

Predicted subcellular locations for the 49,907 Gram-negative proteins			
Accession number	Code in Swiss-Prot database	Uncertain Annotation in Swiss-Prot Database	Predicted subcellular location by Gneg-PLoc
O05011	NQRB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O05012	NQRF_HAEIN	bacterial cell inner membrane (potential).	inner membrane
O05023	Y493_HAEIN		cytoplasm
O05029	ISPD_HAEIN		cytoplasm
O05031	Y737_HAEIN		cytoplasm
O05052	T1RH_HAEIN		cytoplasm
O05069	TRA_HAEIN		cytoplasm
O05070	VGAM_HAEIN		cytoplasm
O05071	Y1494_HAEIN		cytoplasm
O05072	VPG1_HAEIN		cytoplasm
O05073	VPI_HAEIN		cytoplasm
O05074	HLDE_HAEIN		cytoplasm
O05083	Y1698_HAEIN		inner membrane
O05086	Y1721_HAEIN		cytoplasm
O05087	Y1728_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O05104	NADA_EHRCR	cytoplasm (by similarity).	cytoplasm
O05113	Y335_METEX		inner membrane
O05118	KPYK_METEX		cytoplasm
O05131	PBPA_NEIGO	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
O05132	ENGB_NEIGO		cytoplasm
O05137	MASZ_PSEFL	cytoplasm (by similarity).	cytoplasm
O05189	PARA_CAUCR		cytoplasm
O05190	PARB_CAUCR		cytoplasm
O05321	DCDA_PSEFL		cytoplasm
O05322	DAPF_PSEFL	cytoplasm (by similarity).	cytoplasm
O05340	RBFA_STIAU	cytoplasm (potential).	cytoplasm
O05385	RP32_BUCAI		cytoplasm
O05467	MVIN_RHITR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O05485	EPRA1_AERHY		extracellular
O05528	FLII_CAUCR	cytoplasm (potential).	inner membrane
O05542	DHET_GLUOX	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	periplasm
O05543	Y755_GLUOX	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O05544	ADHS_GLUOX	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	periplasm
O05545	IPYR_GLUOX	cytoplasm (potential).	cytoplasm

O05616	VANA_PSEUH		cytoplasm
O05617	VANB_PSEUH		cytoplasm
O05689	CRP_PASMU		cytoplasm
O05700	DNAK_RHOS7		cytoplasm
O05725	ALKB_CAUCR		cytoplasm
O05728	VAPD_HELPY		extracellular
O05730	VDLC_HELPY		cytoplasm
O05732	Y888_HELPY	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
O05927	CYSH_PSEAE	cytoplasm (by similarity).	cytoplasm
O05938	LIFO_PSEWI	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
O05941	LIPA_RHIET	cytoplasm (by similarity).	cytoplasm
O05942	DEGP_RICPR		cytoplasm
O05944	Y218_RICPR		cytoplasm
O05945	Y219_RICPR		extracellular
O05946	PUR7_RICPR		cytoplasm
O05948	FTSY_RICPR	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
O05949	DPO1_RICPR		cytoplasm
O05951	Y673_RICPR		cytoplasm
O05952	KSGA_RICPR		cytoplasm
O05953	MURF_RICPR	cytoplasm (probable).	cytoplasm
O05954	MURE_RICPR	cytoplasm (probable).	cytoplasm
O05955	MFD_RICPR		cytoplasm
O05956	END3_RICPR		cytoplasm
O05957	GLRXA_RICPR		cytoplasm
O05958	Y744_RICPR		cytoplasm
O05959	LIPA_RICPR	cytoplasm (by similarity).	cytoplasm
O05960	Y741_RICPR		inner membrane
O05961	Y740_RICPR		cytoplasm
O05963	Y436_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O05964	RBFA_RICPR	cytoplasm (potential).	cytoplasm
O05965	Y434_RICPR		cytoplasm
O05966	SUCC_RICPR		cytoplasm
O05968	SSRP_RICPR	cytoplasm (potential).	cytoplasm
O05969	DAPA_RICPR	cytoplasm (by similarity).	cytoplasm
O05970	ILVE_RICPR		cytoplasm
O05972	Y028_RICPR		cytoplasm
O05973	UDG_RICPR		cytoplasm
O05974	DPO3A_RICPR	cytoplasm (by similarity).	cytoplasm
O05975	Y785_RICPR		outer membrane
O05976	Y786_RICPR	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
O05977	Y787_RICPR		cytoplasm
O05978	Y788_RICPR		cytoplasm
O05979	Y789_RICPR		cytoplasm
O06430	DNAK_NITEU		cytoplasm
O06431	DNAJ_NITEU	cytoplasm (probable).	cytoplasm

O06432	TONB_NEIGO	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
O06434	EXBD_NEIGO	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
O06465	AHPF_XANCH		cytoplasm
O06522	CDTA_HAEDU	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
O06652	HADDL_PSES4		extracellular
O06659	MSBB2_SHIFL	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
O06663	Y5246_SHIFL		cytoplasm
O06695	VLLY_VIBVU		extracellular
O06702	Y4329_BORBR	periplasm (potential).	periplasm
O06703	BBUR_BORBR		cytoplasm
O06758	FLHA_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O06844	NORC_RHOSH	cell membrane; single-pass membrane protein (by similarity). may be attached to the membrane by a signal- anchor (by similarity).	inner membrane
O06875	SP39_BRUAB	periplasm (probable).	periplasm
O06898	BIOC_ESCVU		cytoplasm
O06899	BIOD_ESCVU		cytoplasm
O06912	FRDC_HELPY	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
O06913	FRDA_HELPY		periplasm
O06914	FRDB_HELPY		cytoplasm
O06922	MDCB_MALRU		cytoplasm
O06961	TDCD_SALTY		cytoplasm
O07025	FLII_HELPY	cytoplasm (potential).	inner membrane
O07026	FLAV_KLEPN		cytoplasm
O07046	SLYD_AERHY	cytoplasm (by similarity).	cytoplasm
O07051	LTAA_AERJA		cytoplasm
O07131	HIS8_METFL		cytoplasm
O07293	BLOI_PSEAE		extracellular
O07299	ALGD_PSESH		cytoplasm
O07308	NODP_RHISB		cytoplasm
O07309	NODQ_RHISB		cytoplasm
O07310	NUOD_RHOCA		cytoplasm
O07315	FUR_RHILV	cytoplasm (by similarity).	cytoplasm
O07350	LIFO_VIBCH	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
O07443	UBIB_PROST	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O07451	HYPF2_RALEU		cytoplasm
O07458	BADR_RHOPA		cytoplasm
O07465	BADM_RHOPA		cytoplasm

O07650	TONB_CAMCO	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
O07674	TNAA_HAEIN		cytoplasm
O07678	PRMA_HELPY	cytoplasm (by similarity).	cytoplasm
O07680	Y1070_HELPY		cytoplasm
O07813	RS6_NEIGO		cytoplasm
O07816	RL9_NEIGO		cytoplasm
O07817	HITA_NEIGO		cytoplasm
O07824	NDOB_PSEFL		cytoplasm
O07898	VIBC_VIBCH		cytoplasm
O07899	VIBE_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O07900	VIBB_VIBCH		cytoplasm
O07948	NUOF_RHOCA		periplasm
O08248	SURE_RHIME	cytoplasm (potential).	periplasm
O08249	PIMT_RHIME	cytoplasm (by similarity).	cytoplasm
O08250	BIOS_RHIME		cytoplasm
O08314	TGT_HELPY		cytoplasm
O08315	GCH2_HELPY		extracellular
O08337	BLA8_ECOLI		extracellular
O08356	DNAJ_RHOS7	cytoplasm (probable).	cytoplasm
O08371	SUCD_RICPR		cytoplasm
O08374	SGAA_HYPME		cytoplasm
O08384	GRPE_NITEU	cytoplasm (probable).	cytoplasm
O08385	HIS1_RHOS4	cytoplasm (by similarity).	cytoplasm
O08400	URE1_BORBR	cytoplasm (by similarity).	cytoplasm
O08429	RP54_VIBAN		cytoplasm
O08430	APHA_SALTI	periplasm (potential).	periplasm
O08437	FKBA_AERHY	periplasm (by similarity).	periplasm
O08444	UDP_KLEAE	cytoplasm (by similarity).	cytoplasm
O09460	PPCK_ANASU	cytoplasm (by similarity).	cytoplasm
O24714	HIS2_KLEPN	cytoplasm (by similarity).	cytoplasm
O24744	RPOD_SHEVI		cytoplasm
O24755	FUR_VIBPA	cytoplasm (by similarity).	cytoplasm
O24840	VANB_ACIAD		cytoplasm
O24849	MTGA_ACIAD	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
O24853	NUSB_HELPY		cytoplasm
O24854	RISB_HELPY		cytoplasm
O24855	CYNT_HELPY		cytoplasm
O24872	BIOD_HELPY		cytoplasm
O24876	Y035_HELPY		inner membrane
O24915	MTNN_HELPY		cytoplasm
O24924	THRC_HELPY		cytoplasm
O24931	LUXS_HELPY		extracellular
O24933	HRCA_HELPY		cytoplasm
O24968	HCPD_HELPY	secreted protein (by similarity).	extracellular
O24970	Y162_HELPY		inner membrane
O24982	LNT_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

O24989	Y189_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O24990	FABI_HELPY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O24991	LPXD_HELPY		cytoplasm
O24993	PLSX_HELPY		cytoplasm
O24994	FABH_HELPY	cytoplasm (probable).	cytoplasm
O24999	MRP_HELPY		cytoplasm
O25004	CDSA_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O25008	ISCS_HELPY		cytoplasm
O25011	MSRAB_HELPY		cytoplasm
O25016	KDSB_HELPY	cytoplasm (potential).	cytoplasm
O25018	Y232_HELPY	bacterial cell membrane; lipid-anchor (probable).	inner membrane
O25021	HCPE_HELPY	secreted protein (potential).	extracellular
O25028	FLGI_HELPY	periplasm (by similarity).	periplasm
O25039	EX7L_HELPY	cytoplasm (by similarity).	cytoplasm
O25042	Y262_HELPY		cytoplasm
O25045	PYRX_HELPY		cytoplasm
O25067	AMIE_HELPY		cytoplasm
O25074	Y303_HELPY		cytoplasm
O25088	TATA_HELPY	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
O25092	FLGH_HELPY	bacterial cell outer membrane (probable).	inner membrane
O25095	LPXK_HELPY		cytoplasm
O25096	NADE_HELPY		cytoplasm
O25097	ILVC_HELPY		cytoplasm
O25098	MIND_HELPY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
O25099	MINE_HELPY		inner membrane
O25101	RUVX_HELPY	cytoplasm (potential).	cytoplasm
O25103	HCPB_HELPY		extracellular
O25114	Y347_HELPY		cytoplasm
O25116	PYRG_HELPY		cytoplasm
O25121	DXS_HELPY		cytoplasm
O25122	LEPA_HELPY		cytoplasm
O25136	DCD_HELPY		extracellular
O25138	Y374_HELPY		cytoplasm
O25149	PRIA_HELPY		cytoplasm
O25151	TPX_HELPY		cytoplasm
O25156	Y395_HELPY		cytoplasm
O25157	Y396_HELPY		cytoplasm
O25165	GUAA_HELPY		cytoplasm
O25170	Y415_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O25225	TYPA_HELPY		cytoplasm

O25235	MRAY_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O25236	MURD_HELPY	cytoplasm (by similarity).	cytoplasm
O25237	Y495_HELPY		cytoplasm
O25242	DPO3B_HELPY	cytoplasm (by similarity).	cytoplasm
O25249	PGBA_HELPY	the plasminogen-binding region is localized on the surface the bacterium (probable).	cytoplasm
O25253	HSLV_HELPY	cytoplasm (by similarity).	cytoplasm
O25254	HSLU_HELPY	cytoplasm (by similarity).	cytoplasm
O25283	ACCA_HELPY	cytoplasm (by similarity).	cytoplasm
O25290	DAPF_HELPY	cytoplasm (by similarity).	cytoplasm
O25291	Y567_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O25294	AMPA_HELPY	cytoplasm (by similarity).	cytoplasm
O25300	LEP_HELPY	bacterial cell membrane; single-pass type ii membrane protein (potential).	inner membrane
O25320	BIOF_HELPY		cytoplasm
O25325	DCUP_HELPY	cytoplasm (probable).	cytoplasm
O25336	DNLJ_HELPY		cytoplasm
O25338	MUTS2_HELPY		cytoplasm
O25340	MURC_HELPY	cytoplasm (probable).	cytoplasm
O25342	ISPG_HELPY		cytoplasm
O25372	GATB_HELPY		cytoplasm
O25376	HEMN_HELPY	cytoplasm (by similarity).	cytoplasm
O25393	GLMU_HELPY	cytoplasm (by similarity).	cytoplasm
O25396	FEOB_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O25411	MRAW_HELPY		cytoplasm
O25424	ASPG_HELPY	cytoplasm (potential).	cytoplasm
O25427	DUS_HELPY		cytoplasm
O25428	TILS_HELPY	cytoplasm (probable).	cytoplasm
O25436	Y736_HELPY		cytoplasm
O25441	Y745_HELPY		cytoplasm
O25443	TRMB_HELPY		cytoplasm
O25455	Y760_HELPY		inner membrane
O25458	FTSY_HELPY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
O25474	LOLA_HELPY	periplasm (by similarity).	periplasm
O25475	SECA_HELPY	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	inner membrane
O25484	RIBB_HELPY		cytoplasm
O25488	ACPS_HELPY	cytoplasm (by similarity).	cytoplasm
O25499	Y823_HELPY		inner membrane
O25502	COAE_HELPY	cytoplasm (by similarity).	cytoplasm
O25503	SPEE_HELPY		cytoplasm
O25505	ENGA_HELPY		cytoplasm
O25506	DBH_HELPY		cytoplasm

O25514	THIE_HELPY		cytoplasm
O25515	THID_HELPY		cytoplasm
O25516	THIM_HELPY		cytoplasm
O25525	GUAC_HELPY		cytoplasm
O25528	GMHA_HELPY	cytoplasm (by similarity).	cytoplasm
O25529	HLDE_HELPY		cytoplasm
O25531	GMHB_HELPY	cytoplasm (by similarity).	cytoplasm
O25534	PGBB_HELPY	the plasminogen-binding region is localized on the surface the bacterium (probable).	cytoplasm
O25536	DUT_HELPY		extracellular
O25537	LPXB_HELPY		cytoplasm
O25544	RUVC_HELPY		cytoplasm
O25549	RUVA_HELPY		cytoplasm
O25551	MVIN_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O25560	HYPB_HELPY		cytoplasm
O25577	CARB_HELPY		cytoplasm
O25578	Y920_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O25581	Y924_HELPY		cytoplasm
O25582	HTPX_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
O25584	SURE_HELPY	cytoplasm (potential).	periplasm
O25587	PTPS_HELPY		extracellular
O25595	ALR_HELPY		cytoplasm
O25597	Y943_HELPY		cytoplasm
O25598	Y944_HELPY		cytoplasm
O25599	Y945_HELPY		cytoplasm
O25603	Y949_HELPY		inner membrane
O25609	LGT_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O25610	Y956_HELPY		cytoplasm
O25613	Y959_HELPY		cytoplasm
O25614	GPDA_HELPY	cytoplasm (probable).	cytoplasm
O25620	VAD2_HELPY		extracellular
O25626	GATC_HELPY		cytoplasm
O25627	BIOA_HELPY		cytoplasm
O25629	FTSA_HELPY		inner membrane
O25654	PPK_HELPY		cytoplasm
O25655	PYRD_HELPY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
O25657	DAPA_HELPY	cytoplasm (by similarity).	cytoplasm
O25664	ISPDF_HELPY		cytoplasm
O25671	FUR_HELPY	cytoplasm (by similarity).	cytoplasm
O25679	FLHF_HELPY		cytoplasm
O25680	HPPK_HELPY		cytoplasm
O25685	Y1044_HELPY		extracellular
O25686	ACSA_HELPY		cytoplasm
O25687	Y1046_HELPY		inner membrane

O25688	RBFA_HELPY	cytoplasm (potential).	cytoplasm
O25690	KHSE_HELPY	cytoplasm (potential).	cytoplasm
O25692	LPXC_HELPY		cytoplasm
O25693	MINC_HELPY		inner membrane
O25698	PANB_HELPY		cytoplasm
O25699	RUVB_HELPY		cytoplasm
O25700	TATB_HELPY	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
O25701	TATC_HELPY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
O25702	QUEA_HELPY	cytoplasm (potential).	cytoplasm
O25703	GIDB_HELPY		cytoplasm
O25716	PYRB_HELPY		cytoplasm
O25728	HCPC_HELPY	secreted protein (potential).	extracellular
O25730	6PGL_HELPY		extracellular
O25731	GLK_HELPY	cytoplasm (by similarity).	cytoplasm
O25748	SLYD_HELPY	cytoplasm (potential).	cytoplasm
O25751	TOLB_HELPY	periplasm (potential).	periplasm
O25754	EXDL1_HELPY	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
O25755	EXBL1_HELPY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O25758	PARB_HELPY		cytoplasm
O25766	TRMD_HELPY	cytoplasm (potential).	cytoplasm
O25767	RIMM_HELPY	cytoplasm (potential).	cytoplasm
O25770	MURG_HELPY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O25773	PROC_HELPY	cytoplasm (by similarity).	cytoplasm
O25775	Y1160_HELPY		inner membrane
O25776	FLAV_HELPY		cytoplasm
O25781	G6PI_HELPY	cytoplasm (by similarity).	cytoplasm
O25788	GLUP_HELPY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
O25797	SOTB_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O25801	DHAS_HELPY		cytoplasm
O25806	RPOBC_HELPY		cytoplasm
O25817	PUR2_HELPY		cytoplasm
O25823	CRCB_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O25826	NUDH_HELPY		cytoplasm
O25827	AK_HELPY		cytoplasm
O25832	Y1234_HELPY	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
O25835	CARA_HELPY		cytoplasm
O25836	AMIF_HELPY		cytoplasm
O25838	Y1240_HELPY	cytoplasm (potential).	inner membrane

O25841	SSB_HELPY		cytoplasm
O25842	Y1247_HELPY		cytoplasm
O25847	SECG_HELPY	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
O25849	NPD_HELPY	cytoplasm (probable).	cytoplasm
O25867	TRPC_HELPY		cytoplasm
O25868	TRPG_HELPY		cytoplasm
O25869	TRPE_HELPY		cytoplasm
O25883	FUMC_HELPY	cytoplasm (by similarity).	cytoplasm
O25888	Y1330_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O25889	Y1331_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O25890	DNAJ_HELPY	cytoplasm (probable).	cytoplasm
O25893	TRMU_HELPY	cytoplasm (by similarity).	cytoplasm
O25895	NADD_HELPY		cytoplasm
O25896	NIKR_HELPY		cytoplasm
O25898	EXBD_HELPY	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
O25899	TONB_HELPY	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
O25901	CORA_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O25903	PLSC_HELPY	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
O25909	NADC_HELPY		cytoplasm
O25910	NADA_HELPY	cytoplasm (by similarity).	cytoplasm
O25911	PSD_HELPY		cytoplasm
O25916	DNAB_HELPY		nucleoid
O25926	CLPX_HELPY		cytoplasm
O25927	LPXA_HELPY	cytoplasm (by similarity).	cytoplasm
O25928	FABZ_HELPY	cytoplasm (by similarity).	cytoplasm
O25936	F16P_HELPY	cytoplasm (potential).	cytoplasm
O25943	RECN_HELPY		cytoplasm
O25944	PPNK_HELPY	cytoplasm (by similarity).	cytoplasm
O25956	BIOB_HELPY		cytoplasm
O25959	QUEF_HELPY	cytoplasm (probable).	cytoplasm
O25961	MIAA_HELPY		cytoplasm
O25963	MURB_HELPY	cytoplasm (probable).	inner membrane
O25966	Y1423_HELPY		cytoplasm
O25970	Y1428_HELPY		cytoplasm
O25971	Y1429_HELPY		cytoplasm
O25972	KSGA_HELPY		cytoplasm
O25982	PPIA_HELPY		cytoplasm
O25983	CSRA_HELPY		cytoplasm
O25984	ISPE_HELPY		cytoplasm
O25985	SSRP_HELPY	cytoplasm (potential).	cytoplasm

O25986	EXBL2_HELPY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O25988	Y1449_HELPY		inner membrane
O25989	OXAA_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O25991	TRME_HELPY		cytoplasm
O26004	ILVE_HELPY		cytoplasm
O26009	KTHY_HELPY		cytoplasm
O26010	COAD_HELPY	cytoplasm (by similarity).	cytoplasm
O26011	PAAD_HELPY		cytoplasm
O26016	EX7S_HELPY	cytoplasm (by similarity).	cytoplasm
O26018	Y1484_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O26024	Y1491_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O26027	MURE_HELPY	cytoplasm (probable).	cytoplasm
O26039	Y1509_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O26043	SELA_HELPY	cytoplasm (by similarity).	cytoplasm
O26051	RECG_HELPY		cytoplasm
O26057	DNAA_HELPY	cytoplasm (by similarity).	cytoplasm
O26060	GLMS_HELPY	cytoplasm (by similarity).	cytoplasm
O26061	THYX_HELPY		cytoplasm
O26066	MFD_HELPY		cytoplasm
O26073	SECF_HELPY	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
O26074	SECD_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O26075	Y1551_HELPY		inner membrane
O26087	ENGB_HELPY		cytoplasm
O26091	RLPA_HELPY	bacterial cell membrane; lipid-anchor (probable).	inner membrane
O26102	PDXJ_HELPY	cytoplasm (probable).	cytoplasm
O26103	PDXA_HELPY	cytoplasm (probable).	cytoplasm
O26106	Y1587_HELPY		inner membrane
O26107	Y1588_HELPY		inner membrane
O30330	FUR_RALME	cytoplasm (by similarity).	cytoplasm
O30332	URED1_RALEU		cytoplasm
O30333	URED2_RALEU		cytoplasm
O30334	URE3_RALEU	cytoplasm (by similarity).	cytoplasm
O30338	UREE_RALEU	cytoplasm (by similarity).	cytoplasm
O30372	PVCC_PSEAE		cytoplasm
O30387	LEP4_MYXXA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O30391	FTSY_NEIMC	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
O30444	Y150_BORPE		inner membrane

O30446	Y148_BORPE	periplasm (potential).	periplasm
O30498	PITX_RHIME		cytoplasm
O30499	PIT_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O30506	AOTP_PSEAE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
O30508	ARUC_PSEAE	cytoplasm (probable).	cytoplasm
O30513	BENK_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O30548	HISZ_PSEST	cytoplasm (probable).	cytoplasm
O30557	AROQ1_PSEAE		cytoplasm
O30569	MTS1_RHIME		cytoplasm
O30611	ICEK_PSESX	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
O30612	CLPP1_MYXXA	cytoplasm (by similarity).	cytoplasm
O30633	RECA_XANOR	cytoplasm (by similarity).	cytoplasm
O30646	PAGO_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O30723	HIS2_RHOCA	cytoplasm (by similarity).	cytoplasm
O30724	HIS6_RHOCA	cytoplasm (by similarity).	cytoplasm
O30725	HIS4_RHOCA	cytoplasm (by similarity).	cytoplasm
O30745	DADA_KLEAE		cytoplasm
O30746	ALR2_KLEAE		cytoplasm
O30762	MIAA_PSEPU		cytoplasm
O30807	MAO1_RHIME		cytoplasm
O30808	MAO2_RHIME		cytoplasm
O30819	BCHI_RHOS4		cytoplasm
O30823	RECR_HAEDU		cytoplasm
O30825	Y322_HAEDU		cytoplasm
O30826	SODM_HAEDU		periplasm
O30827	CITG_HAEDU		cytoplasm
O30847	CLCE_PSESB		cytoplasm
O30862	GRPE_VIBCH	cytoplasm (probable).	cytoplasm
O30867	MUKF_HAEAE	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
O30868	MTH2_HAEAE		cytoplasm
O30869	T2H2_HAEAE		extracellular
O30884	RECA_CAMFE	cytoplasm (by similarity).	cytoplasm
O30911	HSLU_SALTY	cytoplasm (by similarity).	cytoplasm
O30917	SIGE_SALTY	cytoplasm (probable).	cytoplasm
O30919	SOLR2_RALSO		cytoplasm
O30920	SOLI2_RALSO		cytoplasm
O30970	SODF_RHOCA		periplasm
O30971	TNAA_RHOCA		cytoplasm
O30993	FTSQ_RHIME	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
O30994	FTSA_RHIME		inner membrane
O30999	CEST_ECO11	cytoplasm (by similarity).	cytoplasm
O31017	ARCA_RHIET	cytoplasm (potential).	extracellular

O31018	OTCC_RHIET	cytoplasm (probable).	cytoplasm
O31019	ARCC_RHIET	cytoplasm (potential).	cytoplasm
O31020	Y3137_VIBCH		extracellular
O31038	MSUE_PSEAE		cytoplasm
O31047	PURA_EDWIC	cytoplasm (by similarity).	cytoplasm
O31066	CATA_CAUCR		periplasm
O31087	XERC_SERMA	cytoplasm (by similarity).	nucleoid
O31099	SRPA_PSEPU	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
O31100	SRPB_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O31101	SRPC_PSEPU	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
O31151	UVRA_ZYMMO	cytoplasm (by similarity).	cytoplasm
O31158	PRXC_PSEFL		cytoplasm
O31183	NIFH_RHOSH		cytoplasm
O31185	Y4105_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O31186	ADHA_RHIME		cytoplasm
O31196	HUTI_RHIME	cytoplasm (potential).	cytoplasm
O31197	HUTH_RHIME	cytoplasm (potential).	cytoplasm
O31200	HUTI_PSEPU	cytoplasm (potential).	cytoplasm
O31206	XERD_PROMI	cytoplasm (by similarity).	nucleoid
O31207	XERC_PROMI	cytoplasm (by similarity).	nucleoid
O31214	UCRI_CHRVI	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
O31215	CYB_CHRVI		inner membrane
O31216	CY1_CHRVI		inner membrane
O31219	DHAS_LEGPN		cytoplasm
O31222	DRNG_AERHY	secreted protein (potential).	extracellular
O31269	ISCS_AZOVI		cytoplasm
O31285	YREP_BUCTC	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O31288	IBP_BUCTS		cytoplasm
O31289	YREP_BUCTS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O31292	LEU3_BUCTS	cytoplasm (by similarity).	cytoplasm
O31293	LEU2_BUCTS		cytoplasm
O31294	LEUD_BUCTS		cytoplasm
O31381	HEMN_BRAJA	cytoplasm (by similarity).	cytoplasm
O31385	PCAB_BRAJA		cytoplasm
O31395	IRLR_BURPS		cytoplasm
O31396	IRLS_BURPS	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O32323	HSS_RHOVI		cytoplasm
O32347	DKSA_CAUCR		cytoplasm
O32350	PPK_CAMCO		cytoplasm
O32377	RECA_CHRVI	cytoplasm (by similarity).	cytoplasm
O32378	KHSE_METGL		cytoplasm
O32414	ILVC_RHOMO		cytoplasm

O32445	NAGA_VIBCH		cytoplasm
O32464	DNAK_BUCAI		cytoplasm
O32465	DNAJ_BUCAI	cytoplasm (probable).	cytoplasm
O32481	GRPE_LEGPN	cytoplasm (probable).	cytoplasm
O32482	DNAK_LEGPN		cytoplasm
O32483	CAPP_RHOPA		cytoplasm
O32487	PSTB_ENTCL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O32488	PHOU_ENTCL	cytoplasm (by similarity).	cytoplasm
O32513	ENO_DESVM	cytoplasm (by similarity).	extracellular
O32521	PTHC_ERWAM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O32522	PTHB_ERWAM	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O32528	YPDI_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
O32583	THIS_ECOLI		cytoplasm
O32605	CH10_EHRSE	cytoplasm (by similarity).	cytoplasm
O32606	CH60_EHRSE	cytoplasm (by similarity).	cytoplasm
O32613	RL11_HAEDU		cytoplasm
O32614	RL1_HAEDU		cytoplasm
O32620	COPP_HELFE		inner membrane
O32622	Y292_HAEIN		inner membrane
O32712	MDCC_KLEPN	cytoplasm (by similarity).	cytoplasm
O32740	RBL1_RHOCA		cytoplasm
O32741	RBS_RHOCA		cytoplasm
O33367	GYRB_MYXXA		cytoplasm
O33380	TYSY_NEIGO	cytoplasm (by similarity).	cytoplasm
O33395	UVRB_NEIMB	cytoplasm (by similarity).	cytoplasm
O33422	FLISA_PSEAE		flagellar
O33479	ICEV_PSESX	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
O33481	MTP1_PSYTA		cytoplasm
O33499	CH10_PSEST	cytoplasm (by similarity).	cytoplasm
O33500	CH60_PSEST	cytoplasm (by similarity).	cytoplasm
O33505	PQQD_RAHAQ		cytoplasm
O33506	PQQE_RAHAQ		cytoplasm
O33512	GPH_RHOCA		cytoplasm
O33513	CBBY_RHOCA		cytoplasm
O33517	SECD_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O33518	SECF_RHOCA	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
O33519	FMT_RICCN		cytoplasm
O33522	DNAK_RALME		cytoplasm
O33523	FMT_RICFE		cytoplasm
O33525	MDH_RHILV		cytoplasm
O33528	DNAK_RHILE		cytoplasm
O33564	HIS7_RHOS4	cytoplasm (by similarity).	cytoplasm
O33565	HIS5_RHOS4	cytoplasm (by similarity).	cytoplasm

O33568	SECF_RHOS4	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
O33570	CCMA_RHOS4	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O33580	CYSD_RHITR		cytoplasm
O33581	CYSN_RHITR		cytoplasm
O33582	FMT_RICTY		cytoplasm
O33682	Y4338_RHIME		cytoplasm
O33683	Y4337_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O33684	Y4336_RHIME		extracellular
O33685	SYRB1_RHIME		cytoplasm
O33787	YICL_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O33789	WZZE_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O33807	BLC3_SALTY		extracellular
O33817	HCRR_THAAR		cytoplasm
O33818	HCRC_THAAR		cytoplasm
O33819	HCRA_THAAR		periplasm
O33820	HCRB_THAAR		periplasm
O33873	CLPX_YEREN		cytoplasm
O33877	FABA_PSEAE	cytoplasm (by similarity).	cytoplasm
O33915	CISY_RHIME		cytoplasm
O33945	CATR_ACILW		cytoplasm
O33946	CATB1_ACILW		cytoplasm
O33947	CATC1_ACILW		cytoplasm
O33948	CATA1_ACILW		cytoplasm
O33949	CATB2_ACILW		cytoplasm
O33950	CATA2_ACILW		cytoplasm
O33951	CATC2_ACILW		cytoplasm
O33952	UDG8_ECOLI		cytoplasm
O33963	CH60_BARHE	cytoplasm (by similarity).	cytoplasm
O33964	CH60_BARQU	cytoplasm (by similarity).	cytoplasm
O33998	DSRA_CHRVI		cytoplasm
O34011	F16P2_RHOCA		cytoplasm
O34162	HEMN_RALEU	cytoplasm (by similarity).	cytoplasm
O34167	Y565_AGRT5		cytoplasm
O34171	FLII_AGRT5	cytoplasm (potential).	inner membrane
O34201	FLHD_PROMI	cytoplasm (by similarity).	cytoplasm
O34202	FLHC_PROMI		cytoplasm
O34206	KINB_PSEAE	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
O34213	GADH3_PECCY	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	inner membrane
O34214	GADH1_PECCY	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	inner membrane

O34215	GADH2_PECCY	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	cytoplasm
O34238	MVIN_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O34239	RS20_VIBCH		cytoplasm
O34241	DNAK_VIBCH		cytoplasm
O34242	DNAJ_VIBCH	cytoplasm (probable).	cytoplasm
O34246	PAND_WOLSU		cytoplasm
O34247	Y1681_WOLSU		inner membrane
O34258	EXBD2_XANCP	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
O34259	EXBD1_XANCP	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
O34261	TONB_XANCP	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
O34262	WXCX_XANCP		cytoplasm
O34263	SCOB_XANCP		cytoplasm
O34264	SCOA_XANCP		cytoplasm
O34268	TDH_XANCP	cytoplasm (by similarity).	cytoplasm
O34273	TRUB_YEREN		cytoplasm
O34274	RS15_YEREN		cytoplasm
O34275	PNP_YEREN	cytoplasm (by similarity).	cytoplasm
O34291	THIC_RHIET		cytoplasm
O34292	THIO_RHIET		cytoplasm
O34293	THIG_RHIET	cytoplasm (by similarity).	cytoplasm
O34294	THIE_RHIET		cytoplasm
O34295	TTUC5_AGRVI	cytoplasm (by similarity).	cytoplasm
O34296	TTUC3_AGRVI	cytoplasm (by similarity).	cytoplasm
O34370	SERC_NEIMA	cytoplasm (by similarity).	cytoplasm
O34419	RSR1_VIBCH		cytoplasm
O34845	BCHD_RHOS4		cytoplasm
O34955	IPYR_LEGPN	cytoplasm (by similarity).	cytoplasm
O50171	BFR1_MAGMG		cytoplasm
O50172	BFR2_MAGMG		cytoplasm
O50173	IAAH_ENTAG		extracellular
O50174	ASTD_PSEAE		cytoplasm
O50175	ASTB_PSEAE		extracellular
O50177	ASTE_PSEAE		periplasm
O50224	RECG_THIFE		cytoplasm
O50235	CARA_ZYMMO		cytoplasm
O50236	CARB_ZYMMO		cytoplasm
O50237	GREA_ZYMMO		cytoplasm
O50258	NLR_DESGI		inner membrane
O50273	CYSD_PSEAE		cytoplasm
O50274	CYSNC_PSEAE		cytoplasm
O50286	DLDH_VIBPA	cytoplasm (by similarity).	cytoplasm
O50288	ATPA_RICPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O50289	ATPG_RICPR		inner membrane

O50290	ATPB_RICPR		inner membrane
O50468	AROB_NEIGO	cytoplasm (probable).	cytoplasm
O50562	SAHH_RHOSH	cytoplasm (by similarity).	cytoplasm
O50563	RS12_THICU		cytoplasm
O50564	RS7_THICU		cytoplasm
O50580	DT3E_PSECI		cytoplasm
O50584	LTAE_PSEUN		cytoplasm
O50593	ARSA_ACIMU		inner membrane
O50595	ARSC_ACIMU		cytoplasm
O50651	NORC_PARHA	cell membrane; single-pass membrane protein (by similarity). may be attached to the membrane by a signal- anchor (by similarity).	inner membrane
O51798	MMLH_RALEJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O51804	RPOS_VIBCH		cytoplasm
O51809	EXBD_HAEDU	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
O51810	TONB_HAEDU	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
O51824	NRDR_SHIFL		cytoplasm
O51830	TRME_BUCMP		cytoplasm
O51831	CH10_BUCMP	cytoplasm (by similarity).	cytoplasm
O51832	CH60_BUCMP	cytoplasm (by similarity).	cytoplasm
O51859	GYRB_VIBPA		cytoplasm
O51871	ATPE_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
O51873	ATPG_BUCAP		inner membrane
O51874	ATPA_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
O51875	ATPD_BUCAP		inner membrane
O51876	ATPF_BUCAP		inner membrane
O51877	ATPL_BUCAP		inner membrane
O51878	ATP6_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O51879	GIDA_BUCAP	cytoplasm (by similarity).	cytoplasm
O51880	Y583_BUCAP		inner membrane
O51881	ENGA_BUCAP		cytoplasm
O51882	FER_BUCAP		cytoplasm
O51883	HSCA_BUCAP		cytoplasm
O51884	HSCB_BUCAP		inner membrane
O51885	NIFU_BUCAP		cytoplasm
O51886	ISCS_BUCAP		cytoplasm
O51887	ILVD_BUCAP		cytoplasm
O51888	ILVC_BUCAP		cytoplasm
O51889	REP_BUCAP		cytoplasm
O51890	THIO_BUCAP		cytoplasm
O51891	RHO_BUCAP		inner membrane

O51900	FES_DICD3	cytoplasm (by similarity).	cytoplasm
O51926	MURC_BUCAP	cytoplasm (probable).	cytoplasm
O51927	DDL_BUCAP	cytoplasm (by similarity).	cytoplasm
O51928	FTSA_BUCAP		inner membrane
O51930	Y205_BUCAP		inner membrane
O51931	MTNN_BUCAP		cytoplasm
O51940	PVDA_BURCE		cytoplasm
O52042	TONB_VIBCH	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
O52044	EXBD1_VIBCH	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
O52055	SGP3_CHRVI		flagellar
O52057	SGP1_CHRVI		flagellar
O52063	ATZC_PSESD	cytoplasm (potential).	extracellular
O52064	DNAK_PASHA		cytoplasm
O52065	DNAJ_PASHA	cytoplasm (probable).	cytoplasm
O52069	FLIF_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O52070	FLGE_BRUAB		flagellar
O52177	RELA_MYXXA		cytoplasm
O52178	DFRA_MYXXA		cytoplasm
O52179	SGP2_CHRVI		flagellar
O52194	ALGX_AZOVI	periplasm (potential).	periplasm
O52195	ALGL_AZOVI	periplasm (by similarity).	periplasm
O52196	ALGI_AZOVI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O52197	ALGJ_AZOVI	bacterial cell inner membrane; peripheral membrane protein; periplasmic side (by similarity).	periplasm
O52213	LSPA_SERMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
O52222	FLHC_SALTY		cytoplasm
O52236	MFD_MYXXA		cytoplasm
O52248	TTGB_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O52249	ECTA_HALEL		cytoplasm
O52250	ECTB_HALEL		cytoplasm
O52256	FLHF_PSEPU		cytoplasm
O52271	RPOB_RICPR		cytoplasm
O52278	VIRR_AGRTU		inner membrane
O52324	ARNC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O52325	ARNA_SALTY		cytoplasm
O52326	YFBH_SALTY		cytoplasm
O52327	ARNT_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

O52328	YFBW_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O52366	UBIA_PROST	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O52376	DSBA_PSESM	periplasm (by similarity).	periplasm
O52398	RPIA_EDWIC		cytoplasm
O52399	ICIA_EDWIC		cytoplasm
O52400	YGGE_EDWIC		cytoplasm
O52401	MSCS_EDWIC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O52402	ALF_EDWIC		cytoplasm
O52485	PUTA_KLEAE		cytoplasm
O52529	FLJM_CAUCR		flagellar
O52530	FLJN_CAUCR		flagellar
O52531	FLJO_CAUCR		flagellar
O52532	DUSB_SERMA		cytoplasm
O52533	DUSB_PROVU		cytoplasm
O52534	FIS_PROVU		cytoplasm
O52535	CAH_KLEPN	periplasm (potential).	periplasm
O52536	DUSB_KLEPN		cytoplasm
O52538	CAH_ERWCA	periplasm (potential).	periplasm
O52539	DUSB_ERWCA		cytoplasm
O52540	FIS_ERWCA		cytoplasm
O52597	DUT_BRAJA		extracellular
O52618	NODI1_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O52619	NODJ_RHIME	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O52655	APAH_ACTAC		extracellular
O52658	ACP2_PSEAE	cytoplasm (by similarity).	cytoplasm
O52702	MTA1_ACEPA		cytoplasm
O52703	T2A1_ACEPA		extracellular
O52714	TUS_SALTY	cytoplasm (by similarity).	cytoplasm
O52715	TUS_KLEPO	cytoplasm (by similarity).	cytoplasm
O52717	RBTT_KLEPN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O52718	DALT_KLEPN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O52720	DALD_KLEPN		cytoplasm
O52727	LOLB_ACTAC	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
O52728	YA57_ACTAC		inner membrane
O52759	RS4_PSEAE		cytoplasm
O52760	RPOA_PSEAE		cytoplasm
O52761	RL17_PSEAE		cytoplasm
O52762	CATA_PSEAE		periplasm
O52787	PTP_ACIJO		extracellular
O52828	Y1113_ACIAD		inner membrane

O52903	BTUB_CITFR	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
O52908	FLHF_CAMJE		cytoplasm
O52959	FLIC_SALNA		flagellar
O52982	YFJS_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
O54037	VANB_PSEPU		cytoplasm
O54053	GLNB_RHIET		cytoplasm
O54067	LPSL_RHIME		cytoplasm
O54068	UDG_RHIME		cytoplasm
O54070	CYB_RICPR		inner membrane
O54075	PCL_RHOS4		cytoplasm
O54239	FLIF_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O54242	Y649_RHIME		cytoplasm
O54243	FLHB_RHIME	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O54244	FLIG_RHIME	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
O54245	FLIN_RHIME	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
O54246	FLIM_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O54248	FLGF_RHIME		flagellar
O54249	FLII_RHIME	cytoplasm (potential).	inner membrane
O54290	YCDZ_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O54294	CSGD_SALTY		cytoplasm
O54296	RS11_SALTY		cytoplasm
O54297	RS4_SALTY		cytoplasm
O54326	FPG_SALTY		cytoplasm
O54364	RNT_PASHA		cytoplasm
O54367	FIS_PASHA		cytoplasm
O54415	MDCC_ACICA	cytoplasm (by similarity).	cytoplasm
O54418	URE3_ACTPL	cytoplasm (by similarity).	cytoplasm
O54419	URE2_ACTPL	cytoplasm (by similarity).	cytoplasm
O54420	URE1_ACTPL	cytoplasm (by similarity).	cytoplasm
O54422	UREE_ACTPL	cytoplasm (by similarity).	cytoplasm
O54423	UREF_ACTPL		cytoplasm
O54424	UREG_ACTPL		cytoplasm
O54425	URED_ACTPL		cytoplasm
O54426	TYRR_CITBR		cytoplasm
O54435	SACB_RAHAQ		extracellular
O54438	FABG_PSEAE		cytoplasm
O54439	ACP1_PSEAE	cytoplasm (by similarity).	cytoplasm
O54451	RAII_RHIET		cytoplasm
O54452	RAIR_RHIET		cytoplasm
O54453	TRPH_SALTY		cytoplasm
O54458	YDIA_ENTAG		cytoplasm

O54459	AROH_ENTAG		cytoplasm
O54478	PARC_CAUCR	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
O54479	PARE_CAUCR		cytoplasm
O65975	BLC5_SALTY		extracellular
O65976	BLC6_SALTY		extracellular
O66026	CH60_KLEPN	cytoplasm (by similarity).	cytoplasm
O66036	SAT_CHRVI		cytoplasm
O66041	UVRC_SALTY	cytoplasm (by similarity).	cytoplasm
O66108	TYSY_VIBCH	cytoplasm (by similarity).	cytoplasm
O66112	ODPA_ZYMMO		cytoplasm
O66113	ODPB_ZYMMO		cytoplasm
O66114	Y505_ZYMMO		cytoplasm
O66116	Y507_ZYMMO		inner membrane
O66119	ODP2_ZYMMO		inner membrane
O66120	Y511_ZYMMO		cytoplasm
O66132	KHSE_BUCAI	cytoplasm (potential).	cytoplasm
O66143	ARGA_BUCAI	cytoplasm (probable).	cytoplasm
O66164	FTHS_PSEPA		cytoplasm
O66186	SCNB_THITI		extracellular
O66187	SCNA_THITI		extracellular
O66188	SCNC_THITI		cytoplasm
O66392	PHBC_AZOCA	cytoplasm (by similarity).	cytoplasm
O68014	BENM_ACIAD		cytoplasm
O68025	Y166_RHOCA		inner membrane
O68032	SBCB_RHOCA		cytoplasm
O68033	SBCD_RHOCA		cytoplasm
O68038	PHAF_RHOCA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O68040	ACSA_RHOCA		cytoplasm
O68045	DPO3E_RHOCA		cytoplasm
O68068	Y260_RHOCA		inner membrane
O68098	COBK_RHOCA		cytoplasm
O68100	COBM_RHOCA		cytoplasm
O68104	CBIN_RHOCA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O68106	CBIO_RHOCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O68108	COBB_RHOCA		cytoplasm
O68109	ALLA_RHOCA		extracellular
O68126	RS6_RHOCA		cytoplasm
O68127	RS18_RHOCA		cytoplasm
O68128	RL9_RHOCA		cytoplasm
O68129	TIG_RHOCA		cytoplasm
O68141	GID_RHOCA	cytoplasm (by similarity).	cytoplasm
O68191	DNAK_BURPS		cytoplasm
O68215	SACB2_NEIMA		cytoplasm
O68249	RIBB_DEHMU		cytoplasm
O68250	RISB_DEHMU		cytoplasm
O68273	DUSC_RALEU		cytoplasm
O68281	HEXR_PSEAE		cytoplasm
O68282	G6PD_PSEAE		cytoplasm

O68284	MSCL_ERWCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O68286	MSCL_PSEFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O68308	CH10_AERSA	cytoplasm (by similarity).	cytoplasm
O68309	CH60_AERSA	cytoplasm (by similarity).	cytoplasm
O68326	FTSQ_CAUCR	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
O68426	TRPD_BUCDN		cytoplasm
O68427	TRPC_BUCDN		cytoplasm
O68428	TRPB_BUCDN		cytoplasm
O68429	TRPA_BUCDN		cytoplasm
O68433	LEP4_LEGPN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
O68446	PURR_SALTY		cytoplasm
O68451	DBHL_RICRI		cytoplasm
O68452	Y754_RICRI		inner membrane
O68460	HPPA_RHORT	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
O68546	GREA_RHILV		cytoplasm
O68547	LPCC_RHILV		cytoplasm
O68549	LCRR_YEREN		extracellular
O68560	Y4767_PSEAE		cytoplasm
O68561	YOM2_PSEAE		inner membrane
O68564	OMLA_PSEFL	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
O68584	T2H3_HAEAE		extracellular
O68638	RCSF2_PSEAE		inner membrane
O68639	Y523_XYLFA		extracellular
O68662	KDSA_ACTPL	cytoplasm (by similarity).	cytoplasm
O68770	DPO3A_YERPE	cytoplasm (by similarity).	cytoplasm
O68799	MOBA_PSEAE	cytoplasm (by similarity).	cytoplasm
O68822	AMPA_PSEAE	cytoplasm (by similarity).	cytoplasm
O68823	HOLC_PSEAE		cytoplasm
O68826	Y3474_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O68827	RARD_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O68838	GSH1_SALTY		cytoplasm
O68845	RL35_VIBCH		cytoplasm
O68853	NUOB1_RHIME		cytoplasm
O68854	NUOC1_RHIME		inner membrane
O68873	ARGE_MYXXA	cytoplasm (probable).	extracellular
O68874	AQPZ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O68883	CISY_SALTY		cytoplasm
O68901	SODC_SALTY	bacterial cell membrane; lipid-anchor (potential).	periplasm

O68926	BFR_SALTY		cytoplasm
O68927	HOPD_SALTY		inner membrane
O68932	HOPD_ECOLI		inner membrane
O68934	BFD_SERMA		cytoplasm
O68935	BFR_SERMA		cytoplasm
O68937	TBPB_NEIMA	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
O68940	ANFG_RHORU		periplasm
O68943	ANFG_AZOPA		periplasm
O68946	ANFG_AZOMA		periplasm
O68951	VNFG_AZOSA		periplasm
O68964	LEP4_AERSA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
O68965	MI2D_RHIME		cytoplasm
O69014	RNH_ZYMMO	cytoplasm (potential).	extracellular
O69015	KHSE_ZYMMO		cytoplasm
O69049	YEGP_PSEST		outer membrane
O69051	PTXA_PSEST	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O69052	PTXB_PSEST		periplasm
O69053	PTXC_PSEST	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O69054	PTXD_PSEST		cytoplasm
O69060	HTXA_PSEST		cytoplasm
O69061	HTXB_PSEST		periplasm
O69062	HTXC_PSEST	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O69063	HTXD_PSEST	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O69064	HTXE_PSEST	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O69066	HTXG_PSEST		cytoplasm
O69067	HTXH_PSEST		cytoplasm
O69077	AK_PSEAE		cytoplasm
O69078	CSRA_PSEAE		cytoplasm
O69082	COAE_PSEPG	cytoplasm (by similarity).	cytoplasm
O69115	APAH_BURPS		extracellular
O69138	PYP_RHOCA		cytoplasm
O69140	PCL_RHOCA		cytoplasm
O69158	PDXJ_BRAJA	cytoplasm (probable).	cytoplasm
O69159	ACPS_BRAJA	cytoplasm (by similarity).	cytoplasm
O69161	RNC_BRAJA	cytoplasm (by similarity).	cytoplasm
O69162	ERA_BRAJA		cytoplasm
O69169	KTHY_YERPE		cytoplasm
O69170	HOLB_YERPE		cytoplasm
O69177	LON_RHIME	cytoplasm (by similarity).	cytoplasm
O69219	YOR2_AZOVI		inner membrane
O69220	HSCB_AZOVI		inner membrane
O69221	HSCA_AZOVI		cytoplasm
O69232	RPOA_BUCAP		cytoplasm

O69241	HSPD_BRAJA		cytoplasm
O69242	HSPE_BRAJA		cytoplasm
O69280	ECNR_CITFR		cytoplasm
O69289	CH60_CAMJE	cytoplasm (by similarity).	cytoplasm
O69290	MURE_CAMJE	cytoplasm (probable).	cytoplasm
O69294	FUMC_CAMJE	cytoplasm (by similarity).	cytoplasm
O69297	GRPE_CAMJE	cytoplasm (probable).	cytoplasm
O69298	DNAK_CAMJE		cytoplasm
O69300	LON_CAMJE	cytoplasm (by similarity).	cytoplasm
O69301	RS18_CAMJE		cytoplasm
O69302	SSB_CAMJE		cytoplasm
O69395	BLT2_ECOLI		extracellular
O69754	PHZF_PSEAE		cytoplasm
O69761	MSRA_PSEFL		cytoplasm
O69768	AMID_PSEPU		extracellular
O69772	AMPR_PROST	cytoplasm (by similarity).	cytoplasm
O69773	AMPC_PROST	periplasm (by similarity).	periplasm
O69777	YRP2_RHIET		cytoplasm
O69782	DAPA2_RHIME	cytoplasm (by similarity).	cytoplasm
O69786	BETI_RHIME		cytoplasm
O69787	BETC_RHIME		inner membrane
O82850	RS2_PSEAE		cytoplasm
O82852	PYRH_PSEAE		cytoplasm
O82853	RRF_PSEAE	cytoplasm (by similarity).	cytoplasm
O82855	NORM_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O82859	BCSA2_ACEXY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O82860	BCSB2_ACEXY	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
O82861	BCSC2_ACEXY	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
O82868	REGA_RHOSU		cytoplasm
O82922	FINO_ECO57		outer membrane
O82947	CYCR_CHRVI	bacterial cell membrane; lipid-anchor (by similarity).	periplasm
O83046	PCAU_ACIAD		cytoplasm
O84917	RBL2_THIIN		extracellular
O84955	BLOK_PSEAE		extracellular
O84970	NUOB_RHOCA		cytoplasm
O84971	NUOC_RHOCA		inner membrane
O85012	MTDB_METEX	cytoplasm (by similarity).	cytoplasm
O85040	RBL1_THINE		extracellular
O85060	REPA1_BUCAP		nucleoid
O85061	YREP_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O85062	REPA2_BUCAP		nucleoid
O85063	LEU1_BUCAP		cytoplasm
O85064	LEU3_BUCAP	cytoplasm (by similarity).	cytoplasm
O85065	LEU2_BUCAP		cytoplasm
O85066	LEUD_BUCAP		cytoplasm

O85068	YREP_BUCDN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O85070	LEU1_BUCDN		cytoplasm
O85071	LEU3_BUCDN	cytoplasm (by similarity).	cytoplasm
O85072	LEU2_BUCDN		cytoplasm
O85073	LEUD_BUCDN		cytoplasm
O85133	FLIP_RHOSH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O85139	FABH_SALTY	cytoplasm (probable).	cytoplasm
O85140	FABD_SALTY		cytoplasm
O85207	PHAG_PSEPK		cytoplasm
O85213	DNAJ_CAMJE	cytoplasm (probable).	cytoplasm
O85243	SYCT_YEREN		cytoplasm
O85264	RSR2_VIBCH		cytoplasm
O85269	Y6A9_YEREN		inner membrane
O85274	NUOB_PECCC		cytoplasm
O85282	DNAK_EHRSE		cytoplasm
O85290	DAPD_BUCAP	cytoplasm (by similarity).	cytoplasm
O85291	DEGP_BUCAP		cytoplasm
O85292	APBE_BUCAP		cytoplasm
O85293	ILVI_BUCAP		cytoplasm
O85294	ILVH_BUCAP		cytoplasm
O85295	MRAW_BUCAP	cytoplasm (by similarity).	cytoplasm
O85296	FTSL_BUCAP		inner membrane
O85297	FTSI_BUCAP	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
O85298	MURE_BUCAP	cytoplasm (probable).	cytoplasm
O85304	PEPD_SALDU		cytoplasm
O85306	SIGE_SALDU	cytoplasm (probable).	cytoplasm
O85341	WBDQ_ECO57		cytoplasm
O85342	MANC_ECO57		cytoplasm
O85343	MANB_ECO57		cytoplasm
O85347	PYRG_NITEU		cytoplasm
O85348	ENO_NITEU	cytoplasm (by similarity).	extracellular
O85387	RL22_COXBU		cytoplasm
O85388	RS3_COXBU		cytoplasm
O85713	NOEL_RHIFR		cytoplasm
O85718	GLYA_ACIRA	cytoplasm (by similarity).	cytoplasm
O85732	FMT_PSEAE		cytoplasm
O85735	CSRA_SERMA		cytoplasm
O85754	CH60_RICTY	cytoplasm (by similarity).	cytoplasm
O85762	SSUE_PSEPU		cytoplasm
O85764	SSUD_PSEPU		cytoplasm
O85806	FLHD_SERMA	cytoplasm (by similarity).	cytoplasm
O85807	FLHC_SERMA		cytoplasm
O85987	PDXA_SPHAR	cytoplasm (probable).	cytoplasm
O86013	BPHF_SPHAR		cytoplasm
O86028	MUTB_RHIME		cytoplasm
O86029	Y6150_RHIME		cytoplasm
O86033	GLPK_RHIME		cytoplasm
O86034	BDHA_RHIME		cytoplasm
O86046	FLHD_YEREN	cytoplasm (by similarity).	cytoplasm
O86047	FLHC_YEREN		cytoplasm

O86062	NUDC_PSEAE		cytoplasm
O86082	RECX_HERSE	cytoplasm (potential).	cytoplasm
O86088	PBPA_NEICI	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
O86090	PPK_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
O86094	RPOB_LEGPN		cytoplasm
O86110	HSPH_BRAJA		cytoplasm
O86200	MCHI_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O86220	Y148A_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O86221	Y213A_HAEIN		inner membrane
O86222	Y220B_HAEIN		cytoplasm
O86223	Y221A_HAEIN		cytoplasm
O86225	Y485A_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O86226	Y559A_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O86227	Y602A_HAEIN		extracellular
O86228	Y666A_HAEIN		cytoplasm
O86229	Y870A_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O86230	Y976A_HAEIN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
O86231	Y1017_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O86232	Y103A_HAEIN		inner membrane
O86233	Y112B_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O86234	Y1127_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O86235	Y122B_HAEIN		cytoplasm
O86236	Y132A_HAEIN		cytoplasm
O86237	Y138A_HAEIN		cytoplasm
O86239	Y143B_HAEIN		cytoplasm
O86241	Y146A_HAEIN		outer membrane
O86242	Y149A_HAEIN		cytoplasm
O86243	Y156A_HAEIN		cytoplasm
O86422	UDG_PSEAE		cytoplasm
O86428	ILVE_PSEAE		cytoplasm
O86436	AMPA_PSEPU	cytoplasm (by similarity).	cytoplasm
O86447	CALB_PSEUH		cytoplasm
O86459	AAT_RHILP	cytoplasm (by similarity).	cytoplasm
O86464	FIXT_RHIME		cytoplasm

O86988	UMOB_PROMI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
O87005	CHPE_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O87006	AROA_SHIDY	cytoplasm (probable).	cytoplasm
O87007	AROD_SHIDY		cytoplasm
O87014	USG_PSEAE		cytoplasm
O87016	TRUA_PSEAE		cytoplasm
O87077	LGT_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
O87120	CDTA_ACTAC	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
O87131	CHER1_PSEAE		cytoplasm
O87278	STCD_RHIME		cytoplasm
O87320	AATC_RHIME	cytoplasm (by similarity).	cytoplasm
O87322	SFSA_RHIME		cytoplasm
O87326	TRL_HELPY		extracellular
O87340	GMHA_HAEDU	cytoplasm (by similarity).	cytoplasm
O87384	DNAK_VIBHA		cytoplasm
O87385	DNAJ_VIBHA	cytoplasm (probable).	cytoplasm
O87387	SOXD_RHIME	cytoplasm (by similarity).	cytoplasm
O87388	SOXB_RHIME	cytoplasm (by similarity).	cytoplasm
O87389	GLXA_RHIME		cytoplasm
O87390	GLXB_RHIME		cytoplasm
O87391	GLXC_RHIME		cytoplasm
O87392	GLXD_RHIME		cytoplasm
O87393	GLNA3_RHIME	cytoplasm (potential).	cytoplasm
O87394	Y093_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
O87405	GLSA_RHIEC		extracellular
O87406	Y425_NEIG1		inner membrane
O87407	ENGA_NEIG1		cytoplasm
O87408	RDGC_NEIG1		cytoplasm
O87475	DBH_CAUCR		cytoplasm
O87574	YBEY_SALTY		inner membrane
O87579	PBPA_NEILA	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
O87612	CLCE_PSEAE		cytoplasm
O87626	PBPA_NEIFL	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
O87627	NIFN_HERSE		periplasm
O87628	NIFX_HERSE		cytoplasm
O87663	YADU_SALTY		extracellular
O87667	GYRB_CAMJE		cytoplasm
O87705	TIG_CAUCR		cytoplasm
O87706	CLPP_CAUCR	cytoplasm (by similarity).	cytoplasm
O87707	CICA_CAUCR		cytoplasm
O87708	CLPX_CAUCR		cytoplasm
O87715	CHEW_CAUCR	cytoplasm (by similarity).	cytoplasm
O87791	RS15_PSEPU		cytoplasm

O87792	PNP_PSEPU	cytoplasm (by similarity).	cytoplasm
O87796	ALF_PSEST		cytoplasm
O87807	INH_PSETO	periplasm (by similarity).	periplasm
O87816	FDHD_RALEU	cytoplasm (potential).	cytoplasm
O87873	DCH_THAAR		cytoplasm
O87885	RL27_LAWIN		cytoplasm
O87886	RL21_LAWIN		cytoplasm
O87887	CH10_LAWIN	cytoplasm (by similarity).	cytoplasm
O87896	DSRE_CHRVI	cytoplasm (by similarity).	cytoplasm
O87897	DSRF_CHRVI		cytoplasm
O87902	RL7_COXBU		cytoplasm
O87903	RPOB_COXBU		cytoplasm
O87908	SYRA_RHIME		cytoplasm
O87924	GLPK_PSETO		cytoplasm
O87949	TORD_SHEMA	cytoplasm (by similarity).	cytoplasm
O87970	YUKI_YERRU		cytoplasm
O88005	BAF_BORBR		cytoplasm
P00080	CYC2_RHOGL		periplasm
P00081	CYC2_AGRTC		periplasm
P00082	CYC2_RHOVA		periplasm
P00083	CYC2_RHOVI		periplasm
P00084	CYC2_RHOAC		periplasm
P00085	CY550_NITWI		periplasm
P00086	CYC21_RHOFU		periplasm
P00087	CYC21_RHOMO		periplasm
P00088	CYC22_RHOMO		periplasm
P00089	CYC22_RHOFU		periplasm
P00090	CYC21_RHOPA		periplasm
P00091	CYC22_RHOPA		periplasm
P00093	CYC2_RHOPH		periplasm
P00094	CYC2_RHOCA		periplasm
P00096	CY550_PARDE		periplasm
P00097	CYC2_RHOGE		periplasm
P00098	CYC2_RHOTE		periplasm
P00100	CY551_PSEFC		periplasm
P00102	CY551_PSEME		periplasm
P00103	CY551_PSEDE		periplasm
P00104	CY551_AZOVI		periplasm
P00105	C554_PARSP		periplasm
P00121	CYC5_PSEME		periplasm
P00122	CY551_ECTHA		periplasm
P00133	CYC3_DESGI		periplasm
P00135	CYC3_DESSA		periplasm
P00137	CYC3_DESAC		periplasm
P00139	C556_AGRTB		periplasm
P00140	C556_AGRTC		periplasm
P00141	C556_AGRTA		periplasm
P00142	CYCP_RHOGE		periplasm
P00143	CYCP_PARSP		periplasm
P00144	CYCP_RHORT		periplasm
P00145	CYCP_RHOPH		periplasm
P00146	CYCP_RHOSP		periplasm
P00147	CYCP_RHOCA		periplasm
P00148	CYCP_RHOS4		periplasm
P00149	CYCP_RHOPA		periplasm
P00150	C556_RHOPA		periplasm

P00151	CYCP_RHOFU		periplasm
P00152	CYCP_RHOMO		periplasm
P00153	CYCP_RHOTE		periplasm
P00154	CYCP_CHRVI		periplasm
P00183	CPXA_PSEPU		inner membrane
P00194	FER1_RHORU		periplasm
P00207	FER1_RHOPA		periplasm
P00208	FER_CHRVI		periplasm
P00209	FER_DESGI		cytoplasm
P00210	FER1_DESAF		periplasm
P00214	FER1_AZOVI		periplasm
P00259	PUTX_PSEPU		cytoplasm
P00261	HIP_THIRO		periplasm
P00263	HIP_THIPF		periplasm
P00264	HIP_PARSP		periplasm
P00265	HIP_RHOGE		periplasm
P00266	HIP1_RHOTE		periplasm
P00272	RUBR2_PSEOL		cytoplasm
P00323	FLAV_DESVH		cytoplasm
P00324	FLAV_AZOVI		cytoplasm
P00335	RIDH_KLEAE		cytoplasm
P00350	6PGD_ECOLI		cytoplasm
P00363	FRDA_ECOLI		periplasm
P00370	DHE4_ECOLI		cytoplasm
P00382	DYR1_ECOLI		cytoplasm
P00383	DYR21_ECOLI		cytoplasm
P00384	DYR22_ECOLI		cytoplasm
P00386	DHNO_AGRT7		cytoplasm
P00392	MERA_PSEAE		cytoplasm
P00436	PCXA_PSEPU		cytoplasm
P00437	PCXB_PSEPU		cytoplasm
P00438	PHHY_PSEFL		cytoplasm
P00448	SODM_ECOLI		periplasm
P00452	RIR1_ECOLI		cytoplasm
P00458	NIFH_KLEPN		cytoplasm
P00459	NIFH1_AZOVI		cytoplasm
P00460	NIFH_RHIME		cytoplasm
P00461	NIFH_RHILT		cytoplasm
P00462	NIFH_RHIET		cytoplasm
P00463	NIFH_BRASP		cytoplasm
P00466	NIFD_KLEPN		periplasm
P00472	MTE1_ECOLI		cytoplasm
P00473	MTH2_HAEPH		cytoplasm
P00474	MTPS_PROST		cytoplasm
P00484	CAT3_ECOLI		cytoplasm
P00490	PHSM_ECOLI		cytoplasm
P00500	TRPD_ACIAD		cytoplasm
P00547	KHSE_ECOLI	cytoplasm (probable).	cytoplasm
P00551	KKA1_ECOLI		cytoplasm
P00552	KKA2_KLEPN		cytoplasm
P00557	KHYB_ECOLI		cytoplasm
P00561	AK1H_ECOLI		cytoplasm
P00562	AK2H_ECOLI		cytoplasm
P00579	RPOD_ECOLI		cytoplasm
P00582	DPO1_ECOLI		cytoplasm
P00632	ELH2_ACIAD		cytoplasm

P00640	T2PS_PROST		extracellular
P00642	T2E1_ECOLI		extracellular
P00643	T2H2_HAEPH		extracellular
P00645	CEAC_ECOLI		extracellular
P00646	CEA3_ECOLI		extracellular
P00722	BGAL_ECOLI		cytoplasm
P00778	PRLA_LYSEN		cytoplasm
P00801	PRLB_LYSEN		extracellular
P00816	CYNS_ECOLI		cytoplasm
P00861	DCDA_ECOLI		cytoplasm
P00864	CAPP_ECOLI		cytoplasm
P00885	ALKD_PSEPU		cytoplasm
P00887	AROH_ECOLI		cytoplasm
P00888	AROF_ECOLI		cytoplasm
P00892	ILVG_ECOLI		cytoplasm
P00893	ILVI_ECOLI		cytoplasm
P00894	ILVH_ECOLI		cytoplasm
P00895	TRPE_ECOLI		cytoplasm
P00897	TRPE_SERMA		cytoplasm
P00898	TRPE_SALTY		cytoplasm
P00900	TRPG_SERMA		cytoplasm
P00901	TRPG_PSEPU		cytoplasm
P00902	TRPG_ACIAD		cytoplasm
P00903	PABA_ECOLI		cytoplasm
P00904	TRPG_ECOLI		cytoplasm
P00905	TRPG_SALTY		cytoplasm
P00909	TRPC_ECOLI		cytoplasm
P00910	TRPC_SALTY		cytoplasm
P00911	TRPC_ACIAD		cytoplasm
P00914	PHR_ECOLI		cytoplasm
P00926	SDHD_ECOLI		cytoplasm
P00929	TRPA_SALTY		cytoplasm
P00930	TRPA_KLEAE		cytoplasm
P00934	THRC_ECOLI		cytoplasm
P00946	MANA_ECOLI	cytoplasm (probable).	cytoplasm
P00947	SDIS_COMTE		inner membrane
P00948	CATC_PSEPU		cytoplasm
P00968	CARB_ECOLI		cytoplasm
P01555	CHTA_VIBCH		extracellular
P01556	CHTB_VIBCH		extracellular
P02344	DBH_RHIME		cytoplasm
P02348	DBH5_RHILE		cytoplasm
P02358	RS6_ECOLI		cytoplasm
P02359	RS7_ECOLI		cytoplasm
P02393	RL7_DESVM		cytoplasm
P02397	RL7_RHOSH		cytoplasm
P02413	RL15_ECOLI		cytoplasm
P02918	PBPA_ECOLI	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P02966	DESS_MYXXA		extracellular
P02969	FLJJ_CAUCR		flagellar
P02970	FMKB_ECOLI		extracellular
P02971	FMC1_ECOLI		fimbrium
P02973	FMPA_PSEAE		extracellular
P02974	FMM1_NEIGO		extracellular

P02975	FMAA_BACNO		extracellular
P02978	CEA1_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	extracellular
P02984	IMM3_ECOLI		cytoplasm
P02985	IMM1_ECOLI		cytoplasm
P02986	IMMC_ECOLI		cytoplasm
P02988	GENK_ECOLI		cytoplasm
P02989	GENL_ECOLI		extracellular
P03007	DPO3E_ECOLI		cytoplasm
P03008	TNP3_ECOLI		cytoplasm
P03009	TRA2_ECOLI		cytoplasm
P03012	TNR1_ECOLI		cytoplasm
P03013	HIN_SALTY		cytoplasm
P03014	PINE_ECOLI		cytoplasm
P03018	UVRD_ECOLI		cytoplasm
P03023	LACI_ECOLI		cytoplasm
P03024	GALR_ECOLI		cytoplasm
P03027	NIFA_KLEPN		cytoplasm
P03028	NIFA_RHIME		cytoplasm
P03029	NTRC_KLEPN		cytoplasm
P03031	NODD1_RHIME		cytoplasm
P03038	TETR1_ECOLI		cytoplasm
P03039	TETR3_ECOLI		cytoplasm
P03051	ROP_ECOLI		cytoplasm
P03052	KORA2_ECOLI		cytoplasm
P03066	REP2_ECOLI		nucleoid
P03067	PIR_ECOLI		cytoplasm
P03813	YGEA_ECOLI		fimbrium
P03817	MIOC_ECOLI		cytoplasm
P03825	PIOO_ECOLI		extracellular
P03828	INA1_SHIDY		cytoplasm
P03829	INA2_SHIDY		cytoplasm
P03830	INSB1_ECOLI		cytoplasm
P03831	INBD_SHIDY		cytoplasm
P03832	INBN_SHIDY		cytoplasm
P03833	NIFJ_KLEPN		cytoplasm
P03835	INSG_ECOLI		cytoplasm
P03837	INSH_ECOLI		cytoplasm
P03839	YT93_ECOLI		inner membrane
P03842	RNPH_ECOLI		cytoplasm
P03847	COPB2_ECOLI		cytoplasm
P03848	YPRR_ECOLI		extracellular
P03849	YPK6_ECOLI		cytoplasm
P03850	YPB2_ECOLI		cytoplasm
P03851	YPB3_ECOLI		cytoplasm
P03852	YPB1_ECOLI		cytoplasm
P03853	YPB4_ECOLI		extracellular
P03854	COPA1_ECOLI		cytoplasm
P03856	REPE1_ECOLI		cytoplasm
P03869	IPT_AGRT4		cytoplasm
P03961	ATKC_ECOLI	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
P04028	YP1_AGRT4		cytoplasm
P04030	YP6_AGRT4		cytoplasm

P04079	GUAA_ECOLI		cytoplasm
P04129	MERP_SHIFL	periplasm (probable).	inner membrane
P04130	TNR2_ECOLI		cytoplasm
P04131	MERP_PSEAE	periplasm (probable).	inner membrane
P04139	MERC_PSEAE		inner membrane
P04152	UMUC_ECOLI		cytoplasm
P04162	TNPM_ECOLI		cytoplasm
P04168	HIP1_ECTHA		periplasm
P04169	HIP2_ECTHA		periplasm
P04174	DYR_NEIGO		cytoplasm
P04252	BAHG_VITST		cytoplasm
P04335	Yafa_ECOLI		cytoplasm
P04337	MERC_SHIFL		inner membrane
P04340	NODC_RHILV	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P04341	NODC_RHIME	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P04369	C555_METCA		periplasm
P04390	T2E5_ECOLI		extracellular
P04391	OTC1_ECOLI	cytoplasm (probable).	cytoplasm
P04393	MTE5_ECOLI		cytoplasm
P04395	3MG2_ECOLI		cytoplasm
P04419	CEA2_ECOLI		extracellular
P04425	GSHB_ECOLI		cytoplasm
P04479	CEIB_ECOLI		extracellular
P04480	CEA_CITFR	bacterial cell membrane; multi-pass membrane protein (potential).	extracellular
P04482	IMM2_ECOLI		cytoplasm
P04483	TETR2_ECOLI		cytoplasm
P04668	FLAW_KLEPN		cytoplasm
P04680	NODD_RHILT		cytoplasm
P04681	NODD_RHILV		cytoplasm
P04682	NODD_BRASP		cytoplasm
P04685	NODF_RHILV		cytoplasm
P04686	NODF_RHILT		cytoplasm
P04718	RBL2_RHORU		cytoplasm
P04739	FMPO_PSEAE		extracellular
P04740	FMK1_ECOLI		extracellular
P04743	PAPI_ECOLI		cytoplasm
P04787	LEUD1_SALTY		cytoplasm
P04949	FLIC_ECOLI		flagellar
P04951	KDSB_ECOLI	cytoplasm (potential).	cytoplasm
P04953	FMAH_BACNO		extracellular
P04968	THD1_ECOLI		cytoplasm
P04977	TOX1_BORPE		extracellular
P04978	TOX2_BORPE		extracellular
P04979	TOX3_BORPE		extracellular
P04981	TOX5_BORPE		extracellular
P04982	RBSD_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P04993	EX5A_ECOLI		cytoplasm
P04995	EX1_ECOLI		cytoplasm

P05020	PYRC_ECOLI		cytoplasm
P05033	DCLY_HAFAL	cytoplasm (probable).	cytoplasm
P05034	DCHS_MORMO		cytoplasm
P05036	ATPA_RHORU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P05038	ATPB_RHORU		inner membrane
P05041	PABB_ECOLI		cytoplasm
P05050	ALKB_ECOLI		cytoplasm
P05052	APPY_ECOLI		cytoplasm
P05100	3MG1_ECOLI		cytoplasm
P05101	MTE2_ECOLI		cytoplasm
P05102	MTH1_HAEPH		cytoplasm
P05103	MTP7_PSEAE		cytoplasm
P05104	T2P7_PSEAE		extracellular
P05192	BLAC_KLEPN		extracellular
P05193	AMPC_CITFR	periplasm (by similarity).	periplasm
P05194	AROD_ECOLI		cytoplasm
P05301	T2D1_DESDN		extracellular
P05302	MTD1_DESDN		cytoplasm
P05338	RCSA_KLEAE		cytoplasm
P05339	RCSB_KLEAE		cytoplasm
P05340	NIFU_AZOVI		cytoplasm
P05341	NIFS_AZOVI		cytoplasm
P05342	NIFV_AZOVI		cytoplasm
P05343	NIFU_KLEPN		cytoplasm
P05344	NIFS_KLEPN		cytoplasm
P05345	NIFV_KLEPN		cytoplasm
P05351	VIRB2_AGRT9		extracellular
P05356	VIRB6_AGRT9		cytoplasm
P05357	VIRB8_AGRT9		cytoplasm
P05359	VIRBA_AGRT9		cytoplasm
P05361	ALDC_ENTAE		cytoplasm
P05364	AMPC_ENTCL	periplasm (by similarity).	periplasm
P05384	DBHB_PSEAE		cytoplasm
P05404	TFDD1_RALEJ		cytoplasm
P05406	FIXR_BRAJA		cytoplasm
P05407	NIFA_BRAJA		cytoplasm
P05415	GLGC_SALTY		cytoplasm
P05416	GLGA_SALTY		cytoplasm
P05418	CYB_PARDE		inner membrane
P05431	FMM1_NEIMB		extracellular
P05436	ATPG_RHOBL		inner membrane
P05437	ATPD_RHOBL		inner membrane
P05438	ATPD_RHORU		inner membrane
P05439	ATPA_RHOBL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P05440	ATPB_RHOBL		inner membrane
P05441	ATPE_RHOBL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P05442	ATPE_RHORU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P05444	CLPA_RHOBL		cytoplasm

P05445	PRIA_RHORU		cytoplasm
P05446	GLO2_RHOBL		cytoplasm
P05447	YAT4_RHOBL		cytoplasm
P05448	YAT5_RHOBL		cytoplasm
P05449	YAT6_RHOBL		cytoplasm
P05450	YAT7_RHOBL		cytoplasm
P05459	PDXB_ECOLI		cytoplasm
P05521	LUXD_VIBHA		cytoplasm
P05523	FPG_ECOLI		cytoplasm
P05529	MCBF_ECOLI		inner membrane
P05530	MCBG_ECOLI		inner membrane
P05637	APAH_ECOLI		extracellular
P05679	YI32_AGRT7		inner membrane
P05680	YI31_AGRT7		inner membrane
P05682	REPA_AGRRH		cytoplasm
P05683	REPB_AGRRH		nucleoid
P05684	REPC_AGRRH		nucleoid
P05706	PTHA_ECOLI	cytoplasm (probable).	cytoplasm
P05707	SRLD_ECOLI		cytoplasm
P05719	T1SK_ECOLI		cytoplasm
P05789	EREB_ECOLI		cytoplasm
P05791	ILVD_ECOLI		cytoplasm
P05793	ILVC_ECOLI		cytoplasm
P05794	DYR23_ECOLI		cytoplasm
P05804	BGLR_ECOLI		cytoplasm
P05818	BMAE_ECOLI		extracellular
P05819	CEAB_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	extracellular
P05820	CEAM_ECOLI		extracellular
P05822	INSF_ECOLI		cytoplasm
P05823	TNR0_ECOLI		cytoplasm
P05824	RECN_ECOLI		cytoplasm
P05831	INCB_ECOLI		cytoplasm
P05833	REPA_ECOLI		cytoplasm
P05834	MCBA_ECOLI		extracellular
P05835	TRAY3_ECOLI		cytoplasm
P05836	TRAY4_ECOLI		cytoplasm
P05839	EXC_ECOLI		cytoplasm
P05845	TNSE_ECOLI		cytoplasm
P05846	TNSC_ECOLI		cytoplasm
P05847	TTDA_ECOLI		cytoplasm
P05852	GCP_ECOLI		extracellular
P05853	CIT2_ECOLI		cytoplasm
P05989	ILVC_SALTY		cytoplasm
P06008	RCEH_RHOVI		inner membrane
P06117	NIFH_BRAJA		cytoplasm
P06118	NIFH2_AZOCH		cytoplasm
P06120	NIFD_BRASP		periplasm
P06121	NIFD_BRAJA		periplasm
P06122	NIFK_BRASP		cytoplasm
P06123	FDXN_AZOCH		periplasm
P06134	ADA_ECOLI		cytoplasm
P06173	RPOB_SALTY		cytoplasm
P06175	FLIC_SALRU		flagellar
P06176	FLIC_SALCH		flagellar

P06177	FLIC_SALMU		flagellar
P06178	FLIC_SALPA		flagellar
P06179	FLIC_SALTY		flagellar
P06184	PGTA_SALTY	cytoplasm (probable).	cytoplasm
P06185	PGTE_SALTY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P06187	T1S_SALTY		cytoplasm
P06188	ARAB_SALTY		cytoplasm
P06189	ARAA_SALTY		cytoplasm
P06190	ARAD_SALTY		cytoplasm
P06191	ALR2_SALTY		cytoplasm
P06193	PABA_SALTY		cytoplasm
P06194	PABA_KLEAE		cytoplasm
P06195	PABA_SERMA		cytoplasm
P06196	USHA_SALTY	periplasm (by similarity).	periplasm
P06200	PHLC_PSEAE		extracellular
P06203	METJ_SALTY	cytoplasm (by similarity).	cytoplasm
P06204	PYRC_SALTY		cytoplasm
P06218	NTRB_KLEOX		inner membrane
P06219	BGAL_KLEPN		cytoplasm
P06223	RP54_KLEOX		cytoplasm
P06232	NODF1_RHIME		cytoplasm
P06233	NODF2_RHIME		cytoplasm
P06234	NODG_RHIME		cytoplasm
P06235	NOG4_RHIME		cytoplasm
P06236	NODH_RHIME		cytoplasm
P06237	NOH4_RHIME		cytoplasm
P06519	XYLR_PSEPU		cytoplasm
P06524	IPTZ_AGRT7		cytoplasm
P06608	ASPG_ERWCH		cytoplasm
P06610	BTUE_ECOLI	periplasm (probable).	periplasm
P06612	TOP1_ECOLI		cytoplasm
P06615	REDF_ECOLI		cytoplasm
P06617	TR2M_PSESS		cytoplasm
P06618	HYIN_PSESS		extracellular
P06619	IPT_PSESS		cytoplasm
P06621	CBPG_PSES6		extracellular
P06622	XYLE1_PSEPU		cytoplasm
P06627	TRAY1_ECOLI		cytoplasm
P06661	NIFH_THIFE		cytoplasm
P06662	NIFD_THIFE		periplasm
P06665	VIRC1_AGRTU		cytoplasm
P06666	VIRC2_AGRTU		cytoplasm
P06667	VIRD1_AGRTU		extracellular
P06668	VIRD2_AGRTU		extracellular
P06672	PDC_ZYMMO		cytoplasm
P06689	MERD_PSEAE		cytoplasm
P06690	MERE_PSEAE		inner membrane
P06691	TNR5_PSEAE		cytoplasm
P06692	TNR7_ECOLI		cytoplasm
P06694	TNP2_ECOLI		cytoplasm
P06695	TNP5_PSEAE		cytoplasm
P06709	BIRA_ECOLI		cytoplasm
P06710	DPO3X_ECOLI		cytoplasm

P06716	CEIA_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	extracellular
P06717	ELAP_ECOLI		extracellular
P06720	AGAL_ECOLI		cytoplasm
P06736	HLYC_ECOLI	cytoplasm (potential).	cytoplasm
P06755	NODJ_RHILV	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P06758	ADH2_ZYMMO		cytoplasm
P06769	NIFD_RHICP		periplasm
P06770	NIFB_BRAJA		cytoplasm
P06772	NIFL_KLEPN		inner membrane
P06846	EBGR_ECOLI		cytoplasm
P06864	BGA2_ECOLI		cytoplasm
P06959	ODP2_ECOLI		inner membrane
P06961	CCA_ECOLI		cytoplasm
P06962	LYS1_CITFR		outer membrane
P06965	DICC_ECOLI		cytoplasm
P06966	DICA_ECOLI		cytoplasm
P06968	DUT_ECOLI		extracellular
P06970	FAED_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P06983	HEM3_ECOLI		cytoplasm
P06986	HIS8_ECOLI		cytoplasm
P06988	HISX_ECOLI		cytoplasm
P06990	T1SB_ECOLI		cytoplasm
P06991	T1SD_ECOLI		cytoplasm
P06992	KSGA_ECOLI		cytoplasm
P06993	MALT_ECOLI		cytoplasm
P06999	K6PF2_ECOLI		cytoplasm
P07010	REM_ECOLI		cytoplasm
P07013	PRIB_ECOLI		cytoplasm
P07021	YFIB_ECOLI	bacterial cell membrane; lipid-anchor (potential).	outer membrane
P07026	SDIA_ECOLI		cytoplasm
P07047	CPPC_NEIGO		cytoplasm
P07048	CPPA_NEIGO		cytoplasm
P07049	CPPB_NEIGO		nucleoid
P07104	FANA_ECOLI		cytoplasm
P07105	FANB_ECOLI		cytoplasm
P07112	MBA2_ECOLI		cytoplasm
P07113	MOBB2_ECOLI		cytoplasm
P07114	MOBC_ECOLI		cytoplasm
P07115	YPMC_ECOLI		inner membrane
P07116	YPMF_ECOLI		inner membrane
P07165	VIRC1_AGRT5		cytoplasm
P07166	VIRC2_AGRT5		cytoplasm
P07167	VIRAL_AGRTU		inner membrane
P07168	VIRAW_AGRTU		inner membrane
P07169	VIRBB_AGRT5	cytoplasm (probable).	inner membrane
P07175	PARA_AGRTU		cytoplasm
P07206	PULA_KLEPN	bacterial cell membrane; lipid-anchor (probable).	extracellular
P07212	H8_NEIMC		periplasm

P07227	ATPG_RHORU		inner membrane
P07254	CHIA_SERMA		extracellular
P07328	NIFD_AZOVI		periplasm
P07329	NIFK_AZOVI		cytoplasm
P07344	TRPA_PSEAE		cytoplasm
P07345	TRPB_PSEAE		cytoplasm
P07346	ASPA_PSEFL		cytoplasm
P07362	PRIM_SALTY		cytoplasm
P07364	CHER_ECOLI		cytoplasm
P07375	MUCB_ECOLI		cytoplasm
P07383	TPES_PSEPU		cytoplasm
P07386	MBHA_MYXXA		cytoplasm
P07445	SDIS_PSEPU		inner membrane
P07485	FER1_DESDN		periplasm
P07543	VIRE1_AGRTU		cytoplasm
P07604	TYRR_ECOLI		cytoplasm
P07620	PARA_ECOLI		cytoplasm
P07621	PARB_ECOLI		cytoplasm
P07625	CY551_ROSDE		periplasm
P07637	ARO_A_SALTY	cytoplasm (probable).	cytoplasm
P07640	FMQ_MORBO		extracellular
P07641	CAT_PROMI		cytoplasm
P07648	EX5C_ECOLI		cytoplasm
P07649	TRUA_ECOLI		cytoplasm
P07650	TYPH_ECOLI		cytoplasm
P07658	FDHF_ECOLI		periplasm
P07673	INCC2_ECOLI		cytoplasm
P07674	KORB2_ECOLI		cytoplasm
P07676	TRFA_ECOLI		cytoplasm
P07684	EREA_ECOLI		cytoplasm
P07691	PYRF_SALTY		cytoplasm
P07739	LUXB_VIBHA		cytoplasm
P07740	LUXA_VIBHA		cytoplasm
P07748	FIXZ_RHILE		cytoplasm
P07760	RBTR_KLEAE		cytoplasm
P07762	GLGB_ECOLI		extracellular
P07769	BENA_ACIAD		cytoplasm
P07770	BENB_ACIAD		cytoplasm
P07771	BENC_ACIAD		cytoplasm
P07772	BEND_ACIAD		cytoplasm
P07773	CATA_ACIAD		cytoplasm
P07774	CATM_ACIAD		cytoplasm
P07776	Y1443_ACIAD		cytoplasm
P07777	Y1444_ACIAD		cytoplasm
P07779	PQQB_ACICA		cytoplasm
P07780	PQQC_ACICA		cytoplasm
P07781	PQQD_ACICA		cytoplasm
P07782	PQQE_ACICA		cytoplasm
P07801	CHER_SALTY		cytoplasm
P07811	PULA_KLEAE	bacterial cell membrane; lipid-anchor (probable).	extracellular
P07874	ALGA_PSEAE		cytoplasm
P07935	VEGA_MYXXA		cytoplasm
P07989	T1M_SALPO		cytoplasm
P07990	T1S_SALPO		cytoplasm
P08015	FMP3_PSEAE		extracellular

P08058	THIO_RHOSH		cytoplasm
P08061	VIRE3_AGRT5		extracellular
P08063	VIRE1_AGRT5		extracellular
P08080	HEM1_RHIME		inner membrane
P08083	CEAN_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	extracellular
P08084	BNZA_PSEPU		cytoplasm
P08085	BNZB_PSEPU		cytoplasm
P08086	BNZC_PSEPU		cytoplasm
P08087	BNZD_PSEPU		cytoplasm
P08088	BNZE_PSEPU		cytoplasm
P08097	MOBB1_ECOLI		cytoplasm
P08098	MOBA1_ECOLI		cytoplasm
P08127	NAHH_PSEPU		cytoplasm
P08142	ILVB_ECOLI		cytoplasm
P08179	PUR3_ECOLI		cytoplasm
P08189	FIMF_ECOLI		extracellular
P08190	FIMG_ECOLI		extracellular
P08191	FIMH_ECOLI		extracellular
P08192	FOLC_ECOLI		cytoplasm
P08200	IDH_ECOLI		cytoplasm
P08201	NIRB_ECOLI		cytoplasm
P08202	ARAA_ECOLI		cytoplasm
P08203	ARAD_ECOLI		cytoplasm
P08204	ARAB_ECOLI		cytoplasm
P08244	PYRF_ECOLI		cytoplasm
P08245	YCIH_ECOLI		cytoplasm
P08262	HEM1_BRAJA		inner membrane
P08280	RECA_PSEAE	cytoplasm (by similarity).	cytoplasm
P08293	NIFE_AZOVI		cytoplasm
P08301	COXX_PARDE	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P08302	YCO2_PARDE		cytoplasm
P08303	COXZ_PARDE		inner membrane
P08304	YCO4_PARDE		cytoplasm
P08310	CATB_PSEPU		cytoplasm
P08315	FINO2_ECOLI		outer membrane
P08322	TRAE1_ECOLI		cytoplasm
P08332	MERA_SHIFL		cytoplasm
P08337	MUTT_ECOLI		cytoplasm
P08339	YZFA_ECOLI		cytoplasm
P08365	CHPS_ECOLI		cytoplasm
P08368	CREB_ECOLI	cytoplasm (probable).	cytoplasm
P08370	YGDB_ECOLI		cytoplasm
P08371	PPDB_ECOLI		cytoplasm
P08372	PPDC_ECOLI		inner membrane
P08373	MURB_ECOLI	cytoplasm (probable).	inner membrane
P08390	USG_ECOLI		cytoplasm
P08394	EX5B_ECOLI		cytoplasm
P08395	SPPA_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	extracellular
P08409	INSL_ECOLI		cytoplasm
P08421	PYRI_SALTY		cytoplasm

P08455	MTP2_NEIGO		cytoplasm
P08470	PTSBC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P08501	CY1_RHOCA		inner membrane
P08502	CYB_RHOCA		inner membrane
P08504	TNP5_ECOLI		cytoplasm
P08623	RP54_AZOVI		cytoplasm
P08632	NODL_RHILV		cytoplasm
P08633	NODM_RHILV	cytoplasm (by similarity).	cytoplasm
P08634	NODN_RHILV		cytoplasm
P08639	LUXC_VIBHA		cytoplasm
P08654	MERD_SERMA		cytoplasm
P08658	LIP_PSEFR		extracellular
P08660	AK3_ECOLI		cytoplasm
P08664	MERB_SERMA		extracellular
P08690	ARSA1_ECOLI		inner membrane
P08692	ARSC1_ECOLI		cytoplasm
P08694	BPHB_PSEPS		cytoplasm
P08695	BPHC_PSEPS		cytoplasm
P08701	IMMI_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P08702	IMMJ_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P08704	CDGT_KLEOX		extracellular
P08710	FIXX_RHILT		cytoplasm
P08716	HLYBP_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P08717	NIFD_RHOCA		periplasm
P08718	NIFH1_RHOCA		cytoplasm
P08719	NODD2_RHIME		cytoplasm
P08720	NODI_RHILV	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P08737	NIFE_KLEPN		cytoplasm
P08738	NIFN_KLEPN		periplasm
P08811	FER_PSEST		periplasm
P08812	FER3_DESAF		periplasm
P08813	FER1_DESVM		periplasm
P08868	YUAZ_ECOLI		periplasm
P08870	PYRE_SALTY		cytoplasm
P08881	S3AD_KLEPN		cytoplasm
P08888	NODX_RHILV	bacterial cell membrane; multi-pass membrane protein (potential).	cytoplasm
P08956	T1RK_ECOLI		cytoplasm
P08957	T1MK_ECOLI		cytoplasm
P08988	AACC4_SALSP		cytoplasm
P09029	PURK_ECOLI		cytoplasm
P09030	EX3_ECOLI		cytoplasm
P09053	AVTA_ECOLI	cytoplasm (by similarity).	cytoplasm
P09060	ODBA_PSEPU		cytoplasm
P09061	ODBB_PSEPU		cytoplasm

P09062	ODB2_PSEPU		inner membrane
P09099	XYLB_ECOLI		cytoplasm
P09120	HNS_SHIFL		cytoplasm
P09126	HEM4_ECOLI		cytoplasm
P09133	NIFA_AZOCA		cytoplasm
P09134	NIFT_KLEPN		cytoplasm
P09135	NIFY_KLEPN		cytoplasm
P09136	NIFX_KLEPN		cytoplasm
P09137	NIFW_KLEPN		cytoplasm
P09140	LUXA1_PHOLE		cytoplasm
P09141	LUXB1_PHOLE		cytoplasm
P09142	LUXF_PHOLE		cytoplasm
P09147	GALE_ECOLI		cytoplasm
P09148	GAL7_ECOLI		cytoplasm
P09151	LEU1_ECOLI		cytoplasm
P09153	TFAE_ECOLI		cytoplasm
P09154	YMFS_ECOLI		cytoplasm
P09158	SPEE_ECOLI		cytoplasm
P09162	YJAA_ECOLI		inner membrane
P09163	YJAB_ECOLI		cytoplasm
P09165	AERC_AERSO		extracellular
P09181	LYS4_ECOLI		outer membrane
P09182	IMMN_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P09184	VSR_ECOLI		cytoplasm
P09213	SODF_PHOLE		periplasm
P09223	SODF_PSEPU		periplasm
P09347	FDEH_PSEPU		cytoplasm
P09372	GRPE_ECOLI	cytoplasm (probable).	cytoplasm
P09424	MTLD_ECOLI		cytoplasm
P09427	NIFT_AZOVI		cytoplasm
P09431	NTRB_RHOCA		inner membrane
P09432	NTRC_RHOCA		cytoplasm
P09433	RP54_RHOCA		cytoplasm
P09434	NIFA_RHOCA		cytoplasm
P09546	PUTA_ECOLI		cytoplasm
P09549	DEDD_ECOLI		cytoplasm
P09557	DICB_ECOLI		inner membrane
P09570	NIFA_AZOVI		cytoplasm
P09657	RBL1C_RALEU		extracellular
P09658	RBSC_RALEU		cytoplasm
P09745	SHU1_ECOLI		extracellular
P09746	SHU2_ECOLI		extracellular
P09747	SHU3_ECOLI		extracellular
P09748	SHU4_ECOLI		extracellular
P09749	SHU5_ECOLI		extracellular
P09750	SHU6_ECOLI		extracellular
P09751	SHU7_ECOLI		extracellular
P09763	RL11_SERMA		cytoplasm
P09764	RL1_SERMA		cytoplasm
P09772	NIFK_KLEPN		cytoplasm
P09773	OCD_AGRT5		cytoplasm
P09776	VIRB2_AGRTU		extracellular
P09779	VIRB6_AGRTU		cytoplasm
P09781	VIRB8_AGRTU		cytoplasm

P09783	VIRBA_AGRTU		cytoplasm
P09785	PHNA_PSEAE		cytoplasm
P09786	PHNB_PSEAE		cytoplasm
P09787	CY4C_PSEPU		periplasm
P09788	DH4C_PSEPU		cytoplasm
P09795	MTS1_SALIN		cytoplasm
P09796	T2S1_SALIN		extracellular
P09815	ICEN_PSEFL	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P09816	VIRD3_AGRTU		cytoplasm
P09817	VIRD4_AGRTU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P09818	FIXA_RHIME		cytoplasm
P09819	FIXB_RHIME		cytoplasm
P09820	FIXC_RHIME		cytoplasm
P09822	FIXX_RHIME		cytoplasm
P09823	FIXX_RHILE		cytoplasm
P09824	NIFB_RHIME		cytoplasm
P09827	GLNB_RHILV		cytoplasm
P09828	NIFA_RHILE		cytoplasm
P09829	FMM_MORNO		extracellular
P09831	GLTB_ECOLI		cytoplasm
P09832	GLTD_ECOLI		cytoplasm
P09857	THIO_CHRVI		cytoplasm
P09881	IMM8_ECOLI		cytoplasm
P09883	CEA9_ECOLI		extracellular
P09885	KKA6_ACIBA		cytoplasm
P09948	CISY_RICPR		cytoplasm
P09980	REP_ECOLI		cytoplasm
P09984	HLC1_ECOLI	cytoplasm (potential).	cytoplasm
P09985	HLC2_ECOLI	cytoplasm (potential).	cytoplasm
P09996	YIDB_ECOLI		extracellular
POA0Q5	ACPS_NEIMB	cytoplasm (by similarity).	cytoplasm
POA0Q6	ACPS_NEIMC	cytoplasm (by similarity).	cytoplasm
POA0Q7	VDLD_HELPY		cytoplasm
POA0Q8	VDLD_HELPJ		cytoplasm
POA0Q9	MURI_NEIMA		cytoplasm
POA0R0	MURI_NEIMB		cytoplasm
POA0R1	BFRA_NEIMB		cytoplasm
POA0R2	BFRA_NEIGO		cytoplasm
POA0R3	CH10_HELPY	cytoplasm (by similarity).	cytoplasm
POA0R4	CH10_HELPJ	cytoplasm (by similarity).	cytoplasm
POA0R5	CH10_XANAC	cytoplasm (by similarity).	cytoplasm
POA0R6	CH10_XANCP	cytoplasm (by similarity).	cytoplasm
POA0R7	CH10_XANCH	cytoplasm (by similarity).	cytoplasm
POA0R8	EXBD_NEIMA	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
POA0R9	EXBD_NEIMB	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
POA0S0	EXBD_NEIMC	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane

P0A0S1	FLAA_HELPY		flagellar
P0A0S2	FLAA_HELPJ		flagellar
P0A0S3	FLIQ_HELPY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A0S4	FLIQ_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A0S7	FUR_NEIMA	cytoplasm (by similarity).	cytoplasm
P0A0S8	FUR_NEIMB	cytoplasm (by similarity).	cytoplasm
P0A0S9	FUR_NEIMC	cytoplasm (by similarity).	cytoplasm
P0A0T0	G6PI_XANAC	cytoplasm (by similarity).	cytoplasm
P0A0T1	G6PI_XANCI	cytoplasm (by similarity).	cytoplasm
P0A0T2	LGUL_NEIMA		extracellular
P0A0T3	LGUL_NEIMB		extracellular
P0A0T4	GPXA_NEIMA		periplasm
P0A0T5	GPXA_NEIMB		periplasm
P0A0T6	GPXA_NEIMC		periplasm
P0A0T7	HEM1_NEIMA		cytoplasm
P0A0T8	HEM1_NEIMC		cytoplasm
P0A0T9	IHFA_XANCP		cytoplasm
P0A0U0	IHFA_XANAC		cytoplasm
P0A0U1	IHFB_NEIMA		cytoplasm
P0A0U2	IHFB_NEIMB		cytoplasm
P0A0U3	IHFB_NEIGO		cytoplasm
P0A0U4	HYPY_HELPY		cytoplasm
P0A0U5	HYPY_HELPJ		cytoplasm
P0A0U6	KAD_NEIMA	cytoplasm (by similarity).	cytoplasm
P0A0U7	KAD_NEIMB	cytoplasm (by similarity).	cytoplasm
P0A0U8	PBP2_NEIMA	bacterial cell inner membrane (by similarity).	inner membrane
P0A0V0	LPP20_HELPY	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P0A0V1	LPP20_HELPJ	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P0A0V2	OMP4_NEIMA	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A0V4	OHR_XANAC		cytoplasm
P0A0V5	OHR_XANCH		cytoplasm
P0A0V6	HPN_HELPY		cytoplasm
P0A0V7	HPN_HELPJ		cytoplasm
P0A0W0	YAV4_XANCP		extracellular
P0A0W1	YAV4_XANCV		extracellular
P0A0W2	FKBP_NEIMB		cytoplasm
P0A0W3	FKBP_NEIMC		cytoplasm
P0A0W4	RECA_XANAC	cytoplasm (by similarity).	cytoplasm
P0A0W5	RECA_XANCI	cytoplasm (by similarity).	cytoplasm
P0A0W6	RECX_XANAC	cytoplasm (potential).	cytoplasm
P0A0W7	RECX_XANCI	cytoplasm (potential).	cytoplasm
P0A0W8	RL7_NEILA		cytoplasm
P0A0W9	RL7_NEISI		cytoplasm
P0A0X0	RL7_NEIMA		cytoplasm
P0A0X1	RL7_NEIMB		cytoplasm
P0A0X2	RS11_XANCP		cytoplasm
P0A0X3	RS11_XANAC		cytoplasm

P0A0X4	RS12_HELPY		cytoplasm
P0A0X5	RS12_HELPJ		cytoplasm
P0A0X6	RS18_NEIMA		cytoplasm
P0A0X7	RS18_NEIMB		cytoplasm
P0A0X8	RS18_NEIGO		cytoplasm
P0A0X9	RS4_XANCP		cytoplasm
P0A0Y0	RS4_XANAC		cytoplasm
P0A0Y1	RPOA_XANCP		cytoplasm
P0A0Y2	RPOA_XANAC		cytoplasm
P0A0Y4	FBPA_NEIMB	periplasm (by similarity).	periplasm
P0A0Y5	GMHA_NEIMA	cytoplasm (by similarity).	cytoplasm
P0A0Y6	GMHA_NEIMB	cytoplasm (by similarity).	cytoplasm
P0A0Y7	SSRP_NEIMA	cytoplasm (potential).	cytoplasm
P0A0Y8	SSRP_NEIMB	cytoplasm (potential).	cytoplasm
P0A0Y9	Y1593_NEIMA		cytoplasm
P0A0Z0	Y1378_NEIMB		cytoplasm
P0A0Z1	Y030_NEIMA		inner membrane
P0A0Z2	Y228_NEIMB		inner membrane
P0A0Z3	MTGA_NEIMA	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P0A0Z4	MTGA_NEIMB	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P0A0Z5	PBPA_NEIMA	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0A0Z6	PBPA_NEIMB	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0A0Z9	ARGA_PSEPK	cytoplasm (probable).	cytoplasm
P0A100	ARGA_PSEPU	cytoplasm (probable).	cytoplasm
P0A101	PCAJ_PSEPK		cytoplasm
P0A102	PCAJ_PSEPU		cytoplasm
P0A103	ATPZ_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0A104	ATPZ_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0A105	CAPB_PSESM	cytoplasm (by similarity).	cytoplasm
P0A107	NAHC_PSEAE		cytoplasm
P0A108	NAHC_PSEU8		cytoplasm
P0A109	NAHC2_PSEPU		cytoplasm
P0A110	NDOB_PSEPU		cytoplasm
P0A111	NDOB_PSEU8		cytoplasm
P0A112	NDOC_PSEPU		cytoplasm
P0A113	NDOC_PSEU8		cytoplasm
P0A114	CLCD_PSEPU		cytoplasm
P0A115	CLCD_PSESB		cytoplasm
P0A116	DNAA_PSEPK	cytoplasm (by similarity).	cytoplasm
P0A117	DNAA_PSEPU	cytoplasm (by similarity).	cytoplasm
P0A118	PRIM_PSEPK		cytoplasm
P0A119	PRIM_PSEPU		cytoplasm
P0A120	DPO3B_PSEPK	cytoplasm (by similarity).	cytoplasm
P0A121	DPO3B_PSEPU	cytoplasm (by similarity).	cytoplasm

P0A122	FER_PSEPK		periplasm
P0A123	FER_PSEPU		periplasm
P0A124	GIDB_PSEPK		cytoplasm
P0A125	GIDB_PSEPU		cytoplasm
P0A126	IHFA_PSEPK		cytoplasm
P0A127	IHFA_PSEPU		cytoplasm
P0A128	IHFB_PSEPK		cytoplasm
P0A129	IHFB_PSEPU		cytoplasm
P0A130	Y1224_PSEPK		extracellular
P0A131	YOPRL_PSEPU		extracellular
P0A134	TRA6_PSEAE		cytoplasm
P0A135	TRA6_SHIFL		cytoplasm
P0A136	KAD_PSEPK	cytoplasm (by similarity).	cytoplasm
P0A137	KAD_PSEPU	cytoplasm (by similarity).	cytoplasm
P0A138	PAL_PSEPK	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P0A140	OXAA_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A141	OXAA_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A142	NAHE_PSEAE		cytoplasm
P0A143	NAHE_PSEU8		cytoplasm
P0A144	NAHE2_PSEPU		cytoplasm
P0A145	NAHQ_PSEAE		cytoplasm
P0A146	NAHQ_PSEU8		cytoplasm
P0A147	RP5M_PSEPK		cytoplasm
P0A148	RP5M_PSEPU		cytoplasm
P0A149	Y002_PSEPK		cytoplasm
P0A150	YGIDB_PSEPU		cytoplasm
P0A151	PARB_PSEPK		cytoplasm
P0A152	PARB_PSEPU		cytoplasm
P0A153	LEXA1_PSEPK		cytoplasm
P0A154	LEXA_PSEPU		cytoplasm
P0A155	AHPF_PSEPK		cytoplasm
P0A156	AHPF_PSEPU		cytoplasm
P0A157	RL7_PSEPK		cytoplasm
P0A158	RL7_PSEPU		cytoplasm
P0A159	RL20_PSESM		cytoplasm
P0A160	RL20_PSESY		cytoplasm
P0A163	RL35_PSESM		cytoplasm
P0A164	RL35_PSESY		cytoplasm
P0A165	RS21_PSEPK		cytoplasm
P0A166	RS21_PSEPU		cytoplasm
P0A167	RNPA_PSEPK		extracellular
P0A168	RNPA_PSEPU		extracellular
P0A169	NAHB_PSEPU		cytoplasm
P0A170	NAHB_PSEU8		cytoplasm
P0A171	RP54_PSEPK		cytoplasm
P0A172	RP54_PSEPU		cytoplasm
P0A173	TOLB_PSEPK	periplasm (potential).	periplasm
P0A174	TOLB_PSEPU	periplasm (potential).	periplasm
P0A175	TRME_PSEPK		cytoplasm
P0A176	TRME_PSEPU		cytoplasm
P0A177	YCLC_PSEPU	periplasm (potential).	periplasm

P0A179	Y007_PSEPK		inner membrane
P0A180	YU161_PSEPU		inner membrane
P0A181	QADG_PSEPK	periplasm (by similarity).	periplasm
P0A183	MERR_PSEAE		cytoplasm
P0A184	MERR_PSEFL		cytoplasm
P0A185	NDOA_PSEPU		cytoplasm
P0A186	NDOA_PSEU8		cytoplasm
P0A187	AROP_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A188	AROP_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A189	YIFK_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A190	YIFK_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A191	LIVF_SALTY		inner membrane
P0A192	LIVF_SALTI		inner membrane
P0A193	LIVG_SALTY		inner membrane
P0A194	LIVG_SALTI		inner membrane
P0A195	UVRA_SALTY	cytoplasm (by similarity).	cytoplasm
P0A196	UVRA_SALTI	cytoplasm (by similarity).	cytoplasm
P0A197	UBIB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A198	UBIB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A199	ACP2_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A1A0	ACP_SHIDY	cytoplasm (by similarity).	cytoplasm
P0A1A1	YCIA_SALTY		cytoplasm
P0A1A2	YCIA_SALTI		cytoplasm
P0A1A3	ALR1_SALTY		cytoplasm
P0A1A4	ALR1_SALTI		cytoplasm
P0A1A5	ILVE_SALTY		cytoplasm
P0A1A6	ILVE_SALTI		cytoplasm
P0A1A8	CYAA_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1A9	EX3_SALTY		cytoplasm
P0A1B0	EX3_SALTI		cytoplasm
P0A1B1	APAH_SALTY		extracellular
P0A1B2	APAH_SALTI		extracellular
P0A1B3	ARGR_SALTY	cytoplasm (probable).	cytoplasm
P0A1B4	ARGR_SALTI	cytoplasm (probable).	cytoplasm
P0A1B5	AROF_SALTY		cytoplasm
P0A1B6	AROF_SALTI		cytoplasm
P0A1B7	ATPE_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0A1B8	ATPE_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0A1B9	SPAL_SALTY	cytoplasm (potential).	inner membrane
P0A1C0	SPAL_SALTI	cytoplasm (potential).	inner membrane

P0A1C1	SPAL_SHIFL	cytoplasm (potential).	inner membrane
P0A1C2	SPAL_SHISO	cytoplasm (potential).	inner membrane
P0A1C3	ACCA_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1C4	ACCA_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1C5	SUFI_SALTY	periplasm (by similarity).	periplasm
P0A1C6	SUFI_SALTI	periplasm (by similarity).	periplasm
P0A1C7	PDUA_SALTY		cytoplasm
P0A1C8	PDUA_SALTI		cytoplasm
P0A1C9	EUTL_SALTY		cytoplasm
P0A1D0	EUTL_SALTI		cytoplasm
P0A1D1	PDUU_SALTY		cytoplasm
P0A1D2	PDUU_SALTI		cytoplasm
P0A1D3	CH60_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1D4	CH60_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1D5	CH10_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1D6	CH10_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1D7	CLPP_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1D8	CLPP_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1D9	CSPH_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1E0	CSPH_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1E2	HEMN_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1E3	CYSK_SALTY		cytoplasm
P0A1E4	CYSK_SALTI		cytoplasm
P0A1E5	CSGA_SALTY		extracellular
P0A1E6	CSGA_SALTI		extracellular
P0A1E7	CSGA_SALEN		extracellular
P0A1E8	CSGB_SALTY		extracellular
P0A1E9	CSGB_SALEN		extracellular
P0A1F1	DDLA_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1F2	DAPF_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1F3	DAPF_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1F4	YGHB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A1F5	YGHB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A1F6	UDP_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1F7	UDP_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1F8	DHAS_SALTY		cytoplasm
P0A1F9	DHAS_SALTI		cytoplasm
P0A1G0	IMPC1_SALTY		cytoplasm
P0A1G1	IMPC_ECOLI		cytoplasm
P0A1G2	IMPC_SHIFL		cytoplasm
P0A1G3	MSGA_SALTY		extracellular
P0A1G4	MSGA_SALTI		extracellular
P0A1G5	DKSA_SALTY		cytoplasm
P0A1G6	DKSA_SALTI		cytoplasm
P0A1G7	DNAJ_SALTY	cytoplasm (probable).	cytoplasm
P0A1G8	DNAJ_SALTI	cytoplasm (probable).	cytoplasm
P0A1G9	DPO3E_SALTY		cytoplasm
P0A1H0	DPO3E_SALTI		cytoplasm
P0A1H1	DSBI_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane

P0A1H2	DSBI_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0A1H8	EXBB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A1H9	FABZ_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1I0	FABZ_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1I1	PHSB_SALTY		periplasm
P0A1I2	PHSB_SALTI		periplasm
P0A1I3	INVA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0A1I4	INVA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0A1I6	MXIA_SHISO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A1I7	FLGC_SALTY		flagellar
P0A1I8	FLGC_SALTI		flagellar
P0A1I9	FLGD_SALTY		inner membrane
P0A1J0	FLGD_SALTI		inner membrane
P0A1J1	FLGE_SALTY		flagellar
P0A1J2	FLGE_SALTI		flagellar
P0A1J3	FLGG_SALTY		flagellar
P0A1J4	FLGG_SALTI		flagellar
P0A1J5	FLGK_SALTY		flagellar
P0A1J6	FLGK_SALTI		flagellar
P0A1J7	FLGN_SALTY		flagellar
P0A1J8	FLGN_SALTI		flagellar
P0A1K0	FLIG_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0A1K2	FLIJ_SALTI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0A1K3	SPAM_SHIFL		extracellular
P0A1K4	SPAM_SHISO		extracellular
P0A1K5	SPAN_SHIFL	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P0A1K6	SPAN_SHISO	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P0A1K7	SPAO_SALDU		extracellular
P0A1K8	SPAO_SALEN		extracellular
P0A1K9	SPAO_SHIFL		extracellular
P0A1L0	SPAO_SHISO		extracellular
P0A1L1	FLIO_SALTY	bacterial cell membrane; single-pass membrane protein (probable).	inner membrane
P0A1L2	FLIO_SALTI	bacterial cell membrane; single-pass membrane protein (probable).	inner membrane

P0A1L3	SPAP_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1L4	SPAP_SHISO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1L5	FLIQ_SALTY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0A1L6	FLIQ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0A1L7	SPAQ_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1L8	SPAQ_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1L9	SPAQ_SALDU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M0	SPAQ_SALEN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M1	SPAQ_SALGL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M2	SPAQ_SALSE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M3	SPAQ_SALTP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M4	SPAQ_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M5	SPAQ_SHISO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M6	SPAR_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M7	SPAR_SHISO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M8	SPAS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0A1M9	SPAS_SHISO	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0A1N0	SPAK_SALTY		extracellular
P0A1N1	SPAK_SALTI		extracellular
P0A1N2	FLIT_SALTY		inner membrane
P0A1N3	FLIT_SALTI		inner membrane
P0A1N4	FLHE_SALTY		inner membrane

P0A1N5	FLHE_SALTI		inner membrane
P0A1N6	FLAW_SALTY		cytoplasm
P0A1N7	FLAW_SALTI		cytoplasm
P0A1N8	FLGH_SALTY	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
P0A1N9	FLGH_SALTI	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
P0A1P1	G3P1_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1P2	GPDA_SALTY	cytoplasm (probable).	cytoplasm
P0A1P3	GPDA_SALTI	cytoplasm (probable).	cytoplasm
P0A1P4	RFBJ_SALTY		cytoplasm
P0A1P7	GLNA_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1P8	GLRX1_SALTY		cytoplasm
P0A1P9	GLRX1_SALTI		cytoplasm
P0A1Q0	OTSA_SALTY		cytoplasm
P0A1Q1	OTSA_SALTI		cytoplasm
P0A1Q2	LGUL_SALTY		extracellular
P0A1Q3	LGUL_SALTI		extracellular
P0A1Q4	DNAB_SALTY		cytoplasm
P0A1Q5	DNAB_SALTI		cytoplasm
P0A1Q6	HEM1_SALTY		cytoplasm
P0A1Q7	HEM1_SALTI		cytoplasm
P0A1Q8	HEM3_SALTY		cytoplasm
P0A1Q9	HEM3_SALTI		cytoplasm
P0A1R0	HFQ_SALTY		cytoplasm
P0A1R1	HFQ_SALTI		cytoplasm
P0A1R3	HIS6_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1R4	HIS5_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1R5	HIS5_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1R6	DBHA_SALTY		cytoplasm
P0A1R7	DBHA_SALTI		cytoplasm
P0A1R8	DBHB_SALTY		cytoplasm
P0A1R9	DBHB_SALTI		cytoplasm
P0A1S0	IHFA_SALTY		cytoplasm
P0A1S1	IHFA_SALTI		cytoplasm
P0A1S2	HNS_SALTY		cytoplasm
P0A1S3	HNS_SALTI		cytoplasm
P0A1S4	STPA_SALTY		cytoplasm
P0A1S5	STPA_SALTI		cytoplasm
P0A1S6	YAEQ_SALTY		cytoplasm
P0A1S7	YAEQ_SALTI		cytoplasm
P0A1S8	YAJD_SALTY		cytoplasm
P0A1S9	YAJD_SALTI		cytoplasm
P0A1T0	YMDA_SALTY		cytoplasm
P0A1T1	YMDA_SALTI		cytoplasm
P0A1T2	YCED_SALTY		cytoplasm
P0A1T3	YCED_SALTI		cytoplasm
P0A1T4	YDHZ_SALTY		cytoplasm
P0A1T5	YDHZ_SALTI		cytoplasm
P0A1T6	YIFL_SALTY	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0A1T7	YIFL_SALTI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0A1T8	YIGF_SALTY		cytoplasm
P0A1T9	YIGF_SALTI		cytoplasm
P0A1U0	YIGI_SALTY		cytoplasm

P0A1U1	YIGI_SALTI		cytoplasm
P0A1U2	YIGM_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1U3	YIGM_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A1U4	YJAG_SALTY		cytoplasm
P0A1U5	YJAG_SALTI		cytoplasm
P0A1U6	YJGD_SALTY		cytoplasm
P0A1U7	YJGD_SALTI		cytoplasm
P0A1U8	YIIY_SALTY		outer membrane
P0A1U9	YIIY_SALTI		outer membrane
P0A1V0	YTGA_SALTY		extracellular
P0A1V1	YTGA_SALTI		extracellular
P0A1V2	YIS3_SHIFL		cytoplasm
P0A1V3	YIS3_SHISO		cytoplasm
P0A1V5	KAD_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1V6	KPRS_SALTY	cytoplasm (by similarity).	cytoplasm
P0A1V7	KPRS_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1V8	BLO2_SALTY		extracellular
P0A1V9	BLO2_ECOLI		extracellular
P0A1W2	LEP_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A1W3	LEP_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A1W7	LIVK_SALTI	periplasm (by similarity).	periplasm
P0A1W9	BRNQ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A1X0	SLYB_SALTY	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P0A1X1	SLYB_SALTI	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P0A1X2	MXIM_SHIFL	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P0A1X3	MXIM_SHISO	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P0A1X4	LPXD_SALTY		cytoplasm
P0A1X5	LPXD_SALTI		cytoplasm
P0A1X6	AMPM_SALTY		cytoplasm
P0A1X7	AMPM_SALTI		cytoplasm
P0A1X8	MSCL_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A1X9	MSCL_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A1Y0	MUTS_SALTY		cytoplasm
P0A1Y1	MUTS_SALTI		cytoplasm
P0A1Y3	ASRC_SALTI	cytoplasm (by similarity).	cytoplasm
P0A1Y4	NUOG_SALTY		cytoplasm
P0A1Y5	NUOG_SALTI		cytoplasm
P0A1Y6	NUOCD_SALTY		inner membrane

P0A1Y7	NUOCD_SALTI		inner membrane
P0A1Y8	NUOE_SALTY		cytoplasm
P0A1Y9	NUOE_SALTI		cytoplasm
P0A1Z0	LDCI_SALTY	cytoplasm (probable).	cytoplasm
P0A1Z1	LDCI_SALTI	cytoplasm (probable).	cytoplasm
P0A1Z2	SKP_SALTY	periplasm (by similarity).	periplasm
P0A1Z3	SKP_SALTI	periplasm (by similarity).	periplasm
P0A1Z4	PYRB_SALTY		cytoplasm
P0A1Z5	PYRB_SALTI		cytoplasm
P0A1Z6	BFD_SALTY		cytoplasm
P0A1Z7	BFD_SALTI		cytoplasm
P0A1Z8	CSGC_SALTY		periplasm
P0A1Z9	CSGC_SALEN		periplasm
P0A200	CSGE_SALTY		cytoplasm
P0A201	CSGE_SALTI		cytoplasm
P0A202	CSGF_SALTY		extracellular
P0A203	CSGF_SALTI		extracellular
P0A204	CSGG_SALTY	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0A205	CSGG_SALTI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0A206	EUTJ_SALTY		cytoplasm
P0A207	EUTJ_SALTI		cytoplasm
P0A208	EUTP_SALTY		cytoplasm
P0A209	EUTP_SALTI		cytoplasm
P0A210	FLIZ_SALTY		cytoplasm
P0A211	FLIZ_SALTI		cytoplasm
P0A212	GYRI_SALTY	cytoplasm (by similarity).	cytoplasm
P0A213	GYRI_SALTI	cytoplasm (by similarity).	cytoplasm
P0A214	MARB_SALTY		inner membrane
P0A215	MARB_SALTI		inner membrane
P0A216	MERP_SALTI	periplasm (probable).	inner membrane
P0A217	MERP_ENTAG	periplasm (probable).	inner membrane
P0A218	MERP_ENTCL	periplasm (probable).	inner membrane
P0A223	MXIH_SHIFL		inner membrane
P0A224	MXIH_SHISO		inner membrane
P0A225	MXII_SHIFL		inner membrane
P0A226	MXII_SHISO		inner membrane
P0A227	MXIK_SHIFL		extracellular
P0A228	MXIK_SHISO		extracellular
P0A230	NIRD_SALTI	cytoplasm (by similarity).	cytoplasm
P0A233	PGTC_SALTY		inner membrane
P0A234	PGTC_SALTI		inner membrane
P0A235	RFC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A236	RFC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A237	SIRB2_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A238	SIRB2_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A239	SSAP_SALTY		inner membrane

P0A240	SSAP_SALTI		inner membrane
P0A242	SULA_SALTI	bacterial cell inner membrane (by similarity).	inner membrane
P0A243	YSPT_SHIFL		cytoplasm
P0A244	YSPT_SHISO		cytoplasm
P0A245	UBIX_SALTY		cytoplasm
P0A246	UBIX_SALTI		cytoplasm
P0A247	VIRB_SHIFL		cytoplasm
P0A248	VIRB_SHISO		cytoplasm
P0A250	PT1_SALTI	cytoplasm (by similarity).	cytoplasm
P0A251	AHPC_SALTY		cytoplasm
P0A252	AHPC_SALTI		cytoplasm
P0A253	REP12_SALTI		cytoplasm
P0A254	REP12_ECOLI		cytoplasm
P0A255	AAC32_SALSP		cytoplasm
P0A256	AACC3_ENTCL		cytoplasm
P0A258	PLSC_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0A259	PLSX_SALTY		cytoplasm
P0A260	PLSX_SALTI		cytoplasm
P0A262	TSX_SALTI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A265	PPIC_SALTY	cytoplasm (by similarity).	cytoplasm
P0A266	PPIC_SALTI	cytoplasm (by similarity).	cytoplasm
P0A267	GPPA_SALTY		extracellular
P0A268	GPPA_SALTI		extracellular
P0A269	PPX_SALTY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0A270	PPX_SALTI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0A271	HSLV_SALTY	cytoplasm (by similarity).	cytoplasm
P0A272	HSLV_SALTI	cytoplasm (by similarity).	cytoplasm
P0A273	LEXA_SALTY		cytoplasm
P0A274	LEXA_SALTI		cytoplasm
P0A275	MUCA_SALTY		cytoplasm
P0A276	MUCA_ECOLI		cytoplasm
P0A278	XGPT_SALTI	cytoplasm (by similarity).	cytoplasm
P0A279	PSIE_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A280	PSIE_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A281	PTH_SALTY	cytoplasm (by similarity).	cytoplasm
P0A283	PTGA_SALTY	cytoplasm (by similarity).	cytoplasm
P0A284	PTGA_SALTI	cytoplasm (by similarity).	cytoplasm
P0A285	RECO_SALTY		cytoplasm
P0A286	RECO_SALTI		cytoplasm
P0A291	DMA_SALTY		cytoplasm
P0A292	DMA_SALTI		cytoplasm
P0A293	YFCB_SALTY		cytoplasm
P0A294	YFCB_SALTI		cytoplasm

P0A295	RHO_SALTY		inner membrane
P0A296	RHO_SALTI		inner membrane
P0A297	RL10_SALTY		cytoplasm
P0A298	RL10_SALTI		cytoplasm
P0A299	RL7_SALTY		cytoplasm
P0A2A0	RL7_SALTI		cytoplasm
P0A2A1	RL19_SALTY		cytoplasm
P0A2A2	RL19_SALTI		cytoplasm
P0A2A3	RL1_SALTY		cytoplasm
P0A2A4	RL1_SALTI		cytoplasm
P0A2A5	RL28_SALTY		cytoplasm
P0A2A6	RL28_SALTI		cytoplasm
P0A2A7	RL30_SALTY		cytoplasm
P0A2A8	RL30_SALTI		cytoplasm
P0A2A9	RS16_SALTY		extracellular
P0A2B0	RS16_SALTI		extracellular
P0A2B1	RS20_SALTY		cytoplasm
P0A2B2	RS20_SALTI		cytoplasm
P0A2B3	RS7_SALTY		cytoplasm
P0A2B4	RS7_SALTI		cytoplasm
P0A2B5	RIMM_SALTY	cytoplasm (potential).	cytoplasm
P0A2B6	RIMM_SALTI	cytoplasm (potential).	cytoplasm
P0A2B7	RMUC_SALTY		cytoplasm
P0A2B8	RMUC_SALTI		cytoplasm
P0A2B9	RNH_SALTY	cytoplasm (potential).	extracellular
P0A2C0	RNH_SALTI	cytoplasm (potential).	extracellular
P0A2C1	RNH2_SALTY	cytoplasm (potential).	extracellular
P0A2C2	RNH2_SALTI	cytoplasm (potential).	extracellular
P0A2C3	RNPH_SALTY		cytoplasm
P0A2C4	RNPH_SALTI		cytoplasm
P0A2C6	RBSB_SALTI	periplasm (by similarity).	periplasm
P0A2C8	POTD_SALTI	periplasm (by similarity).	periplasm
P0A2C9	FABG_SALTY		cytoplasm
P0A2D0	FABG_SALTI		cytoplasm
P0A2D1	UCPA_SALTY		cytoplasm
P0A2D2	UCPA_SALTI		cytoplasm
P0A2D3	SECE_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2D4	SECE_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2D6	CHEY_SALTI	cytoplasm (by similarity).	cytoplasm
P0A2D7	TYRR_SALTY		cytoplasm
P0A2D8	TYRR_SALTI		cytoplasm
P0A2D9	NTRB_SALTY		inner membrane
P0A2E0	NTRB_SALTI		inner membrane
P0A2E2	GLYA_SALTI	cytoplasm (by similarity).	cytoplasm
P0A2E3	RPOD_SALTY		cytoplasm
P0A2E4	RPOD_SALTI		cytoplasm
P0A2E5	RPOS_SALTY		cytoplasm
P0A2E6	RPOS_SALTI		cytoplasm
P0A2E7	RPOS_SALDU		cytoplasm
P0A2E8	FLIA_SALTY		cytoplasm
P0A2E9	FLIA_SALTI		cytoplasm
P0A2F0	RPOE_SALTY		cytoplasm

P0A2F1	RPOE_SALTI		cytoplasm
P0A2F2	NPD_SALTY	cytoplasm (probable).	cytoplasm
P0A2F3	NPD_SALTI	cytoplasm (probable).	cytoplasm
P0A2F4	SODF_SALTY		periplasm
P0A2F5	SODF_SALTI		periplasm
P0A2F6	SSB_SALTY		cytoplasm
P0A2F7	SSB_SALTI		cytoplasm
P0A2F8	CITN_SALTY	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2F9	CITN_SALTI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2G1	SSRP_SALTY	cytoplasm (potential).	cytoplasm
P0A2G2	SSRP_SALTI	cytoplasm (potential).	cytoplasm
P0A2G4	CIT1_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2G5	NANT_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2G6	NANT_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2G7	YCIH_SALTY		cytoplasm
P0A2G8	YCIH_SALTI		cytoplasm
P0A2G9	CBIF_SALTY		cytoplasm
P0A2H0	CBIF_SALTI		cytoplasm
P0A2H1	CBIE_SALTY		cytoplasm
P0A2H2	CBIE_SALTI		cytoplasm
P0A2H3	TATA_SALTY	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P0A2H4	TATA_SALTI	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P0A2H5	TATE_SALTY	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
P0A2H6	TATE_SALTI	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
P0A2H7	FADA_SALTY	cytoplasm (by similarity).	cytoplasm
P0A2H8	FADA_SALTI	cytoplasm (by similarity).	cytoplasm
P0A2I0	DSBA_SALEN	periplasm (by similarity).	periplasm
P0A2I1	TOP1_SALTY		cytoplasm
P0A2I2	TOP1_SALTI		cytoplasm
P0A2I3	GYRB_SALTY		cytoplasm
P0A2I4	GYRB_SALTI		cytoplasm
P0A2I5	PARE_SALTY		cytoplasm
P0A2I6	PARE_SALTI		cytoplasm
P0A2I8	HISM_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0A2J0	HISQ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2J2	LIVH_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2J3	SAPB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2J4	SAPB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2J5	SAPC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2J6	SAPC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2J7	POTB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2J8	POTB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2K1	TRPB_SALTY		cytoplasm
P0A2K2	TRPB_SALTI		cytoplasm
P0A2K5	UBIE_SALTY		cytoplasm
P0A2K6	UBIE_SALTI		cytoplasm
P0A2K7	GALF_SALTY		cytoplasm
P0A2K8	GALF_SALTI		cytoplasm
P0A2K9	YEAB_SALTY		cytoplasm
P0A2L0	YEAB_SALTI		cytoplasm
P0A2L1	YJBQ_SALTY		inner membrane
P0A2L2	YJBQ_SALTI		inner membrane
P0A2L3	CORC_SALTY		inner membrane
P0A2L4	CORC_SALTI		inner membrane
P0A2L5	MARC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2L6	MARC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2L7	YFJF_SALTY		inner membrane
P0A2L8	YFJF_SALTI		inner membrane
P0A2L9	SIRB1_SALTY		cytoplasm
P0A2M0	SIRB1_SALTI		cytoplasm
P0A2M1	NRDR_SALTY		cytoplasm
P0A2M2	NRDR_SALTI		cytoplasm
P0A2M3	YOA_H_SALTY		inner membrane
P0A2M4	YOA_H_SALTI		inner membrane
P0A2M5	UPP_SALTY		cytoplasm
P0A2M6	UPP_SALTI		cytoplasm
P0A2M7	UXUA_SALTY		cytoplasm
P0A2M8	UXUA_SALTI		cytoplasm
P0A2M9	VRP3_SALTY		extracellular
P0A2N0	VRP3_SALDU		extracellular

P0A2N1	VRP3_SALEN		extracellular
P0A2N2	VSDE_SALTY		extracellular
P0A2N3	VSDE_SALDU		extracellular
P0A2N4	YJEM_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2N5	YJEM_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2N6	ZIPA_SALTY	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
P0A2N7	ZIPA_SALTI	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
P0A2N8	YIHI_SALTY		inner membrane
P0A2N9	YIHI_SALTI		inner membrane
P0A2P0	RHLB_SALTY		cytoplasm
P0A2P1	RHLB_SALTI		cytoplasm
P0A2P4	YHEU_SALTY		inner membrane
P0A2P5	YHEU_SALTI		inner membrane
P0A2P6	XERD_SALTY	cytoplasm (by similarity).	nucleoid
P0A2P7	XERD_SALTI	cytoplasm (by similarity).	nucleoid
P0A2P8	FRUR_SALTY		cytoplasm
P0A2P9	FRUR_SALTI		cytoplasm
P0A2Q0	HDFR_SALTY		cytoplasm
P0A2Q1	HDFR_SALTI		cytoplasm
P0A2Q3	ILVY_SALTI	cytoplasm (by similarity).	cytoplasm
P0A2Q5	METR_SALTI	cytoplasm (by similarity).	cytoplasm
P0A2Q6	MERD_SALTI		cytoplasm
P0A2Q7	MERD_ENTAG		cytoplasm
P0A2Q8	MERR_SALTI		cytoplasm
P0A2Q9	MERR_SHIFL		cytoplasm
P0A2R0	SOXR_SALTY		cytoplasm
P0A2R1	SOXR_SALTI		cytoplasm
P0A2R2	FLHD_SALTY	cytoplasm (by similarity).	cytoplasm
P0A2R3	FLHD_SALTI	cytoplasm (by similarity).	cytoplasm
P0A2R4	RPOC_SALTY		cytoplasm
P0A2R5	RPOC_SALTI		cytoplasm
P0A2R6	DUSB_SALTY		cytoplasm
P0A2R7	DUSB_SALTI		cytoplasm
P0A2R9	CORA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A2S0	LRP_SALTY		cytoplasm
P0A2S1	LRP_KLEAE		cytoplasm
P0A2S2	PDHR_SALTY		cytoplasm
P0A2S3	PDHR_SALTI		cytoplasm
P0A2S4	MARA_SALTY		cytoplasm
P0A2S5	MARA_SALTI		cytoplasm
P0A2S6	MARA_SALEN		cytoplasm
P0A2S7	MXIE_SHIFL		cytoplasm
P0A2S8	MXIE_SHISO		cytoplasm
P0A2S9	RHAS_SALTY	cytoplasm (by similarity).	cytoplasm
P0A2T0	RHAS_SALTI	cytoplasm (by similarity).	cytoplasm
P0A2T1	VIRF_SHIFL		cytoplasm

P0A2T2	VIRF_SHIDY		cytoplasm
P0A2T3	VIRF_SHISO		cytoplasm
P0A2T4	MARR_SALTY		cytoplasm
P0A2T5	MARR_SALTI		cytoplasm
P0A2T6	CRP_SALTY		cytoplasm
P0A2T7	CRP_KLEAE		cytoplasm
P0A2T8	FNR_SALTY	cytoplasm (probable).	cytoplasm
P0A2T9	FNR_SALMS	cytoplasm (probable).	cytoplasm
P0A2U0	OGT_SALTY		cytoplasm
P0A2U1	OGT_SALTI		cytoplasm
P0A2U2	RIMK_SALTY		cytoplasm
P0A2U3	RIMK_SALTI		cytoplasm
P0A2U5	IPGC_SHIDY	cytoplasm (by similarity).	cytoplasm
P0A2V0	CHVA_AGRT5	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A2V1	CHVA_AGRTU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A2V2	OCCP_AGRTU	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0A2V3	OCCP_AGRT4	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0A2W2	ACP_VIBPA	cytoplasm (by similarity).	cytoplasm
P0A2W3	ACP_VIBHA	cytoplasm (by similarity).	cytoplasm
P0A2W4	ACPXL_RHIET	cytoplasm (by similarity).	cytoplasm
P0A2X0	HYIN_AGRTU		extracellular
P0A2X1	HYIN_AGRT4		extracellular
P0A2X2	HYIN_AGRT9		extracellular
P0A2X9	ARGI_BRUME		cytoplasm
P0A2Y0	ARGI_BRUSU		cytoplasm
P0A2Y1	ARGI_BRUAB		cytoplasm
P0A2Y4	AROA_PSES2	cytoplasm (probable).	cytoplasm
P0A2Y5	AROA_ACHSL	cytoplasm (probable).	cytoplasm
P0A308	ATPL_VIBPA		inner membrane
P0A309	ATPL_VIBAL		inner membrane
P0A320	AZUR_BORPE	periplasm (by similarity).	periplasm
P0A322	AZUR_BORPA	periplasm (by similarity).	periplasm
P0A323	CATA_BORPE		periplasm
P0A324	CATA_BORBR		periplasm
P0A325	CATA_BORPA		periplasm
P0A331	FTSA_AGRT5		inner membrane
P0A332	FTSA_AGRTU		inner membrane
P0A339	CH10_BORPE	cytoplasm (by similarity).	cytoplasm
P0A340	CH10_BORBR	cytoplasm (by similarity).	cytoplasm
P0A341	CH10_BORPA	cytoplasm (by similarity).	cytoplasm
P0A342	CH10_BRUME	cytoplasm (by similarity).	cytoplasm
P0A343	CH10_BRUSU	cytoplasm (by similarity).	cytoplasm
P0A344	CH10_BRUAB	cytoplasm (by similarity).	cytoplasm
P0A352	CSPA_BORPE	cytoplasm (by similarity).	cytoplasm
P0A353	CSPA_BORBR	cytoplasm (by similarity).	cytoplasm
P0A354	CSPA_BORPA	cytoplasm (by similarity).	cytoplasm
P0A362	CSPB_YERPE	cytoplasm (by similarity).	cytoplasm
P0A363	CSPB_YEREN	cytoplasm (by similarity).	cytoplasm

P0A388	CYCA_ACEPO	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	periplasm
P0A389	CYCA_ACEEU	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	periplasm
P0A390	NAHF_PSEU8		cytoplasm
P0A391	NAHF_PSEPU		cytoplasm
P0A394	DHLO_AGRT4		cytoplasm
P0A395	DHLO_AGRT9		cytoplasm
P0A396	TFDC_RALEJ		cytoplasm
P0A397	TFDC_BURCE		cytoplasm
P0A3B1	TYP_A_ECOL6		cytoplasm
P0A3B2	TYP_A_ECO27		cytoplasm
P0A3B3	TYP_A_ECO57		cytoplasm
P0A3B4	TYP_A_SHIFL		cytoplasm
P0A3B9	ENTD_ECO57		periplasm
P0A3C0	ENTD_SHIFL		periplasm
P0A3D6	HEMH_BRUME	cytoplasm (by similarity).	extracellular
P0A3D7	HEMH_BRUSU	cytoplasm (by similarity).	extracellular
P0A3D8	HEMH_BRUAB	cytoplasm (by similarity).	extracellular
P0A3E3	FUR_BORPE	cytoplasm (by similarity).	cytoplasm
P0A3E4	FUR_BORBR	cytoplasm (by similarity).	cytoplasm
P0A3E5	FUR_BORPA	cytoplasm (by similarity).	cytoplasm
P0A3F2	GLGA1_AGRT5		cytoplasm
P0A3F3	GLGA1_AGRTU		cytoplasm
P0A3F6	DSDX_ECO57		inner membrane
P0A3F7	DSDX_SHIFL		inner membrane
P0A3F8	VIRBB_AGRTU	cytoplasm (probable).	inner membrane
P0A3F9	VIRBB_AGRT9	cytoplasm (probable).	inner membrane
P0A3G4	DHAA_PSEPV		cytoplasm
P0A3G7	HFQ_BRUME		cytoplasm
P0A3G8	HFQ_BRUSU		cytoplasm
P0A3H3	DBH1_RHILE		cytoplasm
P0A3H4	DBH1_AGRTU		cytoplasm
P0A3I1	IHFB_AGRTU		cytoplasm
P0A3I2	IHFB_AGRT9		cytoplasm
P0A3I3	RT2C_ACTPL	cytoplasm (potential).	cytoplasm
P0A3I4	RT2C_ACTSU	cytoplasm (potential).	cytoplasm
P0A3I5	CYAC_BORPE	cytoplasm (potential).	cytoplasm
P0A3I6	CYAC_BORBR	cytoplasm (potential).	cytoplasm
P0A3I7	CYAC_BORPA	cytoplasm (potential).	cytoplasm
P0A3I8	HRCA_AGRT5		cytoplasm
P0A3I9	HRCA_AGRTU		cytoplasm
P0A3J5	AHYR_AERHY		cytoplasm
P0A3J6	AHYR_AERSA		cytoplasm
P0A3J7	YP7_AGRTU		cytoplasm
P0A3J8	YP7_AGRT4		cytoplasm
P0A3J9	YP7_AGRT9		cytoplasm
P0A3K2	Y915_RICCN		cytoplasm
P0A3K3	YDNAA_RICMO		cytoplasm
P0A3L3	Y4601_AGRT5		cytoplasm
P0A3L4	Y4601_AGRT7		cytoplasm
P0A3L5	IPT_AGRT5		cytoplasm
P0A3L6	IPT_AGRT7		cytoplasm
P0A3L9	KITH_ECO57	cytoplasm (potential).	cytoplasm

P0A3M0	KITH_SHIFL	cytoplasm (potential).	cytoplasm
P0A3M1	BLA5_KLEPN		extracellular
P0A3M2	BLA5_PSEAE		extracellular
P0A3M3	BLAD_KLEPN		extracellular
P0A3M4	BLAD_ENTAE		extracellular
P0A3N4	17KD_RICCN	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P0A3N5	17KD_RICRI	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P0A3P4	LFXD_BRUME		cytoplasm
P0A3P5	LFXD_BRUSU		cytoplasm
P0A3Q2	MGSA_BRUME		cytoplasm
P0A3Q3	MGSA_BRUSU		cytoplasm
P0A3Q4	MGSA_BRUAB		cytoplasm
P0A3Q5	MIAA_AGRT5		cytoplasm
P0A3Q6	MIAA_AGRTU		cytoplasm
P0A3R5	TOX4_BORPE		extracellular
P0A3R6	TOX4_BORPA		extracellular
P0A3R7	CVRA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0A3R8	CVRA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0A3T0	6B1_AGRT4		cytoplasm
P0A3T1	6B1_AGRT9		cytoplasm
P0A3T2	BCSP_BRUME		cytoplasm
P0A3T3	BCSP_BRUAB		cytoplasm
P0A3U2	NIFZ_KLEPN		cytoplasm
P0A3U3	NIFZ_KLEOX		cytoplasm
P0A3U6	PIC1_AGRT5		extracellular
P0A3U7	PIC1_AGRTU		extracellular
P0A3V2	TR2M_AGRT4		cytoplasm
P0A3V3	TR2M_AGRTU		cytoplasm
P0A3V4	TRZD_ENTCL		extracellular
P0A3V5	TRZD_PSESP		extracellular
P0A3V6	VIRB1_AGRTU		extracellular
P0A3V7	VIRB1_AGRT9		extracellular
P0A3V8	VIRB3_AGRTU		cytoplasm
P0A3V9	VIRB3_AGRT9		cytoplasm
P0A3W0	VIRB4_AGRTU		cytoplasm
P0A3W1	VIRB4_AGRT9		cytoplasm
P0A3W2	VIRB5_AGRTU		fimbrium
P0A3W3	VIRB5_AGRT9		fimbrium
P0A3W7	VIRB9_AGRT9	bacterial cell membrane; peripheral membrane protein (by similarity).	extracellular
P0A3W9	VIRE2_AGRT9	plant host: nucleus (by similarity).	cytoplasm
P0A3X0	YMOA_YERPE		cytoplasm
P0A3X1	YMOA_YEREN		cytoplasm
P0A3X2	Y3529_BORPE		cytoplasm
P0A3X3	Y1978_BORBR		cytoplasm
P0A3X4	Y2533_BORPA		cytoplasm
P0A3Y9	NIFM_KLEPN		cytoplasm
P0A3Z0	NIFM_KLEOX		cytoplasm

P0A3Z5	DEGP_BRUSU	periplasm (potential).	periplasm
P0A435	PTKB_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A436	PTKB_ECO57	cytoplasm (by similarity).	cytoplasm
P0A437	PTKB_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A440	PURU_ECOL6		cytoplasm
P0A441	PURU_SHIFL		cytoplasm
P0A448	RECA_BORPE	cytoplasm (by similarity).	cytoplasm
P0A449	RECA_BORBR	cytoplasm (by similarity).	cytoplasm
P0A450	RECA_BORPA	cytoplasm (by similarity).	cytoplasm
P0A468	RL7_BRUME		cytoplasm
P0A469	RL7_BRUSU		cytoplasm
P0A470	RL7_BRUAB		cytoplasm
P0A479	RL20_VIBCH		cytoplasm
P0A480	RL20_VIBMI		cytoplasm
P0A481	RL20_VIBPA		cytoplasm
P0A482	RL20_VIBVU		cytoplasm
P0A4C5	RS4_BORPE		cytoplasm
P0A4C6	RS4_BORBR		cytoplasm
P0A4C7	RS4_BORPA		cytoplasm
P0A4D0	RS6_ECOL6		cytoplasm
P0A4D1	RS6_ECO57		cytoplasm
P0A4D2	RS6_SHIFL		cytoplasm
P0A4D9	RNH2_BRUME	cytoplasm (potential).	extracellular
P0A4E0	RNH2_BRUSU	cytoplasm (potential).	extracellular
P0A4E3	RPOA_AGRT5		cytoplasm
P0A4E4	RPOA_AGRTU		cytoplasm
P0A4E5	RPOA_BORPE		cytoplasm
P0A4E6	RPOA_BORBR		cytoplasm
P0A4E7	RPOA_BORPA		cytoplasm
P0A4F9	OCCT_AGRT4	periplasm (by similarity).	periplasm
P0A4H2	BVGA_BORPE		cytoplasm
P0A4H3	BVGA_BORBR		cytoplasm
P0A4H4	BVGA_BORPA		cytoplasm
P0A4L5	DSBA_ECOL6	periplasm (by similarity).	periplasm
P0A4L6	DSBA_ECO27	periplasm (by similarity).	periplasm
P0A4L7	DSBL_ECOL6	periplasm (by similarity).	periplasm
P0A4L8	DSBL_KLEPN	periplasm (by similarity).	periplasm
P0A4N5	OCCQ_AGRTU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A4N6	OCCQ_AGRT4	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A4P6	USPF_ECOL6		cytoplasm
P0A4P7	USPF_ECO57		cytoplasm
P0A4P8	USPF_SHIFL		cytoplasm
P0A4Q1	YAJC_BRUME		inner membrane
P0A4Q2	YAJC_BRUSU		inner membrane
P0A4R0	Y305_BRUME		cytoplasm
P0A4R1	Y305_BRUSU		cytoplasm
P0A4R2	URE2_BORBR	cytoplasm (by similarity).	cytoplasm
P0A4R3	URE2_BORPA	cytoplasm (by similarity).	cytoplasm
P0A4R4	URED_BORPE		cytoplasm
P0A4R5	URED_BORBR		cytoplasm
P0A4R6	URED_BORPA		cytoplasm
P0A4R7	UREG_BORBR		cytoplasm

P0A4R8	UREG_BORPA		cytoplasm
P0A4R9	UXUA_ECOL6		cytoplasm
P0A4S0	UXUA_SHIFL		cytoplasm
P0A4S7	LFTR_BRUME	cytoplasm (by similarity).	cytoplasm
P0A4S8	LFTR_BRUSU	cytoplasm (by similarity).	cytoplasm
P0A4T3	OCCR_AGRTU		cytoplasm
P0A4T4	OCCR_AGRT4		cytoplasm
P0A4T6	PCAQ_AGRT5		cytoplasm
P0A4T7	PCAQ_AGRTU		cytoplasm
P0A4U0	URE3_BORBR	cytoplasm (by similarity).	cytoplasm
P0A4U1	URE3_BORPA	cytoplasm (by similarity).	cytoplasm
P0A4U4	SLYA_ECOL6		cytoplasm
P0A4U5	SLYA_ECO27		cytoplasm
P0A4U8	DAPD_BORPE	cytoplasm (by similarity).	cytoplasm
P0A4U9	DAPD_BORBR	cytoplasm (by similarity).	cytoplasm
P0A698	UVRA_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A699	UVRA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6A1	UBIB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A6A2	UBIB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A6A4	ACKA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6A5	ACKA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6A6	LEU2_ECOLI		cytoplasm
P0A6A7	LEU2_ECOL6		cytoplasm
P0A6A9	ACP_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6B0	ACP_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6B1	ACP_SALTY	cytoplasm (by similarity).	cytoplasm
P0A6B2	ACP_SALTI	cytoplasm (by similarity).	cytoplasm
P0A6B3	ACP1_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6B4	ALR1_ECOLI		cytoplasm
P0A6B5	ALR1_SHIFL		cytoplasm
P0A6B6	ALR1_SHIBO		cytoplasm
P0A6B7	ISCS_ECOLI	cytoplasm (probable).	cytoplasm
P0A6B8	ISCS_ECOL6		cytoplasm
P0A6B9	ISCS_ECO57		cytoplasm
P0A6C0	ISCS_SHIFL		cytoplasm
P0A6C1	END4_ECOLI		extracellular
P0A6C2	END4_ECO57		extracellular
P0A6C5	ARGA_ECOLI	cytoplasm (probable).	cytoplasm
P0A6C6	ARGA_ECOL6	cytoplasm (probable).	cytoplasm
P0A6C7	ARGA_ECO57	cytoplasm (probable).	cytoplasm
P0A6C8	ARGB_ECOLI	cytoplasm (probable).	cytoplasm
P0A6C9	ARGB_ECO57	cytoplasm (probable).	cytoplasm
P0A6D1	ARGR_ECO57	cytoplasm (probable).	cytoplasm
P0A6D2	ARGR_SHIFL	cytoplasm (probable).	cytoplasm
P0A6D3	AROA_ECOLI	cytoplasm (probable).	cytoplasm
P0A6D4	AROA_ECO57	cytoplasm (probable).	cytoplasm
P0A6D5	YDIB_ECOLI		cytoplasm
P0A6D6	YDIB_ECOL6		cytoplasm
P0A6D7	AROK_ECOLI	cytoplasm (probable).	cytoplasm
P0A6D8	AROK_ECOL6	cytoplasm (probable).	cytoplasm
P0A6D9	AROK_ECO57	cytoplasm (probable).	cytoplasm
P0A6E0	AROK_SHIFL	cytoplasm (probable).	cytoplasm

P0A6E1	AROL_ECOLI	cytoplasm (probable).	cytoplasm
P0A6E2	AROL_ECOL6	cytoplasm (probable).	cytoplasm
P0A6E3	AROL_ECO57	cytoplasm (probable).	cytoplasm
P0A6E4	ASSY_ECOLI	cytoplasm (probable).	cytoplasm
P0A6E5	ASSY_ECOL6	cytoplasm (probable).	cytoplasm
P0A6E7	ATPE_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0A6E8	ATPE_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0A6E9	BIOD2_ECOLI		cytoplasm
P0A6F0	BIOD2_ECO57		cytoplasm
P0A6F1	CARA_ECOLI		cytoplasm
P0A6F2	CARA_ECO57		cytoplasm
P0A6F3	GLPK_ECOLI		cytoplasm
P0A6F4	GLPK_ECO57		cytoplasm
P0A6G3	YGAD_ECOLI		inner membrane
P0A6G4	YGAD_ECOL6		inner membrane
P0A6G5	CITX_ECOLI		cytoplasm
P0A6G6	CITX_ECO57		cytoplasm
P0A6H1	CLPX_ECOLI		cytoplasm
P0A6H2	CLPX_ECOL6		cytoplasm
P0A6H3	CLPX_ECO57		cytoplasm
P0A6H4	CLPX_SHIFL		cytoplasm
P0A6H9	CLS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A6I1	KCY_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6I2	KCY_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6I3	COAA_ECOLI	cytoplasm (probable).	cytoplasm
P0A6I4	COAA_ECOL6	cytoplasm (probable).	cytoplasm
P0A6I5	COAA_ECO57	cytoplasm (probable).	cytoplasm
P0A6I7	COAD_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6I8	COAD_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6I9	COAE_ECOLI	cytoplasm (probable).	cytoplasm
P0A6J0	COAE_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6J1	CYSC_ECOLI		cytoplasm
P0A6J2	CYSC_ECO57		cytoplasm
P0A6J4	CYSZ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A6J6	DADA_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0A6J7	DADA_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0A6J9	DDLA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6K0	DDLA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6K2	DAPF_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6K3	DEF_ECOLI		cytoplasm
P0A6K4	DEF_ECOL6		cytoplasm
P0A6K5	DEF_ECO57		cytoplasm
P0A6K7	DEOB_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6K8	DEOB_ECO57	cytoplasm (by similarity).	cytoplasm

P0A6K9	DEOB_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6L1	DEOC_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6L3	DAPA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6L5	NANA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6L6	NANA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6L7	UXAB_ECOLI		cytoplasm
P0A6L8	UXAB_ECO57		cytoplasm
P0A6L9	HSCB_ECOLI		inner membrane
P0A6M0	HSCB_ECOL6		inner membrane
P0A6M1	HSCB_ECO57		inner membrane
P0A6M3	DSBB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A6M4	DTD_ECOLI	cytoplasm (probable).	cytoplasm
P0A6M5	DTD_ECOL6	cytoplasm (probable).	cytoplasm
P0A6M6	DTD_ECO57	cytoplasm (probable).	cytoplasm
P0A6M7	DTD_SHIFL	cytoplasm (probable).	cytoplasm
P0A6N8	EFPL_ECOLI		cytoplasm
P0A6N9	EFPL_ECOL6		cytoplasm
P0A6P0	EFPL_ECO57		cytoplasm
P0A6P5	ENGA_ECOLI		cytoplasm
P0A6P6	ENGA_ECO57		cytoplasm
P0A6P7	ENGB_ECOLI		cytoplasm
P0A6P8	ENGB_ECOL6		cytoplasm
P0A6Q0	ENO_ECOL6	cytoplasm (by similarity).	extracellular
P0A6Q1	ENO_ECO57	cytoplasm (by similarity).	extracellular
P0A6Q2	ENO_SHIFL	cytoplasm (by similarity).	extracellular
P0A6Q4	FABA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6Q5	FABA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6Q7	FABZ_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6Q8	FABZ_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6Q9	FABZ_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6R1	FABH_ECOL6	cytoplasm (probable).	cytoplasm
P0A6R2	FABH_ECO57	cytoplasm (probable).	cytoplasm
P0A6R3	FIS_ECOLI		cytoplasm
P0A6R4	FIS_ECOL6		cytoplasm
P0A6R5	FIS_ECO57		cytoplasm
P0A6R6	FIS_SALTY		cytoplasm
P0A6R7	FIS_SALTI		cytoplasm
P0A6R8	FIS_SHIFL		cytoplasm
P0A6R9	FIS_SERMA		cytoplasm
P0A6S0	FLGH_ECOLI	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
P0A6S1	FLGH_ECO57	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
P0A6S2	FLGH_SHIFL	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
P0A6S4	FLGI_SHIFL	periplasm (by similarity).	periplasm
P0A6S5	FTSB_ECOLI	inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site.	inner membrane

P0A6S6	FTSB_ECO57	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsI to the division site (by similarity).	inner membrane
P0A6S7	GPDA_ECOLI	cytoplasm (probable).	cytoplasm
P0A6S8	GPDA_ECOL6	cytoplasm (probable).	cytoplasm
P0A6S9	GPDA_ECO57	cytoplasm (probable).	cytoplasm
P0A6T0	GPDA_SHIFL	cytoplasm (probable).	cytoplasm
P0A6T2	G6PI_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6T3	GAL1_ECOLI	cytoplasm (potential).	cytoplasm
P0A6T4	GAL1_ECO57	cytoplasm (potential).	cytoplasm
P0A6T5	GCH1_ECOLI		cytoplasm
P0A6T6	GCH1_ECOL6		cytoplasm
P0A6T7	GCH1_ECO57		cytoplasm
P0A6T8	GCH1_SHIFL		cytoplasm
P0A6T9	GCSH_ECOLI		cytoplasm
P0A6U0	GCSH_ECO57		cytoplasm
P0A6U1	GCSH_SALT1		cytoplasm
P0A6U2	GCSH_SHIFL		cytoplasm
P0A6U3	GIDA_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A6U4	GIDA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6U5	GIDB_ECOLI		cytoplasm
P0A6U6	GIDB_ECOL6		cytoplasm
P0A6U7	GIDB_ECO57		cytoplasm
P0A6U8	GLGA_ECOLI		cytoplasm
P0A6U9	GLGA_ECOL6		cytoplasm
P0A6V0	GLGA_ECO57		cytoplasm
P0A6V1	GLGC_ECOLI		cytoplasm
P0A6V2	GLGC_ECOL6		cytoplasm
P0A6V3	GLGC_ECO57		cytoplasm
P0A6V4	GLGC_SHIFL		cytoplasm
P0A6V6	GLPE_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6V7	GLPE_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A6V9	GLK_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6W0	GLSA2_ECOLI		extracellular
P0A6W1	GLSA2_ECOL6		extracellular
P0A6W2	GLSA2_ECO57		extracellular
P0A6W4	MRAY_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A6W5	GREA_ECOLI		cytoplasm
P0A6W6	GREA_ECOL6		cytoplasm
P0A6W7	GREA_ECO57		cytoplasm
P0A6W8	GREA_SHIFL		cytoplasm
P0A6W9	GSH1_ECOLI		cytoplasm
P0A6X0	GSH1_ECOL6		cytoplasm
P0A6X1	HEM1_ECOLI		cytoplasm
P0A6X2	HEM1_ECO57		cytoplasm
P0A6X3	HFQ_ECOLI		cytoplasm
P0A6X4	HFQ_ECOL6		cytoplasm
P0A6X5	HFQ_ECO57		cytoplasm
P0A6X6	HFQ_SHIFL		cytoplasm
P0A6X7	IHFA_ECOLI		cytoplasm
P0A6X8	IHFA_ECOL6		cytoplasm
P0A6X9	IHFA_ECO57		cytoplasm

P0A6Y0	IHFA_SHIFL		cytoplasm
P0A6Y1	IHFB_ECOLI		cytoplasm
P0A6Y2	IHFB_ECOL6		cytoplasm
P0A6Y3	IHFB_ECO57		cytoplasm
P0A6Y4	IHFB_SHIFL		cytoplasm
P0A6Y6	HSLO_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6Y7	HSLO_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6Y9	DNAK_ECOL6		cytoplasm
P0A6Z0	DNAK_ECO57		cytoplasm
P0A6Z1	HSCA_ECOLI		cytoplasm
P0A6Z2	HSCA_ECO57		cytoplasm
P0A6Z4	HTPG_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A6Z5	HTPG_ECO57	cytoplasm (by similarity).	cytoplasm
P0A6Z6	NIKR_ECOLI		cytoplasm
P0A6Z7	NIKR_ECOL6		cytoplasm
P0A6Z8	NIKR_ECO57		cytoplasm
P0A6Z9	NIKR_SHIFL		cytoplasm
P0A700	HYP_A_ECOLI		cytoplasm
P0A701	HYP_A_ECO57		cytoplasm
P0A702	HYP_A_SHIFL		cytoplasm
P0A703	HYP_B_ECOLI		cytoplasm
P0A704	HYP_B_SHIFL		cytoplasm
P0A711	ISPZ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A713	KDGT_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A714	KDGT_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A716	KDSA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A718	KPRS_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A719	KPRS_ECO57	cytoplasm (by similarity).	cytoplasm
P0A720	KTHY_ECOLI		cytoplasm
P0A721	KTHY_SHIFL		cytoplasm
P0A723	LPXA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A724	LPXA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A725	LPXC_ECOLI		cytoplasm
P0A726	LPXC_ECOL6		cytoplasm
P0A727	LPXC_ECO57		cytoplasm
P0A728	LPXC_SHIFL		cytoplasm
P0A729	YCEF_ECOLI	cytoplasm (potential).	inner membrane
P0A730	YCEF_SHIFL	cytoplasm (potential).	inner membrane
P0A732	MGSA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A733	MGSA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A734	MINE_ECOLI		inner membrane
P0A735	MINE_ECOL6		inner membrane
P0A736	MINE_ECO57		inner membrane
P0A737	MINE_SHIFL		inner membrane
P0A738	MOAC_ECOLI		fimbrium
P0A739	MOAC_ECOL6		cytoplasm
P0A740	MOAC_ECO57		cytoplasm
P0A741	MOAC_SHIFL		cytoplasm

P0A743	MSCL_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A744	MSRA_ECOLI		cytoplasm
P0A745	MSRA_SHIFL		cytoplasm
P0A746	MSRB_ECOLI		cytoplasm
P0A747	MSRB_ECOL6		cytoplasm
P0A748	MSRB_ECO57		cytoplasm
P0A749	MURA_ECOLI	cytoplasm (probable).	inner membrane
P0A750	MURA_ECOL6	cytoplasm (probable).	inner membrane
P0A751	MURA_SHIFL	cytoplasm (probable).	inner membrane
P0A752	NADD_ECOLI		cytoplasm
P0A753	NADD_SHIFL		cytoplasm
P0A754	KEFF_ECOLI		cytoplasm
P0A755	KEFF_ECOL6		cytoplasm
P0A756	KEFG_ECOLI		cytoplasm
P0A757	KEFG_ECOL6		cytoplasm
P0A758	KEFG_ECO57		cytoplasm
P0A759	NAGB_ECOLI		extracellular
P0A760	NAGB_ECO57		extracellular
P0A761	NANE_ECOLI		cytoplasm
P0A762	NANE_ECOL6		cytoplasm
P0A764	NDK_ECO57	cytoplasm (by similarity).	cytoplasm
P0A765	NDK_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A767	RNFA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A768	RNFA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A770	MNTH_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0A771	MNTH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0A772	NRDI_ECOLI		inner membrane
P0A773	NRDI_ECOL6		inner membrane
P0A774	NRDI_ECO57		inner membrane
P0A775	NRDI_SHIFL		inner membrane
P0A776	NUDH_ECOLI		cytoplasm
P0A777	NUDH_ECOL6		cytoplasm
P0A778	NUDH_ECO57		cytoplasm
P0A779	NUDH_SHIFL		cytoplasm
P0A780	NUSB_ECOLI		cytoplasm
P0A781	NUSB_ECOL6		cytoplasm
P0A782	NUSB_ECO57		cytoplasm
P0A783	NUSB_SHIFL		cytoplasm
P0A784	ORN_ECOLI	cytoplasm (potential).	cytoplasm
P0A785	ORN_ECOL6	cytoplasm (potential).	cytoplasm
P0A786	PYRB_ECOLI		cytoplasm
P0A787	PYRB_ECOL6		cytoplasm
P0A788	PYRB_ECO57		cytoplasm
P0A789	PYRB_SHIFL		cytoplasm
P0A790	PAND_ECOLI		cytoplasm
P0A791	PAND_ECOL6		cytoplasm

P0A792	PAND_ECO57		cytoplasm
P0A793	PAND_SHIFL		cytoplasm
P0A795	PDXJ_ECO57	cytoplasm (probable).	cytoplasm
P0A797	K6PF1_ECO57	cytoplasm (by similarity).	cytoplasm
P0A798	K6PF1_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A7A0	PGK_ECOL6	cytoplasm (potential).	cytoplasm
P0A7A1	PGK_SHIFL	cytoplasm (potential).	cytoplasm
P0A7A2	GPMB_ECOLI		cytoplasm
P0A7A3	GPMB_ECO57		cytoplasm
P0A7A4	GPMB_SHIFL		cytoplasm
P0A7A6	PIMT_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7A8	PLSB_ECOL6	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0A7B0	IPYR_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7B2	PPK_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0A7B3	PPNK_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A7B4	PPNK_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A7B6	PROB_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A7B7	PROB_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7B9	HSLV_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A7C0	HSLV_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7C1	HSLV_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A7C2	LEXA_ECOLI		cytoplasm
P0A7C3	LEXA_ECOL6		cytoplasm
P0A7C4	LEXA_ECO57		cytoplasm
P0A7C5	LEXA_SHIFL		cytoplasm
P0A7C6	PEPE_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A7C7	PEPE_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7C9	PSIE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A7D0	PSIE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A7D2	PTH_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A7D3	PTH_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7D5	PURA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A7D6	PURA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7D7	PUR7_ECOLI		cytoplasm
P0A7D8	PUR7_ECOL6		cytoplasm
P0A7D9	PUR7_ECO57		cytoplasm
P0A7E0	PUR7_SHIFL		cytoplasm
P0A7E2	PYRD_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0A7E3	PYRE_ECOLI		cytoplasm
P0A7E4	PYRE_ECOL6		cytoplasm
P0A7E5	PYRG_ECOLI		cytoplasm
P0A7E6	PYRG_ECOL6		cytoplasm
P0A7E7	PYRG_ECO57		cytoplasm
P0A7E8	PYRG_SHIFL		cytoplasm
P0A7E9	PYRH_ECOLI		cytoplasm
P0A7F0	PYRH_ECOL6		cytoplasm

P0A7F1	PYRH_ECO57		cytoplasm
P0A7F2	PYRH_SHIFL		cytoplasm
P0A7F3	PYRI_ECOLI		cytoplasm
P0A7F4	PYRI_ECOL6		cytoplasm
P0A7F5	PYRI_ECO57		cytoplasm
P0A7F6	SPED_ECOLI		cytoplasm
P0A7F7	SPED_ECO57		cytoplasm
P0A7F8	SPED_SHIFL		cytoplasm
P0A7F9	QUEA_ECOLI	cytoplasm (potential).	cytoplasm
P0A7G0	QUEA_ECO57	cytoplasm (potential).	cytoplasm
P0A7G1	QUEA_SHIFL	cytoplasm (potential).	cytoplasm
P0A7G2	RBFA_ECOLI	cytoplasm (potential).	cytoplasm
P0A7G3	RBFA_ECOL6	cytoplasm (potential).	cytoplasm
P0A7G4	RBFA_ECO57	cytoplasm (potential).	cytoplasm
P0A7G5	RBFA_SHIFL	cytoplasm (potential).	cytoplasm
P0A7G7	RECA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A7G8	RECA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7G9	RECA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A7H0	RECF_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A7H1	RECF_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A7H2	RECF_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7H3	RECO_ECOLI		cytoplasm
P0A7H4	RECO_ECOL6		cytoplasm
P0A7H5	RECO_ECO57		cytoplasm
P0A7H6	RECR_ECOLI		cytoplasm
P0A7H7	RECR_ECOL6		cytoplasm
P0A7H8	RECR_ECO57		cytoplasm
P0A7H9	RECR_SHIFL		cytoplasm
P0A7I7	GCH2_ECOLI		cytoplasm
P0A7I8	GCH2_ECO57		cytoplasm
P0A7I9	GCH2_SHIFL		cytoplasm
P0A7J0	RIBB_ECOLI	bacterial cell membrane; peripheral membrane protein (potential).	cytoplasm
P0A7J1	RIBB_ECO57		cytoplasm
P0A7J2	RIBB_SHIFL		cytoplasm
P0A7J3	RL10_ECOLI		cytoplasm
P0A7J4	RL10_ECOL6		cytoplasm
P0A7J5	RL10_ECO57		cytoplasm
P0A7J6	RL10_SHIFL		cytoplasm
P0A7J7	RL11_ECOLI		cytoplasm
P0A7J8	RL11_ECOL6		cytoplasm
P0A7J9	RL11_ECO57		cytoplasm
P0A7K0	RL11_SALTY		cytoplasm
P0A7K1	RL11_SALTI		cytoplasm
P0A7K2	RL7_ECOLI		cytoplasm
P0A7K3	RL7_ECOL6		cytoplasm
P0A7K4	RL7_ECO57		cytoplasm
P0A7K5	RL7_SHIFL		cytoplasm
P0A7K6	RL19_ECOLI		cytoplasm
P0A7K7	RL19_ECOL6		cytoplasm
P0A7K8	RL19_ECO57		cytoplasm
P0A7K9	RL19_SHIFL		cytoplasm
P0A7L0	RL1_ECOLI		cytoplasm
P0A7L1	RL1_ECOL6		cytoplasm
P0A7L2	RL1_ECO57		cytoplasm

P0A7L3	RL20_ECOLI		cytoplasm
P0A7L4	RL20_ECOL6		cytoplasm
P0A7L5	RL20_ECO57		cytoplasm
P0A7L6	RL20_SALTY		cytoplasm
P0A7L7	RL20_SALTI		cytoplasm
P0A7L8	RL27_ECOLI		cytoplasm
P0A7L9	RL27_ECOL6		cytoplasm
P0A7M0	RL27_ECO57		cytoplasm
P0A7M1	RL27_SHIFL		cytoplasm
P0A7M2	RL28_ECOLI		cytoplasm
P0A7M3	RL28_ECOL6		cytoplasm
P0A7M4	RL28_ECO57		cytoplasm
P0A7M5	RL28_SHIFL		cytoplasm
P0A7M6	RL29_ECOLI		cytoplasm
P0A7M7	RL29_ECOL6		cytoplasm
P0A7M8	RL29_ECO57		cytoplasm
P0A7M9	RL31_ECOLI		cytoplasm
P0A7N0	RL31_ECO57		cytoplasm
P0A7N1	RL31B_ECOLI		cytoplasm
P0A7N2	RL31B_ECOL6		cytoplasm
P0A7N3	R31B1_ECO57		cytoplasm
P0A7N4	RL32_ECOLI		cytoplasm
P0A7N5	RL32_ECO57		cytoplasm
P0A7N6	RL32_SALTY		cytoplasm
P0A7N7	RL32_SALTI		cytoplasm
P0A7N8	RL32_SHIFL		cytoplasm
P0A7N9	RL33_ECOLI		cytoplasm
P0A7P0	RL33_ECOL6		cytoplasm
P0A7P1	RL33_ECO57		cytoplasm
P0A7P2	RL33_SALTY		cytoplasm
P0A7P3	RL33_SALTI		cytoplasm
P0A7P4	RL33_SHIFL		cytoplasm
P0A7Q1	RL35_ECOLI		cytoplasm
P0A7Q2	RL35_ECO57		cytoplasm
P0A7Q3	RL35_SALTY		cytoplasm
P0A7Q4	RL35_SALTI		cytoplasm
P0A7Q5	RL35_SHIFL		cytoplasm
P0A7R1	RL9_ECOLI		cytoplasm
P0A7R2	RL9_ECOL6		cytoplasm
P0A7R3	RL9_ECO57		cytoplasm
P0A7R4	RL9_SHIFL		cytoplasm
P0A7R5	RS10_ECOLI		cytoplasm
P0A7R6	RS10_ECOL6		cytoplasm
P0A7R7	RS10_ECO57		cytoplasm
P0A7R8	RS10_SHIFL		cytoplasm
P0A7R9	RS11_ECOLI		cytoplasm
P0A7S0	RS11_ECOL6		cytoplasm
P0A7S1	RS11_ECO57		cytoplasm
P0A7S2	RS11_SHIFL		cytoplasm
P0A7S3	RS12_ECOLI		cytoplasm
P0A7S4	RS12_ECOL6		cytoplasm
P0A7S5	RS12_ECO57		cytoplasm
P0A7S6	RS12_SALTY		cytoplasm
P0A7S7	RS12_SALTI		cytoplasm
P0A7S8	RS12_SHIFL		cytoplasm
P0A7S9	RS13_ECOLI		cytoplasm

P0A7T0	RS13_ECOL6		cytoplasm
P0A7T1	RS13_ECO57		cytoplasm
P0A7T2	RS13_SHIFL		cytoplasm
P0A7T3	RS16_ECOLI		extracellular
P0A7T4	RS16_ECOL6		extracellular
P0A7T5	RS16_ECO57		extracellular
P0A7T6	RS16_SHIFL		extracellular
P0A7T7	RS18_ECOLI		cytoplasm
P0A7T8	RS18_ECOL6		cytoplasm
P0A7T9	RS18_ECO57		cytoplasm
P0A7U0	RS18_PHOLL		cytoplasm
P0A7U1	RS18_SALTI		cytoplasm
P0A7U2	RS18_SHIFL		cytoplasm
P0A7U3	RS19_ECOLI		cytoplasm
P0A7U4	RS19_ECOL6		cytoplasm
P0A7U5	RS19_ECO57		cytoplasm
P0A7U6	RS19_SHIFL		cytoplasm
P0A7U7	RS20_ECOLI		cytoplasm
P0A7U8	RS20_ECOL6		cytoplasm
P0A7U9	RS20_ECO57		cytoplasm
P0A7V0	RS2_ECOLI		cytoplasm
P0A7V1	RS2_ECOL6		cytoplasm
P0A7V2	RS2_ECO57		cytoplasm
P0A7V3	RS3_ECOLI		cytoplasm
P0A7V4	RS3_ECOL6		cytoplasm
P0A7V5	RS3_ECO57		cytoplasm
P0A7V6	RS3_SALTY		cytoplasm
P0A7V7	RS3_SALTI		cytoplasm
P0A7V8	RS4_ECOLI		cytoplasm
P0A7V9	RS4_ECOL6		cytoplasm
P0A7W0	RS4_ECO57		cytoplasm
P0A7W1	RS5_ECOLI		cytoplasm
P0A7W2	RS5_ECOL6		cytoplasm
P0A7W3	RS5_ECO57		cytoplasm
P0A7W4	RS5_SALTY		cytoplasm
P0A7W5	RS5_SALTI		cytoplasm
P0A7W6	RS5_SHIFL		cytoplasm
P0A7W7	RS8_ECOLI		cytoplasm
P0A7W8	RS8_ECOL6		cytoplasm
P0A7W9	RS8_ECO57		cytoplasm
P0A7X0	RS8_SALTY		cytoplasm
P0A7X1	RS8_SALTI		cytoplasm
P0A7X2	RS8_SHIFL		cytoplasm
P0A7X3	RS9_ECOLI		cytoplasm
P0A7X4	RS9_ECOL6		cytoplasm
P0A7X5	RS9_ECO57		cytoplasm
P0A7X6	RIMM_ECOLI	cytoplasm (potential).	cytoplasm
P0A7X7	RIMM_ECOL6	cytoplasm (potential).	cytoplasm
P0A7X8	RIMM_ECO57	cytoplasm (potential).	cytoplasm
P0A7X9	RIMM_SHIFL	cytoplasm (potential).	cytoplasm
P0A7Y1	RNC_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A7Y2	RNC_ECO57	cytoplasm (by similarity).	cytoplasm
P0A7Y3	RNC_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A7Y4	RNH_ECOLI	cytoplasm (potential).	extracellular
P0A7Y5	RNH_ECOL6	cytoplasm (potential).	extracellular
P0A7Y6	RNH_ECO57	cytoplasm (potential).	extracellular

P0A7Y7	RNH_SHIFL	cytoplasm (potential).	extracellular
P0A7Y8	RNPA_ECOLI		extracellular
P0A7Y9	RNPA_SHIFL		extracellular
P0A7Z0	RPIA_ECOLI		cytoplasm
P0A7Z1	RPIA_ECOL6		cytoplasm
P0A7Z2	RPIA_ECO57		cytoplasm
P0A7Z3	RPIA_SHIFL		cytoplasm
P0A7Z4	RPOA_ECOLI		cytoplasm
P0A7Z5	RPOA_ECOL6		cytoplasm
P0A7Z6	RPOA_ECO57		cytoplasm
P0A7Z7	RPOA_SALTY		cytoplasm
P0A7Z8	RPOA_SALTI		cytoplasm
P0A7Z9	RPOA_SHIFL		cytoplasm
P0A800	RPOZ_ECOLI		cytoplasm
P0A801	RPOZ_ECOL6		cytoplasm
P0A802	RPOZ_ECO57		cytoplasm
P0A803	RPOZ_SALTY		cytoplasm
P0A804	RPOZ_SALTI		cytoplasm
P0A806	RRF_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A807	RRF_ECO57	cytoplasm (by similarity).	cytoplasm
P0A808	RRF_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A809	RUVA_ECOLI		cytoplasm
P0A810	RUVA_ECOL6		cytoplasm
P0A811	RUVA_ECO57		cytoplasm
P0A812	RUVB_ECOLI		cytoplasm
P0A813	RUVB_ECO57		cytoplasm
P0A814	RUVC_ECOLI		cytoplasm
P0A815	RUVC_ECOL6		cytoplasm
P0A816	RUVC_ECO57		cytoplasm
P0A822	SELA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A823	SFSA_ECOLI		cytoplasm
P0A824	SFSA_ECO57		cytoplasm
P0A826	GLYA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A827	GLYA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A828	SMG_ECOLI		cytoplasm
P0A829	SMG_SHIFL		cytoplasm
P0A831	DCTA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A832	SSRP_ECOLI	cytoplasm (potential).	cytoplasm
P0A833	SSRP_ECOL6	cytoplasm (potential).	cytoplasm
P0A834	SSRP_ECO57	cytoplasm (potential).	cytoplasm
P0A835	SSRP_SHIFL	cytoplasm (potential).	cytoplasm
P0A836	SUCC_ECOLI		cytoplasm
P0A837	SUCC_ECOL6		cytoplasm
P0A838	SUCC_ECO57		cytoplasm
P0A839	SUCC_SHIFL		cytoplasm
P0A840	SURE_ECOLI	cytoplasm (potential).	periplasm
P0A841	SURE_ECOL6	cytoplasm (potential).	periplasm
P0A842	SURE_ECO57	cytoplasm (potential).	periplasm
P0A843	TATE_ECOLI	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
P0A844	TATE_ECOL6	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane

P0A845	TATE_ECO57	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
P0A846	TATE_SHIFL	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
P0A847	TGT_ECOLI		cytoplasm
P0A848	TGT_ECOL6		cytoplasm
P0A849	TGT_ECO57		cytoplasm
P0A850	TIG_ECOLI		cytoplasm
P0A851	TIG_ECO57		cytoplasm
P0A852	TIG_SHIFL		cytoplasm
P0A853	TNAA_ECOLI		cytoplasm
P0A854	TNAA_SHIFL		cytoplasm
P0A856	TOLB_ECOL6	periplasm (by similarity).	inner membrane
P0A857	TOLB_ECO57	periplasm (by similarity).	inner membrane
P0A858	TPIS_ECOLI		cytoplasm
P0A859	TPIS_ECOL6		cytoplasm
P0A860	TPIS_ECO57		cytoplasm
P0A861	TPIS_SHIFL		cytoplasm
P0A863	TPX_ECOL6	periplasm (by similarity).	periplasm
P0A864	TPX_ECO57	periplasm (by similarity).	periplasm
P0A865	TPX_SHIFL	periplasm (by similarity).	periplasm
P0A866	TPX_SHIDY	periplasm (by similarity).	periplasm
P0A867	TALA_ECOLI	cytoplasm (probable).	cytoplasm
P0A868	TALA_ECOL6	cytoplasm (probable).	cytoplasm
P0A869	TALA_ECO57	cytoplasm (probable).	cytoplasm
P0A870	TALB_ECOLI	cytoplasm (probable).	cytoplasm
P0A871	TALB_ECO57	cytoplasm (probable).	cytoplasm
P0A872	TALB_SHIFL	cytoplasm (probable).	cytoplasm
P0A873	TRMD_ECOLI	cytoplasm (potential).	cytoplasm
P0A874	TRMD_ECOL6	cytoplasm (potential).	cytoplasm
P0A875	TRMD_ECO57	cytoplasm (potential).	cytoplasm
P0A876	TRMD_SHIFL	cytoplasm (potential).	cytoplasm
P0A877	TRPA_ECOLI		cytoplasm
P0A878	TRPA_SHIFL		cytoplasm
P0A879	TRPB_ECOLI		cytoplasm
P0A880	TRPB_SHIFL		cytoplasm
P0A882	TRPR_ECO57	cytoplasm (by similarity).	cytoplasm
P0A883	TRPR_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A885	TYSY_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A886	TYSY_ECO57	cytoplasm (by similarity).	cytoplasm
P0A887	UBIE_ECOLI		cytoplasm
P0A888	UBIE_ECO57		cytoplasm
P0A889	UBIE_SHIFL		cytoplasm
P0A891	TUSA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A892	TUSA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A893	TUSA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A894	YHBJ_ECOLI		cytoplasm
P0A895	YHBJ_ECOL6		cytoplasm
P0A896	YHBJ_ECO57		cytoplasm
P0A897	YHBJ_SHIFL		cytoplasm
P0A898	YBEY_ECOLI		fimbrium
P0A899	YBEY_SHIFL		inner membrane
P0A8A0	YEBC_ECOLI		inner membrane
P0A8A1	YEBC_SHIFL		inner membrane

P0A8A2	YEEN_ECOLI		fimbrium
P0A8A3	YEEN_ECOL6		inner membrane
P0A8A4	YDIA_ECOLI		cytoplasm
P0A8A5	YDIA_ECOL6		cytoplasm
P0A8A6	YDIA_ECO57		cytoplasm
P0A8A7	YDIA_SHIFL		cytoplasm
P0A8A8	YHBC_ECOLI		fimbrium
P0A8A9	YHBC_ECOL6		inner membrane
P0A8B0	YHBC_ECO57		inner membrane
P0A8B1	YHBC_SHIFL		inner membrane
P0A8B2	YFCN_ECOLI		inner membrane
P0A8B3	YFCN_ECOL6		inner membrane
P0A8B4	YFCN_SHIFL		inner membrane
P0A8B5	YBAB_ECOLI		inner membrane
P0A8B6	YBAB_ECOL6		inner membrane
P0A8B7	YBAB_ECO57		inner membrane
P0A8B8	YBAB_SALTY		inner membrane
P0A8B9	YBAB_SALTI		inner membrane
P0A8C0	YBAB_SHIFL		inner membrane
P0A8C1	YBJQ_ECOLI		fimbrium
P0A8C2	YBJQ_ECOL6		inner membrane
P0A8C3	YBJQ_ECO57		inner membrane
P0A8C4	YGFB_ECOLI		inner membrane
P0A8C5	YGFB_ECOL6		inner membrane
P0A8C6	YGFB_ECO57		inner membrane
P0A8C7	YGFB_SHIFL		inner membrane
P0A8C8	YIDD_ECOLI		inner membrane
P0A8C9	YIDD_SALTY		inner membrane
P0A8D0	NRDR_ECOLI		cytoplasm
P0A8D1	NRDR_ECOL6		cytoplasm
P0A8D2	NRDR_ECO57		cytoplasm
P0A8D3	YAII_ECOLI		inner membrane
P0A8D4	YAII_ECOL6		inner membrane
P0A8D5	YAII_ECO57		inner membrane
P0A8D6	YMDB_ECOLI		inner membrane
P0A8D7	YMDB_ECOL6		inner membrane
P0A8D8	YMDB_ECO57		inner membrane
P0A8E0	YFBV_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8E1	YCFP_ECOLI		inner membrane
P0A8E2	YCFP_ECOL6		inner membrane
P0A8E3	YCFP_ECO57		inner membrane
P0A8E4	YCFP_SHIFL		inner membrane
P0A8E5	YAACL_ECOLI		inner membrane
P0A8E6	YAACL_ECO57		inner membrane
P0A8E7	YAJQ_ECOLI		fimbrium
P0A8E8	YAJQ_ECOL6		inner membrane
P0A8E9	YAJQ_ECO57		inner membrane
P0A8F0	UPP_ECOLI		cytoplasm
P0A8F1	UPP_ECOL6		cytoplasm
P0A8F2	UPP_ECO57		cytoplasm
P0A8F3	UPP_SHIFL		cytoplasm
P0A8F5	URK_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A8F6	URK_ECO57	cytoplasm (by similarity).	cytoplasm
P0A8F7	URK_SHIFL	cytoplasm (by similarity).	cytoplasm

P0A8F9	UVRB_ECO57	cytoplasm (by similarity).	cytoplasm
P0A8G1	UVRC_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A8G2	UVRC_ECO57	cytoplasm (by similarity).	cytoplasm
P0A8G3	UXAC_ECOLI		cytoplasm
P0A8G4	UXAC_ECOL6		cytoplasm
P0A8G5	UXAC_SHIFL		cytoplasm
P0A8G6	WRBA_ECOLI		cytoplasm
P0A8G7	WRBA_ECOL6		cytoplasm
P0A8G8	WRBA_SHIFL		cytoplasm
P0A8H0	EX7S_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A8H1	EX7S_ECO57	cytoplasm (by similarity).	cytoplasm
P0A8H2	EX7S_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A8H3	ZUPT_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A8H4	ZUPT_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A8H5	ZUPT_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0A8H6	YIHI_ECOLI		fimbrium
P0A8H7	YIHI_SHIFL		inner membrane
P0A8H8	YACG_ECOLI		cytoplasm
P0A8H9	YACG_ECO57		cytoplasm
P0A8I0	YACG_SHIFL		cytoplasm
P0A8I1	RUVX_ECOLI	cytoplasm (potential).	cytoplasm
P0A8I2	RUVX_ECOL6	cytoplasm (potential).	cytoplasm
P0A8I3	YAAA_ECOLI		inner membrane
P0A8I4	YAAA_SHIFL		inner membrane
P0A8I5	TRMB_ECOLI		cytoplasm
P0A8I6	TRMB_ECO57		cytoplasm
P0A8I7	TRMB_SHIFL		cytoplasm
P0A8I8	YBEA_ECOLI		fimbrium
P0A8I9	YBEA_ECOL6		inner membrane
P0A8J0	YBEA_ECO57		inner membrane
P0A8J1	YBEA_SHIFL		inner membrane
P0A8J2	DNAT_ECOLI		fimbrium
P0A8J3	DNAT_SHIFL		nucleoid
P0A8J4	YBED_ECOLI		fimbrium
P0A8J5	YBED_ECOL6		inner membrane
P0A8J6	YBED_ECO57		inner membrane
P0A8J7	YBED_SHIFL		inner membrane
P0A8J8	RHLB_ECOLI		cytoplasm
P0A8J9	RHLB_ECOL6		cytoplasm
P0A8K0	RHLB_SHIFL		cytoplasm
P0A8K2	PSD_ECOL6		cytoplasm
P0A8K3	PSD_ECO57		cytoplasm
P0A8K4	PSD_SHIFL		cytoplasm
P0A8K5	YAEP_ECOLI		inner membrane
P0A8K6	YAEP_ECO57		inner membrane
P0A8K7	YAEP_SHIFL		inner membrane
P0A8K9	RBN_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0A8L0	RBN_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8L5	YCGN_ECOLI		inner membrane
P0A8L6	YCGN_SHIFL		inner membrane
P0A8L7	YCIU_ECOLI		fimbrium
P0A8L8	YCIU_ECOL6		inner membrane
P0A8L9	YCIU_ECO57		inner membrane
P0A8M6	YEEX_ECOLI		inner membrane
P0A8M7	YEEX_ECOL6		inner membrane
P0A8M8	YEEX_ECO57		inner membrane
P0A8M9	YEEX_SHIFL		inner membrane
P0A8N0	YCBG_ECOLI		inner membrane
P0A8N1	YCBG_ECO57		inner membrane
P0A8N2	YCBG_SHIFL		inner membrane
P0A8N7	SYK3_ECOLI		cytoplasm
P0A8N8	SYK3_ECOL6		cytoplasm
P0A8N9	SYK3_ECO57		cytoplasm
P0A8P0	SYK3_SHIFL		cytoplasm
P0A8P2	LFTR_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A8P3	FETP_ECOLI		cytoplasm
P0A8P4	FETP_ECO57		cytoplasm
P0A8P5	FETP_SHIFL		cytoplasm
P0A8P7	XERC_ECOL6	cytoplasm (by similarity).	nucleoid
P0A8P8	XERD_ECOLI	cytoplasm (by similarity).	nucleoid
P0A8P9	XERD_ECOL6	cytoplasm (by similarity).	nucleoid
P0A8Q1	FRDC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8Q2	FRDC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8Q4	FRDD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8Q5	FRDD_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8Q6	CLPS_ECOLI		cytoplasm
P0A8Q7	CLPS_ECOL6		cytoplasm
P0A8Q8	CLPS_ECO57		cytoplasm
P0A8Q9	CLPS_SHIFL		cytoplasm
P0A8R0	RRAA_ECOLI		cytoplasm
P0A8R1	RRAA_ECOL6		cytoplasm
P0A8R2	RRAA_ECO57		cytoplasm
P0A8R3	RRAA_SHIFL		cytoplasm
P0A8R4	SLYX_ECOLI		inner membrane
P0A8R5	SLYX_ECOL6		inner membrane
P0A8R6	SLYX_ECO57		inner membrane
P0A8R8	YCJF_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8R9	HDFR_ECOLI		cytoplasm
P0A8S0	HDFR_SHIFL		cytoplasm
P0A8S1	ICIA_ECOLI		cytoplasm
P0A8S2	ICIA_ECOL6		cytoplasm

P0A8S3	ICIA_ECO57		cytoplasm
P0A8S4	ICIA_SHIFL		cytoplasm
P0A8S6	USPB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8S7	USPB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8S8	USPB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0A8T0	FLHD_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A8T2	PRMA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A8T3	PRMA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A8T4	PRMA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A8T5	FLIE_ECOLI		flagellar
P0A8T6	FLIE_ECO57		flagellar
P0A8T7	RPOC_ECOLI		cytoplasm
P0A8T8	RPOC_ECO57		cytoplasm
P0A8T9	RPOC_SHIFL		cytoplasm
P0A8U2	YAFD_ECOLI	cytoplasm (potential).	inner membrane
P0A8U3	YAFD_ECOL6	cytoplasm (potential).	inner membrane
P0A8U4	YAFD_ECO57	cytoplasm (potential).	inner membrane
P0A8U5	YAFD_SHIFL	cytoplasm (potential).	inner membrane
P0A8U7	METJ_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A8U8	METJ_ECO57	cytoplasm (by similarity).	cytoplasm
P0A8U9	METJ_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A8V0	RNZ_ECOLI		extracellular
P0A8V1	RNZ_SHIFL		extracellular
P0A8V2	RPOB_ECOLI		cytoplasm
P0A8V3	RPOB_ECOL6		cytoplasm
P0A8V4	RPOB_ECO57		cytoplasm
P0A8V5	RPOB_SHIFL		cytoplasm
P0A8V6	FADR_ECOLI	cytoplasm (potential).	cytoplasm
P0A8V7	FADR_ECOL6	cytoplasm (potential).	cytoplasm
P0A8V8	FADR_ECO57	cytoplasm (potential).	cytoplasm
P0A8V9	FADR_SHIFL	cytoplasm (potential).	cytoplasm
P0A8W0	NANR_ECOLI		cytoplasm
P0A8W1	NANR_SHIFL		cytoplasm
P0A8W2	SLYA_ECOLI		cytoplasm
P0A8W3	SLYA_ECO57		cytoplasm
P0A8W4	SLYA_SHIFL		cytoplasm
P0A8W5	YQGE_ECOLI		inner membrane
P0A8W6	YQGE_ECOL6		inner membrane
P0A8W7	YQGE_SHIFL		inner membrane
P0A8W8	YFBU_ECOLI		inner membrane
P0A8W9	YFBU_ECOL6		inner membrane
P0A8X0	YJGA_ECOLI		cytoplasm
P0A8X1	YJGA_ECO57		cytoplasm
P0A8X2	YCEI_ECOLI	periplasm (potential).	periplasm
P0A8X3	YCEI_ECO57	periplasm (potential).	periplasm
P0A8X4	YCCT_ECOLI		inner membrane
P0A8X5	YCCT_ECOL6		inner membrane
P0A8X6	YCCT_ECO57		inner membrane
P0A8X7	YCCT_SHIFL		inner membrane
P0A8Y1	YJG_ECOLI	cytoplasm (potential).	periplasm

P0A8Y2	YJG_ECO57	cytoplasm (potential).	periplasm
P0A8Y3	YIHX_ECOLI		cytoplasm
P0A8Y4	YIHX_SHIFL		cytoplasm
P0A8Y5	YIDA_ECOLI		periplasm
P0A8Y6	YIDA_ECOL6		periplasm
P0A8Y7	YIDA_SHIFL		periplasm
P0A8Y8	YBDB_ECOLI		cytoplasm
P0A8Y9	YBDB_ECOL6		cytoplasm
P0A8Z0	YCIA_ECOLI		cytoplasm
P0A8Z1	YCIA_ECOL6		cytoplasm
P0A8Z2	YCIA_SHIFL		cytoplasm
P0A8Z3	YBGC_ECOLI		cytoplasm
P0A8Z4	YBGC_ECOL6		cytoplasm
P0A8Z5	YBGC_ECO57		cytoplasm
P0A8Z6	YBGC_SHIFL		cytoplasm
P0A8Z7	YQIA_ECOLI		cytoplasm
P0A8Z8	YQIA_ECOL6		cytoplasm
P0A8Z9	YQIA_ECO57		cytoplasm
P0A900	YQIA_SHIFL		cytoplasm
P0A902	BLC_ECO57	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P0A904	NLPB_SHIFL	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P0A905	SLYB_ECOLI	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P0A906	SLYB_ECO57	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P0A907	SLYB_SHIFL	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P0A909	MIPA_SHIFL	bacterial cell outer membrane (probable).	outer membrane
P0A911	OMPA_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A913	PAL_ECO57	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P0A914	PAL_SHIFL	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P0A916	OMPW_SHIFL	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P0A918	OMPX_ECOL6	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A919	OMPX_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A920	OMPX_SHIFL	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A928	TSX_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A929	TSX_SHIFL	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane

P0A930	WZA_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A931	WZA_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A932	YCCZ_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A933	YCCZ_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A934	YCCZ_SHIFL	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P0A936	MLTA_ECO57	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
P0A937	SMPA_ECOLI	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P0A938	SMPA_ECOL6	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P0A939	SMPA_SHIFL	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P0A941	YAET_ECOL6	bacterial cell outer membrane (by similarity).	outer membrane
P0A942	YAET_ECO57	bacterial cell outer membrane (by similarity).	outer membrane
P0A943	YAET_SHIFL	bacterial cell outer membrane (by similarity).	outer membrane
P0A945	RIMI_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A946	RIMI_ECO57	cytoplasm (by similarity).	cytoplasm
P0A947	RIMI_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A949	RIMJ_ECO57	cytoplasm (by similarity).	cytoplasm
P0A950	RIMJ_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A951	ATDA_ECOLI	cytoplasm (potential).	cytoplasm
P0A952	ATDA_ECO57	cytoplasm (potential).	cytoplasm
P0A954	FABB_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A956	ALKH_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A957	ALKH_ECO57	cytoplasm (by similarity).	cytoplasm
P0A958	ALKH_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A959	YFBQ_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A960	YFBQ_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A961	YFBQ_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A963	ASPG1_ECO57	cytoplasm (by similarity).	cytoplasm
P0A965	CHEW_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A966	CHEW_ECO57	cytoplasm (by similarity).	cytoplasm
P0A967	CHEW_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A969	CSPD_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A970	CSPD_ECO57	cytoplasm (by similarity).	cytoplasm
P0A971	CSPD_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A972	CSPE_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A973	CSPE_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A974	CSPE_ECO57	cytoplasm (by similarity).	cytoplasm
P0A975	CSPE_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A976	CSPF_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A977	CSPF_ECOL6	cytoplasm (by similarity).	cytoplasm

P0A978	CSPG_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A979	CSPG_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A980	CSPG_ECO57	cytoplasm (by similarity).	cytoplasm
P0A981	CSPG_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A982	CSPH_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A983	CSPH_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A984	CSPH_ECO57	cytoplasm (by similarity).	cytoplasm
P0A985	CSPH_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A986	CSPI_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A987	CSPI_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A989	DPO3B_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A990	DPO3B_ECO57	cytoplasm (by similarity).	cytoplasm
P0A991	ALF1_ECOLI	cytoplasm (probable).	cytoplasm
P0A992	ALF1_ECOL6	cytoplasm (probable).	cytoplasm
P0A993	F16P_ECOLI	cytoplasm (potential).	cytoplasm
P0A994	F16P_ECOL6	cytoplasm (potential).	cytoplasm
P0A995	F16P_SHIFL	cytoplasm (potential).	cytoplasm
P0A999	FTNA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9A0	FTNA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9A1	FTNA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9A2	FTNB_ECOLI	cytoplasm (potential).	cytoplasm
P0A9A3	FTNB_ECOL6	cytoplasm (potential).	cytoplasm
P0A9A4	FTNB_ECO57	cytoplasm (potential).	cytoplasm
P0A9A5	FTNB_SHIFL	cytoplasm (potential).	cytoplasm
P0A9B0	FUR_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9B1	FUR_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9B3	G3P1_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9B4	G3P1_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9B5	G3P1_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9B7	E4PD_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9B8	E4PD_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9B9	E4PD_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9C3	GALM_ECOLI	cytoplasm (probable).	cytoplasm
P0A9C4	GALM_SHIFL	cytoplasm (probable).	cytoplasm
P0A9C6	GLNA_ECOL6	cytoplasm (probable).	cytoplasm
P0A9C7	GLNA_ECO57	cytoplasm (probable).	cytoplasm
P0A9C8	GLNA_SHIFL	cytoplasm (probable).	cytoplasm
P0A9D0	GLPX_ECOL6	cytoplasm (probable).	cytoplasm
P0A9D1	GLPX_SHIFL	cytoplasm (probable).	cytoplasm
P0A9D3	GST_ECO57	cytoplasm (probable).	cytoplasm
P0A9D5	CYSE_ECOL6	cytoplasm (probable).	cytoplasm
P0A9D6	CYSE_ECO57	cytoplasm (probable).	cytoplasm
P0A9D7	CYSE_SHIFL	cytoplasm (probable).	cytoplasm
P0A9D9	DAPD_SHIFL	cytoplasm (probable).	cytoplasm
P0A9E1	ARAC_ECO57	cytoplasm (probable).	cytoplasm
P0A9E3	SOXS_ECOL6	cytoplasm (probable).	cytoplasm
P0A9E4	SOXS_ECO57	cytoplasm (probable).	cytoplasm
P0A9E5	FNR_ECOLI	cytoplasm (probable).	cytoplasm
P0A9E6	FNR_ECOL6	cytoplasm (probable).	cytoplasm
P0A9E7	FNR_ECO57	cytoplasm (probable).	cytoplasm
P0A9E8	FNR_SHIFL	cytoplasm (probable).	cytoplasm
P0A9E9	NSR_ECOLI	cytoplasm (probable).	cytoplasm
P0A9F0	NSR_ECO57	cytoplasm (probable).	cytoplasm
P0A9F1	MNTR_ECOLI	cytoplasm (probable).	cytoplasm
P0A9F2	MNTR_ECOL6	cytoplasm (probable).	cytoplasm
P0A9F4	CYSB_ECOL6	cytoplasm (probable).	cytoplasm

P0A9F5	CYSB_ECO57	cytoplasm (probable).	cytoplasm
P0A9F7	GCVA_ECOL6	cytoplasm (probable).	cytoplasm
P0A9F8	GCVA_ECO57	cytoplasm (probable).	cytoplasm
P0A9G0	METR_ECO57	cytoplasm (probable).	cytoplasm
P0A9G1	METR_SHIFL	cytoplasm (probable).	cytoplasm
P0A9G3	NHAR_SHIFL	cytoplasm (probable).	cytoplasm
P0A9G4	CUER_ECOLI	cytoplasm (probable).	cytoplasm
P0A9G5	CUER_SHIFL	cytoplasm (probable).	cytoplasm
P0A9G7	ACEA_ECOL6	cytoplasm (probable).	cytoplasm
P0A9G9	MODE_ECO57	cytoplasm (probable).	cytoplasm
P0A9H0	MODE_SHIFL	cytoplasm (probable).	cytoplasm
P0A9H1	MUG_ECOLI	cytoplasm (potential).	extracellular
P0A9H2	MUG_ECOL6	cytoplasm (potential).	extracellular
P0A9H3	LDCI_ECOLI	cytoplasm (probable).	cytoplasm
P0A9H4	LDCI_ECO57	cytoplasm (probable).	cytoplasm
P0A9H6	BTUR_ECOL6	cytoplasm (probable).	cytoplasm
P0A9H8	CFA_ECOL6	cytoplasm (probable).	cytoplasm
P0A9I0	CHEZ_ECOL6	cytoplasm (probable).	cytoplasm
P0A9I2	CILB_ECOL6	cytoplasm (probable).	cytoplasm
P0A9I4	GCVR_SHIFL	cytoplasm (probable).	cytoplasm
P0A9I5	NAPD_ECOLI	cytoplasm (potential).	fimbrium
P0A9I6	NAPD_ECO57	cytoplasm (potential).	cytoplasm
P0A9I7	NAPD_SHIFL	cytoplasm (potential).	cytoplasm
P0A9I9	NIRD_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9J1	RNG_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9J2	RNG_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9J3	RNG_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9J4	PANE_ECOLI	cytoplasm (potential).	cytoplasm
P0A9J5	PANE_SHIFL	cytoplasm (potential).	cytoplasm
P0A9J7	RBSK_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9J9	PHEA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9K0	PHEA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9K2	PHOH_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9K3	PHOL_ECOLI	cytoplasm (potential).	cytoplasm
P0A9K4	PHOL_ECOL6	cytoplasm (potential).	cytoplasm
P0A9K5	PHOL_ECO57	cytoplasm (potential).	cytoplasm
P0A9K6	PHOL_SHIFL	cytoplasm (potential).	cytoplasm
P0A9K8	PHOU_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9L0	SLYD_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9L1	SLYD_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9L2	SLYD_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9L4	FKBB_SHIFL		cytoplasm
P0A9L6	PPIC_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9L7	PPIC_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9L9	PROC_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9M1	LON_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9M3	HPRT_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9M4	HPRT_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9M6	XGPT_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9M7	XGPT_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9M8	PTA_ECOLI	cytoplasm (potential).	inner membrane
P0A9M9	PTA_SHIFL	cytoplasm (potential).	inner membrane
P0A9N0	PTSO_ECOLI	cytoplasm (probable).	cytoplasm
P0A9N1	PTSO_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9N2	PTSO_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9N3	PTSO_SHIFL	cytoplasm (by similarity).	cytoplasm

P0A9N5	PFLA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9N6	PFLA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9N7	PFLA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9N9	NRDG_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9P1	DLDH_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9P2	DLDH_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9P3	DLDH_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9P5	TRXB_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9P6	DEAD_ECOLI	cytoplasm (probable).	cytoplasm
P0A9P7	DEAD_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9P8	DEAD_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9P9	IDNO_ECOLI	cytoplasm (probable).	cytoplasm
P0A9Q0	IDNO_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9Q1	ARCA_ECOLI	cytoplasm (probable).	cytoplasm
P0A9Q2	ARCA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9Q3	ARCA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9Q4	ARCA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9Q5	ACCD_ECOLI		cytoplasm
P0A9Q6	ACCD_ECO57		cytoplasm
P0A9Q7	ADHE_ECOLI		cytoplasm
P0A9Q8	ADHE_ECO57		cytoplasm
P0A9Q9	DHAS_ECOLI		cytoplasm
P0A9R0	DHAS_ECO57		cytoplasm
P0A9R1	DHAS_SHIFL		cytoplasm
P0A9R2	ESSD_ECOLI		extracellular
P0A9R3	ESSD_ECOL6		extracellular
P0A9R4	FER_ECOLI		cytoplasm
P0A9R5	FER_ECO57		cytoplasm
P0A9R6	FER_SHIFL		cytoplasm
P0A9R7	FTSE_ECOLI		cytoplasm
P0A9R8	FTSE_ECOL6		cytoplasm
P0A9R9	FTSE_ECO57		cytoplasm
P0A9S0	FTSE_SHIFL		cytoplasm
P0A9S1	FUCO_ECOLI		cytoplasm
P0A9S2	FUCO_ECO57		cytoplasm
P0A9S3	GATD_ECOLI		cytoplasm
P0A9S4	GATD_ECO57		cytoplasm
P0A9S5	GLDA_ECOLI		cytoplasm
P0A9S6	GLDA_ECOL6		cytoplasm
P0A9S7	LIVG_ECOLI		inner membrane
P0A9S8	LIVG_ECO57		inner membrane
P0A9S9	LIVG_SHIFL		inner membrane
P0A9T0	SERA_ECOLI		cytoplasm
P0A9T1	SERA_ECOL6		cytoplasm
P0A9T2	SERA_ECO57		cytoplasm
P0A9T3	SERA_SHIFL		cytoplasm
P0A9T4	TAS_ECOLI		cytoplasm
P0A9T5	TAS_SHIFL		cytoplasm
P0A9T6	YBAQ_ECOLI		cytoplasm
P0A9T7	YBAQ_ECOL6		cytoplasm
P0A9T8	YBBA_ECOLI		inner membrane
P0A9T9	YBBA_ECO57		inner membrane
P0A9U0	YBBA_SHIFL		inner membrane
P0A9U1	YBHF_ECOLI		inner membrane
P0A9U2	YBHF_SHIFL		inner membrane
P0A9U3	YBIT_ECOLI		inner membrane

P0A9U4	YBIT_ECOL6		inner membrane
P0A9U5	YBIT_ECO57		inner membrane
P0A9U6	PUUR_ECOLI		cytoplasm
P0A9U7	PUUR_SHIFL		cytoplasm
P0A9U8	YDIO_ECOLI		cytoplasm
P0A9U9	YDIO_ECOL6		cytoplasm
P0A9V0	YDIO_ECO57		cytoplasm
P0A9V2	YHBG_ECOL6		inner membrane
P0A9V3	YHBG_ECO57		inner membrane
P0A9V4	YHBG_SHIFL		inner membrane
P0A9V5	YIAG_ECOLI		cytoplasm
P0A9V6	YIAG_ECOL6		cytoplasm
P0A9V7	YIAG_ECO57		cytoplasm
P0A9V8	YIHU_ECOLI		cytoplasm
P0A9V9	YIHU_SHIFL		cytoplasm
P0A9W0	ULAR_ECOLI	cytoplasm (potential).	cytoplasm
P0A9W1	ULAR_ECOL6	cytoplasm (potential).	cytoplasm
P0A9W2	ULAR_ECO57	cytoplasm (potential).	cytoplasm
P0A9W3	YJJK_ECOLI		inner membrane
P0A9W4	YJJK_ECO57		inner membrane
P0A9W5	YJJK_SHIFL		inner membrane
P0A9W6	YRBA_ECOLI		cytoplasm
P0A9W7	YRBA_ECOL6		cytoplasm
P0A9W8	YRBA_SHIFL		cytoplasm
P0A9W9	YRDA_ECOLI		cytoplasm
P0A9X0	YRDA_SHIFL		cytoplasm
P0A9X1	ZNUC_ECOLI		inner membrane
P0A9X2	ZNUC_ECOL6		inner membrane
P0A9X3	ZNUC_ECO57		inner membrane
P0A9X4	MREB_ECOLI		cytoplasm
P0A9X5	MREB_ECOL6		cytoplasm
P0A9X6	MREB_SALTY		cytoplasm
P0A9X7	MREB_SALTI		cytoplasm
P0A9X8	MREB_SHIFL		cytoplasm
P0A9Y0	CSPA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9Y1	CSPA_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9Y2	CSPA_SALTY	cytoplasm (by similarity).	cytoplasm
P0A9Y3	CSPA_SALTI	cytoplasm (by similarity).	cytoplasm
P0A9Y4	CSPA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0A9Y5	CSPA_SALEN	cytoplasm (by similarity).	cytoplasm
P0A9Y6	CSPC_ECOLI	cytoplasm (by similarity).	cytoplasm
P0A9Y7	CSPC_ECOL6	cytoplasm (by similarity).	cytoplasm
P0A9Y8	CSPC_ECO57	cytoplasm (by similarity).	cytoplasm
P0A9Y9	CSPC_SALTY	cytoplasm (by similarity).	cytoplasm
P0A9Z0	CSPC_SALTI	cytoplasm (by similarity).	cytoplasm
P0A9Z1	GLNB_ECOLI		cytoplasm
P0A9Z2	GLNB_ECOL6		cytoplasm
P0A9Z3	GLNB_ECO57		cytoplasm
P0A9Z4	GLNB_SALTY		cytoplasm
P0A9Z5	GLNB_SALTI		cytoplasm
P0A9Z6	GLNB_SHIFL		cytoplasm
P0A9Z7	BLA2_ECOLI		extracellular
P0A9Z8	BLA2_KLEPN		extracellular
P0A9Z9	BLA2_KLEPO		extracellular
P0AA00	BLA2_SALTY		extracellular
P0AA01	NUSG_ECOL6		cytoplasm

P0AA02	NUSG_SALTY		cytoplasm
P0AA03	NUSG_SALTI		cytoplasm
P0AA05	PTHP_ECOL6	cytoplasm (by similarity).	cytoplasm
P0AA06	PTHP_ECO57	cytoplasm (by similarity).	cytoplasm
P0AA07	PTHP_SALTY	cytoplasm (by similarity).	cytoplasm
P0AA08	PTHP_SALTI	cytoplasm (by similarity).	cytoplasm
P0AA09	PTHP_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AA10	RL13_ECOLI		cytoplasm
P0AA11	RL13_ECOL6		cytoplasm
P0AA12	RL13_ECO57		cytoplasm
P0AA13	RL13_SALTY		cytoplasm
P0AA14	RL13_SALTI		cytoplasm
P0AA15	RL13_SHIFL		cytoplasm
P0AA17	OMPR_ECOL6	cytoplasm (by similarity).	cytoplasm
P0AA18	OMPR_ECO57	cytoplasm (by similarity).	cytoplasm
P0AA19	OMPR_SALTY	cytoplasm (by similarity).	cytoplasm
P0AA20	OMPR_SALTI	cytoplasm (by similarity).	cytoplasm
P0AA21	OMPR_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AA22	EBR_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AA23	EBR_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AA24	EBR_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AA25	THIO_ECOLI		cytoplasm
P0AA26	THIO_ECOL6		cytoplasm
P0AA27	THIO_ECO57		cytoplasm
P0AA28	THIO_SALTY		cytoplasm
P0AA29	THIO_SALTI		cytoplasm
P0AA30	THIO_SHIFL		cytoplasm
P0AA31	YEDF_ECOLI		cytoplasm
P0AA32	YEDF_ECOL6		cytoplasm
P0AA33	YEDF_ECO57		cytoplasm
P0AA34	YEDF_SALTY		cytoplasm
P0AA35	YEDF_SALTI		cytoplasm
P0AA36	YEDF_SHIFL		cytoplasm
P0AA37	RLUA_ECOLI		cytoplasm
P0AA38	RLUA_SHIFL		cytoplasm
P0AA39	RLUC_ECOLI		cytoplasm
P0AA40	RLUC_SHIFL		cytoplasm
P0AA41	TRUC_ECOLI		cytoplasm
P0AA42	TRUC_SHIFL		cytoplasm
P0AA43	RSUA_ECOLI		cytoplasm
P0AA44	RSUA_ECOL6		cytoplasm
P0AA45	RSUA_ECO57		cytoplasm
P0AA46	RSUA_SHIFL		cytoplasm
P0AA48	YEEF_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA49	YFDV_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

P0AA50	YFDV_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AA51	YFDV_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AA52	YFDV_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AA53	YBBK_ECOLI		inner membrane
P0AA54	YBBK_ECOL6		inner membrane
P0AA55	YBBK_ECO57		inner membrane
P0AA56	YBBK_SHIFL		inner membrane
P0AA57	YOBA_ECOLI	periplasm (by similarity).	periplasm
P0AA58	YOBA_ECOL6	periplasm (by similarity).	periplasm
P0AA59	YOBA_SHIFL	periplasm (by similarity).	periplasm
P0AA61	YGHB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA62	YGHB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA64	YQJA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA65	YQJA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA66	YQJA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA68	YBIF_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA69	YBIF_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA71	YEDA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA72	YEDA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA74	YHBE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA75	YHBE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA77	DGOT_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA79	EXUT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AA81	GARP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA83	CODB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA84	YBHO_ECOLI	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P0AA85	YBHO_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0AA87	DSBE_ECO57	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
P0AA88	DSBE_SHIFL	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
P0AA89	YDDV_ECOLI		inner membrane
P0AA90	YDDV_ECO57		inner membrane
P0AA91	YEAY_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AA92	YEAY_ECOL6	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AA94	YPDA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AA95	YACC_ECOLI		cytoplasm
P0AA96	YACC_ECOL6		cytoplasm
P0AA97	YAEQ_ECOLI		cytoplasm
P0AA98	YAEQ_SHIFL		cytoplasm
P0AA99	YAFK_ECOLI		extracellular
P0AAA0	YAFK_ECO57		extracellular
P0AAA2	YAGU_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAA3	YAGZ_ECOLI		fimbrium
P0AAA4	YAGZ_ECO57		fimbrium
P0AAA5	SFA_ECOLI		inner membrane
P0AAA6	SFA_ECOL6		inner membrane
P0AAA8	WZXE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAB0	ZRAP_ECO57	periplasm (by similarity).	periplasm
P0AAB1	ZRAP_SHIFL	periplasm (by similarity).	periplasm
P0AAB2	WZB_ECOLI		extracellular
P0AAB3	WZB_ECO57		extracellular
P0AAB4	UBID_ECOLI	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P0AAB5	UBID_ECOL6	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0AAB6	GALF_ECOLI		cytoplasm

P0AAB7	GALF_ECO57		cytoplasm
P0AAB8	USPD_ECOLI	cytoplasm (by similarity).	cytoplasm
P0AAB9	USPD_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AAC0	USPE_ECOLI	cytoplasm (by similarity).	cytoplasm
P0AAC1	USPE_ECOL6	cytoplasm (by similarity).	cytoplasm
P0AAC2	USPE_ECO57	cytoplasm (by similarity).	cytoplasm
P0AAC3	USPE_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AAC5	YBHL_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAC7	YCCA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAC8	ISCA_ECOLI		cytoplasm
P0AAC9	ISCA_ECOL6		cytoplasm
P0AAD0	ISCA_ECO57		cytoplasm
P0AAD1	ISCA_SHIFL		cytoplasm
P0AAD3	MTR_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAD5	TYRP_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAD7	SDAC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAD9	TDCC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAE1	CYCA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAE3	PROY_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAE4	PROY_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAE6	ARCD_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAE7	ARCD_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAE9	CADB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAF0	CADB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAF2	POTE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAF4	ARAG_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

P0AAF6	ARTP_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAF7	ARTP_ECOL6	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAF8	ARTP_ECO57	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAF9	ARTP_SHIFL	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAG0	DPPD_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAG1	DPPD_ECO57	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAG2	DPPD_SHIFL	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAG3	GLTL_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAG4	GLTL_ECO57	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAG6	MDLB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAG7	MDLB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAG8	MGLA_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAG9	MGLA_SHIFL	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAH1	PSTB_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AAH2	PSTB_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AAH3	PSTB_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AAH4	SAPD_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAH5	SAPD_ECOL6	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane

P0AAH6	SAPD_ECO57	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAH7	SAPD_SHIFL	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAH8	SAPF_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAH9	SAPF_ECOL6	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAI0	SAPF_SHIFL	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P0AAI1	SSUB_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AAI2	SSUB_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AAI4	FTSH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAI5	FABF_ECOLI		cytoplasm
P0AAI6	FABF_ECOL6		cytoplasm
P0AAI7	FABF_ECO57		cytoplasm
P0AAI8	FABF_SHIFL		cytoplasm
P0AAI9	FABD_ECOLI		cytoplasm
P0AAJ0	FABD_ECOL6		cytoplasm
P0AAJ1	YNFG_ECOLI		periplasm
P0AAJ2	YNFG_ECOL6		periplasm
P0AAJ4	FDNH_SHIFL	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAJ6	FDOH_ECOL6	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAJ7	FDOH_SHIFL	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAJ9	HYBA_ECO57	periplasm (by similarity).	periplasm
P0AAK0	HYBA_SHIFL	periplasm (by similarity).	periplasm
P0AAK1	HYCB_ECOLI		periplasm
P0AAK2	HYCB_ECOL6		periplasm
P0AAK3	HYCB_SHIFL		periplasm
P0AAK4	HYDN_ECOLI		periplasm
P0AAK5	HYDN_ECOL6		periplasm
P0AAK6	HYDN_ECO57		periplasm
P0AAK7	NRFC_ECOLI		cytoplasm
P0AAK8	NRFC_ECO57		cytoplasm
P0AAK9	NRFC_SHIFL		cytoplasm
P0AAL0	NAPF_ECOLI		periplasm
P0AAL1	NAPF_ECOL6		periplasm
P0AAL2	NAPF_ECO57		periplasm
P0AAL3	NAPG_ECOLI		periplasm

P0AAL4	NAPG_ECOL6		periplasm
P0AAL5	NAPG_SHIFL		periplasm
P0AAL6	YDHY_ECOLI		periplasm
P0AAL7	YDHY_ECO57		periplasm
P0AAL8	YDHY_SHIFL		periplasm
P0AAL9	YKGY_ECOLI		cytoplasm
P0AAM0	YKGY_ECO57		cytoplasm
P0AAM2	CYBH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAM3	HYPG_ECOLI		cytoplasm
P0AAM4	HYPG_ECOL6		cytoplasm
P0AAM5	HYPG_ECO57		cytoplasm
P0AAM6	HYPG_SHIFL		cytoplasm
P0AAM7	HYBG_ECOLI		cytoplasm
P0AAM8	HYBG_ECOL6		cytoplasm
P0AAM9	HYBG_ECO57		cytoplasm
P0AAN0	HYBG_SHIFL		cytoplasm
P0AAN1	HYBE_ECOLI		fimbrium
P0AAN2	HYBE_ECOL6		cytoplasm
P0AAN3	HYPB_ECOLI		cytoplasm
P0AAN4	HYPB_ECOL6		cytoplasm
P0AAN5	YAIA_ECOLI		periplasm
P0AAN6	YAIA_ECOL6		periplasm
P0AAN7	YAIA_ECO57		cytoplasm
P0AAN8	YAIA_SHIFL		cytoplasm
P0AAN9	YAIB_ECOLI		cytoplasm
P0AAP0	YAIB_ECO57		cytoplasm
P0AAP1	YAIC_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AAP2	YAIC_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AAP3	YAIN_ECOLI		cytoplasm
P0AAP4	YAIN_ECOL6		cytoplasm
P0AAP5	YAIV_ECOLI		cytoplasm
P0AAP6	YAIV_ECO57		cytoplasm
P0AAP8	YAIY_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAP9	YAIY_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAQ0	YAIZ_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AAQ1	YAIZ_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AAQ2	YAJD_ECOLI		cytoplasm
P0AAQ3	YAJD_ECOL6		cytoplasm
P0AAQ4	YAJD_ECO57		cytoplasm
P0AAQ5	YAJD_SHIFL		cytoplasm
P0AAQ6	YBAA_ECOLI		cytoplasm
P0AAQ7	YBAA_ECOL6		cytoplasm

P0AAQ8	YBAA_ECO57		cytoplasm
P0AAQ9	YBAA_SHIFL		cytoplasm
P0AAR0	YBAJ_ECOLI		cytoplasm
P0AAR1	YBAJ_ECOL6		cytoplasm
P0AAR2	YBAJ_ECO57		cytoplasm
P0AAR3	YBAK_ECOLI		cytoplasm
P0AAR4	YBAK_SHIFL		cytoplasm
P0AAR6	YBAN_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAR7	YBAN_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAR8	YBAV_ECOLI		cytoplasm
P0AAR9	YBAV_SHIFL		cytoplasm
P0AAS1	YLAC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAS2	YLAC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAS4	YBBJ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAS5	YLBF_ECOLI		cytoplasm
P0AAS6	YLBF_ECO57		cytoplasm
P0AAS7	YBCJ_ECOLI		cytoplasm
P0AAS8	YBCJ_SHIFL		cytoplasm
P0AAS9	YBDD_ECOLI		cytoplasm
P0AAT0	YBDD_ECOL6		cytoplasm
P0AAT1	YBDD_SHIFL		cytoplasm
P0AAT2	YBDF_ECOLI		cytoplasm
P0AAT3	YBDF_ECO57		cytoplasm
P0AAT4	YBDG_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AAT5	YBDG_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AAT6	YBEB_ECOLI		inner membrane
P0AAT7	YBEB_ECOL6		inner membrane
P0AAT8	YBEB_SHIFL		inner membrane
P0AAT9	YBEL_ECOLI		cytoplasm
P0AAU0	YBEL_ECOL6		cytoplasm
P0AAU1	YBEL_ECO57		cytoplasm
P0AAU2	YBFA_ECOLI		cytoplasm
P0AAU3	YBFA_ECO57		cytoplasm
P0AAU4	YBFA_SHIFL		cytoplasm
P0AAU5	YBFB_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AAU6	YBFB_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AAU7	YBFE_ECOLI		cytoplasm
P0AAU8	YBFE_ECO57		cytoplasm

P0AAU9	YBFE_SHIFL		cytoplasm
P0AAV0	YBGE_ECOLI		inner membrane
P0AAV1	YBGE_ECOL6		inner membrane
P0AAV2	YBGE_ECO57		inner membrane
P0AAV3	YBGE_SHIFL		inner membrane
P0AAV4	YBGJ_ECOLI		cytoplasm
P0AAV5	YBGJ_ECO57		cytoplasm
P0AAV6	YBGS_ECOLI		outer membrane
P0AAV7	YBGS_SHIFL		outer membrane
P0AAV8	YBHH_ECOLI		cytoplasm
P0AAV9	YBHH_ECOL6		cytoplasm
P0AAW0	YBHH_SHIFL		cytoplasm
P0AAW1	YBHP_ECOLI		cytoplasm
P0AAW2	YBHP_ECOL6		cytoplasm
P0AAW3	YBHP_ECO57		cytoplasm
P0AAW4	YBHP_SHIFL		cytoplasm
P0AAW6	YBHQ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAW7	YBHQ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAW8	YBHQ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAX3	YBIJ_ECOLI		fimbrium
P0AAX4	YBIJ_ECOL6		fimbrium
P0AAX5	YBIJ_SHIFL		fimbrium
P0AAX6	YBIM_ECOLI		cytoplasm
P0AAX7	YBIM_ECO57		cytoplasm
P0AAX8	YBIS_ECOLI	periplasm (potential).	periplasm
P0AAX9	YBIS_ECOL6	periplasm (potential).	periplasm
P0AAY0	YBIS_ECO57	periplasm (potential).	periplasm
P0AAY1	BSSR_ECOLI		cytoplasm
P0AAY2	BSSR_ECO57		cytoplasm
P0AAY3	BSSR_SHIFL		cytoplasm
P0AAY4	YBJH_ECOLI		cytoplasm
P0AAY5	YBJH_SHIFL		cytoplasm
P0AAY6	YBJN_ECOLI		inner membrane
P0AAY7	YBJN_ECOL6		inner membrane
P0AAY8	YBJN_ECO57		inner membrane
P0AAY9	YBJN_SHIFL		inner membrane
P0AAZ1	YBJO_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAZ2	YBJO_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAZ3	YBJO_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AAZ4	YCAJ_ECOLI		cytoplasm
P0AAZ5	YCAJ_ECOL6		cytoplasm
P0AAZ6	YCAJ_ECO57		cytoplasm
P0AAZ7	YCAR_ECOLI		cytoplasm
P0AAZ8	YCAR_ECOL6		cytoplasm

P0AAZ9	YCAR_ECO57		cytoplasm
P0AB00	YCAR_SHIFL		cytoplasm
P0AB01	YCBC_ECOLI		inner membrane
P0AB02	YCBC_ECO57		inner membrane
P0AB03	YCBJ_ECOLI		cytoplasm
P0AB04	YCBJ_ECO57		cytoplasm
P0AB05	YCBJ_SHIFL		cytoplasm
P0AB06	YCBK_ECOLI		inner membrane
P0AB07	YCBK_ECOL6		inner membrane
P0AB08	YCBK_ECO57		inner membrane
P0AB09	YCBK_SHIFL		inner membrane
P0AB10	YMBA_ECOLI		inner membrane
P0AB11	YMBA_SHIFL		inner membrane
P0AB13	YCCF_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AB14	YCCJ_ECOLI		cytoplasm
P0AB15	YCCJ_ECOL6		cytoplasm
P0AB16	YCCJ_ECO57		cytoplasm
P0AB17	YCCJ_SHIFL		cytoplasm
P0AB18	TUSE_ECOLI	cytoplasm (probable).	cytoplasm
P0AB19	TUSE_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AB20	YCCV_ECOLI		cytoplasm
P0AB21	YCCV_ECOL6		cytoplasm
P0AB22	YCCV_ECO57		cytoplasm
P0AB23	YCCV_SHIFL		cytoplasm
P0AB24	YCDO_ECOLI		cytoplasm
P0AB25	YCDO_SHIFL		cytoplasm
P0AB26	YCEB_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AB27	YCEB_SHIFL	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AB28	YCED_ECOLI		cytoplasm
P0AB29	YCED_ECOL6		cytoplasm
P0AB30	YCED_ECO57		cytoplasm
P0AB31	YCEK_ECOLI		inner membrane
P0AB32	YCEK_ECO57		inner membrane
P0AB33	BSSS_ECOLI		cytoplasm
P0AB34	BSSS_ECOL6		cytoplasm
P0AB35	YCFJ_ECOLI		outer membrane
P0AB36	YCFJ_ECOL6		outer membrane
P0AB37	YCFJ_SHIFL		outer membrane
P0AB38	YCFM_ECOLI		periplasm
P0AB39	YCFM_ECO57		periplasm
P0AB40	YCFR_ECOLI		outer membrane
P0AB41	YCFR_ECOL6		outer membrane
P0AB42	YCFR_ECO57		outer membrane
P0AB43	YCGL_ECOLI		cytoplasm
P0AB44	YCGL_ECOL6		cytoplasm
P0AB45	YCGL_ECO57		cytoplasm
P0AB46	YMGD_ECOLI		cytoplasm
P0AB47	YMGD_ECOL6		cytoplasm
P0AB48	YMGD_SHIFL		cytoplasm
P0AB49	YCHH_ECOLI		inner membrane
P0AB50	YCHH_ECOL6		inner membrane
P0AB51	YCHH_ECO57		inner membrane

P0AB52	YCHN_ECOLI		cytoplasm
P0AB53	YCHN_ECO57		cytoplasm
P0AB54	YCHN_SHIFL		cytoplasm
P0AB55	YCII_ECOLI		cytoplasm
P0AB56	YCII_ECOL6		cytoplasm
P0AB57	YCII_SHIFL		cytoplasm
P0AB58	YCIM_ECOLI		cytoplasm
P0AB59	YCIM_ECOL6		cytoplasm
P0AB60	YCIM_ECO57		cytoplasm
P0AB61	YCIN_ECOLI		cytoplasm
P0AB62	YCIN_ECOL6		cytoplasm
P0AB63	YCIN_ECO57		cytoplasm
P0AB64	YCIN_SHIFL		cytoplasm
P0AB65	ACYP_ECOLI		extracellular
P0AB66	ACYP_ECO57		extracellular
P0AB68	PNTB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AB69	PNTB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AB70	PNTB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AB71	ALF_ECOLI		cytoplasm
P0AB72	ALF_ECO57		cytoplasm
P0AB73	ALF_SHIFL		cytoplasm
P0AB74	AGAY_ECOLI		cytoplasm
P0AB75	AGAY_ECOL6		cytoplasm
P0AB76	AGAY_ECO57		cytoplasm
P0AB77	KBL_ECOLI		inner membrane
P0AB78	KBL_ECO57		inner membrane
P0AB79	KBL_SHIFL		inner membrane
P0AB80	ILVE_ECOLI		cytoplasm
P0AB81	ILVE_ECOL6		cytoplasm
P0AB82	ILVE_ECO57		cytoplasm
P0AB83	END3_ECOLI		cytoplasm
P0AB84	END3_ECOL6		cytoplasm
P0AB85	APBE_ECOLI	bacterial cell membrane; lipid-anchor (potential).	cytoplasm
P0AB86	APBE_ECO57	bacterial cell membrane; lipid-anchor (potential).	cytoplasm
P0AB87	FUCA_ECOLI		extracellular
P0AB88	FUCA_SHIFL		extracellular
P0AB89	PUR8_ECOLI		cytoplasm
P0AB90	PUR8_ECOL6		cytoplasm
P0AB91	AROG_ECOLI		cytoplasm
P0AB92	AROG_ECO57		cytoplasm
P0AB94	ARSB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AB95	ARSB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AB96	ARSC_ECOLI		cytoplasm
P0AB97	ARSC_SHIFL		cytoplasm

P0AB99	ATP6_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABA1	ATPF_ECOL6	bacterial cell inner membrane (by similarity).	inner membrane
P0ABA2	ATPF_ECO57	bacterial cell inner membrane (by similarity).	inner membrane
P0ABA3	ATPF_SHIFL	bacterial cell inner membrane (by similarity).	inner membrane
P0ABA5	ATPD_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABA7	ATPG_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABA8	ATPG_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABA9	ATPG_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABB1	ATPA_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABB2	ATPA_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABB3	ATPA_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABB4	ATPB_ECOLI		inner membrane
P0ABB5	ATPB_ECOL6		inner membrane
P0ABB6	ATPB_ECO57		inner membrane
P0ABB7	ATPB_SHIFL		inner membrane
P0ABB9	ATMA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABC1	ATPZ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABC2	ATPZ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABC3	HFLC_ECOLI		extracellular
P0ABC4	HFLC_ECOL6		extracellular
P0ABC5	HFLC_ECO57		extracellular
P0ABC6	HFLC_SHIFL		extracellular
P0ABC7	HFLK_ECOLI		inner membrane
P0ABC8	HFLK_ECO57		inner membrane
P0ABD0	BETT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABD1	YEAV_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane

P0ABD2	YEAV_ECO57	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0ABD3	BFR_ECOLI		cytoplasm
P0ABD4	BFR_ECOL6		cytoplasm
P0ABD6	ACCA_ECO57	cytoplasm (by similarity).	cytoplasm
P0ABD7	ACCA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0ABD8	BCCP_ECOLI		cytoplasm
P0ABD9	BCCP_ECOL6		cytoplasm
P0ABE0	BCCP_ECO57		cytoplasm
P0ABE1	BCCP_SHIFL		cytoplasm
P0ABE2	BOLA_ECOLI		cytoplasm
P0ABE3	BOLA_ECOL6		cytoplasm
P0ABE4	BOLA_SHIFL		cytoplasm
P0ABE6	C561_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABE8	C562_ECO57	periplasm (by similarity).	periplasm
P0ABE9	CYNT_ECOLI		cytoplasm
P0ABF0	CYNT_ECO57		cytoplasm
P0ABF1	PCNB_ECOLI		cytoplasm
P0ABF2	PCNB_ECOL6		cytoplasm
P0ABF3	PCNB_ECO57		cytoplasm
P0ABF4	EUTM_ECOLI		cytoplasm
P0ABF5	EUTM_ECOL6		cytoplasm
P0ABF6	CDD_ECOLI		cytoplasm
P0ABF7	CDD_SHIFL		cytoplasm
P0ABF9	PGSA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABG0	PGSA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABG2	CDSA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABG3	CDSA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABG5	FTSW_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABG8	RODA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABG9	RODA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABH0	FTSA_ECOLI		inner membrane
P0ABH1	FTSA_ECOL6		inner membrane
P0ABH2	FTSA_ECO57		inner membrane
P0ABH3	FTSA_SHIFL		inner membrane
P0ABH5	MRED_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0ABH6	MRED_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABH7	CISY_ECOLI		cytoplasm
P0ABH8	CISY_ECOL6		cytoplasm
P0ABH9	CLPA_ECOLI		cytoplasm
P0ABI0	CLPA_ECOL6		cytoplasm
P0ABI1	CLPA_ECO57		cytoplasm
P0ABI5	CORA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABI6	CORA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABI7	CORA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABI9	CYOB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABJ0	CYOB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABJ2	CYOA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABJ4	CYOC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABJ5	CYOC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABJ7	CYOD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABJ8	CYOD_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABK0	CYDA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABK1	CYDA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABK3	CYDB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABK4	CYDB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABK5	CYSK_ECOLI		cytoplasm
P0ABK6	CYSK_ECO57		cytoplasm
P0ABK7	CSGB_ECOLI		extracellular
P0ABK8	CSGB_ECO57		extracellular
P0ABL0	NRFA_ECO57	periplasm (by similarity).	periplasm
P0ABL1	NRFB_ECOLI	periplasm (potential).	periplasm

P0ABL2	NRFB_ECOL6	periplasm (potential).	periplasm
P0ABL3	NAPB_ECOLI	periplasm (potential).	periplasm
P0ABL4	NAPB_SHIFL	periplasm (potential).	periplasm
P0ABL9	CCMB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABM0	CCMB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABM2	CCMC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABM3	CCMC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABM4	CCMC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABM5	CCMD_ECOLI		inner membrane
P0ABM6	CCMD_ECOL6		inner membrane
P0ABM7	CCMD_ECO57		inner membrane
P0ABM8	CCMD_SHIFL		inner membrane
P0ABN0	CCMH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABN2	KDGL_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABN3	KDGL_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABN4	KDGL_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABN6	DCUA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABN7	DCUA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABN8	DCUA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABP0	DCUB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABP1	DCUB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABP2	DCUB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABP4	DCUC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0ABP5	DCUC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABP7	DEDA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABP8	DEOD_ECOLI		cytoplasm
P0ABP9	DEOD_ECO57		cytoplasm
P0ABQ0	COABC_ECOLI		cytoplasm
P0ABQ1	COABC_ECOL6		cytoplasm
P0ABQ2	GARR_ECOLI		cytoplasm
P0ABQ3	GARR_ECOL6		cytoplasm
P0ABQ4	DYR_ECOLI		cytoplasm
P0ABQ5	DYR_ECOL6		cytoplasm
P0ABQ6	DYR_SHIFL		cytoplasm
P0ABQ7	DYR8_ECOLI		cytoplasm
P0ABQ8	DYR8_SHISO		cytoplasm
P0ABR1	DINI_ECOLI		inner membrane
P0ABR2	DINI_ECOL6		inner membrane
P0ABR3	DINI_ECO57		inner membrane
P0ABR4	DINI_SHIFL		inner membrane
P0ABR5	HCAE_ECOLI		cytoplasm
P0ABR6	HCAE_ECO57		cytoplasm
P0ABR7	YEAW_ECOLI		cytoplasm
P0ABR8	YEAW_ECO57		cytoplasm
P0ABR9	MHPB_ECOLI		cytoplasm
P0ABS0	MHPB_ECO57		cytoplasm
P0ABS1	DKSA_ECOLI		cytoplasm
P0ABS2	DKSA_ECOL6		cytoplasm
P0ABS3	DKSA_ECO57		cytoplasm
P0ABS4	DKSA_SHIFL		cytoplasm
P0ABS5	PRIM_ECOLI		cytoplasm
P0ABS6	PRIM_ECO57		cytoplasm
P0ABS7	PRIM_SHIFL		cytoplasm
P0ABS8	HOLE_ECOLI		cytoplasm
P0ABS9	HOLE_ECOL6		cytoplasm
P0ABT0	HOLE_ECO57		cytoplasm
P0ABT1	HOLE_SHIFL		cytoplasm
P0ABT2	DPS_ECOLI		cytoplasm
P0ABT3	DPS_ECO57		cytoplasm
P0ABT4	DPS_SHIFL		cytoplasm
P0ABT5	DUSB_ECOLI		cytoplasm
P0ABT6	DUSB_ECOL6		cytoplasm
P0ABT7	DUSB_ECO57		cytoplasm
P0ABT9	YIJE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABU0	MENB_ECOLI		cytoplasm
P0ABU1	MENB_ECOL6		cytoplasm
P0ABU2	ENGD_ECOLI		cytoplasm
P0ABU3	ENGD_ECO57		cytoplasm
P0ABU4	ENGD_SHIFL		cytoplasm
P0ABU5	ELBB_ECOLI		cytoplasm
P0ABU6	ELBB_SHIFL		cytoplasm

P0ABU8	EXBB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABV0	TOLQ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABV1	TOLQ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ABV3	EXBD_ECOL6	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0ABV4	EXBD_ECO57	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0ABV5	EXBD_SHIFL	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0ABV7	TOLR_ECOL6	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0ABV8	TOLR_ECO57	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0ABV9	TOLR_SHIFL	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0ABW0	HCAC_ECOLI		cytoplasm
P0ABW1	HCAC_ECO57		cytoplasm
P0ABW2	HCAC_SHIFL		cytoplasm
P0ABW3	YFAE_ECOLI		cytoplasm
P0ABW4	YFAE_ECOL6		cytoplasm
P0ABW6	SFMA_ECO57	fimbrium (by similarity).	extracellular
P0ABW7	FMS1_ECOLI		fimbrium
P0ABW8	FMS1_ECOL6		fimbrium
P0ABW9	FLGB_ECOLI		flagellar
P0ABX0	FLGB_ECOL6		flagellar
P0ABX1	FLGB_SHIFL		flagellar
P0ABX2	FLGC_ECOLI		flagellar
P0ABX3	FLGC_ECOL6		flagellar
P0ABX4	FLGC_ECO57		flagellar
P0ABX5	FLGG_ECOLI		flagellar
P0ABX6	FLGG_ECOL6		flagellar
P0ABX7	FLGG_ECO57		flagellar
P0ABX9	FLIL_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABY0	FLIL_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABY1	FLIL_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABY2	FLIT_ECOLI		inner membrane
P0ABY3	FLIT_ECO57		inner membrane
P0ABY4	FLAW_ECOLI		cytoplasm

P0ABY5	FLAW_ECOL6		cytoplasm
P0ABY6	FLAW_ECO57		cytoplasm
P0ABY7	FLHC_ECOLI		cytoplasm
P0ABY8	FLHC_ECOL6		cytoplasm
P0ABY9	FLHC_ECO57		cytoplasm
P0ABZ0	FLHC_SHIFL		cytoplasm
P0ABZ2	FLIG_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABZ3	FLIG_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0ABZ4	KDSC_ECOLI		cytoplasm
P0ABZ5	KDSC_ECO57		cytoplasm
P0ABZ7	SURA_ECOL6	periplasm (by similarity).	periplasm
P0ABZ8	SURA_ECO57	periplasm (by similarity).	periplasm
P0ABZ9	SURA_SHIFL	periplasm (by similarity).	periplasm
P0AC00	FRLB_ECOLI		cytoplasm
P0AC01	FRLB_SHIFL		cytoplasm
P0AC02	YFIO_ECOLI	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P0AC03	YFIO_ECOL6	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P0AC04	YFIO_ECO57	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P0AC06	FLIP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC07	FLIQ_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0AC08	FLIQ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0AC09	FLIQ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0AC10	FLIQ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0AC11	DHP2_ECOLI		cytoplasm
P0AC12	DHP2_SHIFL		cytoplasm
P0AC13	DHPS_ECOLI		cytoplasm
P0AC14	DHPS_ECOL6		cytoplasm
P0AC15	DHPS_SHIFL		cytoplasm
P0AC16	FOLB_ECOLI		cytoplasm
P0AC17	FOLB_ECOL6		cytoplasm
P0AC18	FOLB_SHIFL		cytoplasm
P0AC19	FOLX_ECOLI		cytoplasm
P0AC20	FOLX_ECOL6		cytoplasm
P0AC21	FOLX_ECO57		cytoplasm
P0AC22	FOLX_SHIFL		cytoplasm
P0AC24	FOCA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AC25	FOCA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC27	NIRC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC28	YGFA_ECOLI		cytoplasm
P0AC29	YGFA_ECO57		cytoplasm
P0AC31	FTSX_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC32	FTSX_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC33	FUMA_ECOLI		cytoplasm
P0AC34	FUMA_ECOL6		cytoplasm
P0AC35	TTDB_ECOLI		cytoplasm
P0AC36	TTDB_ECOL6		cytoplasm
P0AC37	TTDB_SHIFL		cytoplasm
P0AC38	ASPA_ECOLI		cytoplasm
P0AC39	ASPA_ECOL6		cytoplasm
P0AC40	ASPA_SHIFL		cytoplasm
P0AC42	DHSA_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AC43	DHSA_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AC45	DHSD_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC46	DHSD_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC47	FRDB_ECOLI		cytoplasm
P0AC48	FRDB_ECOL6		cytoplasm
P0AC49	FRDB_ECO57		cytoplasm
P0AC50	FRDB_SHIFL		cytoplasm
P0AC51	ZUR_ECOLI		cytoplasm
P0AC52	ZUR_SHIFL		cytoplasm
P0AC53	G6PD_ECOLI		cytoplasm
P0AC54	G6PD_ECOL6		cytoplasm
P0AC55	GLNK_ECOLI		cytoplasm
P0AC56	GLNK_ECOL6		cytoplasm
P0AC57	GLNK_ECO57		cytoplasm
P0AC58	GLNK_SHIFL		cytoplasm
P0AC59	GLRX2_ECOLI		cytoplasm
P0AC60	GLRX2_ECOL6		cytoplasm
P0AC61	GLRX2_ECO57		cytoplasm
P0AC62	GLRX3_ECOLI		cytoplasm
P0AC63	GLRX3_ECOL6		cytoplasm
P0AC64	GLRX3_ECO57		cytoplasm
P0AC65	NRDH_ECOLI		cytoplasm
P0AC66	NRDH_ECOL6		cytoplasm
P0AC67	NRDH_ECO57		cytoplasm
P0AC68	NRDH_SHIFL		cytoplasm

P0AC69	YDHD_ECOLI		periplasm
P0AC70	YDHD_ECOL6		periplasm
P0AC71	YDHD_ECO57		periplasm
P0AC72	YDHD_SHIFL		periplasm
P0AC73	EBGC_ECOLI		inner membrane
P0AC74	EBGC_ECOL6		inner membrane
P0AC75	KDTA_ECOLI	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
P0AC76	KDTA_ECOL6	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P0AC77	KDTA_ECO57	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P0AC79	WECA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC80	WECA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC81	LGUL_ECOLI		extracellular
P0AC82	LGUL_ECO57		extracellular
P0AC83	LGUL_SHIFL		extracellular
P0AC84	GLO2_ECOLI		cytoplasm
P0AC85	GLO2_ECO57		cytoplasm
P0AC86	PHSG_ECOLI		cytoplasm
P0AC87	PHSG_SHIFL		cytoplasm
P0AC88	GM4D_ECOLI		cytoplasm
P0AC89	GM4D_ECOL6		cytoplasm
P0AC90	GM4D_ECO57		cytoplasm
P0AC91	GM4D_SHIFL		cytoplasm
P0AC92	GNSA_ECOLI		cytoplasm
P0AC93	GNSA_ECOL6		cytoplasm
P0AC95	Gntp_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC97	Gntu_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AC99	YAAH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ACA0	YAAH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ACA1	YIBF_ECOLI		cytoplasm
P0ACA2	YIBF_ECOL6		cytoplasm
P0ACA3	SSPA_ECOLI		cytoplasm
P0ACA4	SSPA_ECOL6		cytoplasm
P0ACA5	SSPA_ECO57		cytoplasm
P0ACA6	SSPA_SHIFL		cytoplasm
P0ACA7	YLIJ_ECOLI		cytoplasm
P0ACA8	YLIJ_ECOL6		cytoplasm
P0ACA9	YLIJ_SHIFL		cytoplasm
P0ACB0	DNAB_ECOLI		cytoplasm

P0ACB1	DNAB_SHIFL		cytoplasm
P0ACB2	HEM2_ECOLI		cytoplasm
P0ACB3	HEM2_SHIFL		cytoplasm
P0ACB4	HEMG_ECOLI		cytoplasm
P0ACB5	HEMG_ECOL6		cytoplasm
P0ACB6	HEMG_ECO57		cytoplasm
P0ACB8	HEMY_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ACB9	HEMY_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ACC0	HEMY_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ACC1	HEMK_ECOLI		cytoplasm
P0ACC2	HEMK_SHIFL		cytoplasm
P0ACC3	YADR_ECOLI		inner membrane
P0ACC4	YADR_ECOL6		inner membrane
P0ACC5	YADR_ECO57		inner membrane
P0ACC6	YADR_SHIFL		inner membrane
P0ACC8	GLMU_ECO57	cytoplasm (by similarity).	cytoplasm
P0ACC9	WCAB_ECOLI		cytoplasm
P0ACD0	WCAB_ECO57		cytoplasm
P0ACD1	WCAB_SHIFL		cytoplasm
P0ACD2	WCAF_ECOLI		cytoplasm
P0ACD3	WCAF_SHIFL		cytoplasm
P0ACD4	NIFU_ECOLI		cytoplasm
P0ACD5	NIFU_ECOL6		cytoplasm
P0ACD6	NIFU_ECO57		cytoplasm
P0ACD7	NIFU_SHIFL		cytoplasm
P0ACD9	MBHL_SHIFL	bacterial cell membrane; peripheral membrane protein (by similarity).	periplasm
P0ACE1	MBHM_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	periplasm
P0ACE2	MBHM_SHIFL	bacterial cell membrane; peripheral membrane protein (by similarity).	periplasm
P0ACE3	HHA_ECOLI		cytoplasm
P0ACE4	HHA_ECOL6		cytoplasm
P0ACE5	HHA_ECO57		cytoplasm
P0ACE6	HHA_SHIFL		cytoplasm
P0ACE7	YCFF_ECOLI		cytoplasm
P0ACE8	YCFF_ECO57		cytoplasm
P0ACE9	YCFF_SHIFL		cytoplasm
P0ACF0	DBHA_ECOLI		cytoplasm
P0ACF1	DBHA_ECOL6		cytoplasm
P0ACF2	DBHA_ECO57		cytoplasm
P0ACF3	DBHA_SHIFL		cytoplasm
P0ACF4	DBHB_ECOLI		cytoplasm
P0ACF5	DBHB_ECOL6		cytoplasm
P0ACF6	DBHB_ECO57		cytoplasm
P0ACF7	DBHB_SHIFL		cytoplasm
P0ACF8	HNS_ECOLI		cytoplasm

P0ACF9	HNS_ECOL6		cytoplasm
P0ACG0	HNS_ECO57		cytoplasm
P0ACG1	STPA_ECOLI		cytoplasm
P0ACG2	STPA_ECO57		cytoplasm
P0ACG3	STPA_SHIFL		cytoplasm
P0ACG5	HOKC_SHIFL	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0ACG6	HOKD_ECOLI		inner membrane
P0ACG7	HOKD_ECO57		inner membrane
P0ACG8	HSLR_ECOLI		cytoplasm
P0ACG9	HSLR_ECOL6		cytoplasm
P0ACH0	HSLR_ECO57		cytoplasm
P0ACH1	SFSB_ECOLI		cytoplasm
P0ACH2	SFSB_ECOL6		cytoplasm
P0ACH3	SFSB_ECO57		cytoplasm
P0ACH4	SFSB_SHIFL		cytoplasm
P0ACH5	MARA_ECOLI		cytoplasm
P0ACH6	MARA_ECO57		cytoplasm
P0ACH7	MARA_SHIFL		cytoplasm
P0ACH8	MELR_ECOLI		cytoplasm
P0ACH9	MELR_ECOL6		cytoplasm
P0ACI0	ROB_ECOLI		cytoplasm
P0ACI1	ROB_ECO57		cytoplasm
P0ACI2	ROB_SHIFL		cytoplasm
P0ACI3	XYLR_ECOLI		cytoplasm
P0ACI4	XYLR_ECOL6		cytoplasm
P0ACI5	XYLR_ECO57		cytoplasm
P0ACI6	ASNC_ECOLI		cytoplasm
P0ACI7	ASNC_ECOL6		cytoplasm
P0ACI8	ASNC_ECO57		cytoplasm
P0ACI9	ASNC_SHIFL		cytoplasm
P0ACJ0	LRP_ECOLI		cytoplasm
P0ACJ1	LRP_ECOL6		cytoplasm
P0ACJ2	LRP_ECO57		cytoplasm
P0ACJ3	LRP_SHIFL		cytoplasm
P0ACJ4	LRP_ENTAE		cytoplasm
P0ACJ5	YBAO_ECOLI		cytoplasm
P0ACJ6	YBAO_ECOL6		cytoplasm
P0ACJ7	YBAO_ECO57		cytoplasm
P0ACJ8	CRP_ECOLI		cytoplasm
P0ACJ9	CRP_ECOL6		cytoplasm
P0ACK0	CRP_ECO57		cytoplasm
P0ACK1	CRP_SHIFL		cytoplasm
P0ACK2	AGAR_ECOLI		cytoplasm
P0ACK3	AGAR_ECOL6		cytoplasm
P0ACK4	AGAR_ECO57		cytoplasm
P0ACK5	DEOR_ECOLI		cytoplasm
P0ACK6	DEOR_ECOL6		cytoplasm
P0ACK7	DEOR_ECO57		cytoplasm
P0ACK8	FUCR_ECOLI		cytoplasm
P0ACK9	FUCR_SHIFL		cytoplasm
P0ACL0	GLPR_ECOLI		cytoplasm
P0ACL1	GLPR_SHIFL		cytoplasm
P0ACL2	EXUR_ECOLI		cytoplasm
P0ACL3	EXUR_ECO57		cytoplasm

P0ACL4	EXUR_SHIFL		cytoplasm
P0ACL5	GLCC_ECOLI		cytoplasm
P0ACL6	GLCC_ECOL6		cytoplasm
P0ACL7	LLDR_ECOLI		cytoplasm
P0ACL8	LLDR_SHIFL		cytoplasm
P0ACL9	PDHR_ECOLI		cytoplasm
P0ACM0	PDHR_ECOL6		cytoplasm
P0ACM1	PDHR_ECO57		cytoplasm
P0ACM2	YDFH_ECOLI		cytoplasm
P0ACM3	YDFH_ECO57		cytoplasm
P0ACM4	YDFH_SHIFL		cytoplasm
P0ACM5	YEGW_ECOLI		cytoplasm
P0ACM6	YEGW_ECOL6		cytoplasm
P0ACM7	YEGW_ECO57		cytoplasm
P0ACM8	YEGW_SHIFL		cytoplasm
P0ACM9	YIHL_ECOLI		cytoplasm
P0ACN0	YIHL_ECO57		cytoplasm
P0ACN1	YIHL_SHIFL		cytoplasm
P0ACN2	YTFH_ECOLI		cytoplasm
P0ACN3	YTFH_SHIFL		cytoplasm
P0ACN4	ALLR_ECOLI		cytoplasm
P0ACN5	ALLR_ECOL6		cytoplasm
P0ACN6	ALLR_ECO57		cytoplasm
P0ACN7	CYTR_ECOLI		cytoplasm
P0ACN8	CYTR_ECOL6		cytoplasm
P0ACN9	CYTR_ECO57		cytoplasm
P0ACP0	CYTR_SHIFL		cytoplasm
P0ACP1	FRUR_ECOLI		cytoplasm
P0ACP2	FRUR_ECOL6		cytoplasm
P0ACP3	FRUR_ECO57		cytoplasm
P0ACP4	FRUR_SHIFL		cytoplasm
P0ACP5	GNTR_ECOLI		cytoplasm
P0ACP6	GNTR_ECOL6		cytoplasm
P0ACP7	PURR_ECOLI		cytoplasm
P0ACP8	PURR_ECO57		cytoplasm
P0ACP9	PURR_SHIFL		cytoplasm
P0ACQ0	RBSR_ECOLI		cytoplasm
P0ACQ1	RBSR_ECOL6		cytoplasm
P0ACQ2	RBSR_ECO57		cytoplasm
P0ACQ3	RBSR_SHIFL		cytoplasm
P0ACQ4	OXYR_ECOLI		cytoplasm
P0ACQ5	OXYR_ECOL6		cytoplasm
P0ACQ6	OXYR_ECO57		cytoplasm
P0ACQ7	TDCA_ECOLI		cytoplasm
P0ACQ8	TDCA_ECO57		cytoplasm
P0ACQ9	TDCA_SHIFL		cytoplasm
P0ACR0	YBBS_ECOLI		cytoplasm
P0ACR1	YBBS_ECO57		cytoplasm
P0ACR2	YDHB_ECOLI		cytoplasm
P0ACR3	YDHB_ECO57		cytoplasm
P0ACR4	YEIE_ECOLI		cytoplasm
P0ACR5	YEIE_ECOL6		cytoplasm
P0ACR6	YEIE_ECO57		cytoplasm
P0ACR7	YFER_ECOLI		cytoplasm
P0ACR8	YFER_SHIFL		cytoplasm
P0ACR9	MPRA_ECOLI		cytoplasm

P0ACS0	MPRA_ECOL6		cytoplasm
P0ACS1	MPRA_ECO57		cytoplasm
P0ACS2	SOXR_ECOLI		cytoplasm
P0ACS3	SOXR_ECO57		cytoplasm
P0ACS4	SOXR_SHIFL		cytoplasm
P0ACS5	ZNTR_ECOLI		cytoplasm
P0ACS6	ZNTR_ECO57		cytoplasm
P0ACS7	RPIR_ECOLI		cytoplasm
P0ACS8	RPIR_ECOL6		cytoplasm
P0ACS9	ACRR_ECOLI		cytoplasm
P0ACT0	ACRR_ECOL6		cytoplasm
P0ACT1	ACRR_SHIFL		cytoplasm
P0ACT2	ENVR_ECOLI		cytoplasm
P0ACT3	ENVR_ECO57		cytoplasm
P0ACT4	TETR4_ECOLI		cytoplasm
P0ACT5	TETR4_SALOR		cytoplasm
P0ACT6	UIDR_ECOLI		cytoplasm
P0ACT7	UIDR_ECOL6		cytoplasm
P0ACT8	UIDR_ECO57		cytoplasm
P0ACT9	UIDR_SHIFL		cytoplasm
P0ACU0	YBIH_ECOLI		cytoplasm
P0ACU1	YBIH_SHIFL		cytoplasm
P0ACU2	RUTR_ECOLI		cytoplasm
P0ACU3	RUTR_ECOL6		cytoplasm
P0ACU4	RUTR_SHIFL		cytoplasm
P0ACU5	YIJC_ECOLI		cytoplasm
P0ACU6	YIJC_SHIFL		cytoplasm
P0ACU7	YJDC_ECOLI		cytoplasm
P0ACU8	YJDC_ECOL6		cytoplasm
P0ACU9	YJDC_SHIFL		cytoplasm
P0ACV0	HTRB_ECOLI	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P0ACV1	HTRB_ECO57	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P0ACV2	DDG_ECOLI	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P0ACV3	DDG_SHIFL	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P0ACV5	YCIS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ACV6	YCJI_ECOLI		cytoplasm
P0ACV7	YCJI_ECO57		cytoplasm
P0ACV8	YMJA_ECOLI		cytoplasm
P0ACV9	YMJA_SHIFL		cytoplasm
P0ACW0	YDAF_ECOLI		outer membrane
P0ACW1	YDAF_ECO57		outer membrane
P0ACW2	YDBJ_ECOLI		outer membrane
P0ACW3	YDBJ_ECOL6		outer membrane
P0ACW4	YDCA_ECOLI		extracellular
P0ACW5	YDCA_SHIFL		extracellular
P0ACW6	YDCH_ECOLI		cytoplasm

P0ACW7	YDCH_SHIFL		cytoplasm
P0ACW8	YDFA_ECOLI		extracellular
P0ACW9	YDFA_ECO57		outer membrane
P0ACX1	YDGC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ACX2	YDGC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ACX3	YDHR_ECOLI		cytoplasm
P0ACX4	YDHR_SHIFL		cytoplasm
P0ACX5	YDHZ_ECOLI		cytoplasm
P0ACX6	YDHZ_ECOL6		cytoplasm
P0ACX7	YDHZ_ECO57		cytoplasm
P0ACX8	YDHZ_SHIFL		cytoplasm
P0ACX9	YDIE_ECOLI		cytoplasm
P0ACY0	YDIE_ECO57		cytoplasm
P0ACY1	YDJA_ECOLI		cytoplasm
P0ACY2	YDJA_ECO57		cytoplasm
P0ACY3	YEAG_ECOLI		cytoplasm
P0ACY4	YEAG_ECOL6		cytoplasm
P0ACY5	YEAG_ECO57		cytoplasm
P0ACY6	YEAL_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ACY7	YEAL_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ACY8	YEAL_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ACY9	YEBG_ECOLI		cytoplasm
P0ACZ0	YEBG_ECOL6		cytoplasm
P0ACZ1	YEBG_ECO57		cytoplasm
P0ACZ2	ETP_ECOLI		extracellular
P0ACZ3	ETP_ECO57		extracellular
P0ACZ4	EVGA_ECOLI	cytoplasm (probable).	cytoplasm
P0ACZ5	EVGA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0ACZ6	EVGA_ECO57	cytoplasm (by similarity).	cytoplasm
P0ACZ7	EVGA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0ACZ8	CUSR_ECOLI	cytoplasm (probable).	cytoplasm
P0ACZ9	CUSR_ECOL6	cytoplasm (by similarity).	cytoplasm
P0AD00	CUSR_ECO57	cytoplasm (by similarity).	cytoplasm
P0AD01	DCUR_ECOLI	cytoplasm (probable).	cytoplasm
P0AD02	DCUR_ECO57	cytoplasm (probable).	cytoplasm
P0AD04	YEBS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD05	YECA_ECOLI		cytoplasm
P0AD06	YECA_SHIFL		cytoplasm
P0AD07	YECF_ECOLI		flagellar
P0AD08	YECF_ECOL6		flagellar
P0AD09	YECF_SHIFL		cytoplasm
P0AD10	YECJ_ECOLI		inner membrane
P0AD11	YECJ_ECOL6		inner membrane
P0AD12	YEEZ_ECOLI		cytoplasm

P0AD13	YEEZ_ECO57		cytoplasm
P0AD15	YEHU_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD16	YEHU_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD18	YOHC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD20	YOHK_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD21	YEJG_ECOLI		cytoplasm
P0AD22	YEJG_ECOL6		cytoplasm
P0AD23	YEJG_SHIFL		cytoplasm
P0AD24	YEJL_ECOLI		inner membrane
P0AD25	YEJL_ECOL6		inner membrane
P0AD26	YEJL_ECO57		inner membrane
P0AD28	YEJM_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD29	YEJM_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD31	YFCA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD32	YFCA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD33	YFCZ_ECOLI		cytoplasm
P0AD34	YFCZ_ECOL6		cytoplasm
P0AD35	YFDO_ECOLI		nucleoid
P0AD36	YFDO_ECOL6		nucleoid
P0AD37	YFEC_ECOLI		cytoplasm
P0AD38	YFEC_ECOL6		cytoplasm
P0AD39	YFEC_SHIFL		cytoplasm
P0AD40	YPEB_ECOLI		cytoplasm
P0AD41	YPEB_ECO57		cytoplasm
P0AD42	YFHB_ECOLI		inner membrane
P0AD43	YFHB_SHIFL		inner membrane
P0AD44	YFHG_ECOLI		cytoplasm
P0AD45	YFHG_ECO57		cytoplasm
P0AD46	YFHG_SHIFL		cytoplasm
P0AD48	YPHA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AD49	RAIA_ECOLI		cytoplasm
P0AD50	RAIA_ECOL6		cytoplasm
P0AD51	RAIA_ECO57		cytoplasm
P0AD52	RAIA_SHIFL		cytoplasm
P0AD53	YGAC_ECOLI		cytoplasm
P0AD54	YGAC_ECOL6		cytoplasm
P0AD55	YGAC_ECO57		cytoplasm
P0AD56	YGAC_SHIFL		cytoplasm

P0AD57	ISPB_ECOLI		cytoplasm
P0AD58	ISPB_SHIFL		cytoplasm
P0AD60	IVY_ECO57	periplasm (by similarity).	periplasm
P0AD61	KPYK1_ECOLI		cytoplasm
P0AD62	KPYK1_ECO57		cytoplasm
P0AD63	BLA1_ECOLI		extracellular
P0AD64	BLA1_KLEPN		extracellular
P0AD66	PBP2_ECOL6	bacterial cell inner membrane (by similarity).	inner membrane
P0AD67	PBP2_ECO57	bacterial cell inner membrane (by similarity).	inner membrane
P0AD70	AMPH_ECOLI		periplasm
P0AD71	AMPH_ECO57		periplasm
P0AD97	LIVJ_ECOL6	periplasm (by similarity).	periplasm
P0AD98	LIVJ_ECO57	periplasm (by similarity).	periplasm
P0ADA0	BRNQ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADA2	TESA_ECOL6	periplasm (by similarity).	extracellular
P0ADA3	NLPD_ECOLI	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P0ADA4	NLPD_SHIFL	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P0ADA5	YAJG_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P0ADA6	YAJG_ECOL6	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P0ADA7	OSMB_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P0ADA8	OSMB_ECOL6	bacterial cell membrane; lipid-anchor (by similarity).	inner membrane
P0ADA9	OSMB_ECO57	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P0ADB0	OSMB_SHIFL	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P0ADB1	OSME_ECOLI	bacterial cell membrane; lipid-anchor (probable).	outer membrane
P0ADB2	OSME_ECO57	bacterial cell membrane; lipid-anchor (potential).	outer membrane
P0ADB3	OSME_SHIFL	bacterial cell membrane; lipid-anchor (potential).	outer membrane
P0ADC1	RLPB_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P0ADC2	RLPB_ECOL6	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0ADC4	LOLC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADC5	LOLC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADC7	YJGQ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADC8	YJIX_ECOLI		inner membrane
P0ADC9	YJIX_ECOL6		inner membrane

P0ADD0	YJIX_ECO57		inner membrane
P0ADD1	YJIX_SHIFL		inner membrane
P0ADD2	YJJB_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ADD3	YJJB_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ADD4	YJJB_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ADD6	YJJP_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADD7	YJJQ_ECOLI		cytoplasm
P0ADD8	YJJQ_ECO57		cytoplasm
P0ADE2	YTFK_ECOLI		cytoplasm
P0ADE3	YTFK_ECOL6		cytoplasm
P0ADE4	YTFM_ECOLI		outer membrane
P0ADE5	YTFM_ECO57		outer membrane
P0ADE6	YGAU_ECOLI		cytoplasm
P0ADE7	YGAU_ECOL6		cytoplasm
P0ADE8	YGFZ_ECOLI		cytoplasm
P0ADE9	YGFZ_SHIFL		cytoplasm
P0ADF6	EDD_ECOLI		cytoplasm
P0ADF7	EDD_ECO57		cytoplasm
P0ADF8	ILVN_ECOLI		cytoplasm
P0ADF9	ILVN_ECOL6		cytoplasm
P0ADG0	ILVN_ECO57		cytoplasm
P0ADG1	ILVM_ECOLI		cytoplasm
P0ADG2	ILVM_ECOL6		cytoplasm
P0ADG3	ILVM_SHIFL		cytoplasm
P0ADG4	SUHB_ECOLI		cytoplasm
P0ADG5	SUHB_ECOL6		cytoplasm
P0ADG6	SUHB_ECO57		cytoplasm
P0ADG7	IMDH_ECOLI		cytoplasm
P0ADG8	IMDH_ECO57		cytoplasm
P0ADG9	IMDH_SHIFL		cytoplasm
P0ADH0	INSA_ECOLI		cytoplasm
P0ADH1	INSA_SALTI		cytoplasm
P0ADH2	INSA_SHISO		cytoplasm
P0ADH3	INSE_ECOLI		cytoplasm
P0ADH4	INSE_ECO11		cytoplasm
P0ADH5	FIMB_ECOLI		cytoplasm
P0ADH6	FIMB_ECO57		cytoplasm
P0ADH7	FIME_ECOLI		cytoplasm
P0ADH8	FIME_ECOL6		cytoplasm
P0ADH9	FIME_SHIFL		cytoplasm
P0ADI0	PINR_ECOLI		cytoplasm
P0ADI1	PINR_ECOL6		cytoplasm
P0ADI2	TNR3_ECOLI		cytoplasm
P0ADI3	TNR3_KLEPN		cytoplasm
P0ADI4	ENTB_ECOLI		cytoplasm
P0ADI5	ENTB_ECO57		cytoplasm
P0ADI6	ENTB_SHIFL		cytoplasm
P0ADI7	YECD_ECOLI		cytoplasm

P0ADI8	YECD_ECO57		cytoplasm
P0ADI9	YHHN_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ADJ0	YHHN_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ADJ1	YHHN_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ADJ2	YHHN_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ADJ3	YHJR_ECOLI		cytoplasm
P0ADJ4	YHJR_SHIFL		cytoplasm
P0ADJ5	YHJT_ECOLI		cytoplasm
P0ADJ6	YHJT_ECOL6		cytoplasm
P0ADJ7	YHJT_ECO57		cytoplasm
P0ADJ9	YIAA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADK0	YIAF_ECOLI		periplasm
P0ADK1	YIAF_ECOL6		periplasm
P0ADK2	YIAF_ECO57		periplasm
P0ADK3	YIAF_SHIFL		periplasm
P0ADK5	YIAW_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADK6	YIBA_ECOLI		extracellular
P0ADK7	YIBA_ECO57		extracellular
P0ADK8	YIBL_ECOLI		cytoplasm
P0ADK9	YIBL_ECOL6		cytoplasm
P0ADL0	YIBL_ECO57		cytoplasm
P0ADL2	YICM_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADL3	YICN_ECOLI		inner membrane
P0ADL4	YICN_ECOL6		inner membrane
P0ADL5	YICN_ECO57		inner membrane
P0ADL7	YIDG_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADL8	YIDG_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADL9	YIDG_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADM1	YIDH_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADM2	YIDH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0ADM3	YIDH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADM4	YIDQ_ECOLI		cytoplasm
P0ADM5	YIDQ_SHIFL		cytoplasm
P0ADM6	YIDX_ECOLI		inner membrane
P0ADM7	YIDX_SHIFL		inner membrane
P0ADM8	YIEE_ECOLI		cytoplasm
P0ADM9	YIEE_ECOL6		cytoplasm
P0ADN0	VIAA_ECOLI		nucleoid
P0ADN1	VIAA_ECO57		nucleoid
P0ADN2	YIFE_ECOLI		cytoplasm
P0ADN3	YIFE_ECOL6		cytoplasm
P0ADN4	YIFE_ECO57		cytoplasm
P0ADN5	YIFE_SHIFL		cytoplasm
P0ADN6	YIFL_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0ADN7	YIFL_ECOL6	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0ADN8	YIFL_ECO57	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0ADN9	YIFL_SHIFL	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0ADP0	YIGB_ECOLI		cytoplasm
P0ADP1	YIGB_SHIFL		cytoplasm
P0ADP2	YIGI_ECOLI		cytoplasm
P0ADP3	YIGI_ECOL6		cytoplasm
P0ADP4	YIGI_SHIFL		cytoplasm
P0ADP5	YIGM_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ADP6	YIGM_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0ADP7	YIGP_ECOLI		cytoplasm
P0ADP8	YIGP_SHIFL		cytoplasm
P0ADP9	YIHD_ECOLI		cytoplasm
P0ADQ0	YIHD_ECOL6		cytoplasm
P0ADQ1	YIHD_ECO57		cytoplasm
P0ADQ2	YIID_ECOLI		cytoplasm
P0ADQ3	YIID_ECOL6		cytoplasm
P0ADQ4	YIID_ECO57		cytoplasm
P0ADQ5	YIIE_ECOLI		cytoplasm
P0ADQ6	YIIE_SHIFL		cytoplasm
P0ADQ7	YGAM_ECOLI		cytoplasm
P0ADQ8	YGAM_ECOL6		cytoplasm
P0ADQ9	YGAM_ECO57		cytoplasm
P0ADR1	YQAA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADR3	YGDD_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADR4	YGDD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0ADR5	YGDD_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0ADR6	MTFA_ECOLI		cytoplasm
P0ADR7	MTFA_ECO57		cytoplasm
P0ADR8	YGDH_ECOLI		cytoplasm
P0ADR9	YGDH_ECOL6		cytoplasm
P0ADS0	YGDH_ECO57		cytoplasm
P0ADS1	YGDH_SHIFL		cytoplasm
P0ADS2	YGFE_ECOLI		nucleoid
P0ADS3	YGFE_ECOL6		nucleoid
P0ADS4	YGFE_ECO57		nucleoid
P0ADS5	YGFE_SHIFL		nucleoid
P0ADS6	YGGE_ECOLI		cytoplasm
P0ADS7	YGGE_ECO57		cytoplasm
P0ADS8	YGGE_SHIFL		cytoplasm
P0ADS9	YGGN_ECOLI		cytoplasm
P0ADT0	YGGN_ECOL6		cytoplasm
P0ADT1	YGGN_ECO57		cytoplasm
P0ADT2	YGIB_ECOLI		extracellular
P0ADT3	YGIB_ECOL6		extracellular
P0ADT4	YGIB_SHIFL		extracellular
P0ADT5	YGIC_ECOLI		cytoplasm
P0ADT6	YGIC_ECOL6		cytoplasm
P0ADT7	YGIC_ECO57		cytoplasm
P0ADT8	YGIM_ECOLI		inner membrane
P0ADT9	YGIM_ECOL6		inner membrane
P0ADU0	YGIM_ECO57		inner membrane
P0ADU1	YGIM_SHIFL		inner membrane
P0ADU2	YGIN_ECOLI		cytoplasm
P0ADU3	YGIN_ECO57		cytoplasm
P0ADU4	YGIN_SHIFL		cytoplasm
P0ADU5	YGIW_ECOLI	periplasm (potential).	periplasm
P0ADU6	YGIW_ECO57	periplasm (potential).	periplasm
P0ADU7	YQIB_ECOLI		cytoplasm
P0ADU8	YQIB_ECOL6		cytoplasm
P0ADU9	YQIB_ECO57		cytoplasm
P0ADV0	YQIB_SHIFL		cytoplasm
P0ADV1	YHBN_ECOLI	periplasm (potential).	periplasm
P0ADV2	YHBN_ECOL6	periplasm (potential).	periplasm
P0ADV3	YHBN_ECO57	periplasm (potential).	periplasm
P0ADV4	YHBN_SHIFL	periplasm (potential).	periplasm
P0ADV5	YHBW_ECOLI		cytoplasm
P0ADV6	YHBW_ECOL6		cytoplasm
P0ADV7	YRBC_ECOLI		inner membrane
P0ADV8	YRBC_SHIFL		inner membrane
P0ADV9	YRBK_ECOLI		inner membrane
P0ADW0	YRBK_ECOL6		inner membrane
P0ADW1	YRBK_ECO57		inner membrane
P0ADW2	YRBK_SHIFL		inner membrane
P0ADW4	YHCB_ECOL6	bacterial cell inner membrane (by similarity).	inner membrane
P0ADW5	YHCB_ECO57	bacterial cell inner membrane (by similarity).	inner membrane
P0ADW6	YHCC_ECOLI		cytoplasm
P0ADW7	YHCC_ECO57		cytoplasm

P0ADW8	YHEV_ECOLI		cytoplasm
P0ADW9	YHEV_ECOL6		cytoplasm
P0ADX0	YHEV_ECO57		cytoplasm
P0ADX1	YHFA_ECOLI		cytoplasm
P0ADX2	YHFA_ECOL6		cytoplasm
P0ADX3	YHFA_ECO57		cytoplasm
P0ADX4	YHFA_SHIFL		cytoplasm
P0ADX5	YHFG_ECOLI		nucleoid
P0ADX6	YHFG_ECO57		nucleoid
P0ADX7	YHHA_ECOLI		inner membrane
P0ADX8	YHHA_SHIFL		inner membrane
P0ADX9	YHHF_ECOLI		cytoplasm
P0ADY0	YHHF_ECOL6		cytoplasm
P0ADY2	PPID_ECOL6	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	periplasm
P0ADY3	RL14_ECOLI		cytoplasm
P0ADY4	RL14_ECOL6		cytoplasm
P0ADY5	RL14_ECO57		cytoplasm
P0ADY6	RL14_SHIFL		cytoplasm
P0ADY7	RL16_ECOLI		cytoplasm
P0ADY8	RL16_ECOL6		cytoplasm
P0ADY9	RL16_ECO57		cytoplasm
P0ADZ0	RL23_ECOLI		cytoplasm
P0ADZ1	RL23_ECOL6		cytoplasm
P0ADZ2	RL23_ECO57		cytoplasm
P0ADZ3	RL23_SHIFL		cytoplasm
P0ADZ4	RS15_ECOLI		cytoplasm
P0ADZ5	RS15_ECOL6		cytoplasm
P0ADZ6	RS15_SHIFL		cytoplasm
P0ADZ8	YAJC_ECOL6	bacterial cell inner membrane (by similarity).	inner membrane
P0ADZ9	YAJC_ECO57	bacterial cell inner membrane (by similarity).	inner membrane
P0AE00	YAJC_SHIFL	bacterial cell inner membrane (by similarity).	inner membrane
P0AE01	YFHQ_ECOLI		cytoplasm
P0AE02	YFHQ_ECOL6		cytoplasm
P0AE03	YFHQ_SHIFL		cytoplasm
P0AE04	AADB_ECOLI		cytoplasm
P0AE05	AADB1_KLEPN		cytoplasm
P0AE07	ACRA_ECO57	bacterial cell inner membrane; lipid-anchor (by similarity).	inner membrane
P0AE08	AHPC_ECOLI		cytoplasm
P0AE09	AHPC_ECOL6		cytoplasm
P0AE10	AHPC_ECO57		cytoplasm
P0AE11	AHPC_SHIFL		cytoplasm
P0AE12	AMN_ECOLI		cytoplasm
P0AE13	AMN_ECO57		cytoplasm
P0AE15	AMPE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE17	AMPG_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE18	AMPM_ECOLI		cytoplasm

P0AE19	AMPM_ECOL6		cytoplasm
P0AE20	AMPM_ECO57		cytoplasm
P0AE21	AMPM_SHIFL		cytoplasm
P0AE22	APHA_ECOLI	periplasm (potential).	periplasm
P0AE23	APHA_SHIFL	periplasm (potential).	periplasm
P0AE25	ARAE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE27	ARAH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE28	AROM_ECOLI		cytoplasm
P0AE29	AROM_SHIFL		cytoplasm
P0AE32	ARTM_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE33	ARTM_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE35	ARTQ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE36	ARTQ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE37	ASTA_ECOLI		cytoplasm
P0AE38	ASTA_ECO57		cytoplasm
P0AE39	YPDB_ECOLI		cytoplasm
P0AE40	YPDB_ECO57		cytoplasm
P0AE41	YPDB_SHIFL		cytoplasm
P0AE42	YQAE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AE43	YQAE_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AE44	YQAE_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AE46	YTFL_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE47	YTFL_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE48	YTFP_ECOLI		inner membrane
P0AE49	YTFP_ECOL6		inner membrane
P0AE50	YTFP_ECO57		inner membrane
P0AE51	YTFP_SHIFL		inner membrane
P0AE52	BCP_ECOLI		cytoplasm
P0AE53	BCP_ECOL6		cytoplasm
P0AE54	BCP_ECO57		cytoplasm
P0AE55	BCP_SHIFL		cytoplasm
P0AE56	BFD_ECOLI		cytoplasm
P0AE57	BFD_ECOL6		cytoplasm
P0AE58	CAIF_ECOLI		cytoplasm

P0AE59	CAIF_SHIFL		cytoplasm
P0AE60	CEDA_ECOLI		inner membrane
P0AE61	CEDA_ECO57		inner membrane
P0AE62	CEDA_SHIFL		inner membrane
P0AE63	CHAB_ECOLI		cytoplasm
P0AE64	CHAB_ECOL6		cytoplasm
P0AE65	CHAB_ECO57		cytoplasm
P0AE66	CHAB_SHIFL		cytoplasm
P0AE68	CHEY_ECO57	cytoplasm (by similarity).	cytoplasm
P0AE69	CHEY_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AE70	CHPA_ECOLI		cytoplasm
P0AE71	CHPA_ECO57		cytoplasm
P0AE72	CHPR_ECOLI		cytoplasm
P0AE73	CHPR_ECO57		cytoplasm
P0AE75	CITT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE76	COBU_ECOLI		cytoplasm
P0AE77	COBU_ECO57		cytoplasm
P0AE78	CORC_ECOLI		inner membrane
P0AE79	CORC_ECOL6		inner membrane
P0AE80	CORC_ECO57		inner membrane
P0AE81	CORC_SHIFL		inner membrane
P0AE83	CPXA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE84	CPXA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AE86	CPXP_ECOL6	periplasm (by similarity).	periplasm
P0AE88	CPXR_ECOLI	cytoplasm (probable).	cytoplasm
P0AE89	CPXR_ECO57	cytoplasm (probable).	cytoplasm
P0AE90	CPXR_SHIFL	cytoplasm (probable).	cytoplasm
P0AE91	CREA_ECOLI		cytoplasm
P0AE92	CREA_ECOL6		cytoplasm
P0AE93	CREA_ECO57		cytoplasm
P0AE94	CREA_SHIFL		cytoplasm
P0AE95	CSGE_ECOLI		cytoplasm
P0AE96	CSGE_ECOL6		cytoplasm
P0AE97	CSGE_ECO57		cytoplasm
P0AE98	CSGF_ECOLI		extracellular
P0AE99	CSGF_ECOL6		extracellular
P0AEA0	CSGF_ECO57		extracellular
P0AEA1	CSGF_SHIFL		extracellular
P0AEA2	CSGG_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AEA3	CSGG_ECOL6	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AEA4	CSGG_ECO57	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AEA6	CYOE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEA7	CYOE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AEA8	CYSG_ECOLI		cytoplasm
P0AEA9	CYSG_ECO57		cytoplasm
P0AEB1	CYSW_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEB6	YNAI_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEB7	YOAB_ECOLI		periplasm
P0AEB8	YOAB_ECOL6		periplasm
P0AEB9	YOAB_SHIFL		periplasm
P0AEC1	YOAE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEC2	YOAE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEC4	ARCB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEC6	BARA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEC7	BARA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEC9	DCUS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AED1	USPA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0AED2	USPA_ECO57	cytoplasm (by similarity).	cytoplasm
P0AED3	USPA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AED4	USPA_SHISO	cytoplasm (by similarity).	cytoplasm
P0AED5	UVRY_ECOLI	cytoplasm (probable).	cytoplasm
P0AED6	UVRY_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AED7	DAPE_ECOLI		cytoplasm
P0AED8	DAPE_SHIFL		cytoplasm
P0AED9	DCM_ECOLI		cytoplasm
P0AEE0	DCM_ECO57		cytoplasm
P0AEE2	DCRB_SHIFL	periplasm (by similarity).	periplasm
P0AEE3	DEGS_ECOLI	periplasm (potential).	periplasm
P0AEE4	DEGS_ECO57	periplasm (potential).	periplasm
P0AEE6	DGAL_ECOL6	periplasm (by similarity).	periplasm
P0AEE7	DGAL_SHIFL	periplasm (by similarity).	periplasm
P0AEE8	DMA_ECOLI		cytoplasm
P0AEE9	DMA_ECO57		cytoplasm
P0AEF0	DNAC_ECOLI		cytoplasm
P0AEF1	DNAC_ECOL6		cytoplasm
P0AEF2	DNAC_ECO57		cytoplasm
P0AEF3	DNAC_SHIFL		cytoplasm
P0AEF4	DPIA_ECOLI	cytoplasm (potential).	cytoplasm
P0AEF5	DPIA_ECOL6	cytoplasm (potential).	cytoplasm
P0AEF6	DPIA_ECO57	cytoplasm (potential).	cytoplasm
P0AEF7	DPIA_SHIFL	cytoplasm (potential).	cytoplasm

P0AEF9	DPPB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEG0	DPPB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEG2	DPPC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEG3	DPPC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEG5	DSBA_ECO57	periplasm (by similarity).	periplasm
P0AEG7	DSBC_ECO57	periplasm (by similarity).	periplasm
P0AEG8	DSRB_ECOLI		inner membrane
P0AEG9	DSRB_ECOL6		inner membrane
P0AEH0	DSRB_ECO57		inner membrane
P0AEH2	ECFE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P0AEH3	ELAA_ECOLI		cytoplasm
P0AEH4	ELAA_SHIFL		cytoplasm
P0AEH5	ELAB_ECOLI		cytoplasm
P0AEH6	ELAB_ECOL6		cytoplasm
P0AEH7	ELAB_ECO57		cytoplasm
P0AEH9	YJIG_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEI0	YJIG_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEI1	YLEA_ECOLI		cytoplasm
P0AEI2	YLEA_ECOL6		cytoplasm
P0AEI3	YLEA_ECO57		cytoplasm
P0AEI4	YLIG_ECOLI		cytoplasm
P0AEI5	YLIG_ECO57		cytoplasm
P0AEI6	YMFB_ECOLI		extracellular
P0AEI7	YMFB_ECOL6		extracellular
P0AEI8	YMFB_ECO57		extracellular
P0AEI9	YMFB_SHIFL		extracellular
P0AEJ1	EMRB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEJ2	ENTC_ECOLI		cytoplasm
P0AEJ3	ENTC_ECO57		cytoplasm
P0AEJ5	ENVZ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEJ6	EUTB_ECOLI		cytoplasm
P0AEJ7	EUTB_ECO57		cytoplasm
P0AEJ8	EUTN_ECOLI		cytoplasm
P0AEJ9	EUTN_ECOL6		cytoplasm
P0AEK0	EXOX_ECOLI		cytoplasm
P0AEK1	EXOX_ECOL6		cytoplasm
P0AEK2	FABG_ECOLI		cytoplasm
P0AEK3	FABG_SHIFL		cytoplasm

P0AEK5	FABI_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AEK6	FABI_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AEK8	FDNI_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEK9	FDNI_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEL1	FDOI_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEL2	FDOI_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEL3	FEOA_ECOLI		cytoplasm
P0AEL4	FEOA_ECOL6		cytoplasm
P0AEL5	FEOA_ECO57		cytoplasm
P0AEL7	FEPB_ECOL6	periplasm (by similarity).	periplasm
P0AEL8	FIMZ_ECOLI	cytoplasm (probable).	cytoplasm
P0AEL9	FIMZ_ECO57	cytoplasm (probable).	cytoplasm
P0AEM0	FKBX_ECOLI		cytoplasm
P0AEM1	FKBX_ECOL6		cytoplasm
P0AEM2	FKBX_ECO57		cytoplasm
P0AEM3	FKBX_SHIFL		cytoplasm
P0AEM4	FLGM_ECOLI		cytoplasm
P0AEM5	FLGM_SHIFL		cytoplasm
P0AEM6	FLIA_ECOLI		cytoplasm
P0AEM7	FLIA_ECOL6		cytoplasm
P0AEM8	FLIA_ECO57		cytoplasm
P0AEN0	FLIY_ECOL6	periplasm (by similarity).	periplasm
P0AEN1	FRE_ECOLI		cytoplasm
P0AEN2	FRE_ECOL6		cytoplasm
P0AEN3	FRE_ECO57		cytoplasm
P0AEN5	FTSL_ECOL6	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0AEN6	FTSL_ECO57	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0AEN7	FTSL_SHIFL	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0AEN8	FUCU_ECOLI		cytoplasm
P0AEN9	FUCU_ECOL6		cytoplasm
P0AEP0	FUCU_ECO57		cytoplasm
P0AEP2	GALP_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEP3	GALU_ECOLI		cytoplasm
P0AEP4	GALU_ECOL6		cytoplasm
P0AEP5	GALU_ECO57		cytoplasm
P0AEP6	GALU_SHIFL		cytoplasm

P0AEP7	GCL_ECOLI		cytoplasm
P0AEP8	GCL_ECO57		cytoplasm
P0AEP9	GLCD_ECOLI		cytoplasm
P0AEQ0	GLCD_ECOL6		cytoplasm
P0AEQ1	GLCG_ECOLI		cytoplasm
P0AEQ2	GLCG_ECOL6		cytoplasm
P0AEQ4	GLNH_ECOL6	periplasm (by similarity).	periplasm
P0AEQ5	GLNH_ECO57	periplasm (by similarity).	periplasm
P0AEQ7	GLNP_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEQ8	GLNP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEQ9	GLNP_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AER1	GLPF_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AER2	GLPF_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AER4	GLTJ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AER6	GLTK_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AER7	GLTK_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AER9	GLTS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AES0	GSP_ECOLI		cytoplasm
P0AES1	GSP_SHIFL		cytoplasm
P0AES2	GUDH_ECOLI		cytoplasm
P0AES3	GUDH_ECO57		cytoplasm
P0AES4	GYRA_ECOLI		cytoplasm
P0AES5	GYRA_SHIFL		cytoplasm
P0AES6	GYRB_ECOLI		cytoplasm
P0AES7	GYRB_ECO57		cytoplasm
P0AES8	GYRB_SHIFL		cytoplasm
P0AES9	HDEA_ECOLI	periplasm (potential).	periplasm
P0AET0	HDEA_ECO57	periplasm (potential).	periplasm
P0AET1	HDEA_SHIFL	periplasm (potential).	periplasm
P0AET2	HDEB_ECOLI		inner membrane
P0AET3	HDEB_ECOL6		periplasm
P0AET4	HDEB_SHIFL		inner membrane
P0AET6	HDED_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AET7	HDED_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AET8	HDHA_ECOLI		cytoplasm
P0AET9	HDHA_ECO57		cytoplasm
P0AEU1	HISJ_ECOL6	periplasm (by similarity).	periplasm
P0AEU2	HISJ_ECO57	periplasm (by similarity).	periplasm
P0AEU4	HISM_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEU5	HISM_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEU6	HISM_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEU8	SKP_ECOL6	periplasm (by similarity).	periplasm
P0AEU9	SKP_ECO57	periplasm (by similarity).	periplasm
P0AEV0	SKP_SHIFL	periplasm (by similarity).	periplasm
P0AEV1	HNR_ECOLI		cytoplasm
P0AEV2	HNR_ECO57		cytoplasm
P0AEV3	HNR_SHIFL		cytoplasm
P0AEV4	HYCA_ECOLI		cytoplasm
P0AEV5	HYCA_ECO57		cytoplasm
P0AEV6	HYCA_SHIFL		cytoplasm
P0AEV7	HYCH_ECOLI		cytoplasm
P0AEV8	HYCH_ECO57		cytoplasm
P0AEV9	HYCI_ECOLI		extracellular
P0AEW0	HYCI_ECO57		extracellular
P0AEW2	HYFE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEW3	HYFE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEW4	ICC_ECOLI		extracellular
P0AEW5	ICC_ECO57		extracellular
P0AEW6	INGK_ECOLI		cytoplasm
P0AEW7	INGK_ECOL6		cytoplasm
P0AEW8	INGK_ECO57		cytoplasm
P0AEW9	K1PF_ECOLI		cytoplasm
P0AEX0	K1PF_ECOL6		cytoplasm
P0AEX1	K1PF_ECO57		cytoplasm
P0AEX2	K1PF_SHIFL		cytoplasm
P0AEX4	KGTP_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEX5	KPPR_ECOLI		cytoplasm
P0AEX6	KPPR_SHIFL		cytoplasm
P0AEX8	LIVH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEY0	MALE_ECO57	periplasm (by similarity).	periplasm
P0AEY2	MARC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEY3	MAZG_ECOLI		nucleoid
P0AEY4	MAZG_ECO57		nucleoid
P0AEY5	MDAB_ECOLI		cytoplasm

P0AEY6	MDAB_ECOL6		cytoplasm
P0AEY7	MDAB_ECO57		cytoplasm
P0AEY9	MDFA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEZ0	MDFA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AEZ1	METF_ECOLI		cytoplasm
P0AEZ2	METF_SHIFL		cytoplasm
P0AEZ4	MIND_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AEZ5	MIND_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AEZ6	MIND_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AEZ7	MLTD_ECOLI	bacterial cell membrane; lipid-anchor (probable).	extracellular
P0AEZ8	MLTD_ECOL6	bacterial cell membrane; lipid-anchor (probable).	extracellular
P0AEZ9	MOAB_ECOLI		cytoplasm
P0AF00	MOAB_ECO57		cytoplasm
P0AF02	MODB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AF03	MOG_ECOLI		cytoplasm
P0AF04	MOG_ECO57		cytoplasm
P0AF05	MOG_SHIFL		cytoplasm
P0AF07	MOTB_ECO57	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P0AF08	MRP_ECOLI		cytoplasm
P0AF09	MRP_ECOL6		cytoplasm
P0AF10	MTLR_ECOLI		cytoplasm
P0AF11	MTLR_ECOL6		cytoplasm
P0AF12	MTNN_ECOLI		cytoplasm
P0AF13	MTNN_ECOL6		cytoplasm
P0AF14	MTNN_ECO57		cytoplasm
P0AF15	MTNN_SHIFL		cytoplasm
P0AF17	MVIN_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AF18	NAGA_ECOLI		cytoplasm
P0AF19	NAGA_ECO57		cytoplasm
P0AF20	NAGC_ECOLI		cytoplasm
P0AF21	NAGC_ECOL6		cytoplasm
P0AF22	NAGC_ECO57		cytoplasm
P0AF23	NAGC_SHIFL		cytoplasm
P0AF24	NAGD_ECOLI		cytoplasm
P0AF25	NAGD_ECO57		cytoplasm
P0AF26	NARJ_ECOLI		cytoplasm
P0AF27	NARJ_ECO57		cytoplasm
P0AF28	NARL_ECOLI		cytoplasm

P0AF29	NARL_ECOL6		cytoplasm
P0AF30	NARL_ECO57		cytoplasm
P0AF31	NARL_SHIFL		cytoplasm
P0AF33	NARV_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AF34	YIIR_ECOLI		inner membrane
P0AF35	YIIR_ECO57		inner membrane
P0AF36	YIIU_ECOLI		nucleoid
P0AF37	YIIU_ECOL6		nucleoid
P0AF38	YIIU_ECO57		nucleoid
P0AF39	YIIU_SHIFL		nucleoid
P0AF41	YIJD_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AF42	YIJD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AF43	YJBB_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AF44	YJBB_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AF45	YJBE_ECOLI		extracellular
P0AF46	YJBE_ECOL6		extracellular
P0AF47	YJBE_ECO57		extracellular
P0AF48	YJBQ_ECOLI		inner membrane
P0AF49	YJBQ_ECO57		inner membrane
P0AF50	YJBR_ECOLI		cytoplasm
P0AF51	YJBR_ECO57		cytoplasm
P0AF53	YJCD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AF55	YJCH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AF56	YJCO_ECOLI		extracellular
P0AF57	YJCO_ECO57		extracellular
P0AF58	YJCO_SHIFL		extracellular
P0AF59	YJDI_ECOLI		cytoplasm
P0AF60	YJDI_ECO57		cytoplasm
P0AF61	YJDK_ECOLI		cytoplasm
P0AF62	YJDK_ECO57		cytoplasm
P0AF63	YJEB_ECOLI		cytoplasm
P0AF64	YJEB_ECOL6		cytoplasm
P0AF65	YJEB_ECO57		cytoplasm
P0AF66	YJEB_SHIFL		cytoplasm
P0AF67	YJEE_ECOLI		cytoplasm
P0AF68	YJEE_ECO57		cytoplasm
P0AF69	YJEE_SHIFL		cytoplasm
P0AF70	YJEI_ECOLI		periplasm
P0AF71	YJEI_ECOL6		periplasm
P0AF72	YJEI_SHIFL		periplasm

P0AF73	YJET_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AF74	YJET_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AF75	YJET_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AF76	YJFI_ECOLI		inner membrane
P0AF77	YJFI_ECO57		inner membrane
P0AF78	YJFJ_ECOLI		nucleoid
P0AF79	YJFJ_ECOL6		nucleoid
P0AF81	YJFL_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AF82	YJFN_ECOLI		inner membrane
P0AF83	YJFN_ECOL6		inner membrane
P0AF84	YJFN_ECO57		inner membrane
P0AF85	YJFN_SHIFL		inner membrane
P0AF86	YJFY_ECOLI		cytoplasm
P0AF87	YJFY_ECOL6		cytoplasm
P0AF88	YJFY_ECO57		cytoplasm
P0AF89	YJFY_SHIFL		cytoplasm
P0AF90	YJGD_ECOLI		cytoplasm
P0AF91	YJGD_ECO57		cytoplasm
P0AF92	YJGD_SHIFL		cytoplasm
P0AF93	YJGF_ECOLI		cytoplasm
P0AF94	YJGF_ECOL6		cytoplasm
P0AF95	YJGF_SHIFL		cytoplasm
P0AF96	YJGK_ECOLI		inner membrane
P0AF97	YJGK_SHIFL		inner membrane
P0AF99	YJGP_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFA0	YJGP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFA1	YJGP_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFA3	NARX_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFA4	NARX_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFA6	NFRB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFA8	NHAB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFB0	NIKC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AFB2	NLPI_ECOL6	bacterial cell membrane; lipid-anchor (by similarity).	inner membrane
P0AFB3	NLPI_ECO57	bacterial cell membrane; lipid-anchor (by similarity).	inner membrane
P0AFB4	NLPI_SHIFL	bacterial cell membrane; lipid-anchor (by similarity).	inner membrane
P0AFB5	NTRB_ECOLI		inner membrane
P0AFB6	NTRB_ECO57		inner membrane
P0AFB7	NTRB_SHIFL		inner membrane
P0AFB8	NTRC_ECOLI		cytoplasm
P0AFB9	NTRC_ECO57		cytoplasm
P0AFC0	NUDB_ECOLI		cytoplasm
P0AFC1	NUDB_ECO57		cytoplasm
P0AFC2	NUDB_SHIFL		cytoplasm
P0AFC4	NUOA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFC5	NUOA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFC6	NUOA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFC7	NUOB_ECOLI		cytoplasm
P0AFC8	NUOB_ECOL6		cytoplasm
P0AFC9	NUOB_ECO57		cytoplasm
P0AFD0	NUOB_SHIFL		cytoplasm
P0AFD1	NUOE_ECOLI		cytoplasm
P0AFD2	NUOE_ECOL6		cytoplasm
P0AFD3	NUOE_SHIFL		cytoplasm
P0AFD5	NUOH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFD6	NUOI_ECOLI		cytoplasm
P0AFD7	NUOI_ECOL6		cytoplasm
P0AFD8	NUOI_ECO57		cytoplasm
P0AFD9	NUOI_SHIFL		cytoplasm
P0AFE1	NUOJ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFE2	NUOJ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFE3	NUOJ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFE5	NUOK_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFE6	NUOK_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFE7	NUOK_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AFE9	NUOM_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFF1	NUON_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFF3	NUPC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFF5	NUPG_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFF6	NUSA_ECOLI		cytoplasm
P0AFF7	NUSA_ECOL6		cytoplasm
P0AFF8	NUSA_ECO57		cytoplasm
P0AFF9	NUSA_SHIFL		cytoplasm
P0AFG0	NUSG_ECOLI		cytoplasm
P0AFG1	NUSG_ECO57		cytoplasm
P0AFG2	NUSG_SHIFL		cytoplasm
P0AFG3	ODO1_ECOLI		cytoplasm
P0AFG4	ODO1_ECOL6		cytoplasm
P0AFG5	ODO1_ECO57		cytoplasm
P0AFG6	ODO2_ECOLI		inner membrane
P0AFG7	ODO2_ECO57		inner membrane
P0AFG8	ODP1_ECOLI		cytoplasm
P0AFG9	ODP1_ECO57		cytoplasm
P0AFH0	OGT_ECOLI		cytoplasm
P0AFH1	OGT_SHIFL		cytoplasm
P0AFH3	OPPB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFH4	OPPB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFH5	OPPB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFH7	OPPC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFH9	OSMY_ECOL6	periplasm (by similarity).	periplasm
P0AFI0	OXC_ECOLI		cytoplasm
P0AFI1	OXC_ECO57		cytoplasm
P0AFI3	PARC_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P0AFI4	PARC_SHIFL	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P0AFI6	PBP7_SHIFL	periplasm (by similarity).	cytoplasm
P0AFI7	PDXH_ECOLI		cytoplasm
P0AFI8	PDXH_SHIFL		cytoplasm
P0AFI9	PERM_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

P0AFJ0	PERM_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AFJ1	PHNA_ECOLI		cytoplasm
P0AFJ2	PHNA_ECOL6		cytoplasm
P0AFJ3	PHNA_ECO57		cytoplasm
P0AFJ4	PHNA_SHIFL		cytoplasm
P0AFJ6	PHOB_ECO57	cytoplasm (by similarity).	cytoplasm
P0AFJ8	PITA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFJ9	PITA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFK1	PMBA_ECO57	cytoplasm (by similarity).	cytoplasm
P0AFK3	PNUC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFK5	POTB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFK7	POTC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFK8	POTC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFL0	POTD_SHIFL	periplasm (by similarity).	periplasm
P0AFL2	POTI_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFL4	PPIA_ECOL6	periplasm (by similarity).	periplasm
P0AFL5	PPIA_ECO57	periplasm (by similarity).	periplasm
P0AFL7	PPX_ECOL6	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0AFL8	PPX_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0AFM0	PQIA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFM1	PQIA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFM3	PROX_ECO57	periplasm (by similarity).	periplasm
P0AFM4	PSIF_ECOLI		inner membrane
P0AFM5	PSIF_ECOL6		outer membrane
P0AFM7	PSPA_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AFM8	PSPA_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P0AFN0	PSPB_ECOL6	bacterial cell inner membrane (by similarity).	inner membrane

P0AFN1	PSPB_ECO57	bacterial cell inner membrane (by similarity).	inner membrane
P0AFN2	PSPC_ECOLI		cytoplasm
P0AFN3	PSPC_ECOL6		cytoplasm
P0AFN4	PSPC_ECO57		cytoplasm
P0AFN5	PSPC_SHIFL		cytoplasm
P0AFN7	YADH_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFN8	YADH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFN9	YADH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFP1	YADS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFP2	YBAZ_ECOLI		cytoplasm
P0AFP3	YBAZ_SHIFL		cytoplasm
P0AFP4	YBBO_ECOLI		cytoplasm
P0AFP5	YBBO_ECOL6		cytoplasm
P0AFP6	YBGI_ECOLI		extracellular
P0AFP7	YBGI_ECOL6		extracellular
P0AFP8	YBGI_ECO57		extracellular
P0AFQ0	YBHR_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFQ1	YBHR_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFQ3	YBHS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFQ4	YBHS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFQ5	RUTC_ECOLI		cytoplasm
P0AFQ6	RUTC_ECOL6		cytoplasm
P0AFQ7	YCFH_ECOLI		extracellular
P0AFQ8	YCFH_ECOL6		extracellular
P0AFQ9	YCFH_ECO57		extracellular
P0AFR0	YCHK_ECOLI		inner membrane
P0AFR1	YCHK_SHIFL		inner membrane
P0AFR3	YCHM_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFR4	YCIO_ECOLI		inner membrane
P0AFR5	YCIO_ECOL6		inner membrane
P0AFR6	YCIO_SHIFL		inner membrane
P0AFR8	YCJO_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFS0	YDCV_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AFS2	YDEZ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFS3	YDGB_ECOLI		cytoplasm
P0AFS4	YDGB_ECO57		cytoplasm
P0AFS6	YDGG_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFS8	YDIK_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFS9	YEBA_ECOLI		extracellular
P0AFT0	YEBA_ECOL6		extracellular
P0AFT1	YEBA_SHIFL		extracellular
P0AFT3	YECS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFT4	YECS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFT5	YEHT_ECOLI		cytoplasm
P0AFT6	YEHT_ECOL6		cytoplasm
P0AFT7	YEHT_ECO57		cytoplasm
P0AFT8	YEIW_ECOLI		inner membrane
P0AFT9	YEIW_ECOL6		inner membrane
P0AFU1	YEJB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFU2	YFBS_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0AFU3	YFBS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0AFU4	YFHA_ECOLI		cytoplasm
P0AFU5	YFHA_ECO57		cytoplasm
P0AFU6	YIIF_ECOLI		nucleoid
P0AFU7	YIIF_SHIFL		nucleoid
P0AFU8	RISA_ECOLI		cytoplasm
P0AFU9	RISA_SHIFL		cytoplasm
P0AFV1	YIBH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFV2	YHID_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AFV3	YHID_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AFV4	SPR_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AFV5	SPR_ECOL6	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AFV6	SPR_ECO57	bacterial cell membrane; lipid-anchor (potential).	inner membrane

P0AFV7	SPR_SHIFL	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P0AFV8	PSPD_ECOLI		inner membrane
P0AFV9	PSPD_ECO57		inner membrane
P0AFW0	RFAH_ECOLI		cytoplasm
P0AFW1	RFAH_ECO57		cytoplasm
P0AFW2	RMF_ECOLI		cytoplasm
P0AFW3	RMF_ECO57		cytoplasm
P0AFW4	RNK_ECOLI		cytoplasm
P0AFW5	RNK_ECOL6		cytoplasm
P0AFW6	RNK_ECO57		cytoplasm
P0AFW7	RNK_SHIFL		cytoplasm
P0AFW8	ROF_ECOLI		cytoplasm
P0AFW9	ROF_SHIFL		cytoplasm
P0AFX0	RP5M_ECOLI		cytoplasm
P0AFX1	RP5M_ECOL6		cytoplasm
P0AFX2	RP5M_ECO57		cytoplasm
P0AFX3	RP5M_SHIFL		cytoplasm
P0AFX4	RSD_ECOLI		cytoplasm
P0AFX5	RSD_ECO57		cytoplasm
P0AFX6	RSD_SHIFL		cytoplasm
P0AFX7	RSEA_ECOLI		inner membrane
P0AFX8	RSEA_ECOL6		inner membrane
P0AFX9	RSEB_ECOLI	periplasm (potential).	periplasm
P0AFY0	RSEB_ECO57	periplasm (potential).	periplasm
P0AFY1	RSEB_SHIFL	periplasm (potential).	periplasm
P0AFY3	SANA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFY4	SANA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFY5	SANA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFY7	SBMA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFY8	SEQA_ECOLI		cytoplasm
P0AFY9	SEQA_ECOL6		cytoplasm
P0AFZ0	SEQA_SHIFL		cytoplasm
P0AFZ1	SSEB_ECOLI		cytoplasm
P0AFZ2	SSEB_SHIFL		cytoplasm
P0AFZ3	SSPB_ECOLI		cytoplasm
P0AFZ4	SSPB_ECO57		cytoplasm
P0AFZ6	SULA_ECO57	bacterial cell inner membrane (by similarity).	inner membrane
P0AFZ8	TRKH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AFZ9	TRKH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AG01	WZZE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AG02	WZZE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AG03	UBIX_ECOLI		cytoplasm
P0AG04	UBIX_ECO57		cytoplasm
P0AG05	S3AD_ECOLI		cytoplasm
P0AG06	S3AX_SHIFL		cytoplasm
P0AG07	RPE_ECOLI		cytoplasm
P0AG08	RPE_ECOL6		cytoplasm
P0AG09	RPE_ECO57		cytoplasm
P0AG10	RPE_SHIFL		cytoplasm
P0AG11	UMUD_ECOLI		cytoplasm
P0AG12	UMUD_ECO57		cytoplasm
P0AG13	UMUD_SHIFL		cytoplasm
P0AG15	SOHB_SHIFL	bacterial cell inner membrane (by similarity).	extracellular
P0AG16	PUR1_ECOLI		cytoplasm
P0AG17	PUR1_SHIFL		cytoplasm
P0AG18	PUR6_ECOLI		cytoplasm
P0AG19	PUR6_ECO57		cytoplasm
P0AG20	RELA_ECOLI		cytoplasm
P0AG21	RELA_ECOL6		cytoplasm
P0AG22	RELA_ECO57		cytoplasm
P0AG23	RELA_SHIFL		cytoplasm
P0AG24	SPOT_ECOLI		cytoplasm
P0AG25	SPOT_ECO57		cytoplasm
P0AG26	SPOT_SHIFL		cytoplasm
P0AG27	YIBN_ECOLI		cytoplasm
P0AG28	YIBN_ECOL6		cytoplasm
P0AG29	YIBN_ECO57		cytoplasm
P0AG30	RHO_ECOLI		inner membrane
P0AG31	RHO_ECOL6		inner membrane
P0AG32	RHO_ECO57		inner membrane
P0AG33	RHO_SHIFL		inner membrane
P0AG34	RHTB_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AG35	RHTB_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AG36	RHTB_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AG37	RHTB_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AG39	RHTC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AG40	RIBF_ECOLI		cytoplasm
P0AG41	RIBF_ECOL6		cytoplasm
P0AG42	RIBF_ECO57		cytoplasm
P0AG43	RIBF_SHIFL		cytoplasm
P0AG44	RL17_ECOLI		cytoplasm
P0AG45	RL17_ECOL6		cytoplasm
P0AG46	RL17_ECO57		cytoplasm

P0AG47	RL17_SHIFL		cytoplasm
P0AG48	RL21_ECOLI		cytoplasm
P0AG49	RL21_ECO57		cytoplasm
P0AG50	RL21_SHIFL		cytoplasm
P0AG51	RL30_ECOLI		cytoplasm
P0AG52	RL30_ECOL6		cytoplasm
P0AG53	RL30_ECO57		cytoplasm
P0AG54	RL30_SHIFL		cytoplasm
P0AG55	RL6_ECOLI		cytoplasm
P0AG56	RL6_ECOL6		cytoplasm
P0AG57	RL6_ECO57		cytoplasm
P0AG58	RL6_SHIFL		cytoplasm
P0AG59	RS14_ECOLI		cytoplasm
P0AG60	RS14_ECOL6		cytoplasm
P0AG61	RS14_ECO57		cytoplasm
P0AG62	RS14_SHIFL		cytoplasm
P0AG63	RS17_ECOLI		cytoplasm
P0AG64	RS17_ECOL6		cytoplasm
P0AG65	RS17_ECO57		cytoplasm
P0AG66	RS17_SHIFL		cytoplasm
P0AG67	RS1_ECOLI		cytoplasm
P0AG68	RS1_ECOL6		cytoplasm
P0AG69	RS1_ECO57		cytoplasm
P0AG70	RS1_SHIFL		cytoplasm
P0AG71	RMUC_ECOLI		cytoplasm
P0AG72	RMUC_ECO57		cytoplasm
P0AG73	RMUC_SHIFL		cytoplasm
P0AG74	RUS_ECOLI		cytoplasm
P0AG75	RUS_ECOL6		cytoplasm
P0AG76	SBCD_ECOLI		extracellular
P0AG77	SBCD_ECO57		extracellular
P0AG79	SUBI_SHIFL	periplasm (by similarity).	periplasm
P0AG81	UGPB_ECO57	periplasm (by similarity).	periplasm
P0AG83	PSTS_SHIFL	periplasm (by similarity).	periplasm
P0AG84	YGHA_ECOLI		cytoplasm
P0AG85	YGHA_ECO57		cytoplasm
P0AG87	SECB_ECOL6	cytoplasm (by similarity).	cytoplasm
P0AG88	SECB_ECO57	cytoplasm (by similarity).	cytoplasm
P0AG89	SECB_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AG91	SECD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AG92	SECD_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AG94	SECF_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AG95	SECF_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AG97	SECE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AG98	SECE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGA0	SECG_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGA1	SECG_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGA3	SECY_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGA4	SECY_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGA5	SECY_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGA6	UHPA_ECOLI	cytoplasm (probable).	cytoplasm
P0AGA7	UHPA_ECOL6	cytoplasm (probable).	cytoplasm
P0AGA8	UHPA_ECO57	cytoplasm (probable).	cytoplasm
P0AGA9	UHPA_SHIFL	cytoplasm (probable).	cytoplasm
P0AGB0	SERB_ECOLI		cytoplasm
P0AGB1	SERB_ECO57		cytoplasm
P0AGB2	SERB_SHIFL		cytoplasm
P0AGB3	RP32_ECOLI		cytoplasm
P0AGB4	RP32_ECOL6		cytoplasm
P0AGB5	RP32_ECO57		cytoplasm
P0AGB6	RPOE_ECOLI		cytoplasm
P0AGB7	RPOE_ECOL6		cytoplasm
P0AGB8	RPOE_ECO57		cytoplasm
P0AGB9	RPOE_SHIFL		cytoplasm
P0AGC1	UHPT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGC2	UHPT_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGC5	YFHD_ECOLI		periplasm
P0AGC6	YFHD_SHIFL		periplasm
P0AGC8	SMP_SHIFL	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P0AGC9	QACE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AGD0	QACE_KLEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P0AGD2	SODC_ECO57	periplasm (by similarity).	periplasm
P0AGD3	SODF_ECOLI		periplasm
P0AGD4	SODF_ECOL6		periplasm
P0AGD5	SODF_ECO57		periplasm
P0AGD6	SODF_SHIFL		periplasm
P0AGD7	SRP54_ECOLI		cytoplasm
P0AGD8	SRP54_ECOL6		cytoplasm

P0AGD9	SRP54_ECO57		cytoplasm
P0AGE0	SSB_ECOLI		cytoplasm
P0AGE1	SSB_ECOL6		cytoplasm
P0AGE2	SSB_ECO57		cytoplasm
P0AGE3	SSB_SHIFL		cytoplasm
P0AGE5	YGJU_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGE6	YIEF_ECOLI		cytoplasm
P0AGE7	YIEF_ECO57		cytoplasm
P0AGE8	YIEF_SHIFL		cytoplasm
P0AGE9	SUCD_ECOLI		cytoplasm
P0AGF0	SUCD_ECOL6		cytoplasm
P0AGF1	SUCD_ECO57		cytoplasm
P0AGF2	YGDK_ECOLI		cytoplasm
P0AGF3	YGDK_ECO57		cytoplasm
P0AGF5	XYLE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGF6	THD2_ECOLI		cytoplasm
P0AGF7	THD2_ECOL6		cytoplasm
P0AGF8	THD2_ECO57		cytoplasm
P0AGF9	THD2_SHIFL		cytoplasm
P0AGG0	THIL_ECOLI		cytoplasm
P0AGG1	THIL_ECOL6		cytoplasm
P0AGG2	TESB_ECOLI		extracellular
P0AGG3	TESB_ECO57		extracellular
P0AGG5	THIO2_ECOL6	cytoplasm (by similarity).	cytoplasm
P0AGG6	THIO2_ECO57	cytoplasm (by similarity).	cytoplasm
P0AGG7	THIO2_SHIFL	cytoplasm (by similarity).	cytoplasm
P0AGG8	TLDD_ECOLI		cytoplasm
P0AGG9	TLDD_ECOL6		cytoplasm
P0AGH0	TLDD_SHIFL		cytoplasm
P0AGH2	YHHJ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGH4	SAPB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGH6	SAPC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGH7	SAPC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGH9	PSTC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGI0	PSTC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGI2	RBSC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AGI3	RBSC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGI5	XYLH_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGI6	XYLH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGI7	XYLH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGJ2	TRMH_ECOLI	cytoplasm (potential).	cytoplasm
P0AGJ3	TRMH_ECO57	cytoplasm (potential).	cytoplasm
P0AGJ4	TRMH_SHIFL	cytoplasm (potential).	cytoplasm
P0AGJ5	YFIF_ECOLI		cytoplasm
P0AGJ6	YFIF_ECO57		cytoplasm
P0AGJ7	YIBK_ECOLI		cytoplasm
P0AGJ8	YIBK_ECOL6		cytoplasm
P0AGK0	SYU_ECO57	cytoplasm (by similarity).	cytoplasm
P0AGK2	UBIA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGK3	UBIA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGK4	YHBY_ECOLI		inner membrane
P0AGK5	YHBY_ECOL6		inner membrane
P0AGK6	YHBY_ECO57		inner membrane
P0AGK7	YHBY_SHIFL		inner membrane
P0AGK8	YFHP_ECOLI		cytoplasm
P0AGK9	YFHP_ECOL6		cytoplasm
P0AGL0	YFHP_ECO57		cytoplasm
P0AGL1	YFHP_SHIFL		cytoplasm
P0AGL2	TDCF_ECOLI		cytoplasm
P0AGL3	TDCF_ECOL6		cytoplasm
P0AGL4	TDCF_SHIFL		cytoplasm
P0AGL5	YFJG_ECOLI		inner membrane
P0AGL6	YFJG_SHIFL		inner membrane
P0AGL7	YGGJ_ECOLI		cytoplasm
P0AGL8	YGGJ_ECOL6		cytoplasm
P0AGL9	YGGJ_SHIFL		cytoplasm
P0AGM1	YHHT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGM3	YICG_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGM4	YICG_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGM5	SIRB1_ECOLI		extracellular
P0AGM6	SIRB1_ECO57		extracellular
P0AGM8	URAA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P0AGM9	YICE_ECOLI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P0AGN0	YICE_ECOL6	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGN1	YICE_ECO57	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P0AGN2	YICE_SHIFL	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C002	DHP1_ECOLI		cytoplasm
P0C003	DHP1_PSEAE		cytoplasm
P0C018	RL18_ECOLI		cytoplasm
P0C019	RL18_ECOL6		cytoplasm
P0C020	RL18_ECO57		cytoplasm
P0C021	RL18_SHIFL		cytoplasm
P0C037	YAIE_ECOLI		inner membrane
P0C038	YAIE_ECOL6		inner membrane
P0C039	YAIE_ECO57		inner membrane
P0C040	YAIE_SHIFL		inner membrane
P0C055	IBPA_ECO57	cytoplasm (by similarity).	cytoplasm
P0C056	IBPA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0C057	IBPA_SHIFL	cytoplasm (by similarity).	cytoplasm
P0C059	IBPB_ECOL6	cytoplasm (by similarity).	cytoplasm
P0C060	IBPB_SHIFL	cytoplasm (by similarity).	cytoplasm
P0C066	MLTC_ECOLI	bacterial cell outer membrane; lipid-anchor (probable).	extracellular
P0C067	MLTC_ECOL6	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
P0C068	MEPR_PSEPU		cytoplasm
P0C069	MEPA_PSEPU	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P0C070	MEPB_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P0C071	MEPC_PSEPU	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
P0C076	RL31_ECOL6		cytoplasm
P0C077	RELE_ECOLI		cytoplasm
P0C078	RELE_SHIFL		cytoplasm
P0C079	RELB_ECOLI		cytoplasm
P0C080	RELB_SHIFL		cytoplasm
P0C081	LKA2B_PASHA	secreted protein (by similarity).	extracellular
P0C082	LKA2C_PASHA	secreted protein (by similarity).	extracellular
P0C083	LKTA6_PASHA	secreted protein (by similarity).	extracellular
P0C084	LKA7A_PASHA	secreted protein (by similarity).	extracellular
P0C085	LKA7B_PASHA	secreted protein (by similarity).	extracellular

P0C086	LKTB7_PASHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0C087	LKB14_PASHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0C094	SLMA_SHIFL	nucleoid-associated (by similarity).	cytoplasm
P0C0B2	CYSI_BUCAP		cytoplasm
P0C0F6	RPFC_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C0F7	RPFC_XANC8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C0F8	SODM_XANCP		periplasm
P0C0F9	SODM_XANC8		periplasm
P0C0K3	RDOA_ECOLI	cytoplasm (by similarity).	cytoplasm
P0C0K4	RDOA_ECOL6	cytoplasm (by similarity).	cytoplasm
P0C0L3	OSMC_SHIFL	cytoplasm (by similarity).	cytoplasm
P0C0L8	PROP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C0L9	ISCX_ECOLI		cytoplasm
P0C0M0	ISCX_ECO57		cytoplasm
P0C0M1	ISCX_SHIFL		cytoplasm
P0C0R6	ARNA_SALCH		cytoplasm
P0C0R7	RRMJ_ECOLI	cytoplasm (probable).	cytoplasm
P0C0R8	RRMJ_ECO57	cytoplasm (by similarity).	cytoplasm
P0C0R9	RRMJ_ECOL6	cytoplasm (by similarity).	cytoplasm
P0C0S0	RRMJ_SHIFL	cytoplasm (by similarity).	cytoplasm
P0C0S2	MSCS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C0S3	MSCS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C0T3	DSRB_SALTY		inner membrane
P0C0T6	MEPA_SHIFL	periplasm (by similarity).	extracellular
P0C0U4	RIMK_ECOLI		cytoplasm
P0C0U5	RIMK_ECO57		cytoplasm
P0C0V1	DEGP_ECO57	periplasm (by similarity).	periplasm
P0C0W2	RIHC_ECOL6		extracellular
P0C0Y6	PUFQ_RHOSH		cytoplasm
P0C0Y7	RCEH_RHOSH		inner membrane
P0C0Z0	REGB_RHOSH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C0Z2	UVRA_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C0Z3	RL16_BRUAB		cytoplasm
P0C0Z4	RL16_BRUA2		cytoplasm
P0C0Z5	RS15_BRUAB		cytoplasm
P0C0Z6	RS15_BRUA2		cytoplasm
P0C100	AROAB_BRUAB	cytoplasm (probable).	cytoplasm
P0C101	AROQ_BRUAB		cytoplasm

P0C102	CDSA_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C104	FABZ_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C105	GLUP_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P0C106	FUR_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C107	HFQ_BRUAB		cytoplasm
P0C110	LPXA_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C111	LPXD_BRUAB		cytoplasm
P0C112	AQPZ_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C113	LON_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C114	DEGP_BRUAB	periplasm (potential).	periplasm
P0C115	PURA_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C116	MTB1_BRUAB		cytoplasm
P0C117	SECD_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C118	SSB_BRUAB		cytoplasm
P0C119	TPIS2_BRUAB		cytoplasm
P0C120	YAJC_BRUAB		inner membrane
P0C121	URE31_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C122	XERD_BRUAB	cytoplasm (by similarity).	nucleoid
P0C123	SYI_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C124	SYI_BRUA2	cytoplasm (by similarity).	cytoplasm
P0C125	SECB_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C126	SECB_BRUA2	cytoplasm (by similarity).	cytoplasm
P0C144	UREE_BRUAB	cytoplasm (by similarity).	cytoplasm
P0C145	UREE_BRUA2	cytoplasm (by similarity).	cytoplasm
P0C151	RPOB_BURMA		cytoplasm
P0C184	ULAA_SHISS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C186	PNTAA_RHORU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C187	PNTAB_RHORU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C188	PNTB_RHORU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P0C189	CYC2_RHORU		periplasm
P0C193	CH60_SODGL	cytoplasm (by similarity).	cytoplasm
P0C1A6	PELL_ERWCH	secreted protein (by similarity).	extracellular
P0C1A8	PMEA_ERWCH	secreted protein (by similarity).	extracellular
P0C1B2	ULAE_SHIDS		cytoplasm
P0C1B9	ULAF_SHIBS		cytoplasm
P0C1C4	TUSB_SHIBS	cytoplasm (by similarity).	cytoplasm
P0C1D5	URE21_BRUAB	cytoplasm (by similarity).	cytoplasm

P0C1F6	CCME_PSEFL	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
P0C1F7	CCMD_PSEFL	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P10018	YPR2_ECOLI		inner membrane
P10027	YPR3_ECOLI		inner membrane
P10030	PIFC_ECOLI		cytoplasm
P10031	PSIB1_ECOLI		cytoplasm
P10032	PSIB2_ECOLI		cytoplasm
P10045	NRLB_KLEPO		cytoplasm
P10046	DCTD_RHILE	cytoplasm (probable).	cytoplasm
P10051	AAC6_CITDI		cytoplasm
P10054	RL1_PROVU		cytoplasm
P10055	RL11_PROVU		cytoplasm
P10088	TFDA_RALEJ		cytoplasm
P10089	HLYBC_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P10100	RLPA_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P10151	LEUO_ECOLI		cytoplasm
P10172	ASPQ_ACIGL	periplasm (by similarity).	periplasm
P10326	FIXX_BRAJA		cytoplasm
P10331	FIXC_BRAJA		cytoplasm
P10336	NIFN_AZOVI		periplasm
P10342	ISOA_PSEAY		cytoplasm
P10343	YI42_PSEAY		cytoplasm
P10368	HIS7_SALTY	cytoplasm (by similarity).	cytoplasm
P10369	HIS8_SALTY		cytoplasm
P10370	HISX_SALTY		cytoplasm
P10378	ENTE_ECOLI		cytoplasm
P10390	NIFB_KLEPN		cytoplasm
P10392	NIFQ_KLEPN		cytoplasm
P10423	IAP_ECOLI		cytoplasm
P10441	LPXB_ECOLI		cytoplasm
P10442	RNH2_ECOLI	cytoplasm (potential).	extracellular
P10449	FIXB_BRAJA		cytoplasm
P10473	THIO_RHORU		cytoplasm
P10476	GUNA_PSEFL		extracellular
P10480	GCAT_AERHY		cytoplasm
P10484	T1M1_ECOLI		cytoplasm
P10485	T1S1_ECOLI		cytoplasm
P10486	T1R1_ECOLI		cytoplasm
P10487	RCI1_ECOLI		cytoplasm
P10498	PSS_RHILP		cytoplasm
P10503	PUTA_SALTY		cytoplasm
P10512	TRAY2_ECOLI		cytoplasm
P10514	TRAE2_ECOLI		cytoplasm
P10576	NTRC_BRASR		cytoplasm
P10577	NTRC_RHIME		cytoplasm
P10578	NTRB_BRASR		inner membrane
P10624	FER2_DESVM		periplasm

P10640	BEXA_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P10738	ERMB_ECOLI		cytoplasm
P10799	VIRA_AGRT9		inner membrane
P10802	ODP2_AZOVI		inner membrane
P10805	ENVY_ECOLI		cytoplasm
P10880	AROL_ERWCH	cytoplasm (probable).	cytoplasm
P10932	AMIR_PSEAE		cytoplasm
P10952	TETO_CAMJE		cytoplasm
P10958	FIXJ_RHIME	cytoplasm (probable).	cytoplasm
P11003	METF_SALTY		cytoplasm
P11054	FERN_AZOVI		periplasm
P11067	NIFB_AZOVI		cytoplasm
P11068	NIFQ_AZOVI		cytoplasm
P11072	LIT_ECOLI		inner membrane
P11080	TRPB_PSEPU		cytoplasm
P11081	TRPA_PSEPU		cytoplasm
P11091	CYAD_BORPE	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P11092	CYAE_BORPE	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P11122	BPHC_PSEPA		cytoplasm
P11252	RL3_YERPS		cytoplasm
P11256	RS19_YERPS		cytoplasm
P11257	T341_ECOLI		cytoplasm
P11289	YFIL_ECOLI		periplasm
P11290	YIBD_ECOLI		cytoplasm
P11291	YZCX_ECOLI		inner membrane
P11405	T2M1_MORSP		extracellular
P11406	RECA_PROMI	cytoplasm (by similarity).	cytoplasm
P11408	MTM1_MORSP		cytoplasm
P11409	MTP2_PROVU		cytoplasm
P11436	AMIE_PSEAE		cytoplasm
P11437	AVRA_PSESG		extracellular
P11439	TOXA_PSEAE		cytoplasm
P11444	MANR_PSEPU		cytoplasm
P11446	ARGC_ECOLI	cytoplasm (probable).	cytoplasm
P11447	ARLY_ECOLI	cytoplasm (probable).	cytoplasm
P11451	CLCA_PSEPU		cytoplasm
P11452	CLCB_PSEPU		cytoplasm
P11454	ENTF_ECOLI		cytoplasm
P11460	FATB_VIBAN	bacterial cell inner membrane; lipid-anchor (potential).	periplasm
P11519	FLMC1_ECOLI		cytoplasm
P11539	RP32_CITFR		cytoplasm
P11553	FUCK_ECOLI		cytoplasm
P11556	FUCT_ECOLI		inner membrane
P11557	DAMX_ECOLI		inner membrane
P11663	YGGD_ECOLI		cytoplasm
P11664	YGGC_ECOLI		cytoplasm
P11671	GLNB_KLEOX		cytoplasm
P11720	TRPI_PSEAE		cytoplasm
P11731	DYR5_ECOLI		cytoplasm

P11732	CYC5_AZOVI		periplasm
P11759	ALGD_PSEAE		cytoplasm
P11764	FMM2_NEIGO		extracellular
P11797	CHIB_SERMA		extracellular
P11861	NAHC1_PSEPU		cytoplasm
P11864	YHAC_ECOLI		cytoplasm
P11865	YHAB_ECOLI		cytoplasm
P11866	TDCR_ECOLI		cytoplasm
P11868	TDCD_ECOLI		cytoplasm
P11888	ROLA_AGRRH		extracellular
P11895	HOK_ECOLI		inner membrane
P11899	IMMD_ECOLI		cytoplasm
P11901	T421_ECOLI		cytoplasm
P11902	PNDA1_ECOLI		inner membrane
P11904	PARM_ECOLI		flagellar
P11905	STBC_ECOLI		cytoplasm
P11906	STBB_ECOLI		cytoplasm
P11907	YPNR_ECOLI		cytoplasm
P11933	FMAG_BACNO		extracellular
P11954	THD2_SALTY		cytoplasm
P11987	MEMG_METCA		cytoplasm
P11988	BGLB_ECOLI		cytoplasm
P11989	BGLG_ECOLI		cytoplasm
P12008	AROC_ECOLI		cytoplasm
P12009	LIVX_ECOLI		inner membrane
P12015	CYMO_ACISP		cytoplasm
P12033	KPPR1_RHOSH		cytoplasm
P12050	FAND_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P12056	CML_ECOLI		inner membrane
P12057	TRAE1_SALTI		cytoplasm
P12058	TRAL_SALTI	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P12059	TRAY1_SALTI		cytoplasm
P12232	NOD12_BRAJA		cytoplasm
P12233	NOD22_BRAJA		cytoplasm
P12281	MOEA_ECOLI		fimbrium
P12282	MOEB_ECOLI		cytoplasm
P12288	USG_CAUCR		cytoplasm
P12289	TRPF_CAUCR		cytoplasm
P12290	TRPB_CAUCR		cytoplasm
P12291	TRPA_CAUCR		cytoplasm
P12364	T3MO_ECOLI		cytoplasm
P12377	COPD_PSESM	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P12380	HUTC_KLEAE		cytoplasm
P12421	AROA_BORPE	cytoplasm (probable).	cytoplasm
P12580	VANB_PSES9		cytoplasm
P12609	VANA_PSES9		cytoplasm
P12614	BGLS_AGRSA		extracellular
P12626	ANFA_AZOVI		cytoplasm
P12627	VNFA_AZOVI		cytoplasm
P12634	FLAD_CAUCR	periplasm (probable).	flagellar

P12673	CYSZ_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P12677	BIOA_SALTY	cytoplasm (by similarity).	cytoplasm
P12678	BIOB_SALTY		cytoplasm
P12679	PABB_KLEAE		cytoplasm
P12680	PABB_SALTY		cytoplasm
P12691	ALKB_PSEOL	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P12692	RUBR1_PSEOL		cytoplasm
P12693	ALDH_PSEOL		cytoplasm
P12712	FDXN_RHIME		periplasm
P12730	SFAA_ECOL6		extracellular
P12744	LUXB_PHOPO		cytoplasm
P12745	LUXF_PHOPO		cytoplasm
P12746	LUXR_VIBFI		cytoplasm
P12747	LUXI_VIBFI		cytoplasm
P12748	LUXC_VIBFI		cytoplasm
P12779	HIA_RHIFR		cytoplasm
P12780	NOLJ_RHIFR		cytoplasm
P12781	NIFN_RHIME		periplasm
P12784	ACP_RHOSH	cytoplasm (by similarity).	cytoplasm
P12832	CY550_AQUIT		periplasm
P12833	DYR3_SALTY		cytoplasm
P12903	FM12_KLEPN		extracellular
P12983	ATPZ_VIBAL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P12985	ATPA_VIBAL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P12986	ATPB_VIBAL		inner membrane
P12987	ATPD_VIBAL		inner membrane
P12988	ATPE_VIBAL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P12989	ATPF_VIBAL		inner membrane
P12990	ATPG_VIBAL		inner membrane
P12996	BIOB_ECOLI		cytoplasm
P12998	BIOF_ECOLI		cytoplasm
P12999	BIOC_ECOLI		cytoplasm
P13001	BIOH_ECOLI	cytoplasm (by similarity).	cytoplasm
P13009	METH_ECOLI		cytoplasm
P13018	STA_ECOLI		cytoplasm
P13029	CATA_ECOLI		periplasm
P13081	BLE_KLEPN		inner membrane
P13082	STR_KLEPN		cytoplasm
P13111	MERR_SERMA		cytoplasm
P13113	MERP_SERMA	periplasm (probable).	inner membrane
P13118	FLAA2_RHIME		flagellar
P13119	FLAB2_RHIME		flagellar
P13187	DCOA_KLEPN		inner membrane
P13225	VIRF_YEREN		cytoplasm
P13246	AAC31_SALSP		cytoplasm
P13253	FMAD_BACNO		extracellular

P13254	MEGL_PSEPU		cytoplasm
P13295	FIXK_RHIME		cytoplasm
P13364	GYRB_PSEPU		cytoplasm
P13370	TODB_PSEPU		cytoplasm
P13398	NYLA_PSES8		extracellular
P13421	FMA_SERMA		extracellular
P13429	SFAG_ECOL6		extracellular
P13430	SFAS_ECOL6		extracellular
P13431	SFAH_ECOL6		extracellular
P13441	NODP_RHIME		cytoplasm
P13442	NODQ_RHIME		cytoplasm
P13445	RPOS_ECOLI		cytoplasm
P13452	TODA_PSEPU		cytoplasm
P13453	TODE_PSEPU		cytoplasm
P13456	RECF_PSEPU	cytoplasm (potential).	cytoplasm
P13458	SBCC_ECOLI		cytoplasm
P13459	VIRC1_AGRRH		cytoplasm
P13460	VIRC2_AGRRH		cytoplasm
P13461	VIRD1_AGRRH		extracellular
P13462	VIRD2_AGRRH		extracellular
P13463	VIRD3_AGRRH		cytoplasm
P13464	VIRD4_AGRRH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P13476	IMM5_ECOLI		cytoplasm
P13477	IMM6_ECOLI		cytoplasm
P13478	IMMX_ECOLI		cytoplasm
P13479	IMM9_ECOLI		cytoplasm
P13486	Y4066_RHIME		cytoplasm
P13509	CZCC_RALME		inner membrane
P13510	CZCB_RALME		inner membrane
P13511	CZCA_RALME	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P13512	CZCD_RALME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P13514	LSPA_ENTAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P13518	YHDA_ECOLI		inner membrane
P13556	GLNB_RHOCA		cytoplasm
P13606	RES4_ECOLI		cytoplasm
P13627	CY1_PARDE		inner membrane
P13632	DCTD_RHIME	cytoplasm (probable).	cytoplasm
P13650	DHGB_ACICA		extracellular
P13657	MBEC_ECOLI		cytoplasm
P13658	MBEA_ECOLI		cytoplasm
P13659	MBEB_ECOLI		cytoplasm
P13660	MBED_ECOLI		cytoplasm
P13661	BLO1_ECOLI		extracellular
P13669	FARR_ECOLI		cytoplasm
P13694	TNP7_ECOLI		cytoplasm
P13702	MVAA_PSEMV		cytoplasm
P13703	HMGCL_PSEMV		cytoplasm
P13713	FLIC_SERMA		flagellar

P13809	RPI_ECOLI		extracellular
P13810	E2AA_ECOLI		extracellular
P13811	ELBH_ECOLI		extracellular
P13812	E2AB_ECOLI		extracellular
P13835	AVRB_PSESG		extracellular
P13836	AVRC_PSESG		extracellular
P13859	TODD_PSEPU		cytoplasm
P13946	COPA2_ECOLI		cytoplasm
P13959	COPB1_ECOLI		cytoplasm
P13961	REP4_ECOLI		nucleoid
P13964	KLEA2_ECOLI		cytoplasm
P13965	KLEB2_ECOLI		cytoplasm
P13966	KLEC2_ECOLI		nucleoid
P13967	KLED2_ECOLI		cytoplasm
P13970	SRNB_ECOLI		inner membrane
P13971	SOK_ECOLI		extracellular
P13974	YPRA_ECOLI		inner membrane
P13975	PEMI_ECOLI		cytoplasm
P13976	PEMK_ECOLI		cytoplasm
P13981	ARCA_PSEAE	cytoplasm (potential).	extracellular
P13982	ARCC_PSEAE	cytoplasm (potential).	cytoplasm
P13988	TNSA_ECOLI		cytoplasm
P13989	TNSB_ECOLI		cytoplasm
P13991	TNSD_ECOLI		cytoplasm
P14011	IPT_AGRRH		cytoplasm
P14012	ARGI_AGRT5		cytoplasm
P14062	6PGD_SALTY		cytoplasm
P14129	RS1_RHIME		cytoplasm
P14146	PHOP_SALTY	cytoplasm (probable).	cytoplasm
P14147	PHOQ_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P14161	MUTL_SALTY		cytoplasm
P14165	CISY_PSEAE		cytoplasm
P14168	RFBS_SALTI		cytoplasm
P14169	RFBE_SALTI		cytoplasm
P14172	YAMP_RHOCA		extracellular
P14179	GLNB_BRAJA		cytoplasm
P14181	LICA2_HAEIN		cytoplasm
P14182	LICB_HAEIN		inner membrane
P14183	LICC_HAEIN		cytoplasm
P14184	LICD_HAEIN		cytoplasm
P14185	EXOX_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P14190	FMKC_ECOLI		extracellular
P14191	FMKD_ECOLI		extracellular
P14229	T2SM_SERMA		extracellular
P14230	MTSM_SERMA		cytoplasm
P14243	MTC9_CITFR		cytoplasm
P14285	CHRA_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P14286	LUXE_VIBHA		cytoplasm
P14294	TOP3_ECOLI		cytoplasm
P14299	DRAT_RHORU		cytoplasm

P14300	DRAG_RHORU		extracellular
P14301	YDRA_RHORU		cytoplasm
P14303	MUCB_SALTY		cytoplasm
P14307	YSMA_SERMA		cytoplasm
P14310	YFX1_RHILE		cytoplasm
P14311	YFX2_RHILE		cytoplasm
P14312	FIXW_RHILE		periplasm
P14333	IPT2_RALSO		cytoplasm
P14375	ZRAR_ECOLI	cytoplasm (probable).	cytoplasm
P14407	FUMB_ECOLI		cytoplasm
P14489	BLP2_PSEAE		extracellular
P14492	RLX1_SALTY		cytoplasm
P14493	RLX2_SALTY		cytoplasm
P14498	TRAS2_ECOLI		cytoplasm
P14499	X191_ECOLI		cytoplasm
P14504	YP54_ECOLI		cytoplasm
P14505	YP55_ECOLI		cytoplasm
P14508	KKA7_CAMJE		cytoplasm
P14509	KKA8_ECOLI		cytoplasm
P14511	S3AD_SHIFL		cytoplasm
P14565	TRAI1_ECOLI		cytoplasm
P14567	DPO3A_SALTY	cytoplasm (by similarity).	cytoplasm
P14601	PUFQ_RHOCA		cytoplasm
P14633	T2E2_ECOLI		extracellular
P14711	HLYT_VIBHO		extracellular
P14727	YAV1_XANCV		extracellular
P14728	YAV2_XANCV		extracellular
P14729	YAV3_XANCV		extracellular
P14751	MTR1_RHOSH		cytoplasm
P14768	XYNA_PSEFL		cytoplasm
P14774	CYCL_METEX	periplasm (potential).	periplasm
P14801	EXOX_RHILP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P14827	MTEC_ENTCL		cytoplasm
P14829	GYRA_KLEOX		cytoplasm
P14845	CARA_SALTY		cytoplasm
P14846	CARB_SALTY		cytoplasm
P14886	NIFY_AZOVI		cytoplasm
P14887	NIFX_AZOVI		cytoplasm
P14888	NIFW_AZOVI		cytoplasm
P14889	NIFZ_AZOVI		cytoplasm
P14890	NIFM_AZOVI		cytoplasm
P14929	FTSY_NEIGO	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P14930	MSRAB_NEIGO		cytoplasm
P14939	FERV_AZOVI		periplasm
P14940	ADH_RALEU		cytoplasm
P15005	MCRB_ECOLI		cytoplasm
P15006	MCRC_ECOLI		cytoplasm
P15011	ATPZ_RHORU		inner membrane
P15013	ATPF_RHORU		inner membrane
P15014	ATPL_RHORU		inner membrane
P15015	ATPX_RHORU		inner membrane
P15017	DNU4_RHORU		cytoplasm

P15025	ISTA_PSEAE		cytoplasm
P15026	ISTB_PSEAE		cytoplasm
P15032	RECE_ECOLI		cytoplasm
P15033	RACC_ECOLI		cytoplasm
P15038	HELD_ECOLI		cytoplasm
P15042	DNLJ_ECOLI		cytoplasm
P15043	RECQ_ECOLI		cytoplasm
P15047	ENTA_ECOLI		cytoplasm
P15052	NIFK_THIFE		cytoplasm
P15067	GLGX_ECOLI		extracellular
P15068	TRBF_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P15081	GUTM_ECOLI		cytoplasm
P15082	SRLR_ECOLI		cytoplasm
P15109	PCXA_BURCE		cytoplasm
P15110	PCXB_BURCE		cytoplasm
P15111	DHE4_SALTY		cytoplasm
P15255	65KD_ZYMMO		cytoplasm
P15256	21KD_ZYMMO		cytoplasm
P15266	NDK_MYXXA	cytoplasm (by similarity).	cytoplasm
P15275	ALGQ_PSEAE		cytoplasm
P15276	ALGP_PSEAE		cytoplasm
P15286	DIV_ECOLI		inner membrane
P15288	PEPD_ECOLI		cytoplasm
P15332	VNFD_AZOCH		cytoplasm
P15333	VNFG_AZOCH		periplasm
P15334	VNFK_AZOCH		cytoplasm
P15335	NIFH2_AZOVI		cytoplasm
P15345	FLAEY_CAUCR		cytoplasm
P15373	SOHA_ECOLI	cytoplasm (probable).	cytoplasm
P15394	REPA_AGRTU		nucleoid
P15395	TRPE_RHIME		cytoplasm
P15397	ROLT_AGRRH		cytoplasm
P15446	MTH2_HAEP A		cytoplasm
P15483	CS31_ECOLI	periplasm (by similarity).	periplasm
P15484	CS32_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P15488	FMS3_ECOLI		fimbrium
P15493	LIP_VIBCH		extracellular
P15557	PAC1_PSES3		cytoplasm
P15558	PAC2_PSES3		periplasm
P15595	TRAF_AGRT9	periplasm (probable).	periplasm
P15596	YVF3_AGRT9		inner membrane
P15597	VIRF_AGRT9		extracellular
P15639	PUR9_ECOLI		cytoplasm
P15640	PUR2_ECOLI		cytoplasm
P15645	GSPE_KLEPN	cytoplasm (probable).	inner membrane
P15653	IPT_AGRTU		cytoplasm
P15701	KDGT_ERWCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P15713	PHLN_PSEAE		extracellular
P15717	LEU21_SALTY		cytoplasm
P15723	DGTP_ECOLI		cytoplasm

P15727	NODT_RHILV	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P15745	GSPF_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P15746	GSPG_KLEPN		inner membrane
P15747	GSPH_KLEPN		inner membrane
P15748	GSPI_KLEPN		inner membrane
P15749	GSPJ_KLEPN		inner membrane
P15750	GSPK_KLEPN	bacterial cell inner membrane (potential).	inner membrane
P15751	GSPL_KLEPN	bacterial cell inner membrane (probable).	inner membrane
P15752	GSPM_KLEPN	bacterial cell inner membrane (probable).	inner membrane
P15753	GSPN_KLEPN	bacterial cell inner membrane (probable).	inner membrane
P15754	LEP4_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P15770	AROE_ECOLI		cytoplasm
P15795	TOXR_VIBCH		cytoplasm
P15805	VRP3_SALCH		extracellular
P15875	LEU1_SALTY		cytoplasm
P15888	NFNB_SALTY		cytoplasm
P15905	ARSR1_ECOLI		cytoplasm
P15934	FLIH_SALTY		inner membrane
P15939	NODV_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P15940	NODW_BRAJA	cytoplasm (probable).	cytoplasm
P16021	FDXN_RHOCA		periplasm
P16022	FER4_RHOCA		cytoplasm
P16028	MOXJ_METEX	periplasm (potential).	periplasm
P16034	RECA_MAGMG	cytoplasm (by similarity).	cytoplasm
P16095	SDHL_ECOLI		cytoplasm
P16099	DHTM_METME		cytoplasm
P16113	PYP_ECTHA		cytoplasm
P16122	URE1_PROVU	cytoplasm (by similarity).	cytoplasm
P16123	URE2_PROVU	cytoplasm (by similarity).	cytoplasm
P16124	URE3_PROVU	cytoplasm (by similarity).	cytoplasm
P16238	RECA_THIFE	cytoplasm (by similarity).	cytoplasm
P16266	ANFD_AZOVI		cytoplasm
P16267	ANFK_AZOVI		cytoplasm
P16268	ANFG_AZOVI		periplasm
P16269	NIFH3_AZOVI		cytoplasm
P16280	AROC_SALTI		cytoplasm
P16323	FLGF_SALTY		flagellar
P16326	FLGL_SALTY		flagellar
P16384	MIAA_ECOLI		cytoplasm
P16431	HYCE_ECOLI		periplasm
P16432	HYCF_ECOLI		periplasm
P16433	HYCG_ECOLI		cytoplasm
P16437	FLGB_SALTY		flagellar
P16447	LUXG_VIBHA		cytoplasm
P16448	RIBB_VIBHA		cytoplasm

P16454	AIL_YEREN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P16456	SELD_ECOLI		cytoplasm
P16461	LKTC_ACTAC	cytoplasm (potential).	cytoplasm
P16462	LKTA_ACTAC	or: bacterial cell outer membrane; peripheral membrane protein (by similarity). or: secreted protein (by similarity).	extracellular
P16470	RCI2_ECOLI		cytoplasm
P16477	PNDA2_ECOLI		inner membrane
P16528	ICLR_ECOLI		cytoplasm
P16530	PELX_ERWCA		extracellular
P16532	LKB1A_PASHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P16533	LKC1_PASHA	cytoplasm (potential).	cytoplasm
P16534	HLYD_PASHA	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P16536	14KD_RHOSH		inner membrane
P16551	RAFA_ECOLI		cytoplasm
P16553	RAFD_ECOLI		cytoplasm
P16556	NODD_RHILE		cytoplasm
P16565	BCHC_RHOCA		cytoplasm
P16575	BVGS_BORPE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P16588	PROA_VIBAL		extracellular
P16624	17KD_RICPR	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P16625	CH60_RICTS	cytoplasm (by similarity).	cytoplasm
P16626	CH10_RICTS	cytoplasm (by similarity).	cytoplasm
P16630	GUNS_PECCC		extracellular
P16635	LGUL_PSEPU		extracellular
P16640	CAMA_PSEPU		cytoplasm
P16656	YCJE_SALTY		cytoplasm
P16676	CYSA_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P16677	PHNC_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P16678	PHNK_ECOLI		inner membrane
P16679	PHNL_ECOLI		inner membrane
P16681	PHNB_ECOLI		cytoplasm
P16682	PHND_ECOLI	periplasm (potential).	periplasm
P16684	PHNF_ECOLI		cytoplasm
P16685	PHNG_ECOLI		cytoplasm
P16686	PHNH_ECOLI		cytoplasm
P16687	PHNI_ECOLI		cytoplasm
P16688	PHNJ_ECOLI		cytoplasm
P16689	PHNM_ECOLI		extracellular
P16690	PHNN_ECOLI		cytoplasm
P16691	PHNO_ECOLI		cytoplasm

P16692	PHNP_ECOLI		cytoplasm
P16694	YJDA_ECOLI		nucleoid
P16703	CYSM_ECOLI		cytoplasm
P16706	TRPB_ACICA		cytoplasm
P16855	VNFD_AZOVI		cytoplasm
P16856	VNFK_AZOVI		cytoplasm
P16857	VNFG_AZOVI		periplasm
P16897	BLP4_PSEAE		extracellular
P16916	RHSA_ECOLI		inner membrane
P16917	RHSB_ECOLI		inner membrane
P16918	RHSC_ECOLI		inner membrane
P16919	RHSD_ECOLI		inner membrane
P16923	TRPF_ACICA		cytoplasm
P16926	MREC_ECOLI	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P16931	DGDR_BURCE		cytoplasm
P16932	DGDA_BURCE		cytoplasm
P16939	YIS1_SHISO		cytoplasm
P16940	YIS2_SHISO		cytoplasm
P16942	T629_SHISO		cytoplasm
P16943	YIS5_SHISO		cytoplasm
P16944	ISTA_SHISO		cytoplasm
P16966	TTR_PSESZ		cytoplasm
P17051	ALKS_PSEOL		cytoplasm
P17054	CRTI_RHOCA		cytoplasm
P17055	CRTA_RHOCA		cytoplasm
P17056	CRTB_RHOCA		cytoplasm
P17057	CRTK_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P17058	CRTC_RHOCA		cytoplasm
P17059	CRTD_RHOCA		cytoplasm
P17060	CRTE_RHOCA		cytoplasm
P17061	CRTF_RHOCA		cytoplasm
P17086	URE1_PROMI	cytoplasm (by similarity).	cytoplasm
P17087	URE2_PROMI	cytoplasm (by similarity).	cytoplasm
P17088	URE3_PROMI	cytoplasm (by similarity).	cytoplasm
P17089	URED_PROMI		cytoplasm
P17090	UREE_PROMI	cytoplasm (by similarity).	cytoplasm
P17091	UREF_PROMI		cytoplasm
P17109	MEND_ECOLI		cytoplasm
P17115	GUTQ_ECOLI		cytoplasm
P17117	NFSA_ECOLI		cytoplasm
P17127	PTFAH_SALTY	cytoplasm (by similarity).	cytoplasm
P17160	RP5M_AZOVI		cytoplasm
P17161	RP5M_KLEOX		cytoplasm
P17163	YPTO_KLEOX		cytoplasm
P17222	T1SP_ECOLI		cytoplasm
P17223	PRRC_ECOLI		extracellular
P17224	T1RP_ECOLI		cytoplasm
P17239	MERA_THIFE		cytoplasm
P17262	DMPB_PSEUF		cytoplasm
P17263	RP541_RHIME		cytoplasm
P17265	RP5M_RHIME		cytoplasm
P17295	MPC1_RALEU		cytoplasm

P17296	MPC2_RALEU		cytoplasm
P17297	BPHC_PSES1		cytoplasm
P17303	NIFH_AZOBR		cytoplasm
P17410	CHBR_ECOLI		cytoplasm
P17411	CHBF_ECOLI		cytoplasm
P17412	FRDA_WOLSU		inner membrane
P17416	FMZD_BACNO		extracellular
P17417	FMZH_BACNO		extracellular
P17418	FIMCD_BACNO		fimbrium
P17419	FIMCH_BACNO		fimbrium
P17420	FIMDD_BACNO		extracellular
P17421	FIMDH_BACNO		extracellular
P17422	CLPB_BACNO	cytoplasm (probable).	cytoplasm
P17424	RIR4_SALTY		cytoplasm
P17430	ICLR_SALTY		cytoplasm
P17434	NIFB_RHOCA		cytoplasm
P17435	YNF1_RHOCA		extracellular
P17445	BETB_ECOLI		cytoplasm
P17446	BETI_ECOLI		cytoplasm
P17449	VRP1_SALCH		extracellular
P17450	VRP2_SALCH		extracellular
P17452	TOXA_PASMU	cytoplasm (probable).	inner membrane
P17479	RECA_SERMA	cytoplasm (by similarity).	cytoplasm
P17492	REPA_NEIGO		cytoplasm
P17531	RPOD_MYXXA		cytoplasm
P17548	BPHD_PSES1		cytoplasm
P17550	SODM_RALME		periplasm
P17551	CHRA_RALME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P17596	FRDB_WOLSU		cytoplasm
P17611	NODG_AZOBR		cytoplasm
P17738	X192_ECOLI		cytoplasm
P17743	T2C2_HAEIN		extracellular
P17744	MTHC_HAEIN		cytoplasm
P17750	CATA_SALTY		periplasm
P17791	VIRB1_AGRT5		cytoplasm
P17792	VIRB2_AGRT5		extracellular
P17793	VIRB3_AGRT5		inner membrane
P17794	VIRB4_AGRT5		cytoplasm
P17795	VIRB5_AGRT5		fimbrium
P17796	VIRB6_AGRT5		cytoplasm
P17798	VIRB8_AGRT5		cytoplasm
P17799	VIRB9_AGRT5	bacterial cell membrane; peripheral membrane protein (by similarity).	extracellular
P17800	VIRBA_AGRT5		cytoplasm
P17802	MUTY_ECOLI		cytoplasm
P17811	COLY_YERPE	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P17822	FMAB_BACNO		extracellular
P17823	FMA1_BACNO		extracellular
P17824	FMA2_BACNO		extracellular
P17825	FMAE_BACNO		extracellular
P17826	FMAF_BACNO		extracellular

P17827	FMAI_BACNO		extracellular
P17834	FIMBI_BACNO		fimbrium
P17836	FMK1_PSEAE		extracellular
P17837	FMCD_PSEAE		extracellular
P17838	FMP1_PSEAE		extracellular
P17845	CYSI_SALTY		cytoplasm
P17846	CYSI_ECOLI		cytoplasm
P17853	CYSH_SALTY	cytoplasm (by similarity).	cytoplasm
P17857	PROA_SERMA	cytoplasm (by similarity).	cytoplasm
P17862	NODC_RHILO	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P17888	PRIA_ECOLI		cytoplasm
P17899	FLBD_CAUCR		cytoplasm
P17908	TRAK5_ECOLI		cytoplasm
P17909	TRAJ6_ECOLI		cytoplasm
P17910	TRAK4_ECOLI		cytoplasm
P17942	LSPA_PSEFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P17952	MURC_ECOLI	cytoplasm (probable).	cytoplasm
P17974	GUN2_RALSO	bacterial cell membrane; lipid-anchor (probable).	extracellular
P17976	PSIA2_ECOLI		cytoplasm
P17983	YIA1_RHISP		inner membrane
P17984	YIA2_RHISP		cytoplasm
P17985	YIA3_RHISP		cytoplasm
P17986	YIA_RHISP		inner membrane
P17993	UBIG_ECOLI		cytoplasm
P17994	YFAA_ECOLI		periplasm
P17998	CEAD_ECOLI		extracellular
P17999	CEA6_ECOLI		extracellular
P18002	IMMM_ECOLI		cytoplasm
P18005	YUBP_ECOLI		cytoplasm
P18009	IPA4_SHIFL		extracellular
P18014	IPA7_SHIFL		extracellular
P18021	REDM_ECOLI		cytoplasm
P18022	SSBP_ECOLI		cytoplasm
P18023	REPZ_ECOLI		nucleoid
P18032	ARTA_ECOLI		inner membrane
P18076	RECA_METFL	cytoplasm (by similarity).	cytoplasm
P18079	HEM1_RHOCA		inner membrane
P18082	FER2_RHOCA		periplasm
P18086	FLAV_DESSA		cytoplasm
P18103	FANC_ECOLI		extracellular
P18127	ICEN_XANCT	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P18128	YPC2_ECOLI		cytoplasm
P18133	PNCB_ECOLI	cytoplasm (by similarity).	cytoplasm
P18148	PSIA1_ECOLI		cytoplasm
P18152	MP13_FRATH		inner membrane
P18196	MINC_ECOLI		inner membrane
P18204	IAAL_PSESS		cytoplasm
P18205	YIAL_PSESS		cytoplasm
P18245	FLAB_CAMCO		flagellar

P18251	BLA1_ENTCL		extracellular
P18275	ARCD_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P18299	LUXA_KRYAS		cytoplasm
P18310	SSBF_ECOLI		cytoplasm
P18314	URE1_KLEAE	cytoplasm (by similarity).	cytoplasm
P18315	URE2_KLEAE	cytoplasm (by similarity).	cytoplasm
P18316	URE3_KLEAE	cytoplasm (by similarity).	cytoplasm
P18318	UREF_KLEAE		cytoplasm
P18319	UREG_KLEAE		cytoplasm
P18335	ARGD_ECOLI	cytoplasm (probable).	cytoplasm
P18351	YPC1_ECOLI		cytoplasm
P18352	YPC3_ECOLI		cytoplasm
P18353	TRBJ_ECOLI		inner membrane
P18355	YPFU_ECOLI		cytoplasm
P18390	YJJA_ECOLI		extracellular
P18393	YBDZ_ECOLI		cytoplasm
P18539	AMPC_SERMA	periplasm (by similarity).	periplasm
P18540	VIRA_AGRT5		inner membrane
P18561	SYRM_RHIME		cytoplasm
P18591	VIRD1_AGRT5		extracellular
P18592	VIRD2_AGRT5		extracellular
P18593	VIRD3_AGRT5		cytoplasm
P18594	VIRD4_AGRT5	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P18607	FPRA_RHOCA		inner membrane
P18641	IMPA_SALTY		cytoplasm
P18642	IMPB_SALTY		cytoplasm
P18767	NDVA_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P18769	FRZE_MYXXA		inner membrane
P18774	FM12_PSEAE		extracellular
P18776	DMSB_ECOLI		periplasm
P18778	TERA_ALCSP		cytoplasm
P18779	TERB_ALCSP		cytoplasm
P18781	TERD_ALCSP		cytoplasm
P18782	TERE_ALCSP		cytoplasm
P18785	HIS5_AZOBR	cytoplasm (by similarity).	cytoplasm
P18787	HIS7_AZOBR	cytoplasm (by similarity).	cytoplasm
P18789	CISY_COXBU		cytoplasm
P18790	HLYD_ACTAC	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P18797	MMOB_METCA		cytoplasm
P18798	MEMB_METCA		cytoplasm
P18809	FINQ_ECOLI		cytoplasm
P18811	MALI_ECOLI		cytoplasm
P18818	HNS_PROVU		cytoplasm
P18843	NADE_ECOLI		cytoplasm
P18896	IGRA_PSES2		cytoplasm
P18913	FLJK_CAUCR		flagellar
P18914	FLJL_CAUCR		flagellar
P18952	PA1_SERLI		extracellular

P18953	GRCA_SERLI		cytoplasm
P18954	PHLB_SERLI		cytoplasm
P18955	HNS_SERMA		cytoplasm
P18957	UVP1_ECOLI		cytoplasm
P19055	NIFE_RHOCA		cytoplasm
P19056	RCEH_RHOCA		inner membrane
P19059	DMPC_PSEUF		cytoplasm
P19066	NIFD_RHISN		periplasm
P19067	NIFK_RHISN		cytoplasm
P19068	NIFH_RHISN		cytoplasm
P19071	PARP_ECOLI		cytoplasm
P19076	DMPD_PSEUF		cytoplasm
P19077	NIFN_RHOCA		periplasm
P19078	NIFX_RHOCA		cytoplasm
P19175	RPOB_PSEPU		cytoplasm
P19176	RPOC_PSEPU		cytoplasm
P19197	LUXD1_PHOLU		cytoplasm
P19247	VVHA_VIBVU		extracellular
P19248	VVHB_VIBVU		extracellular
P19249	HLY1_VIBPA		extracellular
P19250	HLY2_VIBPA		extracellular
P19257	PIV_MORLA		cytoplasm
P19264	EUTB_SALTY		cytoplasm
P19265	EUTC_SALTY		cytoplasm
P19317	NARW_ECOLI		cytoplasm
P19323	FHLA_ECOLI		cytoplasm
P19372	ACP_RHIME	cytoplasm (by similarity).	cytoplasm
P19381	TRBH_ECOLI	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P19390	BEXB1_HAEIN	bacterial cell inner membrane; multi- pass membrane protein (potential).	inner membrane
P19391	BEXB2_HAEIN	bacterial cell inner membrane; multi- pass membrane protein (potential).	inner membrane
P19421	CH60_COXBU	cytoplasm (by similarity).	cytoplasm
P19422	CH10_COXBU	cytoplasm (by similarity).	cytoplasm
P19423	CBEP_COXBU		inner membrane
P19433	RPSB_MYXXA		cytoplasm
P19441	NODS_RHISN		cytoplasm
P19449	BCSA1_ACEXY	bacterial cell inner membrane; multi- pass membrane protein (potential).	inner membrane
P19450	BCSC1_ACEXY	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P19451	BCSD1_ACEXY		inner membrane
P19480	AHPF_SALTY		cytoplasm
P19485	CYA1_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	cytoplasm
P19487	GUNA_XANCP		extracellular
P19520	YAV5_XANCV		extracellular
P19528	FMAJ_BACNO		extracellular
P19543	NIFJ_ENTAG		cytoplasm

P19572	DCDA_PSEAE		cytoplasm
P19592	ISS_ECOLI		inner membrane
P19624	PDXA_ECOLI	cytoplasm (probable).	cytoplasm
P19636	EUTC_ECOLI		cytoplasm
P19650	AAC6_KLEPN		cytoplasm
P19685	SODF_COXBU		periplasm
P19688	AROA_YEREN	cytoplasm (probable).	cytoplasm
P19689	SERC_YEREN	cytoplasm (by similarity).	cytoplasm
P19690	RECA_BURCE	cytoplasm (by similarity).	cytoplasm
P19704	T1SA_ECOLI		cytoplasm
P19705	T1SE_ECOLI		cytoplasm
P19729	DMPK_PSEUF		cytoplasm
P19730	DMPL_PSEUF		cytoplasm
P19731	DMPM_PSEUF		cytoplasm
P19732	DMPN_PSEUF		cytoplasm
P19733	DMPO_PSEUF		cytoplasm
P19734	DMPP_PSEUF		cytoplasm
P19763	INSA_SHIFL		cytoplasm
P19765	INSB_SHIFL		cytoplasm
P19766	INSB_SHISO		cytoplasm
P19767	INSA7_ECOLI		cytoplasm
P19768	INSJ_ECOLI		inner membrane
P19769	INSK_ECOLI		cytoplasm
P19770	YI5C_ECOLI		inner membrane
P19776	INSC_ECOLI		cytoplasm
P19777	INSD_ECOLI		cytoplasm
P19778	YI23_ECOLI		inner membrane
P19779	YI24_ECOLI		inner membrane
P19786	AROA_SALTI	cytoplasm (probable).	cytoplasm
P19816	RFAI_SALTY		cytoplasm
P19817	RFAJ_SALTY		cytoplasm
P19828	ANGR_VIBAN		cytoplasm
P19829	SAST_VIBAN		extracellular
P19833	LIP1_MORS1		extracellular
P19839	LUXA1_PHOLU		cytoplasm
P19840	LUXB1_PHOLU		cytoplasm
P19844	NOSF_PSEST	bacterial cell membrane; peripheral membrane protein (potential).	inner membrane
P19871	3BHD_COMTE		cytoplasm
P19906	NTRB_VIBAL		inner membrane
P19907	LUXA_VIBFI		cytoplasm
P19908	LUXB_VIBFI		cytoplasm
P19910	PYRB_SERMA		cytoplasm
P19911	F16P1_RALEU		cytoplasm
P19912	F16P2_RALEU		cytoplasm
P19913	DCML_HYDPS		periplasm
P19914	DCMM_HYDPS		cytoplasm
P19915	DCMS_HYDPS		cytoplasm
P19919	DCML_OLICA		periplasm
P19920	DCMM_OLICA		cytoplasm
P19921	DCMS_OLICA		cytoplasm
P19922	HYIN2_BRAJA		extracellular
P19923	KPPR2_RALEU		cytoplasm
P19924	KPPR1_RALEU		cytoplasm
P19925	ENTD_ECOLI		periplasm

P19930	HYAD_ECOLI		extracellular
P19931	HYAE_ECOLI		cytoplasm
P19932	HYAF_ECOLI		cytoplasm
P19936	PYRI_SERMA		cytoplasm
P20083	PARE_ECOLI		cytoplasm
P20085	MOBL_THIFE		cytoplasm
P20086	MOBS_THIFE		cytoplasm
P20087	YML1_THIFE		extracellular
P20088	YML2_THIFE		cytoplasm
P20089	YML3_THIFE		cytoplasm
P20090	YMS1_THIFE		inner membrane
P20092	AAC6_SERMA		cytoplasm
P20098	RCSA_ERWAM		cytoplasm
P20099	BISC_ECOLI		periplasm
P20102	MERD_SHIFL		cytoplasm
P20343	CYSX_ECOLI		cytoplasm
P20356	REPJ_ECOLI		cytoplasm
P20368	ADH1_ZYMMO		cytoplasm
P20370	DC4C_ACIAD		cytoplasm
P20371	PCXA_ACIAD		cytoplasm
P20372	PCXB_ACIAD		cytoplasm
P20401	OPGH_PSESY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P20402	ROB1_AGRRH		extracellular
P20403	ROC1_AGRRH		extracellular
P20418	DESR_DESVH		inner membrane
P20442	DNAK_CAUCR		cytoplasm
P20469	ICEA_PANAN	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P20506	THD1_SALTY		cytoplasm
P20574	TRPD_PSEAE		cytoplasm
P20575	TRPD_PSEPU		cytoplasm
P20576	TRPG_PSEAE		cytoplasm
P20577	TRPC_PSEAE		cytoplasm
P20578	TRPC_PSEPU		cytoplasm
P20579	TRPE_PSEPU		cytoplasm
P20580	TRPE_PSEAE		cytoplasm
P20581	Y1000_PSEAE		cytoplasm
P20582	PQSD_PSEAE	cytoplasm (probable).	cytoplasm
P20586	PHHY_PSEAE		cytoplasm
P20588	T2F1_HAEIN		extracellular
P20589	MTH3_HAEAE		cytoplasm
P20590	MTH1_HAEIN		cytoplasm
P20605	FIC_ECOLI		cytoplasm
P20621	NIFK_BRAJA		cytoplasm
P20624	FER3_RHOCA		periplasm
P20629	NIFQ_RHOCA		cytoplasm
P20657	FMI_MORBO		extracellular
P20665	PIV_MORBO		cytoplasm
P20666	TFPB_MORBO		inner membrane
P20669	NODD_AZOCA		cytoplasm
P20707	ODO1_AZOVI		cytoplasm
P20708	ODO2_AZOVI		inner membrane
P20713	ATSA_KLEAE	periplasm (by similarity).	periplasm

P20714	ATSB_KLEAE	cytoplasm (by similarity).	cytoplasm
P20725	GSPB_KLEPN		inner membrane
P20751	FIC_SALTY		cytoplasm
P20753	PPIA_SALTY	periplasm (by similarity).	periplasm
P20861	FANG_ECOLI		extracellular
P20862	FANH_ECOLI		fimbrium
P20901	CISY_ACEAC		cytoplasm
P20902	CISY_ACIAN		cytoplasm
P20906	MDLC_PSEPU		cytoplasm
P20921	FRDB_PROVU		cytoplasm
P20922	FRDA_PROVU		periplasm
P20923	FRDC_PROVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P20924	FRDD_PROVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P20926	HYBF_PROVU		cytoplasm
P20927	YFRC_PROVU		cytoplasm
P20928	SUGE_PROVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P20932	MDLB_PSEPU		cytoplasm
P20960	NRLA_ALCFA		cytoplasm
P21152	RECA_NEIGO	cytoplasm (by similarity).	cytoplasm
P21156	CYSD_ECOLI		cytoplasm
P21159	PDXH_MYXXA		cytoplasm
P21161	DCMA_METDI	cytoplasm (probable).	cytoplasm
P21165	PEPQ_ECOLI		extracellular
P21169	DCOR_ECOLI		cytoplasm
P21177	FADB_ECOLI		cytoplasm
P21178	CEA1_SHISO	bacterial cell membrane; multi-pass membrane protein (potential).	extracellular
P21179	CATE_ECOLI	cytoplasm (probable).	periplasm
P21184	FLICA_PSEAE		flagellar
P21189	DPO2_ECOLI		cytoplasm
P21193	GLNB_AZOBR		cytoplasm
P21207	LCRH_YERPE		cytoplasm
P21211	YLC6_YEREN		inner membrane
P21258	OGL_DICD3	periplasm (potential).	periplasm
P21267	GSA_SALTY	cytoplasm (potential).	cytoplasm
P21295	FLAF_CAUCR		inner membrane
P21296	FLBA_CAUCR		extracellular
P21297	FLBT_CAUCR		cytoplasm
P21302	OTC_NEIGO	cytoplasm (probable).	cytoplasm
P21308	LUXR_VIBHA		cytoplasm
P21309	LUXD1_PHOLE		cytoplasm
P21310	HUTH_PSEPU	cytoplasm (potential).	cytoplasm
P21311	DMA7_ECOLI		cytoplasm
P21312	YR72_ECOLI		inner membrane
P21313	YR73_ECOLI		inner membrane
P21314	YR75_ECOLI		inner membrane
P21315	YR7A_ECOLI		inner membrane
P21316	YR7B_ECOLI		inner membrane
P21318	YR7D_ECOLI		inner membrane

P21319	YR7E_ECOLI		inner membrane
P21320	DINIR_ECOLI		cytoplasm
P21321	YR7G_ECOLI		inner membrane
P21322	YR7H_ECOLI		inner membrane
P21323	YR7I_ECOLI		cytoplasm
P21324	YR7J_ECOLI		inner membrane
P21325	RT67_ECOLI		extracellular
P21337	TETR5_ECOLI		cytoplasm
P21361	YCIG_ECOLI		extracellular
P21362	YCIF_ECOLI		cytoplasm
P21363	YCIE_ECOLI		cytoplasm
P21367	YCAC_ECOLI		cytoplasm
P21369	PNCA_ECOLI		cytoplasm
P21394	XYLA_PSEPU		cytoplasm
P21395	XYLM_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P21408	FBPA_SERMA	periplasm (probable).	periplasm
P21409	FBPB_SERMA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P21410	FBPC_SERMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P21413	FM98_ECOLI		extracellular
P21418	YDFC_ECOLI		cytoplasm
P21423	YAC4_KLEPN		inner membrane
P21424	TNR4_KLEPN		cytoplasm
P21437	YGGF_ECOLI		cytoplasm
P21454	VRP2_SALTY		extracellular
P21455	VRP1_SALTY		extracellular
P21482	DSBB1_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P21499	RNR_ECOLI		cytoplasm
P21507	SRMB_ECOLI		cytoplasm
P21514	YAHA_ECOLI		cytoplasm
P21515	ACPH_ECOLI		cytoplasm
P21559	P34_RICRI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P21578	LUXY_VIBFI		cytoplasm
P21599	KPYK2_ECOLI		cytoplasm
P21622	ILVH_SALTY		cytoplasm
P21629	BRAF_PSEAE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P21630	BRAG_PSEAE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P21631	SUMT_PSEDE		cytoplasm
P21632	COBB_PSEDE		cytoplasm
P21633	COBC_PSEDE	cytoplasm (by similarity).	cytoplasm
P21634	COBD_PSEDE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

P21635	COBE_PSEDE		cytoplasm
P21636	COBF_PSEDE		cytoplasm
P21637	COBG_PSEDE		cytoplasm
P21638	COBH_PSEDE		cytoplasm
P21639	COBI_PSEDE		cytoplasm
P21640	COBJ_PSEDE		cytoplasm
P21645	LPXD_ECOLI		cytoplasm
P21647	MRKC_KLEPN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P21649	MRKE_KLEPN		cytoplasm
P21683	CRTB_PANAN		cytoplasm
P21684	CRTE_PANAN		cytoplasm
P21685	CRTI_PANAN		cytoplasm
P21686	CRTX_PANAN		cytoplasm
P21687	CRTY_PANAN		cytoplasm
P21688	CRTZ_PANAN		cytoplasm
P21689	TRPE_PSESS		cytoplasm
P21693	DBPA_ECOLI	cytoplasm (probable).	cytoplasm
P21725	TKTC_RALEU		extracellular
P21726	TKTP_RALEU		extracellular
P21763	T2R1_RHOSH		extracellular
P21829	YBHA_ECOLI		inner membrane
P21863	FKBX_PSEFL		cytoplasm
P21866	KDPE_ECOLI	cytoplasm (probable).	cytoplasm
P21867	RAFR_ECOLI		cytoplasm
P21877	ACSA1_ACEXY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P21893	RECJ_ECOLI		cytoplasm
P21907	G6PD_ZYMMO		cytoplasm
P21908	GLK_ZYMMO	cytoplasm (by similarity).	cytoplasm
P21909	EDD_ZYMMO		cytoplasm
P21920	COBK_PSEDE		cytoplasm
P21921	COBL_PSEDE		cytoplasm
P21922	COBM_PSEDE		cytoplasm
P22008	PROC_PSEAE	cytoplasm (by similarity).	cytoplasm
P22037	ATMC_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P22038	TRMA_SALTY		cytoplasm
P22076	DESR_DESDE		inner membrane
P22088	LIP_BURCE		extracellular
P22089	LIFO1_BURCE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
P22095	TRPA_VIBPA		cytoplasm
P22096	TRPD_VIBPA		cytoplasm
P22097	TRPB1_VIBPA		cytoplasm
P22098	TRPC_VIBPA		cytoplasm
P22099	TRPE_VIBPA		cytoplasm
P22101	TRPG_VIBPA		cytoplasm
P22104	TCTD_SALTY		cytoplasm
P22106	ASNB_ECOLI		cytoplasm
P22118	GYRB_NEIGO		cytoplasm
P22186	MRAZ_ECOLI		inner membrane

P22188	MURE_ECOLI	cytoplasm (probable).	cytoplasm
P22235	BEXB3_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P22236	BEXD_HAEIN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P22251	FLA2_CAMJE		flagellar
P22252	FLB2_CAMJE		flagellar
P22253	PNCB_SALTY	cytoplasm (by similarity).	cytoplasm
P22255	CYSQ_ECOLI	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (probable).	cytoplasm
P22256	GABT_ECOLI		cytoplasm
P22260	CLP_XANCP		cytoplasm
P22299	ARO_A_SALGL	cytoplasm (probable).	cytoplasm
P22305	DNAJ_CAUCR	cytoplasm (probable).	cytoplasm
P22308	REPY_ECOLI		cytoplasm
P22333	ADD_ECOLI		extracellular
P22359	HTPG_VIBCH	cytoplasm (by similarity).	cytoplasm
P22390	BLAC_CITDI		extracellular
P22391	BLO1_KLEOX		extracellular
P22493	UMUD_SALTY		cytoplasm
P22494	UMUC_SALTY		cytoplasm
P22519	CVAA_ECOLI	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P22521	IMMV_ECOLI		cytoplasm
P22525	YCBB_ECOLI		inner membrane
P22537	NOLA_BRAJA		cytoplasm
P22558	IMM1_SHISO		cytoplasm
P22564	RIHC_ECOLI		extracellular
P22566	MAUJ_PARDE		cytoplasm
P22567	ARGA_PSEAE	cytoplasm (probable).	cytoplasm
P22586	FLIO_ECOLI	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P22606	FLGH_CAUCR	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
P22609	PILC_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P22615	CAT2_ECOLI		cytoplasm
P22616	CAT2_HAEIN		cytoplasm
P22634	MURI_ECOLI		cytoplasm
P22635	PCYA_PSEPA		cytoplasm
P22636	PCYB_PSEPA		cytoplasm
P22643	DHLA_XANAU		cytoplasm
P22645	YDH1_XANAU		cytoplasm
P22706	TRAI2_ECOLI		cytoplasm
P22707	FINO_ECOLI		outer membrane
P22709	TRAX1_ECOLI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane

P22710	TRAX2_ECOLI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P22713	GAL1_SALTY	cytoplasm (potential).	cytoplasm
P22714	GAL7_SALTY		cytoplasm
P22715	GALE_SALTY		cytoplasm
P22731	LIVF_ECOLI		inner membrane
P22751	PELX_ERWCH		extracellular
P22759	BFR_AZOVI		cytoplasm
P22773	HUTC_PSEPU		cytoplasm
P22780	F16P2_RHOSH		cytoplasm
P22782	CAT_CAMCO		cytoplasm
P22787	YIFB_ECOLI		cytoplasm
P22824	SCRK_VIBAL		cytoplasm
P22825	PTSBC_VIBAL	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P22834	YIDD_PROMI		inner membrane
P22835	RNPA_PROMI		extracellular
P22837	DNAA_PROMI	cytoplasm (by similarity).	cytoplasm
P22838	DPO3B_PROMI	cytoplasm (by similarity).	cytoplasm
P22839	RECF_PROMI	cytoplasm (potential).	cytoplasm
P22848	5NTD_VIBPA	bacterial cell outer membrane; lipid-anchor (probable).	periplasm
P22849	RBL1A_CHRVI		cytoplasm
P22850	RBS1_CHRVI		cytoplasm
P22859	RBL1B_CHRVI		cytoplasm
P22860	RBS2_CHRVI		cytoplasm
P22862	ESTE_PSEFL		cytoplasm
P22867	MMOD_METCA		cytoplasm
P22868	MMOC_METCA		cytoplasm
P22869	MEMA_METCA		cytoplasm
P22871	CRTI_ESCVU		cytoplasm
P22872	CRTB_ESCVU		cytoplasm
P22873	CRTE_ESCVU		cytoplasm
P22881	RP54_RHISN		cytoplasm
P22882	17KD_RICTY	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P22899	MOBC_THIFE		cytoplasm
P22900	MOBD_THIFE		cytoplasm
P22901	MOBE_THIFE		cytoplasm
P22902	YME1_THIFE		cytoplasm
P22903	YME2_THIFE		cytoplasm
P22904	YME3_THIFE		cytoplasm
P22905	MERC_THIFE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P22921	NIFH_RHORU		cytoplasm
P22930	BEXC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P22990	RIBF_PSEFL		cytoplasm
P22993	PARC4_ECOLI		cytoplasm
P22994	YPR5_ECOLI		cytoplasm
P22995	PARD4_ECOLI		cytoplasm
P22996	PAA4_ECOLI		cytoplasm

P22997	PAB4_ECOLI		extracellular
P23001	FLAV_AZOCH		cytoplasm
P23003	TRMA_ECOLI		cytoplasm
P23010	KPPR2_RHOSH		cytoplasm
P23011	RBL_XANFL		extracellular
P23012	RBS_XANFL		cytoplasm
P23013	CFXQ_XANFL		cytoplasm
P23014	F16P_XANFL		cytoplasm
P23015	KPPR_XANFL		cytoplasm
P23030	XYNB_PSEFL		extracellular
P23070	RT86_ECOLI		cytoplasm
P23071	RT65_MYXXA		cytoplasm
P23072	RT16_MYXXA		cytoplasm
P23099	XYLX_PSEPU		cytoplasm
P23100	XYLY_PSEPU		cytoplasm
P23101	XYLZ_PSEPU		cytoplasm
P23102	XYLL_PSEPU		cytoplasm
P23103	FERX_PSEPU		cytoplasm
P23105	XYLG_PSEPU		cytoplasm
P23106	XYLF_PSEPU		cytoplasm
P23107	XYLJ_PSEPU		cytoplasm
P23113	LUXC_PHOLU		cytoplasm
P23119	NIFM_AZOCH		cytoplasm
P23120	NIFS_AZOCH		cytoplasm
P23121	NIFU_AZOCH		cytoplasm
P23122	NIFV_AZOCH		cytoplasm
P23123	NIFW_AZOCH		cytoplasm
P23124	NIFZ_AZOCH		cytoplasm
P23133	TODF_PSEPU		cytoplasm
P23134	CYB_RHORU		inner membrane
P23135	CY1_RHORU		inner membrane
P23138	YPE1_RHORU		cytoplasm
P23139	YPE2_RHORU		cytoplasm
P23145	NIFP_AZOCH		cytoplasm
P23146	LUXA2_PHOLU		cytoplasm
P23147	LUXB2_PHOLU		cytoplasm
P23148	LUXD2_PHOLU		cytoplasm
P23177	YNIW_AZOCH		extracellular
P23181	AACC1_PSEAE		cytoplasm
P23182	CIR1_CITFR		extracellular
P23183	CIR2_CITFR		extracellular
P23184	MCBB_ECOLI	cytoplasm (potential).	cytoplasm
P23185	MCBC_ECOLI	cytoplasm (potential).	cytoplasm
P23186	MCBD_ECOLI	cytoplasm (potential).	cytoplasm
P23190	NODD3_RHIME		cytoplasm
P23191	T2M2_MORBO		extracellular
P23192	MTM2_MORBO		cytoplasm
P23205	Y2024_PSEAE		cytoplasm
P23221	FIXJ_BRAJA	cytoplasm (probable).	cytoplasm
P23234	DCIP_ENTCL		cytoplasm
P23240	ALDH_VIBCH		extracellular
P23247	DHAS_VIBCH		cytoplasm
P23256	MALY_ECOLI		cytoplasm
P23262	NHG1_PSEPU		cytoplasm
P23263	FERN_PSEPU		cytoplasm
P23300	AMOA_AERHY		cytoplasm

P23302	IHFA_SERMA		cytoplasm
P23303	IHFB_SERMA		cytoplasm
P23305	YIGA_ECOLI		inner membrane
P23315	TRPE_ACICA		cytoplasm
P23325	ARPA_ECOLI		cytoplasm
P23331	KITH_ECOLI	cytoplasm (potential).	cytoplasm
P23354	K1PF_XANCP		cytoplasm
P23364	CAT4_AGRT5		cytoplasm
P23367	MUTL_ECOLI		cytoplasm
P23386	K1PF_RHOCA		cytoplasm
P23388	PTFAX_RHOCA	cytoplasm (potential).	cytoplasm
P23460	LHG2_RHOCA		inner membrane
P23461	PUCD_RHOCA		inner membrane
P23476	TCPB_VIBCH		outer membrane
P23481	HYFA_ECOLI		cytoplasm
P23484	FECI_ECOLI		cytoplasm
P23522	GARL_ECOLI		cytoplasm
P23524	GLXK2_ECOLI		cytoplasm
P23538	PPSA_ECOLI		cytoplasm
P23587	FLMC2_ECOLI		cytoplasm
P23621	PHOR_PSEAE		inner membrane
P23657	T2P2_PROVU		extracellular
P23679	GLND_SALTY		cytoplasm
P23702	LKTB_ACTAC	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P23703	NODI_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P23716	NOLE_RHILP	periplasm (potential).	periplasm
P23717	NOLP_RHILP		cytoplasm
P23718	NODD1_RHILP		cytoplasm
P23719	NODD2_RHILP		cytoplasm
P23720	NODD3_RHILP		cytoplasm
P23744	SODF_METJ		periplasm
P23747	ALGB_PSEAE		cytoplasm
P23774	FAPR_ECOLI		cytoplasm
P23831	SAMA_SALTY		cytoplasm
P23832	SAMB_SALTY		cytoplasm
P23835	TETO_CAMCO		cytoplasm
P23836	PHOP_ECOLI	cytoplasm (probable).	cytoplasm
P23839	YICC_ECOLI		nucleoid
P23840	DIND_ECOLI		cytoplasm
P23841	XAPR_ECOLI		cytoplasm
P23842	YFEA_ECOLI		inner membrane
P23845	CYSN_ECOLI		cytoplasm
P23857	PSPE_ECOLI		cytoplasm
P23862	PRIC_ECOLI		cytoplasm
P23873	HIPB_ECOLI		cytoplasm
P23874	HIPA_ECOLI		fimbrium
P23878	FEPC_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P23882	FMT_ECOLI		cytoplasm
P23883	PUUC_ECOLI		cytoplasm
P23890	CADC_ECOLI		cytoplasm

P23893	GSA_ECOLI	cytoplasm (potential).	cytoplasm
P23898	NLPC_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P23908	ARGE_ECOLI	cytoplasm (probable).	extracellular
P23909	MUTS_ECOLI		cytoplasm
P23917	MAK_ECOLI		cytoplasm
P23924	MGLA_SALTY	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P23926	ANR_PSEAE		cytoplasm
P23954	BLO2_KLEOX		extracellular
P23995	LCRH_YERPS		cytoplasm
P24019	TAGA_VIBCH	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P24038	NIRT_PSEST	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P24039	NIRC_PSEST	periplasm (by similarity).	periplasm
P24042	REP5_ECOLI		cytoplasm
P24060	GLYA_BRAJA		cytoplasm
P24069	HAD1_PSEUC		cytoplasm
P24070	HAD2_PSEUC		cytoplasm
P24074	RECA_RHILV	cytoplasm (by similarity).	cytoplasm
P24112	PLYD_ERWCA		cytoplasm
P24113	LUXA_PHOPO		cytoplasm
P24126	FUSA_BURCE	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P24127	FUSB_BURCE		extracellular
P24128	FUSC_BURCE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P24129	FUSD_BURCE		inner membrane
P24130	FUSE_BURCE		cytoplasm
P24144	NODJ_RHILT	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P24145	NODT_RHILT	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P24148	YNB1_RHIET		cytoplasm
P24149	YNB2_RHIET		outer membrane
P24151	NODC_RHILP	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P24162	ECHH_RHOCA		cytoplasm
P24169	DCOS_ECOLI		cytoplasm
P24173	RFAC_ECOLI		cytoplasm
P24174	MANC_ECOLI		cytoplasm
P24175	MANB_ECOLI		cytoplasm
P24178	YFFB_ECOLI		cytoplasm
P24180	ACRE_ECOLI	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P24182	ACCC_ECOLI		cytoplasm
P24186	FOLD_ECOLI		cytoplasm
P24188	YCEA_ECOLI		inner membrane
P24192	HYPD_ECOLI		extracellular
P24193	HYPE_ECOLI		cytoplasm

P24197	YGID_ECOLI		cytoplasm
P24200	MCRA_ECOLI		extracellular
P24202	MRR_ECOLI		cytoplasm
P24203	YJIA_ECOLI		cytoplasm
P24205	MSBB_ECOLI	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P24211	RHSE_ECOLI		cytoplasm
P24215	UXUA_ECOLI		cytoplasm
P24218	INTD_ECOLI		cytoplasm
P24220	PYRF_MYXXA		cytoplasm
P24224	ACPS_ECOLI	cytoplasm (probable).	cytoplasm
P24228	PBP4_ECOLI	periplasm (potential).	inner membrane
P24230	RECG_ECOLI		cytoplasm
P24238	YEBB_ECOLI		cytoplasm
P24240	ASCB_ECOLI		cytoplasm
P24242	ASCG_ECOLI		cytoplasm
P24252	YBGA_ECOLI		cytoplasm
P24254	YZPU_ECOLI		cytoplasm
P24255	RP54_ECOLI		cytoplasm
P24273	LUXG_VIBFI		cytoplasm
P24290	FNRN_RHILV		cytoplasm
P24324	Y352_HAEIN		cytoplasm
P24325	GALE_HAEIN		cytoplasm
P24326	Y350_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P24398	VRP4_SALTY		extracellular
P24418	VRP1_SALDU		extracellular
P24419	VRP2_SALDU		extracellular
P24421	VSDF_SALDU		extracellular
P24423	YVNF_AZOCH		cytoplasm
P24426	NIFA_RHILT		cytoplasm
P24427	NIFB_RHILT		cytoplasm
P24466	CPXC_AGRTU		cytoplasm
P24467	CPXD_AGRTU		cytoplasm
P24484	LIP2_MORS1		extracellular
P24497	AROA_KLEPN	cytoplasm (probable).	cytoplasm
P24508	SCRR_VIBAL		cytoplasm
P24509	YSCR_VIBAL		cytoplasm
P24517	RADA_SALTY		cytoplasm
P24520	PNUC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P24536	T402_BURCE		cytoplasm
P24537	YI22_BURCE		inner membrane
P24538	YI23_BURCE		inner membrane
P24542	RECA_METCL	cytoplasm (by similarity).	cytoplasm
P24543	RECA_RHIET	cytoplasm (by similarity).	cytoplasm
P24546	T2A1_ACICA		extracellular
P24550	RCP_VIBCH		cytoplasm
P24554	RADA_ECOLI		cytoplasm
P24555	PTRB_ECOLI		cytoplasm
P24559	PILT_PSEAE	cytoplasm (probable).	inner membrane
P24562	Y394_PSEAE		cytoplasm
P24575	TRA6_BURCE		cytoplasm

P24576	YI62_BURCE		inner membrane
P24577	YI71_BURCE		cytoplasm
P24578	YI72_BURCE		inner membrane
P24579	YI73_BURCE		inner membrane
P24580	YI74_BURCE		cytoplasm
P24581	MTNX_NEILA		cytoplasm
P24582	MTN3_NEILA		cytoplasm
P24584	KPSM5_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P24607	T801_PSESH		inner membrane
P24617	T2P2_NEIGO		extracellular
P24640	LIP3_MORS1		extracellular
P24670	AROD_SALTI		cytoplasm
P24672	RBL_ALVHS		extracellular
P24682	RBS_ALVHS		cytoplasm
P24687	GSPH_ERWCH		inner membrane
P24688	GSPI_ERWCH		inner membrane
P24689	GSPJ_ERWCH		inner membrane
P24690	GSPK_ERWCH	bacterial cell inner membrane (potential).	inner membrane
P24693	YHBG_THIFE		inner membrane
P24695	RP54_THIFE		cytoplasm
P24696	YHUT_PSEPU		cytoplasm
P24703	DAPB_COXBU	cytoplasm (by similarity).	cytoplasm
P24718	RECF_ACTPL	cytoplasm (potential).	cytoplasm
P24735	AMPC_PSEAE	periplasm (by similarity).	periplasm
P24791	PME2_RALSO		extracellular
P24900	RECF_SALTY	cytoplasm (potential).	cytoplasm
P24908	TRPO_PSEAE		cytoplasm
P24931	RUBY_DESVH	cytoplasm (potential).	cytoplasm
P25015	YNIX_RHOCA		cytoplasm
P25016	HYIN2_AGRVI		extracellular
P25017	TR2N_AGRVI		cytoplasm
P25018	IPT2_AGRVI		cytoplasm
P25019	6B2_AGRVI		cytoplasm
P25026	PRXC_PSEPY		cytoplasm
P25060	GSPL_PSEAE	bacterial cell inner membrane (probable).	inner membrane
P25061	GSPM_PSEAE	bacterial cell inner membrane (probable).	inner membrane
P25077	MDH_SALTY		cytoplasm
P25078	PHR_SALTY		cytoplasm
P25081	MANA_SALTY	cytoplasm (probable).	cytoplasm
P25082	LUXP_PHOPO		cytoplasm
P25084	LASR_PSEAE		cytoplasm
P25177	GLMM_HELPY		cytoplasm
P25178	LSPA_HELPY	bacterial cell membrane; multi-pass membrane protein (by similarity).	extracellular
P25195	NODM2_RHIME	cytoplasm (by similarity).	cytoplasm
P25196	NOLF_RHIME		inner membrane
P25197	NOLG_RHIME		cytoplasm
P25200	NODN_RHIME		cytoplasm
P25201	MTA1_ACICA		cytoplasm
P25237	T2K1_KLEPN		extracellular

P25238	MTK1_KLEPN		cytoplasm
P25239	T257_ECOLI		extracellular
P25240	MT57_ECOLI		cytoplasm
P25254	Y392_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P25275	LIP_PSES5		extracellular
P25276	LIFO_PSES5	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
P25282	MTG1_HAEGA		cytoplasm
P25283	MTG2_HAEGA		cytoplasm
P25287	Y57M_ECOLI		cytoplasm
P25313	NIFD_AZOBR		periplasm
P25314	NIFK_AZOBR		cytoplasm
P25315	NIFY_AZOBR		cytoplasm
P25316	YNIF_AZOBR		cytoplasm
P25393	CFAD_ECOLI		cytoplasm
P25397	TEHB_ECOLI	cytoplasm (potential).	cytoplasm
P25438	YI61_XANCV		cytoplasm
P25468	PYRD_SALTY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P25516	ACON1_ECOLI		cytoplasm
P25519	HFLX_ECOLI		cytoplasm
P25524	CODA_ECOLI		extracellular
P25526	GABD_ECOLI		cytoplasm
P25531	RADC_ECOLI		cytoplasm
P25534	UBIH_ECOLI		cytoplasm
P25535	VISC_ECOLI		cytoplasm
P25536	YHDE_ECOLI	cytoplasm (potential).	inner membrane
P25539	RIBD_ECOLI		cytoplasm
P25543	IRGB_VIBCH		cytoplasm
P25544	RBCR_CHRVI		cytoplasm
P25545	CBBR_XANFL		cytoplasm
P25547	GBPR_AGR TU		cytoplasm
P25549	ASLA_ECOLI	periplasm (potential).	periplasm
P25550	ASLB_ECOLI		cytoplasm
P25552	GPPA_ECOLI		extracellular
P25553	ALDA_ECOLI		cytoplasm
P25664	YZPK_ECOLI		cytoplasm
P25665	METE_ECOLI		cytoplasm
P25666	HTRL_ECOLI		inner membrane
P25728	YGBA_ECOLI		cytoplasm
P25731	CSOB_ECOLI		fimbrium
P25732	CFAA_ECOLI		fimbrium
P25733	CFAC_ECOLI		outer membrane
P25734	CFAE_ECOLI		fimbrium
P25738	MSYB_ECOLI		inner membrane
P25740	RFAG_ECOLI		cytoplasm
P25741	RFAP_ECOLI		inner membrane
P25742	RFAQ_ECOLI		cytoplasm
P25743	YCHE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

P25744	MDTG_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P25746	YCFC_ECOLI		inner membrane
P25747	YEIB_ECOLI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P25748	GALS_ECOLI		cytoplasm
P25749	CH10_BUCAI	cytoplasm (by similarity).	cytoplasm
P25750	CH60_BUCAI	cytoplasm (by similarity).	cytoplasm
P25756	GIDA_PSEPU	cytoplasm (by similarity).	cytoplasm
P25762	SSB_SERMA		cytoplasm
P25772	YICF_ECOLI		cytoplasm
P25841	SODC_HAEIN	periplasm (potential).	periplasm
P25842	SODC_HAEPA	periplasm (potential).	periplasm
P25852	ZRAR_SALTY	cytoplasm (probable).	cytoplasm
P25885	Y382_RHIME		inner membrane
P25888	RHLE_ECOLI		cytoplasm
P25889	YEIA_ECOLI		cytoplasm
P25893	Y383_RHIME		outer membrane
P25894	YGGG_ECOLI		extracellular
P25906	YDBC_ECOLI		cytoplasm
P25907	YDBD_ECOLI		cytoplasm
P25924	CYSG_SALTY		cytoplasm
P25926	NIRC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P25927	BIGA_SALTY		extracellular
P25938	C554_THINE		periplasm
P25967	CH60_BRUAB	cytoplasm (by similarity).	cytoplasm
P25969	CH101_RHOSH	cytoplasm (by similarity).	cytoplasm
P26004	CH60_AMOPS	cytoplasm (by similarity).	cytoplasm
P26005	CH10_AMOPS	cytoplasm (by similarity).	cytoplasm
P26024	NODC_BRAJA	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P26025	NODJ_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P26026	NODS_BRAJA		cytoplasm
P26027	NODU_BRAJA		inner membrane
P26050	NODI_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P26157	YPU1_RHOCA		cytoplasm
P26158	YPU2_RHOCA		cytoplasm
P26159	YPU3_RHOCA		cytoplasm
P26160	YPU4_RHOCA		cytoplasm
P26161	YPU5_RHOCA		cytoplasm
P26162	BCHH_RHOCA		cytoplasm
P26163	BCHB_RHOCA		cytoplasm
P26164	BCHN_RHOCA		cytoplasm
P26165	BCHF_RHOCA		cytoplasm
P26166	YBCC_RHOCA		cytoplasm
P26167	CRTJ_RHOCA		cytoplasm
P26168	BCHE_RHOCA		cytoplasm

P26169	BCHJ_RHOCA		cytoplasm
P26170	BCHG_RHOCA	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P26171	BCH2_RHOCA		inner membrane
P26172	BCHP_RHOCA		cytoplasm
P26173	IDI_RHOCA	cytoplasm (by similarity).	cytoplasm
P26174	BCHO_RHOCA		cytoplasm
P26175	BCHD_RHOCA		cytoplasm
P26176	YPUM_RHOCA		cytoplasm
P26177	BCHX_RHOCA		cytoplasm
P26178	BCHY_RHOCA	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P26179	BCHZ_RHOCA		cytoplasm
P26189	ADA_SALTY		cytoplasm
P26194	CH60_LEGMI	cytoplasm (by similarity).	cytoplasm
P26195	CH10_LEGMI	cytoplasm (by similarity).	cytoplasm
P26206	ARBB_ERWCH		cytoplasm
P26207	PTV3B_ERWCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P26211	ARBG_ERWCH		cytoplasm
P26218	YIEC_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (potential).	outer membrane
P26236	BCHM_RHOCA		cytoplasm
P26237	BCHL_RHOCA		cytoplasm
P26239	BCHI_RHOCA		cytoplasm
P26242	DXS_RHOCA		cytoplasm
P26248	NIFH1_AZOCH		cytoplasm
P26251	NIFH1_AZOCA		cytoplasm
P26252	NIFH2_AZOCA		cytoplasm
P26264	CYSQ_SALTY	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (probable).	cytoplasm
P26275	ALGR_PSEAE		cytoplasm
P26276	ALGC_PSEAE		cytoplasm
P26278	CYCR_ROSDE		periplasm
P26281	HPPK_ECOLI		cytoplasm
P26316	HRPS_PSESH		cytoplasm
P26318	FIMY_SALTY		fimbrium
P26319	FIMZ_SALTY	cytoplasm (probable).	cytoplasm
P26340	MANC_SALTY		cytoplasm
P26341	MANB_SALTY		cytoplasm
P26344	RECA_ERWCA	cytoplasm (by similarity).	cytoplasm
P26346	RECA_PROVU	cytoplasm (by similarity).	cytoplasm
P26348	RECA_VIBAN	cytoplasm (by similarity).	cytoplasm
P26353	HMP_SALTY		cytoplasm
P26365	AMIB_ECOLI		extracellular
P26366	AMIB_SALTY		extracellular
P26388	WCAL_SALTY		cytoplasm
P26389	WCAM_SALTY		inner membrane
P26391	RFBB_SALTY		cytoplasm
P26392	RFBD_SALTY		cytoplasm
P26393	RMLA_SALTY		cytoplasm

P26394	RFBC_SALTY		cytoplasm
P26395	RFBI_SALTY		cytoplasm
P26396	RFBF_SALTY		cytoplasm
P26397	RFBG_SALTY		cytoplasm
P26398	RFBH_SALTY		inner membrane
P26400	RFBX_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P26401	RFBV_SALTY		cytoplasm
P26402	RFBU_SALTY		inner membrane
P26403	RFBN_SALTY		cytoplasm
P26404	RFBM_SALTY		cytoplasm
P26405	RFBK_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P26408	HUPR_RHOCA	cytoplasm (probable).	cytoplasm
P26409	HYPB_RHOCA		cytoplasm
P26410	HYPD_RHOCA		cytoplasm
P26411	HYPD_RHOCA		extracellular
P26412	HYPE_RHOCA		cytoplasm
P26416	FLIK_SALTY		flagellar
P26420	SCRK_KLEPN		cytoplasm
P26462	FLIE_SALTY		flagellar
P26464	DPO3B_SALTY	cytoplasm (by similarity).	cytoplasm
P26465	FLII_SALTY	cytoplasm (potential).	inner membrane
P26469	RFAC_SALTY		cytoplasm
P26472	RFAY_SALTY		cytoplasm
P26473	RFMZ_SALTY		inner membrane
P26480	RPOD_PSEAE		cytoplasm
P26481	NIFW_AZOCA		cytoplasm
P26482	FIXA_AZOCA		cytoplasm
P26483	FIXB_AZOCA		cytoplasm
P26484	FIXC_AZOCA		cytoplasm
P26485	FIXX_AZOCA		cytoplasm
P26486	YFXX_AZOCA		cytoplasm
P26487	FIXJ_AZOCA	cytoplasm (probable).	cytoplasm
P26488	FIXK_AZOCA		cytoplasm
P26492	FLAV_DESDE		cytoplasm
P26494	PHAA_PSEOL		cytoplasm
P26495	PHAB_PSEOL		extracellular
P26496	PHAC_PSEOL		cytoplasm
P26501	ISOA_PSEUM		cytoplasm
P26503	EXOB_RHIME		cytoplasm
P26504	LIPA_PSEFL		extracellular
P26505	HEM1_AGRRD		inner membrane
P26506	NIFE_BRAJA		cytoplasm
P26507	NIFN_BRAJA		periplasm
P26508	NOLC_RHIFR		inner membrane
P26608	FLIS_ECOLI		flagellar
P26609	FLIS_SALTY		flagellar
P26616	MAO1_ECOLI		cytoplasm
P26646	YHDH_ECOLI		cytoplasm
P26649	GLGS_ECOLI		cytoplasm
P26688	CBHE_COXBU	cytoplasm (potential).	cytoplasm
P26720	HIS4_AZOBR	cytoplasm (by similarity).	cytoplasm
P26721	HIS6_AZOBR	cytoplasm (by similarity).	cytoplasm

P26722	HIS2_AZOBR	cytoplasm (by similarity).	cytoplasm
P26723	YHI1_AZOBR		cytoplasm
P26724	YHIT_AZOBR		cytoplasm
P26761	RTX1D_ACTPL	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P26762	BVGS_BORBR	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P26838	CAT4_ECOLI		cytoplasm
P26841	CAT4_PSEAE		cytoplasm
P26879	CH10_LEGPN	cytoplasm (by similarity).	cytoplasm
P26918	BLAB_AERHY	periplasm (probable).	periplasm
P26919	T2A1_AZOBR		extracellular
P26922	TRPG_AZOBR		cytoplasm
P26924	TRPD_AZOBR		cytoplasm
P26938	TRPC_AZOBR		cytoplasm
P26943	YTR1_AZOBR		cytoplasm
P26945	IAAT_AZOBR		cytoplasm
P26950	CAF1R_YERPE		cytoplasm
P26977	PUR2_SALTY		cytoplasm
P26978	PUR9_SALTY		cytoplasm
P26979	RP54_SALTY		cytoplasm
P26983	RP5M_SALTY		cytoplasm
P26984	SCRK_SALTY		cytoplasm
P26993	EXSA_PSEAE		cytoplasm
P26994	EXSB_PSEAE		cytoplasm
P26995	EXSC_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P27017	AMIC_PSEAE		periplasm
P27027	PLYD_PSEMA		cytoplasm
P27030	RHAB_SALTY		cytoplasm
P27031	RHAA_SALTY	cytoplasm (by similarity).	cytoplasm
P27032	GUNY_DICD3		extracellular
P27033	GUNC_PSEFL		extracellular
P27034	BGLS_AGRTU	cytoplasm (probable).	periplasm
P27053	FLAA_CAMCO		flagellar
P27098	TCBC_PSESQ		cytoplasm
P27099	TCBD_PSESQ		cytoplasm
P27100	TCBE_PSESQ		cytoplasm
P27101	TCBF_PSESQ		cytoplasm
P27102	TCBR_PSESQ		cytoplasm
P27103	YTCB_PSESQ	periplasm (potential).	periplasm
P27126	RFAS_ECOLI		inner membrane
P27127	RFAB_ECOLI		cytoplasm
P27128	RFAI_ECOLI		cytoplasm
P27129	RFAJ_ECOLI		cytoplasm
P27135	RHAT_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P27136	TFDE1_RALEJ		cytoplasm
P27137	TFDF1_RALEJ		cytoplasm
P27138	TFDB_RALEJ		cytoplasm
P27186	REPB_THIFE		cytoplasm
P27187	TRAA_ECOLI		cytoplasm

P27188	TRAB4_ECOLI		cytoplasm
P27189	TRC4_ECOLI		cytoplasm
P27190	TRC5_ECOLI		cytoplasm
P27192	TRAD5_ECOLI		cytoplasm
P27193	TRAD4_ECOLI		cytoplasm
P27219	PTSBC_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P27237	OPDA_SALTY		extracellular
P27238	YFED_ECOLI		cytoplasm
P27240	RFAY_ECOLI		cytoplasm
P27241	RFAY_ECOLI		inner membrane
P27245	MARR_ECOLI		cytoplasm
P27247	PLSX_ECOLI		cytoplasm
P27248	GCST_ECOLI		cytoplasm
P27249	GLND_ECOLI		cytoplasm
P27250	YJGB_ECOLI		cytoplasm
P27253	SBM_ECOLI		cytoplasm
P27273	FDHB_WOLSU		periplasm
P27294	INAA_ECOLI		inner membrane
P27296	DING_ECOLI		cytoplasm
P27297	BAX_ECOLI		cytoplasm
P27298	OPDA_ECOLI		extracellular
P27300	LPXK_ECOLI		cytoplasm
P27302	TKT1_ECOLI		extracellular
P27305	GLUQ_ECOLI		cytoplasm
P27342	HFAA_CAUCR		extracellular
P27343	HFAB_CAUCR		cytoplasm
P27345	MUTS_AZOVI		cytoplasm
P27353	MEMA_METTR		cytoplasm
P27354	MEMB_METTR		cytoplasm
P27355	MEMG_METTR		cytoplasm
P27356	MMOB_METTR		cytoplasm
P27375	HTRC_ECOLI		cytoplasm
P27394	FDXN_BRAJA		periplasm
P27422	DYR7_ECOLI		cytoplasm
P27431	YCFD_ECOLI		cytoplasm
P27434	YFGA_ECOLI		cytoplasm
P27461	YLPA_YEREN	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P27474	YOPQ_YEREN	secreted protein (probable).	extracellular
P27488	RCSA_ERWST		cytoplasm
P27498	DYR_KLEAE		cytoplasm
P27504	PQQB_KLEPN		cytoplasm
P27505	PQQC_KLEPN		cytoplasm
P27506	PQQD_KLEPN		cytoplasm
P27507	PQQE_KLEPN		cytoplasm
P27508	PQQF_KLEPN		extracellular
P27550	ACSA_ECOLI		cytoplasm
P27640	GREA_RICPR		cytoplasm
P27644	PGLR_AGRTU		extracellular
P27649	HUPD_RHILV		extracellular
P27651	HUPF_RHILV		cytoplasm
P27667	UHPA_SALTY	cytoplasm (probable).	cytoplasm
P27688	FMA0_BACNO		extracellular
P27689	FMA3_BACNO		extracellular

P27690	FMA6_BACNO		extracellular
P27691	FMA7_BACNO		extracellular
P27696	ILVB_KLEPN		cytoplasm
P27713	NIFA_HERSE		cytoplasm
P27714	NIFB_HERSE		cytoplasm
P27717	LEP4_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P27722	KHSE_SERMA	cytoplasm (potential).	cytoplasm
P27725	AK1H_SERMA		cytoplasm
P27735	THRC_SERMA		cytoplasm
P27745	ACOA_RALEU		cytoplasm
P27746	ACOB_RALEU		cytoplasm
P27747	ACOC_RALEU		cytoplasm
P27748	ACOX_RALEU	bacterial membrane (potential).	inner membrane
P27749	YAC7_RALEU		cytoplasm
P27763	NHAB_PSECL		cytoplasm
P27764	NHAA_PSECL		extracellular
P27765	AMID_PSECL		extracellular
P27829	WECC_ECOLI		cytoplasm
P27830	RFFG_ECOLI		cytoplasm
P27832	RFFC_ECOLI		inner membrane
P27833	RFFA_ECOLI		inner membrane
P27835	WZYE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P27836	WECG_ECOLI		cytoplasm
P27838	CYAY_ECOLI		inner membrane
P27840	YIGE_ECOLI		inner membrane
P27842	YIGF_ECOLI		inner membrane
P27848	YIGL_ECOLI		cytoplasm
P27862	YIGZ_ECOLI		fimbrium
P27865	RECA_RHIME	cytoplasm (by similarity).	cytoplasm
P27873	MAS2_AGRRH		cytoplasm
P27874	MAS11_AGRRH		cytoplasm
P27875	AGS_AGRRH		cytoplasm
P27887	XYLE_PSEAE		cytoplasm
P27897	CELA_ACEXY		cytoplasm
P27905	FIMBX_BACNO		fimbrium
P27906	FMAX_BACNO		extracellular
P27994	F16P_RHOSH		cytoplasm
P27995	ALF1_RHOSH		cytoplasm
P27997	RBL1_RHOSH		cytoplasm
P27998	RBS1_RHOSH		cytoplasm
P28029	HLY3_VIBPA		extracellular
P28030	HLY_VIBMI		extracellular
P28031	HLY1_VIBHO		extracellular
P28036	DHET_ACEPO	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	periplasm
P28043	SSB2_ECOLI		cytoplasm
P28044	SSB7_ECOLI		cytoplasm
P28045	SSBR_ECOLI		cytoplasm
P28046	SSB_PROMI		cytoplasm
P28080	DBHA_VIBPR		cytoplasm

P28149	HUPG_RHILV		cytoplasm
P28150	HUPH_RHILV		cytoplasm
P28151	RUBL_RHILV		cytoplasm
P28152	HUPJ_RHILV		cytoplasm
P28153	HUPK_RHILV		cytoplasm
P28154	HYPB_RHILV		cytoplasm
P28155	HYPB_RHILV		cytoplasm
P28156	HYPF_RHILV		cytoplasm
P28157	HYPF_RHILV		cytoplasm
P28157	HYPC_RHILV		cytoplasm
P28183	SAHH_RHOCA	cytoplasm (by similarity).	cytoplasm
P28224	YDHA_ECOLI		cytoplasm
P28243	YAIB_ESCFE		cytoplasm
P28247	YZBC_ECOLI		cytoplasm
P28248	DCD_ECOLI		extracellular
P28249	ASMA_ECOLI	periplasm (potential).	cytoplasm
P28266	NODL_RHIME		cytoplasm
P28267	NOLR_RHIME		cytoplasm
P28269	OAPT_PSEPU		cytoplasm
P28304	QOR_ECOLI		cytoplasm
P28305	PABC_ECOLI		cytoplasm
P28306	YCEG_ECOLI		cytoplasm
P28307	CSGA_ECOLI		extracellular
P28355	RECJ_SALTY		cytoplasm
P28369	RFH_ECOLI		cytoplasm
P28577	DCHS_ENTAE		cytoplasm
P28578	DCHS_KLEPL		cytoplasm
P28579	FLAV_ENTAG		cytoplasm
P28585	BLC1_ECOLI		extracellular
P28595	PYRG_AZOBR		cytoplasm
P28602	HEMH_BRAJA	cytoplasm (by similarity).	extracellular
P28603	NODP_AZOBR		cytoplasm
P28604	NODQ_AZOBR		cytoplasm
P28607	ARS_ALTCA	periplasm (potential).	periplasm
P28613	RP5M_RALEU		cytoplasm
P28614	ACOR_RALEU		cytoplasm
P28615	RP54_RALEU		cytoplasm
P28630	HOLA_ECOLI		cytoplasm
P28631	HOLB_ECOLI		cytoplasm
P28632	HOLD_ECOLI		cytoplasm
P28634	YAEB_ECOLI		inner membrane
P28635	METQ_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P28638	YHDJ_ECOLI		cytoplasm
P28696	YAAI_ECOLI		flagellar
P28697	HTGA_ECOLI		cytoplasm
P28719	DNLJ_ZYMMO		cytoplasm
P28720	TGT_ZYMMO		cytoplasm
P28721	GLTF_ECOLI		inner membrane
P28722	YHCA_ECOLI	periplasm (by similarity).	periplasm
P28787	NTRC_PROVU		cytoplasm
P28788	NTRB_PROVU		inner membrane
P28790	FADA_PSEFR	cytoplasm (by similarity).	cytoplasm
P28793	FADB_PSEFR		cytoplasm
P28796	TNAA_PROVU		cytoplasm
P28808	LCRF_YERPE		cytoplasm
P28809	MMSR_PSEAE		cytoplasm

P28810	MMSA_PSEAE		cytoplasm
P28811	MMSB_PSEAE		cytoplasm
P28812	Y3568_PSEAE		cytoplasm
P28815	TETC_ECOLI		cytoplasm
P28816	TETD_ECOLI		cytoplasm
P28836	YOR3_YEREN		inner membrane
P28861	FENR_ECOLI		cytoplasm
P28895	RBL1A_THIFE		extracellular
P28896	RBS1_THIFE		cytoplasm
P28901	FENR_SHIFL		cytoplasm
P28903	NRDD_ECOLI		cytoplasm
P28905	HOLC_ECOLI		cytoplasm
P28911	YHHH_ECOLI		cytoplasm
P28912	YHHI_ECOLI		cytoplasm
P28915	YBFC_ECOLI		cytoplasm
P28916	YBFD_ECOLI		cytoplasm
P28917	YDCC_ECOLI		cytoplasm
P29010	YDFD_ECOLI		cytoplasm
P29012	ALR2_ECOLI		cytoplasm
P29013	YCGB_ECOLI		cytoplasm
P29039	XPSM_XANCP		cytoplasm
P29041	GSPD_XANCP	bacterial cell outer membrane (probable).	inner membrane
P29208	MENC_ECOLI		extracellular
P29217	YCEH_ECOLI		cytoplasm
P29236	LUXC2_PHOLE		cytoplasm
P29237	LUXG_PHOLE		cytoplasm
P29238	LUXA2_PHOLE		cytoplasm
P29239	LUXB2_PHOLE		cytoplasm
P29246	RECA_AZOVI	cytoplasm (by similarity).	cytoplasm
P29248	FLIA_PSEAE		cytoplasm
P29267	HOXA_RALEU	cytoplasm (probable).	cytoplasm
P29271	ALF2_RHOSH		cytoplasm
P29277	TKT_RHOSH		extracellular
P29278	RBL2_RHOSH		cytoplasm
P29280	SLMA_HAEIN	nucleoid-associated (by similarity).	cytoplasm
P29281	CRP_HAEIN		cytoplasm
P29285	Y6063_BRAJA		inner membrane
P29286	FIXK_BRAJA		cytoplasm
P29334	LUXE_PHOLE		cytoplasm
P29363	THRC_PSEAE		cytoplasm
P29364	KHSE_PSEAE		cytoplasm
P29365	DHOM_PSEAE		cytoplasm
P29367	FINO3_ECOLI		outer membrane
P29368	YPT1_ECOLI		cytoplasm
P29369	AGMR_PSEAE		cytoplasm
P29370	Y1977_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P29399	MARI_ALTSP		cytoplasm
P29431	OXAA_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P29432	Y015_BUCAP		inner membrane
P29433	RNPA_BUCAP		extracellular

P29434	DNAA_BUCAP	cytoplasm (by similarity).	cytoplasm
P29435	GYRB_BUCAP		cytoplasm
P29439	DPO3B_BUCAP	cytoplasm (by similarity).	cytoplasm
P29440	DNAA_SERMA	cytoplasm (by similarity).	cytoplasm
P29444	XYLB_KLEPN		cytoplasm
P29480	TCPT_VIBCH	cytoplasm (probable).	inner membrane
P29485	TCPV_VIBCH		cytoplasm
P29487	TCPE_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P29488	TCPF_VIBCH	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P29489	TCPH_VIBCH		inner membrane
P29490	TCPQ_VIBCH		cytoplasm
P29491	TCPD_VIBCH		inner membrane
P29537	T2H1_HAEPA		extracellular
P29538	MTH1_HAEPA		cytoplasm
P29680	DCUP_ECOLI	cytoplasm (probable).	cytoplasm
P29698	ENTF_SHIFL		cytoplasm
P29740	Y1013_PHOPR		inner membrane
P29741	DING_PHOPR		cytoplasm
P29744	FLGL_ECOLI		flagellar
P29745	PEPT_ECOLI	cytoplasm (by similarity).	extracellular
P29768	NANH_SALTY		extracellular
P29769	NUC_SHIFL		extracellular
P29770	YNU1_SHIFL		cytoplasm
P29771	YNU2_SHIFL		cytoplasm
P29772	OSA_SHIFL		cytoplasm
P29806	AADB2_KLEPN		cytoplasm
P29807	AACC2_ACIBA		cytoplasm
P29808	AACC3_PSEAE		cytoplasm
P29811	CYC2_PSEAE		cytoplasm
P29848	CYSM_SALTY		cytoplasm
P29895	MAUD_PARDE		inner membrane
P29896	MAUE_PARDE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P29897	MAUF_PARDE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P29899	CYCL_PARDE	periplasm (potential).	periplasm
P29900	MOXJ_PARDE	periplasm (potential).	periplasm
P29903	MOXZ_PARDE		cytoplasm
P29904	MOXX_PARDE	cytoplasm (probable).	cytoplasm
P29905	MOXY_PARDE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P29906	BIRA_PARDE		cytoplasm
P29907	YNQ1_PARDE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P29908	YNQ2_PARDE		cytoplasm
P29909	YNQ3_PARDE		cytoplasm
P29910	YNQ4_PARDE		cytoplasm

P29911	YNQ5_PARDE	bacterial membrane; peripheral membrane protein (probable).	cytoplasm
P29912	YNQ6_PARDE	bacterial membrane; peripheral membrane protein (probable).	cytoplasm
P29913	NQO1_PARDE		periplasm
P29914	NQO2_PARDE		cytoplasm
P29915	NQO3_PARDE		cytoplasm
P29916	NQO4_PARDE		cytoplasm
P29917	NQO5_PARDE		inner membrane
P29918	NQO6_PARDE		cytoplasm
P29921	NQO9_PARDE		cytoplasm
P29929	COBN_PSEDE	cytoplasm (potential).	cytoplasm
P29931	COBP_PSEDE		cytoplasm
P29932	COBQ_PSEDE		cytoplasm
P29933	COBS_PSEDE	cytoplasm (potential).	cytoplasm
P29934	COBT_PSEDE	cytoplasm (potential).	cytoplasm
P29935	COBU_PSEDE		cytoplasm
P29936	COBV_PSEDE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P29937	COBW_PSEDE		cytoplasm
P29938	YCBV_PSEDE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P29939	YCB6_PSEDE	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P29940	YCB7_PSEDE	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P29941	YCB8_PSEDE	bacterial membrane (probable).	inner membrane
P29942	YCB9_PSEDE	bacterial membrane (probable).	inner membrane
P29943	YCB1_PSEDE		cytoplasm
P29944	YCB2_PSEDE		inner membrane
P29945	KDGL_PSEDE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P29946	CBIA_SALTY		cytoplasm
P29953	Y2983_RHIME		inner membrane
P29954	MANA_RHIME		cytoplasm
P29955	XANA_XANCP		cytoplasm
P29956	XANB_XANCP		cytoplasm
P29957	AMY_PSEHA		cytoplasm
P29959	CCMA_RHOCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P29960	CCMB_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P29961	CCMC_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane

P29962	YHEA_RHOCA		cytoplasm
P29963	CCMD_RHOCA		inner membrane
P29968	XOXF_PARDE		periplasm
P30011	NADC_ECOLI		cytoplasm
P30012	NADC_SALTY		cytoplasm
P30014	RNT_ECOLI		cytoplasm
P30015	LHR_ECOLI		cytoplasm
P30103	PRIM_RICPR		cytoplasm
P30126	LEUD_ECOLI		cytoplasm
P30128	GREB_ECOLI		cytoplasm
P30130	FIMD_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P30131	HYPF_ECOLI		cytoplasm
P30136	THIC_ECOLI		cytoplasm
P30137	THIE_ECOLI		cytoplasm
P30138	THIF_ECOLI		cytoplasm
P30140	THIH_ECOLI		cytoplasm
P30143	YAAJ_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P30147	HYI_ECOLI		cytoplasm
P30176	YBIA_ECOLI		cytoplasm
P30177	YBIB_ECOLI		cytoplasm
P30178	YBIC_ECOLI	cytoplasm (potential).	cytoplasm
P30192	YCHG_ECOLI		cytoplasm
P30235	YEIC_ECOLI		cytoplasm
P30297	1A1D_PSES0		cytoplasm
P30332	RP54_BRAJA		cytoplasm
P30333	RP55_BRAJA		cytoplasm
P30334	RP5M_BRAJA		cytoplasm
P30347	LIGF_PSEPA		cytoplasm
P30417	FKBZ_PSEAE		cytoplasm
P30534	CYAY_YERIN		inner membrane
P30663	NIFL_AZOVI		inner membrane
P30667	NIFA_AZOBR		cytoplasm
P30718	CH60_PSEAE	cytoplasm (by similarity).	cytoplasm
P30720	CH10_PSEAE	cytoplasm (by similarity).	cytoplasm
P30739	TNPT_PSEPU		cytoplasm
P30744	SDHM_ECOLI		cytoplasm
P30745	MOAA_ECOLI		cytoplasm
P30748	MOAD_ECOLI		fimbrium
P30749	MOAE_ECOLI		fimbrium
P30750	METN_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P30752	LOLB_SALTY	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P30753	ISPE_SALTY		cytoplasm
P30773	T2X1_XANCC		extracellular
P30774	MTX1_XANCC		cytoplasm
P30778	RUBR_AZOVI		cytoplasm
P30779	CH60_AGRT5	cytoplasm (by similarity).	cytoplasm
P30780	CH10_AGRT5	cytoplasm (by similarity).	cytoplasm
P30781	HOXT_AZOVI		cytoplasm
P30787	IHFA_RHOCA		cytoplasm

P30788	RL32_RHOCA		cytoplasm
P30789	PLSX_RHOCA		cytoplasm
P30790	FABH_RHOCA	cytoplasm (probable).	cytoplasm
P30794	Y1243_ZYMMO		outer membrane
P30795	Y1242_ZYMMO		cytoplasm
P30797	HUPK_RHOCA		cytoplasm
P30798	GPMA_ZYMMO		cytoplasm
P30799	DDH_ZYMMO		cytoplasm
P30813	NODK_BRASP		cytoplasm
P30819	NADC_PSEAE		cytoplasm
P30820	NIGY_DESVH	cytoplasm (probable).	cytoplasm
P30843	BASR_ECOLI	cytoplasm (potential).	cytoplasm
P30851	RNR_SHIFL		cytoplasm
P30852	SMF_ECOLI		cytoplasm
P30855	EVGS_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P30859	ARTI_ECOLI	periplasm (by similarity).	periplasm
P30860	ARTJ_ECOLI	periplasm (by similarity).	periplasm
P30863	DKGB_ECOLI	cytoplasm (by similarity).	cytoplasm
P30864	YAFC_ECOLI		cytoplasm
P30866	YAFE_ECOLI		cytoplasm
P30870	GLNE_ECOLI		cytoplasm
P30871	YGIF_ECOLI		cytoplasm
P30877	AGAL_SALTY		cytoplasm
P30896	BLA3_KLEPN		extracellular
P30897	BLAC_PROMI		extracellular
P30903	DHAS_NEIMB		cytoplasm
P30958	MFD_ECOLI		cytoplasm
P30959	CCMD_BRAJA		inner membrane
P30960	CYCY_BRAJA	periplasm (probable).	periplasm
P30961	ISPZ_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P30962	CCMC_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P30963	CCMA_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P30964	CCMB_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P30967	PH4H_CHRVO		cytoplasm
P30979	YBEF_ECOLI		cytoplasm
P31002	IMDH_ACICA		cytoplasm
P31004	FERX_ACICA		cytoplasm
P31011	TPL_ENTAG	cytoplasm (probable).	cytoplasm
P31012	TPL_ESCIN		cytoplasm
P31013	TPL_CITFR		cytoplasm
P31019	PHEB_PSEUE		cytoplasm
P31020	PHEA_PSEUE		cytoplasm
P31032	T2M4_NEIGO		extracellular
P31033	MTM4_NEIGO		cytoplasm

P31048	CLS_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P31049	YLP3_PSEPU		cytoplasm
P31057	PANB_ECOLI		cytoplasm
P31058	YADC_ECOLI		extracellular
P31060	MODF_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P31061	NOHA_ECOLI		cytoplasm
P31062	NOHB_ECOLI		cytoplasm
P31063	YEDD_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P31068	FLIH_ECOLI		inner membrane
P31073	DYR_CITFR		cytoplasm
P31074	DYR_ENTAE		cytoplasm
P31075	PSRA_WOLSU		periplasm
P31076	PSRB_WOLSU		cytoplasm
P31077	PSRC_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P31078	PETP_RHOCA		cytoplasm
P31079	PETR_RHOCA		cytoplasm
P31108	SODF_LEGPH		periplasm
P31120	GLMM_ECOLI		cytoplasm
P31121	MARB_ECOLI		inner membrane
P31126	YDEE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P31129	YDEH_ECOLI		inner membrane
P31130	YDEI_ECOLI		cytoplasm
P31131	YDEJ_ECOLI		inner membrane
P31134	POTG_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P31142	THTM_ECOLI	cytoplasm (probable).	cytoplasm
P31242	LAMB_KLEPN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P31294	CH60_HAEDU	cytoplasm (by similarity).	cytoplasm
P31296	CH10_HAEDU	cytoplasm (by similarity).	cytoplasm
P31339	ASPA_AERSA		extracellular
P31433	YICH_ECOLI		cytoplasm
P31434	YICI_ECOLI		cytoplasm
P31441	ADEC_ECOLI		cytoplasm
P31443	YIDF_ECOLI		cytoplasm
P31447	YIDJ_ECOLI		inner membrane
P31449	YIDL_ECOLI		cytoplasm
P31450	GLVG_ECOLI		cytoplasm
P31453	YIDP_ECOLI		cytoplasm
P31455	YIDR_ECOLI		extracellular
P31456	YIDS_ECOLI		cytoplasm
P31459	DGOK_ECOLI		cytoplasm
P31460	DGOR_ECOLI		cytoplasm
P31463	YIDZ_ECOLI		cytoplasm
P31467	YIEH_ECOLI		cytoplasm

P31469	YIEJ_ECOLI		inner membrane
P31470	YIEK_ECOLI		cytoplasm
P31471	YIEL_ECOLI		cytoplasm
P31475	YIEP_ECOLI		cytoplasm
P31485	YPCP_YEREN		inner membrane
P31490	YERA_YEREN		cytoplasm
P31491	YERA_YERPE		cytoplasm
P31494	URE1_YEREN	cytoplasm (by similarity).	cytoplasm
P31495	URE2_YEREN	cytoplasm (by similarity).	cytoplasm
P31496	URE3_YEREN	cytoplasm (by similarity).	cytoplasm
P31516	HEMP_YEREN		cytoplasm
P31517	HEMS_YEREN		cytoplasm
P31519	SKP_YEREN	periplasm (by similarity).	periplasm
P31520	SKP_YERPS	periplasm (by similarity).	periplasm
P31521	P47K_PSECL		cytoplasm
P31522	PSAA_YERPE		extracellular
P31527	PSAC_YERPE	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P31545	YCDB_ECOLI		cytoplasm
P31551	CAID_ECOLI		cytoplasm
P31552	CAIC_ECOLI		cytoplasm
P31574	FIXB_ECOLI		cytoplasm
P31585	GSPG_ERWCH		inner membrane
P31586	GSPG_PECCC		inner membrane
P31587	GSPH_PECCC		inner membrane
P31588	GSPI_PECCC		inner membrane
P31589	GSPJ_PECCC		inner membrane
P31592	GLNA3_RHILP		cytoplasm
P31638	ACSA_RALEU		cytoplasm
P31660	PRPC_ECOLI		cytoplasm
P31663	PANC_ECOLI		cytoplasm
P31665	YADD_ECOLI		cytoplasm
P31666	YADE_ECOLI		cytoplasm
P31667	YHGA_ECOLI		cytoplasm
P31668	PAT_ALCFA		cytoplasm
P31677	OTSA_ECOLI		cytoplasm
P31678	OTSB_ECOLI		cytoplasm
P31679	YAAU_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P31698	GSPC1_ERWCH	bacterial cell inner membrane (probable).	inner membrane
P31699	GSPC_PECCC	bacterial cell inner membrane (probable).	inner membrane
P31700	GSPD1_ERWCH	bacterial cell outer membrane (probable).	inner membrane
P31701	GSPD_PECCC	bacterial cell outer membrane (probable).	inner membrane
P31702	GSPE1_ERWCH	cytoplasm (probable).	inner membrane
P31703	GSPE_PECCC	cytoplasm (probable).	inner membrane
P31704	GSPF_ERWCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane

P31705	GSPF_PECCC	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P31706	GSPK_PECCC	bacterial cell inner membrane (potential).	inner membrane
P31707	GSPL_ERWCH		inner membrane
P31708	GSPL_PECCC	bacterial cell inner membrane (probable).	inner membrane
P31709	GSPM_PECCC	bacterial cell inner membrane (probable).	inner membrane
P31711	LEP4_ERWCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P31728	METQ_HAEIN	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
P31733	GSPG_AERHY		inner membrane
P31734	GSPG_XANCP		inner membrane
P31735	GSPH_AERHY		inner membrane
P31736	GSPH_XANCP		inner membrane
P31737	GSPI_AERHY		inner membrane
P31738	GSPI_XANCP		cytoplasm
P31739	GSPJ_AERHY		inner membrane
P31740	GSPJ_XANCP		cytoplasm
P31741	GSPE_AERHY	cytoplasm (probable).	inner membrane
P31742	GSPE_XANCP	cytoplasm (probable).	inner membrane
P31743	GSPF_AERHY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P31744	GSPF_XANCP	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P31758	FRZG_MYXXA		cytoplasm
P31759	FRZF_MYXXA		cytoplasm
P31760	GSPK_AERHY	bacterial cell inner membrane (potential).	inner membrane
P31764	GAL7_HAEIN		cytoplasm
P31765	GALM_HAEIN	cytoplasm (probable).	cytoplasm
P31766	GALR_HAEIN		cytoplasm
P31767	GAL1_HAEIN	cytoplasm (potential).	cytoplasm
P31768	COMA_HAEIN		cytoplasm
P31769	COMB_HAEIN		cytoplasm
P31770	COMC_HAEIN		cytoplasm
P31771	COMD_HAEIN		cytoplasm
P31772	COME_HAEIN	bacterial cell outer membrane (probable).	outer membrane
P31773	COMF_HAEIN		cytoplasm
P31774	Y433_HAEIN		inner membrane
P31776	PBPA_HAEIN	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P31777	Y441_HAEIN		cytoplasm
P31778	HRPB_RALSO		cytoplasm
P31780	GSPD_AERHY	bacterial cell outer membrane (probable).	inner membrane
P31781	RL13_HAESO		cytoplasm
P31782	RS9_HAESO		cytoplasm

P31784	SSPA_HAESO		cytoplasm
P31802	NARP_ECOLI		cytoplasm
P31806	YJEF_ECOLI		fimbrium
P31808	YCIK_ECOLI		cytoplasm
P31811	Y847_HAEIN		cytoplasm
P31812	TRMA_HAEIN		cytoplasm
P31825	YFIC_ECOLI		cytoplasm
P31827	YDDB_ECOLI		outer membrane
P31828	PQQL_ECOLI		extracellular
P31850	TBLA_PSESZ		cytoplasm
P31851	TABA_PSESZ		cytoplasm
P31852	TABB_PSESZ		cytoplasm
P31876	HYDD_WOLSU		periplasm
P31877	HYDE_WOLSU		nucleoid
P31878	HUPK_AZOVI		cytoplasm
P31879	HYP A_AZOVI		cytoplasm
P31880	HYPB_AZOVI		cytoplasm
P31881	HYPC_AZOVI		cytoplasm
P31882	HYPD_AZOVI		extracellular
P31894	COOF_RHORU		periplasm
P31895	COOH_RHORU		cytoplasm
P31897	COOC_RHORU		cytoplasm
P31899	HOXL_RALEU		cytoplasm
P31900	HYPC_RALEU		cytoplasm
P31901	HYP A_RALEU		cytoplasm
P31902	HYPB_RALEU		cytoplasm
P31903	HYPD_RALEU		extracellular
P31904	HYPD2_BRAJA		extracellular
P31905	HYPE_RALEU		cytoplasm
P31906	HYPE_BRAJA		cytoplasm
P31907	HOXX_BRAJA		cytoplasm
P31908	HOXA_BRAJA	cytoplasm (probable).	cytoplasm
P31909	HOXM_RALEU		extracellular
P31910	HOXO_RALEU		cytoplasm
P31911	HOXQ_RALEU		cytoplasm
P31912	RUBR_RALEU		cytoplasm
P31913	HOXT_RALEU		cytoplasm
P31914	HOXV_RALEU		cytoplasm
P31957	NODU_RHISN		inner membrane
P31961	EDD_PSEAE		cytoplasm
P31979	NUOF_ECOLI		periplasm
P31991	YDCD_ECOLI		cytoplasm
P31992	YDCE_ECOLI		cytoplasm
P32000	PRIM_BUCAP		cytoplasm
P32001	RPOD_BUCAP		cytoplasm
P32002	SECB_BUCAP	cytoplasm (by similarity).	cytoplasm
P32003	CYSE_BUCAP	cytoplasm (by similarity).	cytoplasm
P32008	NODD2_RHITR		cytoplasm
P32014	CTRB_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P32015	CTRC_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane

P32016	CTRD_NEIMB	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P32021	EFE_PSESH		cytoplasm
P32051	YDEK_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P32053	INTA_ECOLI		cytoplasm
P32056	NUDD_ECOLI		cytoplasm
P32057	WCAI_ECOLI		cytoplasm
P32090	MUTT_PROVU		cytoplasm
P32104	ACCR_AGRT5		cytoplasm
P32106	YIBG_ECOLI		extracellular
P32108	YIBI_ECOLI		inner membrane
P32109	YIBJ_ECOLI		cytoplasm
P32125	MOBB_ECOLI		cytoplasm
P32128	YIHF_ECOLI		inner membrane
P32132	TYPA_ECOLI		cytoplasm
P32134	YIHM_ECOLI		cytoplasm
P32136	YIHO_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P32138	YIHQ_ECOLI		cytoplasm
P32139	YIHR_ECOLI		cytoplasm
P32140	YIHS_ECOLI		cytoplasm
P32141	YIHT_ECOLI		cytoplasm
P32143	YIHV_ECOLI		cytoplasm
P32144	YIHW_ECOLI		cytoplasm
P32151	YIIG_ECOLI		outer membrane
P32152	FRVR_ECOLI		cytoplasm
P32153	FRVX_ECOLI		extracellular
P32155	PTFA_ECOLI	cytoplasm (probable).	inner membrane
P32156	YIIL_ECOLI		cytoplasm
P32157	YIIM_ECOLI		cytoplasm
P32160	YIIQ_ECOLI		extracellular
P32162	YIIS_ECOLI		cytoplasm
P32167	YIIX_ECOLI		cytoplasm
P32170	RHAA_ECOLI	cytoplasm (probable).	cytoplasm
P32171	RHAB_ECOLI		cytoplasm
P32176	FDOG_ECOLI	cytoplasm (potential).	periplasm
P32177	FDHD_ECOLI	cytoplasm (probable).	cytoplasm
P32180	HSLU_PASHA	cytoplasm (by similarity).	cytoplasm
P32181	LAPB_PASHA		inner membrane
P32202	LPXD_RICRI		cytoplasm
P32203	LPXD_YEREN		cytoplasm
P32204	FABZ_RICRI	cytoplasm (by similarity).	cytoplasm
P32205	FABZ_YEREN	cytoplasm (by similarity).	cytoplasm
P32265	NFXB_PSEAE		cytoplasm
P32326	URER_ECOLI		cytoplasm
P32427	PCAB_PSEPU		cytoplasm
P32662	GPH_ECOLI		cytoplasm
P32664	NUDC_ECOLI		cytoplasm
P32668	YIJF_ECOLI		inner membrane
P32669	FSAB_ECOLI	cytoplasm (probable).	cytoplasm
P32670	PTFX2_ECOLI	cytoplasm (potential).	cytoplasm
P32674	PFLD_ECOLI	cytoplasm (by similarity).	cytoplasm
P32675	PFLC_ECOLI	cytoplasm (by similarity).	cytoplasm

P32676	PTFB3_ECOLI	cytoplasm (potential).	inner membrane
P32677	YIJO_ECOLI		cytoplasm
P32680	YJAG_ECOLI		cytoplasm
P32681	YJAH_ECOLI		cytoplasm
P32684	RLUF_ECOLI		cytoplasm
P32685	YJBD_ECOLI		cytoplasm
P32687	YJBF_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P32688	YJBG_ECOLI		cytoplasm
P32689	YJBH_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P32690	YJBI_ECOLI		cytoplasm
P32693	YJBL_ECOLI		cytoplasm
P32694	YJBM_ECOLI		cytoplasm
P32695	DUSA_ECOLI		cytoplasm
P32696	YJBO_ECOLI		inner membrane
P32700	YJCB_ECOLI		inner membrane
P32701	YJCC_ECOLI		inner membrane
P32703	YJCE_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P32704	YJCF_ECOLI		cytoplasm
P32711	NRFF_ECOLI	periplasm (probable).	periplasm
P32712	NRFG_ECOLI		extracellular
P32714	MDTP_ECOLI	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
P32715	MDTO_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P32716	MDTN_ECOLI	bacterial cell inner membrane; single-pass type ii membrane protein (potential).	inner membrane
P32717	YJCS_ECOLI		extracellular
P32718	ALSK_ECOLI		cytoplasm
P32719	ALSE_ECOLI		cytoplasm
P32721	ALSA_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P32725	RECA_RHOS4	cytoplasm (by similarity).	cytoplasm
P32757	YHOX_RALEU		cytoplasm
P32820	TBTA_ALTSM		inner membrane
P32823	CHIA_ALTSM		extracellular
P32890	ELBP_ECOLI		extracellular
P32920	DCUP_RHOS4	cytoplasm (probable).	cytoplasm
P32966	UVRC_PSEFL	cytoplasm (by similarity).	cytoplasm
P32967	GACA_PSEFL		cytoplasm
P33006	CPXL_PSESP		inner membrane
P33007	TERPB_PSESP		cytoplasm
P33008	ALDH_PSESP		cytoplasm
P33009	TERPA_PSESP		cytoplasm
P33010	TERPD_PSESP		cytoplasm
P33014	YEED_ECOLI		cytoplasm
P33018	YEIG_ECOLI		cytoplasm
P33020	YEII_ECOLI		cytoplasm
P33022	RIHB_ECOLI		extracellular
P33025	YEIN_ECOLI		cytoplasm

P33026	SETB_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P33027	SETB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P33029	YEIQ_ECOLI		cytoplasm
P33030	YEIR_ECOLI		cytoplasm
P33037	RECA_ENTAG	cytoplasm (by similarity).	cytoplasm
P33038	MURA_ENTCL	cytoplasm (probable).	inner membrane
P33086	FUR_YERPE	cytoplasm (by similarity).	cytoplasm
P33087	FUR_VIBCH	cytoplasm (by similarity).	cytoplasm
P33109	ASPA_SERMA		cytoplasm
P33117	FUR_VIBVU	cytoplasm (by similarity).	cytoplasm
P33129	HTRE_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P33156	RECA_AGRT5	cytoplasm (by similarity).	cytoplasm
P33160	FDH_PSESR		cytoplasm
P33164	PDR_BURCE		cytoplasm
P33180	TYRO_RHIME		cytoplasm
P33195	GCSP_ECOLI		cytoplasm
P33208	NOLB_RHIFR		cytoplasm
P33209	NOLT_RHIFR	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P33210	NOLU_RHIFR		cytoplasm
P33211	NOLV_RHIFR		cytoplasm
P33212	NOLW_RHIFR		inner membrane
P33213	NOLX_RHIFR		cytoplasm
P33214	YNOL_RHIFR		cytoplasm
P33216	MTLK_RHOSH		cytoplasm
P33217	NOLK_AZOCA		inner membrane
P33221	PURT_ECOLI		cytoplasm
P33222	YJFC_ECOLI		cytoplasm
P33224	AIDB_ECOLI		cytoplasm
P33228	RECT_ECOLI		cytoplasm
P33229	LAR_ECOLI		cytoplasm
P33230	YDAC_ECOLI		fimbrium
P33232	LLDD_ECOLI		cytoplasm
P33234	ADIY_ECOLI		cytoplasm
P33235	FLGK_ECOLI		flagellar
P33340	YEHA_ECOLI		extracellular
P33341	YEHB_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P33342	YEHC_ECOLI	periplasm (by similarity).	periplasm
P33343	YEHD_ECOLI		extracellular
P33344	YEHE_ECOLI		outer membrane
P33345	MOLR_ECOLI		cytoplasm
P33346	YEHI_ECOLI		cytoplasm
P33347	YEHK_ECOLI		cytoplasm
P33348	YEHL_ECOLI		cytoplasm
P33349	YEHM_ECOLI		inner membrane
P33352	YEHP_ECOLI		inner membrane
P33353	YEHQ_ECOLI		cytoplasm

P33354	YEHR_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P33355	YEHS_ECOLI		cytoplasm
P33358	MLRA_ECOLI		cytoplasm
P33360	YEHX_ECOLI		inner membrane
P33362	YEHZ_ECOLI		extracellular
P33368	YOHF_ECOLI		cytoplasm
P33369	MDTQ_ECOLI	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
P33371	DUSC_ECOLI		cytoplasm
P33387	SEFB_SALEN	periplasm (by similarity).	periplasm
P33388	SEFC_SALEN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P33394	RRF1_DESVH		cytoplasm
P33395	RRF2_DESVH		cytoplasm
P33397	HFC1_HAEIN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P33408	MYFC_YEREN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P33409	FIMB_BORPE	periplasm (by similarity).	periplasm
P33410	FIMC_BORPE	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P33451	RPOD_RICPR		cytoplasm
P33452	RPOD_AGRT5		cytoplasm
P33548	IPGB_SHIFL		extracellular
P33553	BFPA_ECO27		fimbrium
P33554	PPDA_ECOLI		cytoplasm
P33566	LEP4_NEIGO	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P33570	TKT2_ECOLI		extracellular
P33590	NIKA_ECOLI	periplasm (probable).	periplasm
P33593	NIKD_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P33594	NIKE_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P33595	YABN_ECOLI		inner membrane
P33596	RECX_ECOLI	cytoplasm (potential).	inner membrane
P33634	YFIE_ECOLI		cytoplasm
P33637	IMM0_ECOLI		cytoplasm
P33639	PILS_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P33640	RLUD_PSEAE		cytoplasm
P33641	Y9F5_PSEAE	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P33642	Y4548_PSEAE		cytoplasm
P33643	RLUD_ECOLI		cytoplasm
P33644	YFIH_ECOLI		cytoplasm
P33647	CHPB_ECOLI		cytoplasm

P33663	Y4543_PSEAE		cytoplasm
P33666	YDBA_ECOLI		extracellular
P33667	SELU_ECOLI		cytoplasm
P33668	YBBC_ECOLI		cytoplasm
P33669	YBBD_ECOLI		inner membrane
P33678	HIP2_RHOTE		periplasm
P33683	CLPX_AZOVI		cytoplasm
P33696	EXON_RHIME		cytoplasm
P33698	EXOP_RHIME	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P33699	EXOT_RHIME	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P33701	EXOV_RHIME		cytoplasm
P33728	OCD_RHIME		cytoplasm
P33772	AMIA_SALTY		extracellular
P33781	FMS5_ECOLI		fimbrium
P33792	YCS5_ECOLI		cytoplasm
P33880	CARI_PECCC		cytoplasm
P33881	EAGI_ENTAG		cytoplasm
P33882	EXPI_PECCC		extracellular
P33883	LASI_PSEAE		cytoplasm
P33898	G3P3_ECOLI	cytoplasm (potential).	cytoplasm
P33901	NUOF_SALTY		periplasm
P33904	TRAI_AGRTU		cytoplasm
P33905	TRAR_AGRTU		cytoplasm
P33906	DEAD_KLEPN	cytoplasm (probable).	cytoplasm
P33907	TRAI_AGRT5		cytoplasm
P33909	TRAR_AGRVI		cytoplasm
P33910	SUCP_AGRVI		cytoplasm
P33913	YEJA_ECOLI		inner membrane
P33916	YEJF_ECOLI		inner membrane
P33919	YEJH_ECOLI		cytoplasm
P33924	YEJO_ECOLI	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P33931	CCMA_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P33937	NAPA_ECOLI	periplasm (potential).	periplasm
P33940	MQO_ECOLI		cytoplasm
P33941	YOJI_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P33968	YLGX_VIBFI		cytoplasm
P33973	PKN1_MYXXA		cytoplasm
P33976	Y911_CAUCR		cytoplasm
P33977	FLBY_CAUCR		inner membrane
P33980	CHEL_CAUCR		inner membrane
P33982	YHBG_AZOCA		inner membrane
P33983	RP54_ACIGB		cytoplasm
P33984	RP54_AZOCA		cytoplasm
P33985	RP542_RHIME		cytoplasm
P33986	MURA_ACIGB	cytoplasm (probable).	inner membrane
P33987	RP5M_ACIGB		cytoplasm

P33989	YRPM_ACIGB		cytoplasm
P33990	SQHC_ZYMMO	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P33997	ALPA_ECOLI		cytoplasm
P34002	AROB_PSEAE	cytoplasm (probable).	cytoplasm
P34003	AROK_PSEAE	cytoplasm (probable).	cytoplasm
P34026	GSPK_XANCP	bacterial cell inner membrane (potential).	inner membrane
P34027	GSPL_XANCP	bacterial cell inner membrane (probable).	inner membrane
P34037	HIS8_ZYMMO		cytoplasm
P34200	FLIP_PECCC	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P34201	FLIQ_PECCC	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P34202	FLIR_PECCC	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P34209	YDCF_ECOLI		cytoplasm
P34211	YUAR_ECOLI		cytoplasm
P34716	RECA_ACIFA	cytoplasm (by similarity).	cytoplasm
P34719	T2M1_MORBO		extracellular
P34720	MT1A_MORBO		cytoplasm
P34721	MT1B_MORBO		cytoplasm
P34749	HOFQ_ECOLI	bacterial cell outer membrane (probable).	outer membrane
P34750	PILQ_PSEAE	bacterial cell outer membrane (probable).	outer membrane
P34816	TRPA_PSESY		cytoplasm
P34817	TRPB_PSESY		cytoplasm
P34818	TRPI_PSESY		cytoplasm
P34879	MTS2_SHISO		cytoplasm
P34880	T2S2_SHISO		extracellular
P34939	CH60_RHILV	cytoplasm (by similarity).	cytoplasm
P35076	FIMA_BORPE		extracellular
P35077	FHAC_BORPE		outer membrane
P35112	NOCR_AGRT7		cytoplasm
P35113	NOCM_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P35114	OCCM_AGRTU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P35116	NOCP_AGRT5	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P35118	NOCQ_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P35120	NOCT_AGRT5	periplasm (probable).	periplasm
P35327	LUXS_VIBF1		cytoplasm
P35328	LUXJ_VIBF1		cytoplasm
P35340	AHPF_ECOLI		cytoplasm

P35469	CH601_RHIME	cytoplasm (by similarity).	cytoplasm
P35470	CH602_RHIME	cytoplasm (by similarity).	cytoplasm
P35471	CH605_RHIME	cytoplasm (by similarity).	cytoplasm
P35473	CH101_RHIME	cytoplasm (by similarity).	cytoplasm
P35474	CH105_RHIME	cytoplasm (by similarity).	cytoplasm
P35530	SPAK_SHIFL		extracellular
P35540	RPOS_SHIFL		cytoplasm
P35633	FLA1_BARBA		flagellar
P35640	NUDH_BARBA		cytoplasm
P35641	IALB_BARBA		extracellular
P35642	RS7_EIKCO		cytoplasm
P35645	ECPA_EIKCO		extracellular
P35646	ECPB_EIKCO		extracellular
P35647	HAG1_EIKCO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P35648	HAG2_EIKCO	secreted protein (potential).	cytoplasm
P35649	YHA2_EIKCO		cytoplasm
P35651	HPAP_RALSO		cytoplasm
P35652	HRCU_RALSO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P35654	HRPI_ERWAM	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P35655	HRPI_PSESY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P35656	HRCV_RALSO	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P35667	GSHB_ANACE		cytoplasm
P35671	INVE_SALTY	probably associated with a bacterial membrane-localized protein.	outer membrane
P35672	INVG_SALTY	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P35673	GALE_ERWAM		cytoplasm
P35695	BLO7_ECOLI		extracellular
P35755	FBPA_HAEIN	periplasm (probable).	periplasm
P35806	FLGE_CAUCR		flagellar
P35818	GSPD_PSEAE	bacterial cell outer membrane; peripheral membrane protein (potential).	inner membrane
P35841	DCRA_DESVH		inner membrane
P35861	CH602_BRAJA	cytoplasm (by similarity).	cytoplasm
P35863	CH102_BRAJA	cytoplasm (by similarity).	cytoplasm
P35864	CH103_BRAJA	cytoplasm (by similarity).	cytoplasm
P35884	TRA2_THIFE		cytoplasm
P35887	DNAA_CAUCR	cytoplasm (by similarity).	cytoplasm
P35890	DNAA_RHIME	cytoplasm (by similarity).	cytoplasm
P35891	DNAA_SALTY	cytoplasm (by similarity).	cytoplasm
P36219	WTSA_ERWST		cytoplasm
P36223	GLND_AZOVI		cytoplasm
P36234	DHGY_HYPME		cytoplasm

P36243	RL19_SERMA		cytoplasm
P36244	TRMD_SERMA	cytoplasm (potential).	cytoplasm
P36245	TRMD_SALTY	cytoplasm (potential).	cytoplasm
P36247	RL7_LIBAS		cytoplasm
P36248	RL1_LIBAS		cytoplasm
P36249	RL10_LIBAS		cytoplasm
P36250	RL11_LIBAS		cytoplasm
P36267	GGT_PSEUA	periplasm (by similarity).	cytoplasm
P36433	T2H2_HAEP		extracellular
P36434	VSH2_HAEP		extracellular
P36547	EUTR_ECOLI		cytoplasm
P36548	AMIA_ECOLI		extracellular
P36555	YJDB_SALTY	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P36556	BASR_SALTY		cytoplasm
P36557	BASS_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P36558	HOBH_ECOLI		cytoplasm
P36560	ASR_ECOLI		periplasm
P36562	COBT_ECOLI		cytoplasm
P36566	SMTA_ECOLI		cytoplasm
P36568	BIOA_SERMA		cytoplasm
P36569	BIOB_SERMA		cytoplasm
P36570	BIOF_SERMA		cytoplasm
P36571	BIOC_SERMA		cytoplasm
P36572	BIOD_SERMA		cytoplasm
P36634	SAPA_SALTY	periplasm (probable).	periplasm
P36636	SAPD_SALTY	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P36638	SAPF_SALTY	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P36640	ATMA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P36641	PILC_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P36643	FMWC_PSEPU		extracellular
P36644	COAE_PSEPU	cytoplasm (by similarity).	cytoplasm
P36645	HOFB_ECOLI		inner membrane
P36647	PPDD_ECOLI		fimbrium
P36661	YCCE_ECOLI		cytoplasm
P36667	WBBL_ECOLI		cytoplasm
P36673	TRER_ECOLI		cytoplasm
P36674	TRER_SALTY		cytoplasm
P36675	YHDL_ECOLI		cytoplasm
P36677	YHDN_ECOLI		cytoplasm
P36678	GSPM_ECOLI	bacterial cell inner membrane (probable).	inner membrane
P36680	YACF_ECOLI		inner membrane
P36682	YACH_ECOLI		inner membrane
P36683	ACON2_ECOLI		cytoplasm

P36684	Y703_HAESO		inner membrane
P36771	LRHA_ECOLI		cytoplasm
P36879	YADG_ECOLI		inner membrane
P36881	YADI_ECOLI	cytoplasm (potential).	inner membrane
P36928	YEGD_ECOLI		cytoplasm
P36929	RSMB_ECOLI		cytoplasm
P36930	GATR_ECOLI		cytoplasm
P36938	PGM_ECOLI		cytoplasm
P36943	EAEH_ECOLI		extracellular
P36979	YFGB_ECOLI		cytoplasm
P36995	CSPB_ECOLI	cytoplasm (by similarity).	cytoplasm
P36999	RRMA_ECOLI		cytoplasm
P37001	CRCA_ECOLI		outer membrane
P37003	YBFG_ECOLI		cytoplasm
P37005	LAST_ECOLI		cytoplasm
P37007	YAGA_ECOLI		cytoplasm
P37008	YAGB_ECOLI		cytoplasm
P37009	FBPC_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P37013	NORR_ECOLI		cytoplasm
P37014	YFAD_ECOLI		cytoplasm
P37016	YADK_ECOLI		extracellular
P37017	YADL_ECOLI		extracellular
P37018	YADM_ECOLI		extracellular
P37019	CLCA_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P37024	HRPB_ECOLI		cytoplasm
P37025	LIGT_ECOLI		cytoplasm
P37029	Y1755_BRAJA		inner membrane
P37030	NIFS_BRAJA		cytoplasm
P37032	ACON_LEGPH		cytoplasm
P37033	Y1689_LEGPH		cytoplasm
P37034	Y1691_LEGPH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P37047	CDAR_ECOLI		cytoplasm
P37049	YAEI_ECOLI		extracellular
P37050	YADN_ECOLI		extracellular
P37051	PURU_ECOLI		cytoplasm
P37052	YCHJ_ECOLI		fimbrium
P37056	YAEF_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P37057	OGRK_ECOLI		cytoplasm
P37076	SCRR_KLEPN		cytoplasm
P37077	SCRR_SALTY		cytoplasm
P37078	SORC_KLEPN		cytoplasm
P37079	SORD_KLEPN		cytoplasm
P37080	PTRA_KLEPN	cytoplasm (potential).	inner membrane
P37082	PTRC_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P37083	PTRD_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane

P37084	SORE_KLEPN		cytoplasm
P37093	GSPE_VIBCH	cytoplasm (probable).	inner membrane
P37094	PILF_NEIGO	cytoplasm (probable).	inner membrane
P37095	PEPB_ECOLI	cytoplasm (by similarity).	cytoplasm
P37097	FER5_RHOCA		cytoplasm
P37098	FER2_CAUCR		cytoplasm
P37099	F16P_NITVU		cytoplasm
P37100	KPPR_NITVU		cytoplasm
P37126	CGLH_XANMA		cytoplasm
P37127	AEGA_ECOLI		cytoplasm
P37128	YFFH_ECOLI		extracellular
P37132	YOPD_YEREN		flagellar
P37133	Y4551_YERPS		cytoplasm
P37143	DHOM_METGL		cytoplasm
P37144	DHON_METGL		cytoplasm
P37145	THRC_METGL		cytoplasm
P37146	RIR4_ECOLI		cytoplasm
P37168	MVIM_SALTY		cytoplasm
P37169	MVIN_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P37174	YBAK_SALTY		cytoplasm
P37177	PT1P_ECOLI	cytoplasm (probable).	cytoplasm
P37178	PT1P_SALTY	cytoplasm (probable).	cytoplasm
P37182	HYBD_ECOLI		extracellular
P37188	PTKB_ECOLI	cytoplasm (potential).	cytoplasm
P37191	GATZ_ECOLI		cytoplasm
P37192	GATY_ECOLI		cytoplasm
P37195	DCTR_ECOLI		cytoplasm
P37197	YHJA_ECOLI		periplasm
P37226	MDH_PHOPR		cytoplasm
P37246	INSI_ECOLI		cytoplasm
P37248	TRA8_RALME		cytoplasm
P37305	HOKA_ECOLI		inner membrane
P37306	ARCC_ECOLI	cytoplasm (potential).	cytoplasm
P37309	ARSR_ECOLI		cytoplasm
P37313	DPPF_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P37321	BLE1_PSEAE		extracellular
P37322	BLC3_PSEAE		extracellular
P37323	BLA4_KLEPN		extracellular
P37325	YBCH_ECOLI		cytoplasm
P37326	INTS_ECOLI		cytoplasm
P37332	BPHF_BURCE		cytoplasm
P37333	BPHA_BURCE		cytoplasm
P37334	BPHE_BURCE		cytoplasm
P37335	YBPA_BURCE		cytoplasm
P37336	YBPE_BURCE		inner membrane
P37337	BPHG_BURCE		cytoplasm
P37338	YGAE_ECOLI		cytoplasm
P37339	YGAF_ECOLI		inner membrane
P37342	YJJI_ECOLI		fimbrium
P37344	PSPF_ECOLI		cytoplasm
P37348	YECE_ECOLI		cytoplasm
P37349	DHAM_ECOLI		cytoplasm

P37351	RPIB_ECOLI		cytoplasm
P37352	HPCE_ECOLI		extracellular
P37353	MENE_ECOLI		cytoplasm
P37355	YFBB_ECOLI		cytoplasm
P37369	SODF_BORPE		periplasm
P37388	XYLG_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P37402	PANE_SALTY	cytoplasm (potential).	cytoplasm
P37408	HEMH_SALTY	cytoplasm (by similarity).	extracellular
P37409	BTUB_SALTY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P37411	GYRA_SALTY		cytoplasm
P37413	META_SALTY	cytoplasm (by similarity).	cytoplasm
P37414	YTL1_SALTY		cytoplasm
P37415	YTL2_SALTY		cytoplasm
P37416	BIRA_SALTY		cytoplasm
P37417	MURB_SALTY	cytoplasm (probable).	inner membrane
P37418	MENE_SALTY		cytoplasm
P37419	KBL_SALTY		inner membrane
P37421	RFAF_SALTY		cytoplasm
P37422	INVH_SALCH	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P37424	LRP_KLEPN		cytoplasm
P37425	LRP_SERMA		cytoplasm
P37426	RIR1_SALTY		cytoplasm
P37427	RIR2_SALTY		cytoplasm
P37430	NUSA_SALTY		cytoplasm
P37431	UBIG_SALTY		cytoplasm
P37432	OMPF_SALTY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P37439	PTGCB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P37440	UCPA_ECOLI		cytoplasm
P37443	YCAI_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P37444	TRPR_SALTY	cytoplasm (by similarity).	cytoplasm
P37449	PDUB_SALTY		cytoplasm
P37450	DHAB_SALTY		cytoplasm
P37452	LEXA_PSEAE		cytoplasm
P37457	WECG_SALTY		cytoplasm
P37458	WZYE_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P37459	SINR_SALTY		cytoplasm
P37461	ZRAS_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P37462	ALKB_SALTY		cytoplasm
P37463	CIT2_SALTY		cytoplasm
P37586	METH_SALTY		cytoplasm
P37587	FLIU_SALMU		inner membrane

P37588	FLIV_SALMU		inner membrane
P37589	PMRD_SALTY		inner membrane
P37590	PMRD_ECOLI		inner membrane
P37593	NARU_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P37594	SMVA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P37595	ASGX_ECOLI		cytoplasm
P37600	PHSA_SALTY		periplasm
P37602	PHSC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P37610	TAUD_ECOLI		cytoplasm
P37613	YHHK_ECOLI		cytoplasm
P37614	YHHL_ECOLI		inner membrane
P37615	YHHM_ECOLI		inner membrane
P37623	ACPT_ECOLI		cytoplasm
P37624	YHIH_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P37626	YHII_ECOLI		inner membrane
P37627	YHIJ_ECOLI		cytoplasm
P37629	YHIL_ECOLI		cytoplasm
P37631	YHIN_ECOLI		cytoplasm
P37634	YHIR_ECOLI		fimbrium
P37635	YHIS_ECOLI		cytoplasm
P37636	MDTE_ECOLI	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P37639	GADX_ECOLI		cytoplasm
P37640	YHJB_ECOLI		cytoplasm
P37641	YHJC_ECOLI		cytoplasm
P37645	YHJG_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P37646	YHJH_ECOLI		inner membrane
P37647	KDGK_ECOLI		cytoplasm
P37648	YHJJ_ECOLI	periplasm (potential).	extracellular
P37649	YHJK_ECOLI		inner membrane
P37650	BCSC_ECOLI		outer membrane
P37653	BCSA_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P37655	YHJQ_ECOLI		inner membrane
P37657	YHJS_ECOLI		cytoplasm
P37659	YHJU_ECOLI		inner membrane
P37663	YHJY_ECOLI		outer membrane
P37664	YIAC_ECOLI		cytoplasm
P37665	YIAD_ECOLI	bacterial cell inner membrane; multi-pass membrane protein; lipid-anchor (probable).	outer membrane
P37666	TKRA_ECOLI	cytoplasm (probable).	cytoplasm
P37671	YIAJ_ECOLI		cytoplasm
P37672	DLGD_ECOLI	cytoplasm (potential).	cytoplasm
P37673	YIAL_ECOLI		inner membrane

P37675	YIAN_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P37677	LYXK_ECOLI		cytoplasm
P37678	SGBH_ECOLI		cytoplasm
P37679	SGBU_ECOLI		cytoplasm
P37680	SGBE_ECOLI		cytoplasm
P37681	YIAT_ECOLI	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P37682	YIAU_ECOLI		cytoplasm
P37685	ALDB_ECOLI		cytoplasm
P37686	ADH2_ECOLI		cytoplasm
P37689	GPMI_ECOLI		cytoplasm
P37690	YIBP_ECOLI		extracellular
P37691	YIBQ_ECOLI		inner membrane
P37692	RFAF_ECOLI		cytoplasm
P37697	CCPA_ACEXY		cytoplasm
P37716	BCSB1_ACEXY	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
P37718	ACSC_ACEXY	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P37719	ACSD_ACEXY		inner membrane
P37722	YAFA_SALTY		cytoplasm
P37723	OSMB_SALTY	bacterial cell membrane; lipid-anchor (by similarity).	inner membrane
P37724	MIAA_SALTY		cytoplasm
P37725	SUMT_PSEFL		cytoplasm
P37728	KDGR_ERWCH		cytoplasm
P37731	MODB_AZOVI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P37732	MODC1_AZOVI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P37733	MODE_AZOVI		cytoplasm
P37736	FUR_VIBAN	cytoplasm (by similarity).	cytoplasm
P37737	FATC_VIBAN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P37738	FATD_VIBAN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P37740	DCTR_RHOCA	cytoplasm (probable).	cytoplasm
P37741	MANC7_ECOLI		cytoplasm
P37742	RFBK7_ECOLI		cytoplasm
P37743	CATA_RHOCA		periplasm
P37744	RMLA1_ECOLI		cytoplasm
P37745	RFBC_ECOLI		cytoplasm
P37747	GLF_ECOLI		cytoplasm
P37749	YEFG_ECOLI	cytoplasm (potential).	cytoplasm
P37750	WBBJ_ECOLI		cytoplasm

P37751	YEFI_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	cytoplasm
P37753	MANC9_ECOLI		cytoplasm
P37754	6PGD9_ECOLI		cytoplasm
P37755	RFBK9_ECOLI		cytoplasm
P37756	6PGD_SHIFL		cytoplasm
P37757	YDDE_ECOLI		cytoplasm
P37759	RFBB1_ECOLI		cytoplasm
P37760	RFBD1_ECOLI		cytoplasm
P37761	RFBB_NEIGO		cytoplasm
P37762	RMLA_NEIGO		cytoplasm
P37763	RFBC_NEIGO		cytoplasm
P37765	RLUB_ECOLI		cytoplasm
P37766	YDIF_ECOLI		cytoplasm
P37767	YFHH_ECOLI		cytoplasm
P37769	KDUD_ECOLI		cytoplasm
P37771	YHFG_SALTY		nucleoid
P37773	MPL_ECOLI		cytoplasm
P37774	YECC_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P37775	WCAM_SHIFL		inner membrane
P37776	GALF_SHIFL		cytoplasm
P37777	RFBB_SHIFL		cytoplasm
P37778	RFBD_SHIFL		cytoplasm
P37779	RMLA1_SHIFL		cytoplasm
P37780	RFBC_SHIFL		cytoplasm
P37781	RFBE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P37782	RFBF_SHIFL		cytoplasm
P37783	RFBG_SHIFL		cytoplasm
P37784	RFC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P37785	GTRA_SHIFL	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P37786	RFBJ_SHIFL		cytoplasm
P37787	Y2094_SHIFL		cytoplasm
P37788	Y2093_SHIFL		inner membrane
P37789	Y2092_SHIFL		inner membrane
P37790	YK9B_SHIFL		cytoplasm
P37791	UDG_SHIFL		cytoplasm
P37793	HIS2_SHIFL	cytoplasm (by similarity).	cytoplasm
P37794	CHBG_ECOLI		inner membrane
P37798	ACCC_PSEAE		cytoplasm
P37799	BCCP_PSEAE		cytoplasm
P37827	FLIP_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P37828	PRO2_XANCG	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P37858	RECA_YERPE	cytoplasm (by similarity).	cytoplasm

P37860	RECX_PSEAE	cytoplasm (potential).	cytoplasm
P37861	RECX_PSEFL	cytoplasm (potential).	cytoplasm
P37862	RECX_PSEPU	cytoplasm (potential).	cytoplasm
P37864	RECX_LEGPN	cytoplasm (potential).	cytoplasm
P37866	RECX_THIFE	cytoplasm (potential).	cytoplasm
P37867	RECX_YERPE	cytoplasm (potential).	cytoplasm
P37868	PEFC_SALTY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P37893	AMPN_CAUCR		extracellular
P37894	PLEC_CAUCR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P37895	Y2483_CAUCR		inner membrane
P37903	USPF_ECOLI		cytoplasm
P37906	PUUB_ECOLI		cytoplasm
P37909	YBGD_ECOLI		extracellular
P37922	FIMI_SALTY		extracellular
P37923	FIMC_SALTY	periplasm (potential).	periplasm
P37924	FIMD_SALTY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P37925	FIMH_SALTY		extracellular
P37926	FIMF_SALTY		extracellular
P37928	FIMW_SALTY		cytoplasm
P37929	HRPL_PSESY		cytoplasm
P37930	HRPR_PSESY		cytoplasm
P37931	HRPS_PSESY		cytoplasm
P37972	CNRA_RALME	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P37973	CNRB_RALME		inner membrane
P37974	CNRC_RALME		inner membrane
P37975	CNRR_RALME		cytoplasm
P37978	CNRH_RALME		cytoplasm
P37982	IHFA_DICD3		cytoplasm
P37983	IHFB_DICD3		cytoplasm
P37986	G6PD_DICD3		cytoplasm
P38035	RTCR_ECOLI		cytoplasm
P38036	YGCB_ECOLI		inner membrane
P38038	CYSJ_ECOLI		cytoplasm
P38039	CYSJ_SALTY		cytoplasm
P38051	MENF_ECOLI		cytoplasm
P38052	SFMF_ECOLI		extracellular
P38054	CUSA_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P38097	YEGE_ECOLI		inner membrane
P38098	CARA_PSEAE		cytoplasm
P38099	CARA_PSEST		cytoplasm
P38100	CARB_PSEAE		cytoplasm
P38102	Y4757_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P38103	DAPB_PSEAE	cytoplasm (by similarity).	cytoplasm
P38104	RSPA_ECOLI		cytoplasm

P38105	RSPB_ECOLI		cytoplasm
P38107	MUCA_PSEAE		inner membrane
P38108	MUCB_PSEAE	periplasm (potential).	periplasm
P38371	MLPA_MYXXA	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P38393	KIL_ECOLI		inner membrane
P38394	YDAE_ECOLI		cytoplasm
P38441	ACE_VIBCH		inner membrane
P38442	ZOT_VIBCH		extracellular
P38443	Y1460_VIBCH		cytoplasm
P38481	YCHJ_SHIFL		inner membrane
P38488	CREA_PSEPU		cytoplasm
P38489	NFNB_ECOLI		cytoplasm
P38506	EX9_ECOLI		cytoplasm
P38521	YGGL_ECOLI		cytoplasm
P38524	HIP2_ECTVA		periplasm
P38569	PGM_ACEXY		cytoplasm
P38587	CY551_ECTHL		periplasm
P38589	HIP_RHOGL		periplasm
P38684	TORR_ECOLI	cytoplasm (probable).	cytoplasm
P38941	HIP1_ECTVA		periplasm
P38974	ETFA_PARDE		cytoplasm
P38975	ETFB_PARDE		cytoplasm
P39049	ALXM_PHOS4		extracellular
P39160	UXUB_ECOLI		cytoplasm
P39161	UXUR_ECOLI		cytoplasm
P39163	CHAC_ECOLI		cytoplasm
P39165	YCHO_ECOLI		extracellular
P39167	TPX_VIBCH		cytoplasm
P39173	YEAD_ECOLI		cytoplasm
P39176	ERFK_ECOLI	periplasm (potential).	periplasm
P39177	USPG_ECOLI		cytoplasm
P39187	YTFJ_ECOLI	periplasm (potential).	periplasm
P39197	PTA_PARDE	cytoplasm (potential).	inner membrane
P39198	AROC_VIBAN		cytoplasm
P39199	YFCB_ECOLI		cytoplasm
P39206	CAIE_ECOLI		cytoplasm
P39208	IDNK_ECOLI		cytoplasm
P39212	INN2_ECOLI		cytoplasm
P39213	INSN_SHIDY		cytoplasm
P39220	YABP_ECOLI		cytoplasm
P39221	YABQ_ECOLI		cytoplasm
P39263	YFCC_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P39264	FIMI_ECOLI		extracellular
P39267	YJCZ_ECOLI		cytoplasm
P39274	YJDJ_ECOLI		cytoplasm
P39279	YJEJ_ECOLI		cytoplasm
P39280	YJEK_ECOLI		cytoplasm
P39283	YJEN_ECOLI		cytoplasm
P39285	YJEP_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P39286	ENGC_ECOLI		cytoplasm
P39288	YJES_ECOLI		periplasm

P39293	YJFK_ECOLI		cytoplasm
P39295	YJFM_ECOLI		outer membrane
P39297	YJFO_ECOLI	bacterial cell membrane; lipid- anchor (potential).	inner membrane
P39298	YJFP_ECOLI		cytoplasm
P39300	ULAG_ECOLI	cytoplasm (probable).	extracellular
P39304	ULAD_ECOLI		cytoplasm
P39305	ULAE_ECOLI		cytoplasm
P39306	ULAF_ECOLI		cytoplasm
P39308	YJFZ_ECOLI		inner membrane
P39309	YTFA_ECOLI		cytoplasm
P39310	YTFB_ECOLI		inner membrane
P39315	YTFG_ECOLI		cytoplasm
P39317	YTFI_ECOLI		cytoplasm
P39321	YTFN_ECOLI		cytoplasm
P39332	YJGH_ECOLI		fimbrium
P39333	YJGI_ECOLI		cytoplasm
P39334	YJGJ_ECOLI		cytoplasm
P39336	YJGL_ECOLI		cytoplasm
P39337	YJGM_ECOLI		cytoplasm
P39342	YJGR_ECOLI		cytoplasm
P39343	IDNR_ECOLI		cytoplasm
P39346	IDND_ECOLI		cytoplasm
P39347	INTB_ECOLI		cytoplasm
P39349	YJGX_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P39351	YJGZ_ECOLI		cytoplasm
P39352	YJHB_ECOLI	bacterial cell inner membrane; multi- pass membrane protein (potential).	inner membrane
P39353	YJHC_ECOLI		cytoplasm
P39354	YJHD_ECOLI		cytoplasm
P39355	YJHE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P39356	YJHU_ECOLI		cytoplasm
P39357	YJHF_ECOLI	bacterial cell inner membrane; multi- pass membrane protein (potential).	inner membrane
P39358	YJHG_ECOLI		cytoplasm
P39359	YJHH_ECOLI		cytoplasm
P39360	YJHI_ECOLI		cytoplasm
P39361	SGCR_ECOLI		cytoplasm
P39362	SGCE_ECOLI		cytoplasm
P39363	SGCA_ECOLI	cytoplasm (probable).	cytoplasm
P39364	SGCQ_ECOLI		cytoplasm
P39366	SGCX_ECOLI		extracellular
P39367	YJHP_ECOLI		inner membrane
P39368	YJHQ_ECOLI		cytoplasm
P39369	YJHR_ECOLI		cytoplasm
P39370	YJHS_ECOLI		periplasm
P39371	YJHT_ECOLI		extracellular
P39374	YJIC_ECOLI		cytoplasm
P39375	YJID_ECOLI		cytoplasm
P39376	YJIE_ECOLI		cytoplasm

P39379	YJIH_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P39380	KPTA_ECOLI		cytoplasm
P39381	YJIJ_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P39382	YJIK_ECOLI		inner membrane
P39383	YJIL_ECOLI		cytoplasm
P39384	YJIM_ECOLI		cytoplasm
P39385	YJIN_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P39387	YJIP_ECOLI		cytoplasm
P39389	YJIR_ECOLI		cytoplasm
P39390	YJIS_ECOLI		cytoplasm
P39391	YJIT_ECOLI		cytoplasm
P39393	YJIV_ECOLI		cytoplasm
P39394	YJIW_ECOLI		cytoplasm
P39399	YJJM_ECOLI		cytoplasm
P39400	YJJN_ECOLI		cytoplasm
P39404	BGLJ_ECOLI		cytoplasm
P39406	RSMC_ECOLI		cytoplasm
P39407	YJJU_ECOLI		cytoplasm
P39408	YJJV_ECOLI		cytoplasm
P39409	YJJW_ECOLI		cytoplasm
P39410	YJJJ_ECOLI		cytoplasm
P39411	YJJX_ECOLI		inner membrane
P39432	YJJX_SALTY		inner membrane
P39434	SLT_SALTY	periplasm (potential).	periplasm
P39448	TRKA_VIBAL		inner membrane
P39450	ADH3_PASPI	cytoplasm (potential).	cytoplasm
P39451	ADHP_ECOLI		cytoplasm
P39452	RIR3_ECOLI		cytoplasm
P39459	NASD_KLEOX	bacterial cell membrane; peripheral membrane protein (potential).	inner membrane
P39669	GLGC_AGRTU		cytoplasm
P39671	PGM_AGRTU		cytoplasm
P39693	RECJ_DICD3		cytoplasm
P39700	NLPD_SALDU	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P39701	COBC_SALTY		cytoplasm
P39818	CSPJ_SALTY	cytoplasm (by similarity).	cytoplasm
P39829	GARD_ECOLI		cytoplasm
P39831	YDFG_ECOLI		cytoplasm
P39834	YGIL_ECOLI		extracellular
P39836	YFEH_ECOLI		inner membrane
P39838	YOJN_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P39849	XYLB_PSEPU		cytoplasm
P39874	YBEM_ECOLI		cytoplasm
P39879	Y2026_PSEAE		inner membrane
P39897	MTRR_NEIGO		cytoplasm
P39901	YBFI_ECOLI		cytoplasm

P39902	SPRT_ECOLI	cytoplasm (potential).	extracellular
P39906	AABA_BACNO	periplasm (potential).	periplasm
P39907	LPSA_BACNO		inner membrane
P39915	AROA2_BURPS	cytoplasm (probable).	cytoplasm
P39916	TRXB_COXBU	cytoplasm (by similarity).	cytoplasm
P39917	LOLA_COXBU	periplasm (by similarity).	periplasm
P39918	Y1189_COXBU		cytoplasm
P40117	RPEC_RALEU		cytoplasm
P40118	CBXXC_RALEU		cytoplasm
P40119	CBBYC_RALEU		cytoplasm
P40128	CYAY_ERWCH		inner membrane
P40131	FLGA_SALTY	periplasm (probable).	flagellar
P40137	CYAA_STIAU	bacterial cell membrane; multi-pass membrane protein (potential).	cytoplasm
P40138	CYAB_STIAU		cytoplasm
P40149	OXC_OXAFO		cytoplasm
P40191	PDXK_ECOLI		cytoplasm
P40192	PDXK_SALTY		cytoplasm
P40193	PTSJ_SALTY		cytoplasm
P40194	YFEJ_SALTY		cytoplasm
P40195	YFEK_SALTY		cytoplasm
P40196	YFEL_SALTY		inner membrane
P40294	YSCO_YERPS		extracellular
P40296	YSCQ_YERPS		inner membrane
P40330	BVGS_BORPA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P40390	PGM_NEIGO		cytoplasm
P40391	PGM_NEIMB		cytoplasm
P40428	HYPC_RHOCA		cytoplasm
P40431	YVN1_AZOVI		cytoplasm
P40432	YVN2_AZOVI		cytoplasm
P40591	HOXM_AZOVI		extracellular
P40592	HOXL_AZOVI		cytoplasm
P40593	HOXO_AZOVI		cytoplasm
P40594	HOXQ_AZOVI		cytoplasm
P40595	HYPE_AZOVI		cytoplasm
P40596	HYPF_AZOVI		cytoplasm
P40597	HOXV_AZOVI		cytoplasm
P40598	HYPD_RHILV		extracellular
P40599	HYPE_RHILV		cytoplasm
P40604	YTRP_PSEPU		extracellular
P40605	HFLK_VIBPA		inner membrane
P40606	HFLC_VIBPA		extracellular
P40607	PURA_VIBPA	cytoplasm (by similarity).	cytoplasm
P40609	HMP_VIBPA		cytoplasm
P40610	Y2808_VIBPA		inner membrane
P40611	RNR_VIBPA		cytoplasm
P40612	SPAM_SALTY		extracellular
P40613	SPAN_SALTY	cytoplasm (potential).	extracellular
P40676	SLYA_SALTY		cytoplasm
P40677	YJAB_SALTY		cytoplasm
P40680	ERFK_SALTY	periplasm (potential).	periplasm
P40685	CHRR_RHOS4		cytoplasm
P40687	TOP3_SALTY		cytoplasm

P40699	SPAO_SALTY		extracellular
P40700	SPAP_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P40701	SPAR_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P40702	SPAS_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P40709	YEJM_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P40711	YAEJ_ECOLI		cytoplasm
P40713	SCRK_ECOLI		cytoplasm
P40720	FUMA_SALTY		cytoplasm
P40722	SOPD_SALTY		inner membrane
P40723	MURI_SALTY		cytoplasm
P40724	RECQ_SALTY		cytoplasm
P40727	FLHB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P40730	YJBB_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P40731	GCP_SALTY		extracellular
P40732	ARGD_SALTY	cytoplasm (probable).	cytoplasm
P40733	DGTP_SALTY		cytoplasm
P40734	FHLA_SALTY		cytoplasm
P40783	QOR_SALTY		cytoplasm
P40784	YCDD_SALTY		periplasm
P40788	YDDE_SALTY		cytoplasm
P40790	POTA_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P40800	YGIK_SALTY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P40810	ILVD_SALTY		cytoplasm
P40811	ILVI_SALTY		cytoplasm
P40813	T1M_SALTY		cytoplasm
P40814	T3MO_SALTY		cytoplasm
P40815	T3RE_SALTY		extracellular
P40816	HEMK_SALTY		cytoplasm
P40817	DPAL_SALTY		cytoplasm
P40819	YHHV_SALTY		cytoplasm
P40822	YCEB_SALTY	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P40824	YIBR_SALTY		cytoplasm
P40827	NLPD_SALTY	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P40852	GPHC_RALEU		cytoplasm
P40853	GPHP_RALEU		cytoplasm
P40860	CYSA_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

P40861	PROA_SALTY	cytoplasm (by similarity).	cytoplasm
P40865	RHAR_SALTY	cytoplasm (by similarity).	cytoplasm
P40874	MTOX_ECOLI		cytoplasm
P40876	YCBF_ECOLI	periplasm (by similarity).	periplasm
P40882	Y3753_PSEAE		cytoplasm
P40883	PCHR_PSEAE		cytoplasm
P40947	SSB_PSEAE		cytoplasm
P41030	GALS_SALTY		cytoplasm
P41032	CYST_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P41033	PPCK_SALTY	cytoplasm (by similarity).	cytoplasm
P41039	YBII_ECOLI		cytoplasm
P41065	TRAR_ECOLI		cytoplasm
P41066	TRAK1_ECOLI	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P41067	TRAB1_ECOLI		cytoplasm
P41068	TRAP_ECOLI		cytoplasm
P41070	TRBD_ECOLI		cytoplasm
P41072	TRBG_ECOLI		cytoplasm
P41077	PNTAA_RICPR		inner membrane
P41078	Y860_RICPR		inner membrane
P41079	RECA_RICPR	cytoplasm (by similarity).	cytoplasm
P41080	GYRA_RICPR		cytoplasm
P41081	RS7_RICPR		cytoplasm
P41082	RS12_RICPR		cytoplasm
P41087	Y125_RICPR		inner membrane
P41107	RL10_BRUAB		cytoplasm
P41120	RS15_PHOLU		cytoplasm
P41142	ARCA_PSEPU	cytoplasm (potential).	extracellular
P41167	ATPA_THIFE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P41168	ATPB_THIFE		inner membrane
P41169	ATPG_THIFE		inner membrane
P41170	ATPD_THIFE		inner membrane
P41171	ATPE_THIFE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P41172	ATPF_THIFE		inner membrane
P41173	ATPL_THIFE		inner membrane
P41184	RPOB_BUCAP		cytoplasm
P41185	RPOC_BUCAP		cytoplasm
P41186	RS4_BUCAP		cytoplasm
P41188	RL7_BUCAP		cytoplasm
P41189	RL7_LIBAF		cytoplasm
P41191	RL10_LIBAF		cytoplasm
P41257	SYI_CAMJE	cytoplasm (by similarity).	cytoplasm
P41278	RL23_YEREN		cytoplasm
P41302	LUXD_PHOPO		cytoplasm
P41393	GLND_KLEOX		cytoplasm
P41396	DAPD_ACTPL	cytoplasm (by similarity).	cytoplasm
P41399	DHAS_BORPE		cytoplasm
P41407	AZOR_ECOLI		cytoplasm
P41409	RIHA_ECOLI		extracellular
P41442	GSPG_ECOLI		inner membrane

P41443	GSPH_ECOLI		inner membrane
P41501	HRPK_PSESY		extracellular
P41503	NTRB_RHILP		inner membrane
P41504	DUS_RHILP		cytoplasm
P41513	GYRA_ERWCA		cytoplasm
P41560	IDH1_COLMA		cytoplasm
P41561	IDH2_COLMA		cytoplasm
P41576	6PGD_KLEPN		cytoplasm
P41773	LIPB_PSEFL		extracellular
P41780	APBE_SALTY	bacterial cell membrane; lipid- anchor (potential).	cytoplasm
P41782	HILD_SALTY		cytoplasm
P41783	PRGH_SALTY		inner membrane
P41784	PRGI_SALTY		inner membrane
P41785	PRGJ_SALTY		inner membrane
P41786	PRGK_SALTY	bacterial cell outer membrane; lipid- anchor (probable).	inner membrane
P41789	NTRC_SALTY		cytoplasm
P41790	EUTD_SALTY		inner membrane
P41791	EUTM_SALTY		cytoplasm
P41792	EUTN_SALTY		cytoplasm
P41793	EUTE_SALTY		cytoplasm
P41795	EUTG_SALTY		cytoplasm
P41796	EUTH_SALTY	bacterial cell inner membrane; multi- pass membrane protein (by similarity).	inner membrane
P41850	GSPM_AERHY	bacterial cell inner membrane (probable).	inner membrane
P41851	GSPM_VIBCH	bacterial cell inner membrane (probable).	inner membrane
P41852	GSPN_AERHY	bacterial cell inner membrane (potential).	inner membrane
P42030	YNDH_RHOCA		cytoplasm
P42031	NUOI_RHOCA		cytoplasm
P42033	HUPD_AZOCH		extracellular
P42034	HUPE_AZOCH		cytoplasm
P42179	BKDR_PSEPU		cytoplasm
P42191	PRSK_ECOLI		extracellular
P42206	GUDH_PSEPU		cytoplasm
P42216	KPSU5_ECOLI	cytoplasm (potential).	cytoplasm
P42217	KPSC5_ECOLI	cytoplasm (potential).	cytoplasm
P42218	KPSS5_ECOLI		cytoplasm
P42233	KDGD_PSEPU		cytoplasm
P42257	PILJ_PSEAE	bacterial cell inner membrane; multi- pass membrane protein (potential).	inner membrane
P42269	HPC_C_ECOLI		cytoplasm
P42270	HPCG_ECOLI		cytoplasm
P42272	FLIC1_PROMI		flagellar
P42273	FLIC2_PROMI		flagellar
P42373	DNAK_BURCE		cytoplasm
P42374	DNAK_RHIME		cytoplasm
P42378	RP32_PSEAE		cytoplasm
P42381	DNAJ_COXBU	cytoplasm (probable).	cytoplasm
P42382	CH60_EHRCH	cytoplasm (by similarity).	cytoplasm
P42383	CH60_HELPY	cytoplasm (by similarity).	cytoplasm

P42385	CH60_NEIMB	cytoplasm (by similarity).	cytoplasm
P42386	CH10_EHRCH	cytoplasm (by similarity).	cytoplasm
P42387	TRPE_BUCAP		cytoplasm
P42388	TRPG_BUCAP		cytoplasm
P42389	TRPA_BUCAP		cytoplasm
P42391	TRPB_BUCAP		cytoplasm
P42392	TRPD_BUCAP		cytoplasm
P42393	TRPC_BUCAP		cytoplasm
P42394	Y270_BUCAP	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P42395	RLUB_BUCAP		cytoplasm
P42396	Y265_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P42397	ISPZ_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P42398	Y263_BUCAP		cytoplasm
P42428	TFDD2_RALEJ		cytoplasm
P42438	RECA_ACIAD	cytoplasm (by similarity).	cytoplasm
P42440	RECA_CAMJE	cytoplasm (by similarity).	cytoplasm
P42445	RECA_HELPY	cytoplasm (by similarity).	cytoplasm
P42447	RECA_RHOCA	cytoplasm (by similarity).	cytoplasm
P42448	ENO_CAMJE	cytoplasm (by similarity).	extracellular
P42453	RUBR_ACIAD		cytoplasm
P42468	ATPB_BURCE		inner membrane
P42469	ATPB_STIAU		inner membrane
P42470	ATPB_WOLSU		inner membrane
P42501	KPSE1_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P42502	KSF1_ECOLI		cytoplasm
P42503	DCUP_RHOCA	cytoplasm (probable).	cytoplasm
P42504	HEM2_RHOCA		cytoplasm
P42505	HVRA_RHOCA		cytoplasm
P42506	YAHC_RHOCA		cytoplasm
P42507	HVRB_RHOCA		cytoplasm
P42508	REGA_RHOCA		cytoplasm
P42513	FPTB_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P42514	Y4219_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P42535	PCPB_SPHCR		cytoplasm
P42588	OAT_ECOLI		cytoplasm
P42589	YGJH_ECOLI		cytoplasm
P42591	YGJJ_ECOLI		outer membrane
P42592	YGJK_ECOLI		periplasm
P42593	FADH_ECOLI		cytoplasm
P42596	RSMD_ECOLI		cytoplasm
P42597	YGJP_ECOLI		cytoplasm
P42598	YGJQ_ECOLI		inner membrane
P42599	YGJR_ECOLI		inner membrane
P42604	UXAA_ECOLI		cytoplasm

P42615	YQJB_ECOLI		inner membrane
P42616	YQJC_ECOLI		cytoplasm
P42620	YQJG_ECOLI		cytoplasm
P42624	YHAK_ECOLI		cytoplasm
P42625	YHAL_ECOLI		cytoplasm
P42626	YHAM_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P42630	TDCG_ECOLI		cytoplasm
P42632	TDCE_ECOLI	cytoplasm (by similarity).	cytoplasm
P42641	YHBZ_ECOLI		cytoplasm
P42693	PPIA_ACIAD	periplasm (probable).	periplasm
P42710	FIXU_RHILT		cytoplasm
P42711	FDXN_RHILT		periplasm
P42713	YSCQ_YERPE		inner membrane
P42719	PYRE_RHILT		cytoplasm
P42720	SCRK_RHILT		cytoplasm
P42721	RBL1P_RALEU		extracellular
P42722	CFXR_RALEU		cytoplasm
P42724	TFXB_RHILT	cytoplasm (potential).	cytoplasm
P42725	TFXC_RHILT		cytoplasm
P42726	TFXD_RHILT	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P42727	TFXE_RHILT		cytoplasm
P42728	TFXF_RHILT		cytoplasm
P42729	TFXG_RHILT		cytoplasm
P42786	PIP_NEIGO	cytoplasm (probable).	cytoplasm
P42805	ISPE_PSEAE		cytoplasm
P42807	HEM1_PSEAE		cytoplasm
P42808	HEM1_XANCH		cytoplasm
P42810	Y4667_PSEAE		cytoplasm
P42812	LOLB_PSEAE	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P42822	URE23_HELHE	cytoplasm (by similarity).	cytoplasm
P42823	URE1_HELHE	cytoplasm (by similarity).	cytoplasm
P42868	URED_YEREN		cytoplasm
P42869	UREE_YEREN	cytoplasm (by similarity).	cytoplasm
P42870	UREF_YEREN		cytoplasm
P42871	UREG_YEREN		cytoplasm
P42872	TONB_HAEIN	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
P42878	Y2474_RHIME		periplasm
P42879	Y2472_RHIME		cytoplasm
P42880	UREY_RHIME		cytoplasm
P42885	URE1_RHIME	cytoplasm (by similarity).	cytoplasm
P42886	URE2_RHIME	cytoplasm (by similarity).	cytoplasm
P42887	URE3_RHIME	cytoplasm (by similarity).	cytoplasm
P42888	URED_RHIME		cytoplasm
P42903	AGAZ_ECOLI		cytoplasm
P42904	PTPB2_ECOLI	cytoplasm (potential).	cytoplasm
P42906	AGAA_ECOLI		cytoplasm
P42907	AGAS_ECOLI		cytoplasm
P42909	PTPB1_ECOLI	cytoplasm (potential).	cytoplasm

P42912	AGAI_ECOLI		cytoplasm
P42913	YRAH_ECOLI		extracellular
P42914	YRAI_ECOLI	periplasm (by similarity).	periplasm
P42915	YRAJ_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P43010	PNTB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P43011	Y1364_HAEIN		cytoplasm
P43012	TOP1_HAEIN		cytoplasm
P43013	ACPD_HAEIN		extracellular
P43015	HILA_SALTY		cytoplasm
P43016	HILA_SALTI		cytoplasm
P43017	IAGB_SALTY		extracellular
P43018	IAGB_SALTI		extracellular
P43019	SODM_SALTY		periplasm
P43020	YIIZ_SALTY		periplasm
P43085	PPCK_RHIME	cytoplasm (by similarity).	cytoplasm
P43086	PPCK_RHISN	cytoplasm (by similarity).	cytoplasm
P43089	HEM1_PARDE		inner membrane
P43109	VEXB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P43110	VEXC_SALTI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P43111	VEXD_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P43112	VEXE_SALTI		cytoplasm
P43126	FRE_VIBFI		cytoplasm
P43127	FRE_VIBHA		cytoplasm
P43129	FRE_PHOLU		cytoplasm
P43312	SODF_HELPY		periplasm
P43313	NAPA_HELPY		cytoplasm
P43319	YRAK_ECOLI		extracellular
P43329	HRPA_ECOLI		cytoplasm
P43334	PH4H_PSEAE		cytoplasm
P43335	PHS_PSEAE		cytoplasm
P43337	YEAB_ECOLI		cytoplasm
P43340	YCAK_ECOLI		cytoplasm
P43343	MUG_SERMA	cytoplasm (potential).	extracellular
P43387	DCMA_METS1	cytoplasm (probable).	cytoplasm
P43413	HEMH_YEREN	cytoplasm (by similarity).	extracellular
P43418	T2G1_HAEGA		extracellular
P43459	PERA_ECO27		cytoplasm
P43460	CSVR_ECOLI		cytoplasm
P43463	AARP_PROST		cytoplasm
P43464	AGGR_ECOLI		cytoplasm
P43474	PERB_ECO27		cytoplasm
P43475	PERC_ECO27		cytoplasm
P43476	PERD_ECO27		cytoplasm
P43498	FRZA_MYXXA		cytoplasm
P43499	FRZB_MYXXA		cytoplasm
P43501	PILH_PSEAE		cytoplasm

P43502	PILI_PSEAE		cytoplasm
P43503	XYLC_PSEPU		cytoplasm
P43505	MTRC_NEIGO	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P43518	GLNA_RHOSH	cytoplasm (by similarity).	cytoplasm
P43519	GLNB_RHOSH		cytoplasm
P43525	GCH2_AZOBR		extracellular
P43528	E2BA_ECOLI		extracellular
P43529	E2BB_ECOLI		extracellular
P43530	ELAH_ECOLI		extracellular
P43533	FLGN_ECOLI		flagellar
P43660	LPFA_SALTY		extracellular
P43661	LPFB_SALTY	periplasm (by similarity).	periplasm
P43662	LPFC_SALTY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P43663	LPFD_SALTY		extracellular
P43664	LPFE_SALTY		extracellular
P43666	YHJW_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P43667	YGAH_ECOLI		inner membrane
P43669	PRC_SALTY	bacterial cell inner membrane; peripheral membrane protein; periplasmic side (by similarity).	cytoplasm
P43671	PQIB_ECOLI		inner membrane
P43672	UUP_ECOLI		inner membrane
P43674	YCAL_ECOLI		extracellular
P43700	GYRA_HAEIN		cytoplasm
P43701	GYRB_HAEIN		cytoplasm
P43702	PARC_HAEIN	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P43703	PARE_HAEIN		cytoplasm
P43704	TOP3_HAEIN		cytoplasm
P43705	RECA_HAEIN	cytoplasm (by similarity).	cytoplasm
P43706	RECX_HAEIN	cytoplasm (potential).	cytoplasm
P43707	FTN1_HAEIN	cytoplasm (by similarity).	cytoplasm
P43708	FTN2_HAEIN	cytoplasm (by similarity).	cytoplasm
P43709	ACP_HAEIN	cytoplasm (by similarity).	cytoplasm
P43711	FABH_HAEIN	cytoplasm (probable).	cytoplasm
P43712	FABD_HAEIN		cytoplasm
P43713	FABG_HAEIN		cytoplasm
P43714	ATPA_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P43715	ATPB_HAEIN		inner membrane
P43716	ATPG_HAEIN		inner membrane
P43717	ATPD_HAEIN		inner membrane
P43718	ATPE_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P43720	ATPF_HAEIN		inner membrane
P43721	ATPL_HAEIN		inner membrane
P43722	DBH_HAEIN		cytoplasm

P43723	IHFA_HAEIN		cytoplasm
P43724	IHFB_HAEIN		cytoplasm
P43725	SODM_HAEIN		periplasm
P43727	TPIS_HAEIN		cytoplasm
P43728	ERA_HAEIN		cytoplasm
P43729	LEPA_HAEIN	bacterial cell membrane (by similarity). periplasm (by similarity).	cytoplasm
P43730	TRME_HAEIN		cytoplasm
P43731	UNG_HAEIN	cytoplasm (by similarity).	extracellular
P43732	GRPE_HAEIN	cytoplasm (probable).	cytoplasm
P43733	CH60_HAEIN	cytoplasm (by similarity).	cytoplasm
P43734	CH10_HAEIN	cytoplasm (by similarity).	cytoplasm
P43735	DNAJ_HAEIN	cytoplasm (probable).	cytoplasm
P43736	DNAK_HAEIN		cytoplasm
P43737	RPOA_HAEIN		cytoplasm
P43738	RPOB_HAEIN		cytoplasm
P43739	RPOC_HAEIN		cytoplasm
P43740	RPOZ_HAEIN		cytoplasm
P43741	DPO1_HAEIN		cytoplasm
P43742	DNAA_HAEIN	cytoplasm (by similarity).	cytoplasm
P43743	DPO3A_HAEIN	cytoplasm (by similarity).	cytoplasm
P43744	DPO3B_HAEIN	cytoplasm (by similarity).	cytoplasm
P43745	DPO3E_HAEIN		cytoplasm
P43746	DPO3X_HAEIN		cytoplasm
P43747	HOLA_HAEIN		cytoplasm
P43748	HOLB_HAEIN		cytoplasm
P43749	HOLC_HAEIN		cytoplasm
P43750	HOLD_HAEIN		cytoplasm
P43751	PFLA_HAEIN	cytoplasm (by similarity).	cytoplasm
P43752	NRDD_HAEIN		cytoplasm
P43753	PFLB_HAEIN	cytoplasm (by similarity).	cytoplasm
P43754	RIR1_HAEIN		cytoplasm
P43755	RIR2_HAEIN		cytoplasm
P43756	FOCA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P43757	TKT_HAEIN		extracellular
P43758	DKSA_HAEIN		cytoplasm
P43759	TRPA_HAEIN		cytoplasm
P43760	TRPB_HAEIN		cytoplasm
P43761	TRPE_HAEIN		cytoplasm
P43763	PROB_HAEIN	cytoplasm (by similarity).	cytoplasm
P43764	GCP_HAEIN		extracellular
P43766	RPOD_HAEIN		cytoplasm
P43767	RECF_HAEIN	cytoplasm (potential).	cytoplasm
P43768	TOLQ_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P43769	TOLR_HAEIN	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P43770	UDP_HAEIN	cytoplasm (by similarity).	cytoplasm
P43772	HSLV_HAEIN	cytoplasm (by similarity).	cytoplasm
P43773	HSLU_HAEIN	cytoplasm (by similarity).	cytoplasm
P43774	6PGD_HAEIN		cytoplasm

P43775	FOLC_HAEIN		cytoplasm
P43776	DHPS_HAEIN		cytoplasm
P43777	HPPK_HAEIN		cytoplasm
P43778	ACCD_HAEIN		cytoplasm
P43779	TFOX_HAEIN		cytoplasm
P43780	BOLA_HAEIN		cytoplasm
P43782	Y162_HAEIN	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P43785	THIO_HAEIN		cytoplasm
P43786	Y1159_HAEIN		extracellular
P43787	THIX_HAEIN		inner membrane
P43788	TRXB_HAEIN	cytoplasm (by similarity).	cytoplasm
P43789	PSD_HAEIN	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P43790	Y159_HAEIN		cytoplasm
P43791	DYR_HAEIN		cytoplasm
P43792	DUT_HAEIN		extracellular
P43793	DHE4_HAEIN		cytoplasm
P43795	GLNB_HAEIN		cytoplasm
P43796	GLGC_HAEIN		cytoplasm
P43797	DAPA_HAEIN	cytoplasm (by similarity).	cytoplasm
P43798	GPDA_HAEIN	cytoplasm (probable).	cytoplasm
P43800	GLPB_HAEIN		cytoplasm
P43802	NDK_HAEIN	cytoplasm (by similarity).	cytoplasm
P43803	SECA_HAEIN	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	inner membrane
P43804	SECY_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P43805	SECE_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P43806	ENO_HAEIN	cytoplasm (by similarity).	extracellular
P43807	RNH_HAEIN	cytoplasm (potential).	extracellular
P43808	RNH2_HAEIN	cytoplasm (potential).	extracellular
P43809	RECG_HAEIN		cytoplasm
P43810	ISPZ_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P43811	SPOT_HAEIN		cytoplasm
P43812	PYRF_HAEIN		cytoplasm
P43813	DNLJ_HAEIN		cytoplasm
P43824	SYI_HAEIN	cytoplasm (by similarity).	cytoplasm
P43826	SYK3_HAEIN		cytoplasm
P43834	SYV_HAEIN	cytoplasm (by similarity).	cytoplasm
P43836	SYV_HAEIN	cytoplasm (by similarity).	cytoplasm
P43837	MAO2_HAEIN		cytoplasm
P43841	HNS_HAEIN		cytoplasm
P43842	PNTA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P43843	ASPG2_HAEIN	periplasm (by similarity).	periplasm
P43845	PUR2_HAEIN		cytoplasm

P43846	PUR3_HAEIN		cytoplasm
P43847	PUR4_HAEIN	cytoplasm (by similarity).	cytoplasm
P43848	PUR5_HAEIN	cytoplasm (by similarity).	cytoplasm
P43849	PUR6_HAEIN		cytoplasm
P43850	PURK_HAEIN		cytoplasm
P43851	PUR7_HAEIN		cytoplasm
P43852	PUR9_HAEIN		cytoplasm
P43853	HIS1_HAEIN	cytoplasm (by similarity).	cytoplasm
P43854	PUR1_HAEIN		cytoplasm
P43855	PYRE_HAEIN		cytoplasm
P43857	UPP_HAEIN		cytoplasm
P43858	TRPD_HAEIN		cytoplasm
P43861	LEU1_HAEIN		cytoplasm
P43862	SMF_HAEIN		cytoplasm
P43863	K6PF_HAEIN	cytoplasm (by similarity).	cytoplasm
P43865	LONH_HAEIN		cytoplasm
P43866	GCH1_HAEIN		cytoplasm
P43867	CLPP_HAEIN	cytoplasm (by similarity).	cytoplasm
P43868	HEMH_HAEIN	cytoplasm (by similarity).	extracellular
P43869	PROC_HAEIN	cytoplasm (by similarity).	cytoplasm
P43870	T2D3_HAEIN		extracellular
P43871	MTH3_HAEIN		cytoplasm
P43872	ACCA_HAEIN	cytoplasm (by similarity).	cytoplasm
P43873	ACCC_HAEIN		cytoplasm
P43874	BCCP_HAEIN		cytoplasm
P43875	AROC_HAEIN		cytoplasm
P43876	AROE_HAEIN		cytoplasm
P43877	AROQ_ACTPL		cytoplasm
P43878	AROQ_HAEIN		cytoplasm
P43879	AROB_HAEIN	cytoplasm (probable).	cytoplasm
P43880	AROK_HAEIN	cytoplasm (probable).	cytoplasm
P43881	GREA_HAEIN		cytoplasm
P43882	GREB_HAEIN		cytoplasm
P43885	SERA_HAEIN		cytoplasm
P43886	CYSE_HAEIN	cytoplasm (by similarity).	cytoplasm
P43887	LPXA_HAEIN	cytoplasm (by similarity).	cytoplasm
P43888	LPXD_HAEIN		cytoplasm
P43889	GLMU_HAEIN	cytoplasm (by similarity).	cytoplasm
P43890	PYRH_HAEIN		cytoplasm
P43892	KCY1_HAEIN	cytoplasm (by similarity).	cytoplasm
P43893	KCY2_HAEIN	cytoplasm (by similarity).	cytoplasm
P43898	HEM6_PSEAE	cytoplasm (by similarity).	cytoplasm
P43899	Y463_HAEIN		cytoplasm
P43903	QOR_PSEAE		cytoplasm
P43904	AROE_PSEAE		cytoplasm
P43907	Y983_HAEIN		cytoplasm
P43908	Y984_HAEIN		inner membrane
P43910	SELA_HAEIN	cytoplasm (by similarity).	cytoplasm
P43911	SELD_HAEIN		cytoplasm
P43912	TRMD_HAEIN	cytoplasm (potential).	cytoplasm
P43913	EX7L_HAEIN	cytoplasm (by similarity).	cytoplasm
P43914	EX7S_HAEIN	cytoplasm (by similarity).	cytoplasm
P43915	NUSA_HAEIN		cytoplasm
P43916	NUSG_HAEIN		cytoplasm
P43919	GLND_HAEIN		cytoplasm
P43920	CAPP_HAEIN		cytoplasm

P43921	PTHP_HAEIN	cytoplasm (by similarity).	cytoplasm
P43923	PPCK_HAEIN	cytoplasm (by similarity).	cytoplasm
P43924	KPYK_HAEIN		cytoplasm
P43929	Y040_HAEIN		cytoplasm
P43930	Y042_HAEIN		cytoplasm
P43931	Y043_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P43932	Y056_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P43933	Y073_HAEIN		cytoplasm
P43934	Y074_HAEIN		cytoplasm
P43935	Y077_HAEIN		cytoplasm
P43936	Y080_HAEIN		cytoplasm
P43937	Y082_HAEIN		cytoplasm
P43938	Y083_HAEIN		cytoplasm
P43939	Y094_HAEIN		inner membrane
P43940	Y096_HAEIN		inner membrane
P43941	Y100_HAEIN		cytoplasm
P43942	Y101_HAEIN		cytoplasm
P43943	Y109_HAEIN		cytoplasm
P43947	Y120_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P43948	MPL_HAEIN		cytoplasm
P43951	Y131_HAEIN		inner membrane
P43952	Y134_HAEIN		inner membrane
P43953	Y149_HAEIN		inner membrane
P43954	Y152_HAEIN		cytoplasm
P43955	NQRA_HAEIN		inner membrane
P43957	NQRC_HAEIN	bacterial cell inner membrane (potential).	inner membrane
P43958	NQRD_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P43960	Y173_HAEIN		inner membrane
P43961	Y178_HAEIN		cytoplasm
P43963	Y205_HAEIN		cytoplasm
P43965	Y217_HAEIN		cytoplasm
P43966	Y228_HAEIN		inner membrane
P43967	Y233_HAEIN		cytoplasm
P43969	PERM_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P43971	Y243_HAEIN		nucleoid
P43972	Y246_HAEIN		periplasm
P43973	Y256_HAEIN		cytoplasm
P43974	Y258_HAEIN		cytoplasm
P43975	Y275_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P43976	Y278_HAEIN		cytoplasm
P43977	Y279_HAEIN		cytoplasm
P43979	Y291_HAEIN		inner membrane
P43980	Y304_HAEIN		inner membrane

P43981	RUVX_HAEIN	cytoplasm (potential).	cytoplasm
P43982	Y310_HAEIN		cytoplasm
P43984	Y318_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P43985	Y319_HAEIN		cytoplasm
P43987	Y326_HAEIN		cytoplasm
P43988	Y366_HAEIN		extracellular
P43989	Y370_HAEIN		inner membrane
P43990	Y388_HAEIN		cytoplasm
P43991	Y389_HAEIN		inner membrane
P43992	Y391_HAEIN		cytoplasm
P43993	Y392_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P43994	Y395_HAEIN		inner membrane
P43995	Y420_HAEIN		cytoplasm
P43996	Y421_HAEIN		inner membrane
P43997	Y449_HAEIN		cytoplasm
P43998	Y451_HAEIN		cytoplasm
P43999	Y453_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44000	Y466_HAEIN		cytoplasm
P44001	HISX_HAEIN		cytoplasm
P44002	Y476_HAEIN		cytoplasm
P44003	Y487_HAEIN		cytoplasm
P44004	Y488_HAEIN		cytoplasm
P44005	Y489_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44007	LUXS_HAEIN		extracellular
P44009	APHA_HAEIN	periplasm (potential).	periplasm
P44010	Y507_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44011	Y523_HAEIN		cytoplasm
P44012	RN26_HAEIN		extracellular
P44013	Y552_HAEIN		cytoplasm
P44014	Y554_HAEIN		cytoplasm
P44016	Y561_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44017	Y577_HAEIN		cytoplasm
P44018	Y585_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44019	Y586_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44020	Y589_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44021	Y592_HAEIN		cytoplasm

P44023	Y594_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44025	Y627_HAEIN		cytoplasm
P44026	Y633_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44027	Y636_HAEIN		cytoplasm
P44028	Y650_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44029	Y653_HAEIN		cytoplasm
P44030	Y659_HAEIN		cytoplasm
P44031	Y660_HAEIN		cytoplasm
P44033	Y665_HAEIN		cytoplasm
P44034	Y666_HAEIN		cytoplasm
P44035	FTSB_HAEIN	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
P44036	Y677_HAEIN		cytoplasm
P44037	Y688_HAEIN		cytoplasm
P44038	Y698_HAEIN		outer membrane
P44039	TRUD_HAEIN		cytoplasm
P44040	Y704_HAEIN		inner membrane
P44041	Y711_HAEIN		inner membrane
P44042	VACJ_HAEIN	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P44043	Y725_HAEIN		inner membrane
P44044	Y731_HAEIN		cytoplasm
P44045	Y732_HAEIN		inner membrane
P44046	LPXH_HAEIN	cytoplasm (by similarity).	cytoplasm
P44047	Y742_HAEIN		cytoplasm
P44048	FETP_HAEIN		cytoplasm
P44049	MLTC_HAEIN	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
P44050	Y762_HAEIN		extracellular
P44051	ATOE_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44052	Y787_HAEIN		cytoplasm
P44053	Y804_HAEIN		cytoplasm
P44054	Y806_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44055	DXR_HAEIN		cytoplasm
P44056	Y825_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44057	SMPA_HAEIN	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P44058	Y842_HAEIN		cytoplasm
P44060	Y850_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane

P44061	Y854_HAEIN		cytoplasm
P44062	Y857_HAEIN		cytoplasm
P44063	Y866_HAEIN		inner membrane
P44064	Y869_HAEIN		cytoplasm
P44066	Y871_HAEIN		cytoplasm
P44067	Y874_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44068	Y882_HAEIN		cytoplasm
P44069	Y886_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44070	Y902_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44072	Y907_HAEIN		cytoplasm
P44073	Y908_HAEIN		extracellular
P44074	Y912_HAEIN		cytoplasm
P44075	Y925_HAEIN		cytoplasm
P44076	Y926_HAEIN		cytoplasm
P44077	Y930_HAEIN		extracellular
P44078	Y931_HAEIN		cytoplasm
P44079	Y938_HAEIN		inner membrane
P44080	Y939_HAEIN		inner membrane
P44081	Y940_HAEIN		inner membrane
P44082	Y941_HAEIN		inner membrane
P44083	RUMB_HAEIN		cytoplasm
P44084	Y960_HAEIN		cytoplasm
P44085	Y966_HAEIN		cytoplasm
P44087	Y974_HAEIN		inner membrane
P44088	Y977_HAEIN		cytoplasm
P44089	Y996_HAEIN		inner membrane
P44090	Y997_HAEIN		outer membrane
P44092	PPID_HAEIN	bacterial cell membrane; single-pass type ii membrane protein (potential).	periplasm
P44093	Y1011_HAEIN		cytoplasm
P44094	Y1014_HAEIN		cytoplasm
P44095	Y1016_HAEIN		inner membrane
P44096	Y1034_HAEIN		inner membrane
P44097	Y1036_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44098	Y1037_HAEIN		cytoplasm
P44099	Y1038_HAEIN		cytoplasm
P44101	Y1043_HAEIN		periplasm
P44102	Y1044_HAEIN		cytoplasm
P44103	Y1048_HAEIN		periplasm
P44104	Y1054_HAEIN		cytoplasm
P44105	T3RH_HAEIN		extracellular
P44106	Y1058_HAEIN		cytoplasm
P44107	Y1063_HAEIN		cytoplasm
P44110	Y1074_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44112	Y1099_HAEIN		inner membrane

P44113	ZIPA_HAEIN	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
P44114	Y1126_HAEIN		inner membrane
P44115	Y1128_HAEIN		inner membrane
P44116	Y1162_HAEIN		cytoplasm
P44117	Y1168_HAEIN		inner membrane
P44118	Y1169_HAEIN		cytoplasm
P44119	SPRT_HAEIN	cytoplasm (potential).	extracellular
P44120	Y1176_HAEIN		extracellular
P44121	DNLI_HAEIN		cytoplasm
P44123	PTPS_HAEIN		extracellular
P44124	Y1191_HAEIN		inner membrane
P44126	Y1202_HAEIN		inner membrane
P44127	Y1205_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44128	Y1215_HAEIN		cytoplasm
P44129	Y1222_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44130	Y1223_HAEIN		cytoplasm
P44131	Y1235_HAEIN		cytoplasm
P44132	Y1236_HAEIN		inner membrane
P44133	Y1241_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44134	Y1244_HAEIN		cytoplasm
P44135	Y1246_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44136	Y1248_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44137	Y1249_HAEIN		inner membrane
P44138	Y1250_HAEIN		cytoplasm
P44139	Y1253_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44140	Y1254_HAEIN		cytoplasm
P44141	Y1255_HAEIN		cytoplasm
P44142	Y1256_HAEIN		cytoplasm
P44143	Y1257_HAEIN		cytoplasm
P44144	Y1265_HAEIN		cytoplasm
P44145	Y1266_HAEIN		inner membrane
P44150	Y1273_HAEIN		cytoplasm
P44151	Y1280_HAEIN		cytoplasm
P44152	T1SH_HAEIN		cytoplasm
P44153	QUEF_HAEIN	cytoplasm (probable).	cytoplasm
P44154	Y1292_HAEIN		cytoplasm
P44156	Y1293_HAEIN		cytoplasm
P44157	PGPA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44158	Y1310_HAEIN		cytoplasm
P44159	Y1316_HAEIN		cytoplasm

P44160	Y1317_HAEIN		cytoplasm
P44161	Y1323_HAEIN		inner membrane
P44162	Y1326_HAEIN		cytoplasm
P44164	SIXA_HAEIN		extracellular
P44165	Y1340_HAEIN		inner membrane
P44166	Y1341_HAEIN		cytoplasm
P44167	Y1351_HAEIN		cytoplasm
P44168	Y1355_HAEIN		inner membrane
P44169	Y1375_HAEIN		cytoplasm
P44170	Y1376_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44171	Y1386_HAEIN		cytoplasm
P44172	Y1394_HAEIN		inner membrane
P44173	Y1395_HAEIN		cytoplasm
P44174	Y1396_HAEIN		cytoplasm
P44175	Y1399_HAEIN		cytoplasm
P44176	TRPH_HAEIN		cytoplasm
P44177	Y1402_HAEIN		cytoplasm
P44178	Y1403_HAEIN		cytoplasm
P44180	Y1405_HAEIN		cytoplasm
P44181	Y1406_HAEIN		cytoplasm
P44183	Y1409_HAEIN		cytoplasm
P44184	Y1410_HAEIN		cytoplasm
P44185	Y1413_HAEIN		inner membrane
P44186	Y1414_HAEIN		inner membrane
P44187	Y1415_HAEIN		extracellular
P44188	Y1416_HAEIN		inner membrane
P44189	Y1418_HAEIN		cytoplasm
P44190	Y1419_HAEIN		cytoplasm
P44191	Y1420_HAEIN		cytoplasm
P44192	Y1421_HAEIN		cytoplasm
P44193	Y1422_HAEIN		cytoplasm
P44194	Y1423_HAEIN		cytoplasm
P44195	USPE_HAEIN	cytoplasm (by similarity).	cytoplasm
P44196	Y1427_HAEIN		cytoplasm
P44197	TRUC_HAEIN		cytoplasm
P44198	Y1446_HAEIN		cytoplasm
P44199	Y1450_HAEIN		inner membrane
P44201	Y1452_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44202	Y1454_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44203	Y1456_HAEIN		inner membrane
P44204	Y1458_HAEIN		cytoplasm
P44205	Y1469_HAEIN		cytoplasm
P44206	Y1472_HAEIN		periplasm
P44207	Y1476_HAEIN		cytoplasm
P44208	Y1479_HAEIN		cytoplasm
P44209	Y1480_HAEIN		cytoplasm
P44210	Y1482_HAEIN		cytoplasm
P44211	Y1484_HAEIN		cytoplasm
P44212	Y1485_HAEIN		cytoplasm
P44213	Y1486_HAEIN		cytoplasm

P44214	Y1487_HAEIN		cytoplasm
P44215	Y1489_HAEIN		cytoplasm
P44216	Y1490_HAEIN		cytoplasm
P44217	Y1492_HAEIN		outer membrane
P44218	Y1493_HAEIN		cytoplasm
P44219	Y1495_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44220	Y1496_HAEIN		inner membrane
P44221	Y1497_HAEIN		cytoplasm
P44222	Y1498_HAEIN		inner membrane
P44223	VG27_HAEIN		cytoplasm
P44224	VG28_HAEIN		cytoplasm
P44225	VG29_HAEIN		cytoplasm
P44226	VPF_HAEIN		cytoplasm
P44227	VPT_HAEIN		cytoplasm
P44228	VG35_HAEIN		cytoplasm
P44230	VG36_HAEIN		cytoplasm
P44231	VG37_HAEIN		cytoplasm
P44232	VG38_HAEIN		cytoplasm
P44233	VPL_HAEIN		cytoplasm
P44234	VPM_HAEIN		cytoplasm
P44235	VG41_HAEIN		cytoplasm
P44236	VG42_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44238	VG45_HAEIN		cytoplasm
P44239	VG46_HAEIN		inner membrane
P44240	VG47_HAEIN		cytoplasm
P44241	VG48_HAEIN		cytoplasm
P44242	VPS_HAEIN		extracellular
P44243	Y1523_HAEIN		cytoplasm
P44244	Y1524_HAEIN		cytoplasm
P44246	Y1534_HAEIN		cytoplasm
P44247	Y1536_HAEIN		cytoplasm
P44248	Y1543_HAEIN		cytoplasm
P44249	Y1546_HAEIN		cytoplasm
P44250	LOLE_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44251	Y1552_HAEIN		cytoplasm
P44252	LOLC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44253	Y1560_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44254	Y1562_HAEIN		cytoplasm
P44255	Y1563_HAEIN		cytoplasm
P44256	Y1564_HAEIN		cytoplasm
P44258	Y1569_HAEIN		cytoplasm
P44259	Y1570_HAEIN		cytoplasm
P44260	Y1571_HAEIN		cytoplasm
P44262	Y1581_HAEIN		cytoplasm

P44263	Y1586_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44264	Y1593_HAEIN		cytoplasm
P44265	Y1594_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44266	Y1595_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44267	Y1599_HAEIN		cytoplasm
P44268	Y1600_HAEIN		inner membrane
P44269	Y1601_HAEIN		periplasm
P44270	Y1602_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44271	Y1603_HAEIN		cytoplasm
P44272	Y1605_HAEIN		inner membrane
P44273	Y1620_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44274	Y1621_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44275	Y1622_HAEIN		inner membrane
P44276	Y1624_HAEIN		fimbrium
P44277	Y1625_HAEIN		inner membrane
P44278	Y1626_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44279	Y1631_HAEIN		cytoplasm
P44280	Y1637_HAEIN		cytoplasm
P44282	Y1651_HAEIN		cytoplasm
P44284	Y1666_HAEIN		inner membrane
P44285	Y1667_HAEIN		inner membrane
P44286	PROQ_HAEIN	cytoplasm (potential).	inner membrane
P44287	Y1671_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44288	Y1672_HAEIN		inner membrane
P44289	Y1680_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44290	Y1681_HAEIN		inner membrane
P44291	RNFG_HAEIN	bacterial cell inner membrane (potential).	inner membrane
P44292	Y1701_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44293	Y1709_HAEIN	periplasm (potential).	periplasm
P44296	Y1718_HAEIN		flagellar
P44297	Y1724_HAEIN		inner membrane
P44298	Y1730_HAEIN		cytoplasm
P44299	Y1731_HAEIN		cytoplasm

P44300	Y1736_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44301	Y1737_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44302	Y1738_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44303	AROG_HAEIN		cytoplasm
P44305	RIMI_HAEIN	cytoplasm (by similarity).	cytoplasm
P44306	RNPA_HAEIN		extracellular
P44307	RRF_HAEIN	cytoplasm (by similarity).	cytoplasm
P44308	NADR_HAEIN		cytoplasm
P44309	KITH_HAEIN	cytoplasm (potential).	cytoplasm
P44310	KGUA_HAEIN	cytoplasm (by similarity).	cytoplasm
P44311	G6PD_HAEIN		cytoplasm
P44312	G6PI_HAEIN	cytoplasm (by similarity).	cytoplasm
P44313	FOLD_HAEIN		cytoplasm
P44314	ARLY_HAEIN	cytoplasm (probable).	cytoplasm
P44315	ASSY_HAEIN	cytoplasm (probable).	cytoplasm
P44316	DCDA_HAEIN		cytoplasm
P44317	DCOR_HAEIN		cytoplasm
P44318	EX3_HAEIN		cytoplasm
P44319	END3_HAEIN		cytoplasm
P44320	MUTY_HAEIN		cytoplasm
P44321	3MGA_HAEIN		cytoplasm
P44324	ASPA_HAEIN		cytoplasm
P44325	CDD_HAEIN		cytoplasm
P44326	RIBD_HAEIN		cytoplasm
P44327	HIS7_HAEIN	cytoplasm (by similarity).	cytoplasm
P44328	KPRS_HAEIN	cytoplasm (by similarity).	cytoplasm
P44329	RBSR_HAEIN		cytoplasm
P44330	K1PF_HAEIN		cytoplasm
P44331	RBSK_HAEIN	cytoplasm (by similarity).	cytoplasm
P44332	CYSQ_HAEIN	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	cytoplasm
P44333	SUHB_HAEIN		cytoplasm
P44334	IMDH_HAEIN		cytoplasm
P44335	GUAA_HAEIN		cytoplasm
P44336	SERC_HAEIN	cytoplasm (by similarity).	cytoplasm
P44337	ASNC_HAEIN		cytoplasm
P44338	ASNA_HAEIN	cytoplasm (by similarity).	cytoplasm
P44339	Y1171_HAEIN		cytoplasm
P44340	HIS5_HAEIN	cytoplasm (by similarity).	cytoplasm
P44341	PYRG_HAEIN		cytoplasm
P44342	RL1_HAEIN		cytoplasm
P44343	RL2_HAEIN		cytoplasm
P44344	RL3_HAEIN		cytoplasm
P44345	RL4_HAEIN		cytoplasm
P44346	RL5_HAEIN		cytoplasm
P44347	RL6_HAEIN		cytoplasm
P44348	RL7_HAEIN		cytoplasm
P44349	RL9_HAEIN		cytoplasm

P44350	RL10_HAEIN		cytoplasm
P44351	RL11_HAEIN		cytoplasm
P44353	RL15_HAEIN		cytoplasm
P44354	RL16_HAEIN		cytoplasm
P44355	RL17_HAEIN		cytoplasm
P44356	RL18_HAEIN		cytoplasm
P44357	RL19_HAEIN		cytoplasm
P44358	RL20_HAEIN		cytoplasm
P44359	RL21_HAEIN		cytoplasm
P44360	RL22_HAEIN		cytoplasm
P44361	RL23_HAEIN		cytoplasm
P44362	RL24_HAEIN		cytoplasm
P44363	RL27_HAEIN		cytoplasm
P44364	RL28_HAEIN		cytoplasm
P44365	RL29_HAEIN		cytoplasm
P44366	RL30_HAEIN		cytoplasm
P44367	RL31_HAEIN		cytoplasm
P44368	RL32_HAEIN		cytoplasm
P44369	RL33_HAEIN		cytoplasm
P44371	RS2_HAEIN		cytoplasm
P44372	RS3_HAEIN		cytoplasm
P44373	RS4_HAEIN		cytoplasm
P44374	RS5_HAEIN		cytoplasm
P44375	RS6_HAEIN		cytoplasm
P44376	RS7_HAEIN		cytoplasm
P44377	RS8_HAEIN		cytoplasm
P44379	RS11_HAEIN		cytoplasm
P44380	RS13_HAEIN		cytoplasm
P44381	RS14_HAEIN		cytoplasm
P44382	RS16_HAEIN		cytoplasm
P44383	RS17_HAEIN		cytoplasm
P44386	RS21_HAEIN		cytoplasm
P44387	RL13_HAEIN		cytoplasm
P44388	RS9_HAEIN		cytoplasm
P44389	RS15_HAEIN		cytoplasm
P44390	CATA_HAEIN	cytoplasm (probable).	periplasm
P44391	URE1_HAEIN	cytoplasm (by similarity).	cytoplasm
P44392	URE2_HAEIN	cytoplasm (by similarity).	cytoplasm
P44393	URE3_HAEIN	cytoplasm (by similarity).	cytoplasm
P44394	UREE_HAEIN	cytoplasm (by similarity).	cytoplasm
P44395	UREF_HAEIN		cytoplasm
P44396	UREG_HAEIN		cytoplasm
P44397	UREH_HAEIN		cytoplasm
P44399	FUCK_HAEIN		cytoplasm
P44400	GLPK_HAEIN		cytoplasm
P44401	XYLB_HAEIN		cytoplasm
P44402	PRMA_HAEIN	cytoplasm (by similarity).	cytoplasm
P44403	CLPB_HAEIN	cytoplasm (probable).	cytoplasm
P44404	RP32_HAEIN		cytoplasm
P44405	DDL_HAEIN	cytoplasm (by similarity).	cytoplasm
P44407	MSBA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44408	REC2_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane

P44409	SSB_HAEIN		cytoplasm
P44410	UVRA_HAEIN	cytoplasm (by similarity).	cytoplasm
P44411	BCP_HAEIN		cytoplasm
P44413	T2D2_HAEIN		extracellular
P44414	MTHD_HAEIN		cytoplasm
P44416	OAPB_HAEIN		cytoplasm
P44417	DEOD_HAEIN		cytoplasm
P44418	OXYR_HAEIN		cytoplasm
P44419	GLNE_HAEIN		cytoplasm
P44421	AMPM_HAEIN		cytoplasm
P44422	BIOF_HAEIN		cytoplasm
P44423	HIS81_HAEIN		cytoplasm
P44424	KDGL_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44426	BIOA_HAEIN		cytoplasm
P44427	MDH_HAEIN		cytoplasm
P44428	FER_HAEIN		cytoplasm
P44429	ALF_HAEIN		cytoplasm
P44430	DEOC_HAEIN	cytoplasm (by similarity).	cytoplasm
P44431	DMA_HAEIN		cytoplasm
P44432	FABI_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P44433	RLUC_HAEIN		cytoplasm
P44434	HIS2_HAEIN	cytoplasm (by similarity).	cytoplasm
P44435	HIS4_HAEIN	cytoplasm (by similarity).	cytoplasm
P44436	HIS6_HAEIN	cytoplasm (by similarity).	cytoplasm
P44437	HFQ_HAEIN		cytoplasm
P44438	LEUD_HAEIN		cytoplasm
P44439	PCNB_HAEIN		cytoplasm
P44440	RNB_HAEIN	cytoplasm (by similarity).	cytoplasm
P44441	RNC_HAEIN	cytoplasm (by similarity).	cytoplasm
P44444	RNPH_HAEIN		cytoplasm
P44445	RLUD_HAEIN		cytoplasm
P44446	LCFH_HAEIN		cytoplasm
P44447	Y003_HAEIN		inner membrane
P44449	FDHD_HAEIN	cytoplasm (probable).	cytoplasm
P44451	FDXI_HAEIN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P44452	FDHE_HAEIN	cytoplasm (by similarity).	cytoplasm
P44453	RSMC_HAEIN		cytoplasm
P44454	LEP_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44455	GRCA_HAEIN		cytoplasm
P44458	CITXG_HAEIN		cytoplasm
P44461	CITD_HAEIN	cytoplasm (by similarity).	cytoplasm
P44462	CITC_HAEIN		cytoplasm
P44463	LIPA_HAEIN	cytoplasm (by similarity).	cytoplasm
P44464	LIPB_HAEIN	cytoplasm (by similarity).	cytoplasm
P44465	Y028_HAEIN		inner membrane
P44468	RODA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P44469	PBP2_HAEIN	bacterial cell inner membrane (by similarity).	inner membrane
P44470	Y033_HAEIN		inner membrane
P44471	Y034_HAEIN		inner membrane
P44472	Y035_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44474	MREB_HAEIN		cytoplasm
P44475	MREC_HAEIN	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P44476	MRED_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44477	Y044_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44478	Y045_HAEIN	periplasm (potential).	periplasm
P44479	PHNA_HAEIN		cytoplasm
P44480	ALKH_HAEIN	cytoplasm (by similarity).	cytoplasm
P44481	Y048_HAEIN		cytoplasm
P44482	KDGK_HAEIN		cytoplasm
P44483	Y050_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44484	Y051_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44487	UXUR_HAEIN		cytoplasm
P44488	UXUA_HAEIN		cytoplasm
P44489	UVRC_HAEIN	cytoplasm (by similarity).	cytoplasm
P44490	KDSB_HAEIN	cytoplasm (potential).	cytoplasm
P44491	LPXK_HAEIN		cytoplasm
P44492	Y065_HAEIN		cytoplasm
P44493	AMIB_HAEIN		extracellular
P44494	MUTL_HAEIN		cytoplasm
P44495	MIAA_HAEIN		cytoplasm
P44496	RECN_HAEIN		cytoplasm
P44497	PPNK_HAEIN	cytoplasm (by similarity).	cytoplasm
P44498	TESB_HAEIN		extracellular
P44499	PPIB_HAEIN	cytoplasm (by similarity).	cytoplasm
P44500	Y081_HAEIN		extracellular
P44501	DDH_HAEIN		cytoplasm
P44503	THRC_HAEIN		cytoplasm
P44504	KHSE_HAEIN	cytoplasm (potential).	cytoplasm
P44505	AKH_HAEIN		cytoplasm
P44506	Y090_HAEIN		cytoplasm
P44507	GLXK_HAEIN		cytoplasm
P44509	Y093_HAEIN		cytoplasm
P44513	FBPC2_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P44514	DAPE_HAEIN		cytoplasm
P44515	Y103_HAEIN		cytoplasm
P44516	HTPG_HAEIN	cytoplasm (by similarity).	cytoplasm
P44518	SRP54_HAEIN		cytoplasm

P44520	Y108_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44523	TDHA_HAEIN	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P44524	Y116_HAEIN		cytoplasm
P44526	ZNUA_HAEIN	periplasm (by similarity).	periplasm
P44528	PGSA_HAEIN	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P44529	IPYR_HAEIN	cytoplasm (by similarity).	cytoplasm
P44530	Y125_HAEIN		inner membrane
P44531	FBPC1_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P44533	URK_HAEIN	cytoplasm (by similarity).	cytoplasm
P44534	DCD_HAEIN		extracellular
P44535	SOTB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44536	ENGA_HAEIN		cytoplasm
P44537	NAGA_HAEIN		cytoplasm
P44538	NAGB_HAEIN		extracellular
P44540	Y143_HAEIN		cytoplasm
P44541	NANK_HAEIN		cytoplasm
P44542	SIAP_HAEIN	periplasm (probable).	periplasm
P44543	SIAT1_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44544	Y148_HAEIN		outer membrane
P44545	HFLC_HAEIN		extracellular
P44546	HFLK_HAEIN		inner membrane
P44550	APBE_HAEIN	bacterial cell membrane; lipid-anchor (potential).	cytoplasm
P44551	TRMU_HAEIN	cytoplasm (by similarity).	cytoplasm
P44552	Y175_HAEIN		cytoplasm
P44553	Y177_HAEIN	bacterial cell membrane; lipid-anchor (by similarity).	outer membrane
P44554	Y182_HAEIN		cytoplasm
P44555	Y183_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44556	Y184_HAEIN		cytoplasm
P44557	ADH3_HAEIN	cytoplasm (potential).	cytoplasm
P44558	Y186_HAEIN		cytoplasm
P44560	TATC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44561	FUR_HAEIN	cytoplasm (by similarity).	cytoplasm
P44562	FLAV_HAEIN		cytoplasm
P44564	SEQA_HAEIN		cytoplasm
P44565	MENE_HAEIN		cytoplasm
P44566	MEPA_HAEIN	periplasm (potential).	extracellular

P44567	MSBB_HAEIN	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P44568	RIMM_HAEIN	cytoplasm (potential).	cytoplasm
P44569	5NTD_HAEIN		periplasm
P44571	GCH2_HAEIN		extracellular
P44572	Y213_HAEIN	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P44573	OPDA_HAEIN		extracellular
P44577	Y219_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44578	ARCB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44579	Y223_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44580	Y224_HAEIN		cytoplasm
P44581	NHAA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44583	Y227_HAEIN		inner membrane
P44585	NLPI_HAEIN	bacterial cell membrane; lipid-anchor (by similarity).	inner membrane
P44586	DEAD_HAEIN	cytoplasm (probable).	cytoplasm
P44587	Y232_HAEIN		periplasm
P44588	Y235_HAEIN		cytoplasm
P44589	ARSC_HAEIN		cytoplasm
P44590	SECF_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44591	SECD_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44592	Y241_HAEIN		inner membrane
P44593	Y242_HAEIN		cytoplasm
P44594	TGT_HAEIN		cytoplasm
P44595	QUEA_HAEIN	cytoplasm (potential).	cytoplasm
P44596	HAP1_HAEIN	secreted protein (potential).	extracellular
P44598	HAM1_HAEIN		extracellular
P44600	HXUC1_HAEIN	bacterial cell outer membrane (by similarity).	outer membrane
P44603	Y266_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44604	NARQ_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44605	MURB_HAEIN	cytoplasm (probable).	inner membrane
P44606	DUSC_HAEIN		cytoplasm
P44608	RBN_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44609	Y277_HAEIN		inner membrane

P44610	Y281_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44611	Y282_HAEIN		cytoplasm
P44612	MEND_HAEIN		cytoplasm
P44613	MENF_HAEIN		cytoplasm
P44614	MTR_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44615	SDAC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44617	Y293_HAEIN	cytoplasm (probable).	cytoplasm
P44618	METJ_HAEIN	cytoplasm (by similarity).	cytoplasm
P44619	RHO_HAEIN		inner membrane
P44620	LEP4_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44621	HOFC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44622	HOFB_HAEIN		inner membrane
P44623	PPDD_HAEIN		extracellular
P44624	AMPD_HAEIN	cytoplasm (by similarity).	extracellular
P44626	LNT_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44627	Y303_HAEIN		cytoplasm
P44628	RDGC_HAEIN	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
P44629	HCAT_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44630	XERD_HAEIN	cytoplasm (by similarity).	nucleoid
P44631	RUVB_HAEIN		cytoplasm
P44632	RUVA_HAEIN		cytoplasm
P44633	RUVC_HAEIN		cytoplasm
P44634	Y315_HAEIN		inner membrane
P44635	NUDB_HAEIN		cytoplasm
P44638	LGUL_HAEIN		extracellular
P44639	RNT_HAEIN		cytoplasm
P44640	Y325_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44641	Y329_HAEIN		cytoplasm
P44642	RECO_HAEIN		cytoplasm
P44643	RUMA_HAEIN		cytoplasm
P44644	RELA_HAEIN		cytoplasm
P44645	MOG_HAEIN		cytoplasm
P44646	Y338_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44647	PRIA_HAEIN		cytoplasm
P44648	TRMB_HAEIN		cytoplasm
P44649	Y341_HAEIN		cytoplasm

P44650	NAPF_HAEIN		periplasm
P44651	NAPD_HAEIN	cytoplasm (potential).	cytoplasm
P44652	NAPG_HAEIN		periplasm
P44653	NAPH_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	periplasm
P44654	NAPB_HAEIN	periplasm (potential).	periplasm
P44656	Y354_HAEIN		inner membrane
P44658	Y357_HAEIN		cytoplasm
P44659	Y358_HAEIN		cytoplasm
P44660	Y359_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44661	Y360_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44662	Y361_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P44664	PBP7_HAEIN	periplasm (potential).	cytoplasm
P44665	Y365_HAEIN		cytoplasm
P44667	ISPG_HAEIN		cytoplasm
P44668	ISCX_HAEIN		cytoplasm
P44669	HSCA_HAEIN		cytoplasm
P44670	Y374_HAEIN		cytoplasm
P44672	ISCA_HAEIN		cytoplasm
P44675	Y379_HAEIN		cytoplasm
P44676	Y380_HAEIN		cytoplasm
P44677	TOLB_HAEIN	periplasm (potential).	periplasm
P44678	TOLA_HAEIN	bacterial cell inner membrane; single-pass type ii membrane protein (potential).	inner membrane
P44679	Y386_HAEIN		cytoplasm
P44680	Y387_HAEIN		cytoplasm
P44681	ENGD_HAEIN		cytoplasm
P44682	PTH_HAEIN	cytoplasm (by similarity).	cytoplasm
P44683	Y396_HAEIN		cytoplasm
P44684	ADPP_HAEIN		cytoplasm
P44685	ICC_HAEIN		extracellular
P44686	Y400_HAEIN		cytoplasm
P44687	OGT_HAEIN		cytoplasm
P44688	MUTH_HAEIN	cytoplasm (by similarity).	extracellular
P44689	TILS_HAEIN	cytoplasm (probable).	cytoplasm
P44690	PDXY_HAEIN		cytoplasm
P44691	ZNUB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44692	ZNUC_HAEIN		inner membrane
P44693	Y409_HAEIN		extracellular
P44697	THID_HAEIN		cytoplasm
P44699	Y418_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44700	Y419_HAEIN		cytoplasm
P44701	SRMB_HAEIN		cytoplasm
P44702	Y423_HAEIN		cytoplasm

P44703	Y424_HAEIN		cytoplasm
P44705	FADR_HAEIN	cytoplasm (potential).	cytoplasm
P44706	NHAB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44707	DSBB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44708	GLMS_HAEIN	cytoplasm (by similarity).	cytoplasm
P44709	Y431_HAEIN		cytoplasm
P44710	NUDC_HAEIN		cytoplasm
P44711	Y442_HAEIN		inner membrane
P44712	RECR_HAEIN		cytoplasm
P44713	SECG_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44714	PTFBC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44715	PTFAH_HAEIN	cytoplasm (by similarity).	cytoplasm
P44717	Y452_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44718	Y454_HAEIN		extracellular
P44719	KTHY_HAEIN		cytoplasm
P44720	Y457_HAEIN		cytoplasm
P44721	SURA_HAEIN	periplasm (potential).	periplasm
P44722	PYRR_HAEIN		cytoplasm
P44723	MAZG_HAEIN		cytoplasm
P44725	RPIA_HAEIN		cytoplasm
P44726	Y467_HAEIN		nucleoid
P44727	TYRPA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44728	GIDB_HAEIN		cytoplasm
P44731	POTD2_HAEIN	periplasm (potential).	periplasm
P44732	ORDL_HAEIN		cytoplasm
P44733	RMUC_HAEIN		cytoplasm
P44734	RBSD_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P44735	RBSA_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P44736	RBSC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44737	RBSB_HAEIN	periplasm (by similarity).	periplasm
P44738	RRAA_HAEIN		cytoplasm
P44739	MENA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44740	Y510_HAEIN		inner membrane
P44741	TEHA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P44742	Y519_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44743	Y520_HAEIN		cytoplasm
P44744	Y521_HAEIN		cytoplasm
P44746	Y527_HAEIN		periplasm
P44747	TYRPB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44748	PRIB_HAEIN		cytoplasm
P44749	KSGA_HAEIN		cytoplasm
P44751	APAH_HAEIN		extracellular
P44754	HSLR_HAEIN		cytoplasm
P44755	GPH_HAEIN		cytoplasm
P44756	RPE_HAEIN		cytoplasm
P44757	Y570_HAEIN		cytoplasm
P44758	PRX5_HAEIN		cytoplasm
P44759	SLYX_HAEIN		inner membrane
P44760	FKBY_HAEIN		cytoplasm
P44761	Y575_HAEIN		cytoplasm
P44762	TUSD_HAEIN	cytoplasm (by similarity).	cytoplasm
P44763	GIDA_HAEIN	cytoplasm (by similarity).	cytoplasm
P44764	CN16_HAEIN	periplasm (by similarity).	periplasm
P44765	ABGA_HAEIN		extracellular
P44766	PEPE_HAEIN	cytoplasm (by similarity).	cytoplasm
P44768	POTE_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44769	ARCC_HAEIN	cytoplasm (potential).	cytoplasm
P44770	OTCC_HAEIN	cytoplasm (probable).	cytoplasm
P44771	Y597_HAEIN		cytoplasm
P44772	HEMY_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44773	Y603_HAEIN		inner membrane
P44774	Y607_HAEIN		cytoplasm
P44776	FUCP_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44777	FUCA_HAEIN		extracellular
P44778	FUCU_HAEIN		cytoplasm
P44779	FUCI_HAEIN	cytoplasm (by similarity).	cytoplasm
P44780	FUCR_HAEIN		cytoplasm
P44781	RAPA_HAEIN		cytoplasm
P44782	RLUA_HAEIN		cytoplasm
P44783	GLPG_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44784	GLPR_HAEIN		cytoplasm
P44785	METN_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P44786	DEF_HAEIN		cytoplasm
P44787	FMT_HAEIN		cytoplasm
P44788	RSMB_HAEIN		cytoplasm

P44789	MSCL_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44790	RPOE_HAEIN		cytoplasm
P44791	RSEA_HAEIN		inner membrane
P44792	RSEB_HAEIN	periplasm (potential).	periplasm
P44793	COAA_HAEIN	cytoplasm (probable).	cytoplasm
P44794	DUSA_HAEIN		cytoplasm
P44795	HGP1_HAEIN	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P44796	Y638_HAEIN		inner membrane
P44797	PUR8_HAEIN		cytoplasm
P44798	TORZ_HAEIN	periplasm (by similarity).	periplasm
P44799	TORY_HAEIN	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P44800	PLDB_HAEIN	bacterial cell inner membrane (by similarity).	inner membrane
P44801	DHAS_HAEIN		cytoplasm
P44803	MDAB_HAEIN		cytoplasm
P44804	REP_HAEIN		cytoplasm
P44805	COAD_HAEIN	cytoplasm (by similarity).	cytoplasm
P44806	KDTA_HAEIN	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P44807	Y656_HAEIN		inner membrane
P44808	Y658_HAEIN		inner membrane
P44809	HGP2_HAEIN	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P44811	GLPX_HAEIN	cytoplasm (by similarity).	cytoplasm
P44812	Y668_HAEIN		cytoplasm
P44813	MIOC_HAEIN		cytoplasm
P44814	DTD_HAEIN	cytoplasm (probable).	cytoplasm
P44815	ISPF_HAEIN		extracellular
P44817	PEPD_HAEIN		cytoplasm
P44819	GLPE_HAEIN	cytoplasm (by similarity).	cytoplasm
P44821	ILVY_HAEIN	cytoplasm (by similarity).	cytoplasm
P44822	ILVC_HAEIN		cytoplasm
P44826	GLPF_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44827	RLUE_HAEIN		cytoplasm
P44828	PPX_HAEIN	bacterial cell membrane; peripheral membrane protein (potential).	cytoplasm
P44830	SLYD_HAEIN	cytoplasm (potential).	cytoplasm
P44831	Y700_HAEIN		cytoplasm
P44832	Y703_HAEIN		inner membrane
P44834	MUTS_HAEIN		cytoplasm
P44836	HGP3_HAEIN	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P44837	TIG_HAEIN		cytoplasm
P44838	CLPX_HAEIN		cytoplasm

P44839	Y719_HAEIN		cytoplasm
P44840	HTPX_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P44841	TUSA_HAEIN	cytoplasm (by similarity).	cytoplasm
P44842	Y722_HAEIN		inner membrane
P44843	TRKH_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44844	Y724_HAEIN		inner membrane
P44845	NARP_HAEIN		cytoplasm
P44846	OSTA_HAEIN	bacterial cell outer membrane (by similarity).	outer membrane
P44847	SUFI_HAEIN	periplasm (potential).	periplasm
P44848	PLSC_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P44851	ILVD_HAEIN		cytoplasm
P44854	Y744_HAEIN		cytoplasm
P44855	DCUB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44856	DHNA_HAEIN	bacterial membrane (by similarity).	inner membrane
P44857	PLSB_HAEIN	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P44858	LEXA_HAEIN		cytoplasm
P44861	Y753_HAEIN		inner membrane
P44862	Y754_HAEIN		cytoplasm
P44863	Y755_HAEIN		cytoplasm
P44864	Y756_HAEIN		extracellular
P44865	GPMA_HAEIN		cytoplasm
P44866	RIBB_HAEIN		cytoplasm
P44868	Y766_HAEIN		cytoplasm
P44869	Y767_HAEIN		cytoplasm
P44870	FTSY_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P44871	FTSE_HAEIN		cytoplasm
P44872	FTSX_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44873	ATOB_HAEIN	cytoplasm (potential).	cytoplasm
P44874	ATOA_HAEIN		cytoplasm
P44875	ATOD_HAEIN		cytoplasm
P44876	Y775_HAEIN		cytoplasm
P44877	HSLO_HAEIN	cytoplasm (by similarity).	cytoplasm
P44878	GALU_HAEIN		cytoplasm
P44879	CSRA_HAEIN		cytoplasm
P44880	USPA_HAEIN	cytoplasm (by similarity).	cytoplasm
P44881	AMPP_HAEIN		cytoplasm
P44882	Y817_HAEIN		inner membrane
P44883	DGAL_HAEIN	periplasm (by similarity).	periplasm

P44884	MGLA_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P44885	MGLC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44886	Y827_HAEIN		cytoplasm
P44887	Y828_HAEIN		cytoplasm
P44888	SLT_HAEIN	periplasm (potential).	periplasm
P44889	TRPR_HAEIN	cytoplasm (by similarity).	cytoplasm
P44890	MTGA_HAEIN	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P44891	FRDD_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44892	FRDC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44893	FRDB_HAEIN		cytoplasm
P44894	FRDA_HAEIN		periplasm
P44895	CPXR_HAEIN	cytoplasm (probable).	cytoplasm
P44896	NDPA_HAEIN	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
P44897	Y840_HAEIN		inner membrane
P44898	Y841_HAEIN		inner membrane
P44899	MOBA_HAEIN	cytoplasm (by similarity).	cytoplasm
P44900	Y845_HAEIN		cytoplasm
P44901	Y849_HAEIN		inner membrane
P44902	MOBB_HAEIN		cytoplasm
P44903	Y852_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44904	Y855_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44905	Y858_HAEIN		cytoplasm
P44906	RLMB_HAEIN		cytoplasm
P44907	RNR_HAEIN		cytoplasm
P44908	Y862_HAEIN		inner membrane
P44909	PDXH_HAEIN		cytoplasm
P44910	TYP A_HAEIN		cytoplasm
P44914	RFFG_HAEIN		cytoplasm
P44915	Y877_HAEIN		cytoplasm
P44916	ISPB_HAEIN		cytoplasm
P44917	Y883_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44918	ARCA_HAEIN	cytoplasm (probable).	cytoplasm
P44919	DSBD_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44920	COAE_HAEIN	cytoplasm (by similarity).	cytoplasm
P44921	Y891_HAEIN		cytoplasm
P44922	RHLB_HAEIN		cytoplasm

P44923	Y893_HAEIN		cytoplasm
P44927	EMRB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P44928	EMRA_HAEIN	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P44930	LGT_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44931	TADA_HAEIN		cytoplasm
P44932	MUTT_HAEIN		cytoplasm
P44933	KEFX_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44936	Y918_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P44937	CDSA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44938	UPPS_HAEIN		inner membrane
P44940	Y929_HAEIN		cytoplasm
P44941	Y933_HAEIN		cytoplasm
P44942	NRFF_HAEIN	periplasm (probable).	periplasm
P44943	NRFX_HAEIN	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (potential).	periplasm
P44944	NRFE_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44945	EX5C_HAEIN		cytoplasm
P44946	NRDR_HAEIN		cytoplasm
P44947	DEGS_HAEIN	periplasm (potential).	periplasm
P44948	FPG_HAEIN		cytoplasm
P44951	DAT_HAEIN		cytoplasm
P44952	RADC_HAEIN		cytoplasm
P44953	COABC_HAEIN		cytoplasm
P44954	Y956_HAEIN		inner membrane
P44955	NAGZ_HAEIN	cytoplasm (by similarity).	extracellular
P44956	Y961_HAEIN		cytoplasm
P44957	RIBF_HAEIN		cytoplasm
P44958	MVIN_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44959	RS20_HAEIN		cytoplasm
P44960	MENB_HAEIN		cytoplasm
P44961	MENC_HAEIN		extracellular
P44963	PANF_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44965	DUSB_HAEIN		cytoplasm
P44966	FIS_HAEIN		cytoplasm
P44967	SSRP_HAEIN	cytoplasm (potential).	cytoplasm
P44968	LEU2_HAEIN		cytoplasm

P44970	TBP1_HAEIN	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P44971	TBP2_HAEIN	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P44972	Y1000_HAEIN		inner membrane
P44973	OXAA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44974	Y1005_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P44975	LSPA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P44976	ISPH_HAEIN		cytoplasm
P44978	Y1009_HAEIN		cytoplasm
P44979	Y1010_HAEIN		cytoplasm
P44985	THIP_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44986	THIQ_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P44987	BIOB_HAEIN		cytoplasm
P44988	SGBH_HAEIN		cytoplasm
P44989	SGBE_HAEIN		cytoplasm
P44990	SGBU_HAEIN		cytoplasm
P44991	LYXK_HAEIN		cytoplasm
P44992	Y1028_HAEIN	periplasm (probable).	periplasm
P44993	Y1029_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44994	Y1030_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P44995	DLGD_HAEIN	cytoplasm (potential).	cytoplasm
P44996	Y1032_HAEIN		cytoplasm
P44997	SERB_HAEIN		cytoplasm
P44998	CORA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P44999	T2D5_HAEIN		extracellular
P45000	MTH5_HAEIN		cytoplasm
P45002	DMSC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45003	DMSB_HAEIN		periplasm
P45004	DMSA_HAEIN	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	periplasm
P45008	Y1052_HAEIN		cytoplasm
P45011	LPXB_HAEIN		cytoplasm
P45012	FABZ_HAEIN	cytoplasm (by similarity).	cytoplasm

P45014	NRFD_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45015	NRFC_HAEIN		cytoplasm
P45016	NRFB_HAEIN	periplasm (potential).	periplasm
P45017	NRFA_HAEIN	periplasm (by similarity).	periplasm
P45018	HRPA_HAEIN		cytoplasm
P45019	Y1073_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45020	CYOB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45021	CYOA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45022	Y1078_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45023	Y1079_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45024	Y1080_HAEIN	periplasm (potential).	periplasm
P45025	MURA_HAEIN	cytoplasm (probable).	inner membrane
P45026	Y1082_HAEIN		cytoplasm
P45028	Y1084_HAEIN		cytoplasm
P45029	Y1085_HAEIN		cytoplasm
P45030	Y1086_HAEIN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P45031	Y1087_HAEIN		inner membrane
P45032	CCMA_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P45033	CCMB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P45034	CCMC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P45035	CCMD_HAEIN		inner membrane
P45036	CCME_HAEIN	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
P45037	CCMF_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45038	DSBE_HAEIN	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
P45039	CYSZ_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45040	CYSK_HAEIN		cytoplasm

P45042	RFAF_HAEIN		cytoplasm
P45043	XYLR_HAEIN		cytoplasm
P45045	XYLH_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45046	XYLG_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45047	XYLF_HAEIN	periplasm (by similarity).	periplasm
P45048	HLDD_HAEIN		cytoplasm
P45049	COMM_HAEIN		cytoplasm
P45051	OPPF_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45052	OPPD_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45053	OPPC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45054	OPPB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45055	TAL_HAEIN	cytoplasm (probable).	cytoplasm
P45056	MRAZ_HAEIN		inner membrane
P45057	MRAW_HAEIN		cytoplasm
P45058	FTSL_HAEIN	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P45059	FTSI_HAEIN	bacterial cell inner membrane (potential).	inner membrane
P45060	MURE_HAEIN	cytoplasm (probable).	cytoplasm
P45061	MURF_HAEIN	cytoplasm (probable).	cytoplasm
P45062	MRAY_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45063	MURD_HAEIN	cytoplasm (by similarity).	cytoplasm
P45064	FTSW_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45065	MURG_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P45066	MURC_HAEIN	cytoplasm (probable).	cytoplasm
P45067	FTSQ_HAEIN	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P45068	FTSA_HAEIN		inner membrane
P45070	LPXC_HAEIN		cytoplasm
P45071	Y1146_HAEIN		cytoplasm
P45072	PTSN_HAEIN		cytoplasm
P45073	Y1148_HAEIN		inner membrane
P45074	Y1149_HAEIN	periplasm (potential).	periplasm
P45075	Y1150_HAEIN		inner membrane
P45076	Y1151_HAEIN		cytoplasm
P45077	PMBA_HAEIN	cytoplasm (potential).	cytoplasm

P45079	Y1154_HAEIN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P45080	NRDG_HAEIN	cytoplasm (by similarity).	cytoplasm
P45081	CYDC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45082	CYDD_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45083	Y1161_HAEIN		cytoplasm
P45085	Y1165_HAEIN		cytoplasm
P45088	OPA_HAEIN		outer membrane
P45089	ARTM_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45090	ARTQ_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45091	ARTI_HAEIN	periplasm (by similarity).	periplasm
P45092	ARTP_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45093	GMHA_HAEIN	cytoplasm (by similarity).	cytoplasm
P45094	DPPF_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45095	DPPD_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45096	DPPB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45097	Y1189_HAEIN		cytoplasm
P45100	MTFA_HAEIN		cytoplasm
P45101	SUCC_HAEIN		cytoplasm
P45102	SUCD_HAEIN		cytoplasm
P45103	Y1198_HAEIN		inner membrane
P45104	RLUB_HAEIN		cytoplasm
P45105	CYSB_HAEIN	cytoplasm (by similarity).	cytoplasm
P45106	Y1201_HAEIN		cytoplasm
P45107	PTA_HAEIN	cytoplasm (potential).	inner membrane
P45108	CVPA_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45110	ARGR_HAEIN	cytoplasm (probable).	cytoplasm
P45111	DSBC_HAEIN	periplasm (by similarity).	periplasm
P45112	RECJ_HAEIN		cytoplasm
P45113	MTNN_HAEIN		cytoplasm
P45114	Y1217_HAEIN	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P45116	Y1225_HAEIN		cytoplasm
P45117	URAA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P45118	ODP2_HAEIN		inner membrane
P45119	ODP1_HAEIN		cytoplasm
P45120	MGSA_HAEIN		cytoplasm
P45121	PROA_HAEIN	cytoplasm (by similarity).	cytoplasm
P45122	Y1240_HAEIN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P45123	BCR_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45124	RSUA_HAEIN		cytoplasm
P45125	UVRB_HAEIN	cytoplasm (by similarity).	cytoplasm
P45127	Y1252_HAEIN		inner membrane
P45128	MFD_HAEIN		cytoplasm
P45129	HTOA_HAEIN	periplasm (potential).	periplasm
P45134	TEHB_HAEIN	cytoplasm (potential).	cytoplasm
P45135	MRP_HAEIN		cytoplasm
P45138	Y1282_HAEIN		inner membrane
P45141	RBFA_HAEIN	cytoplasm (potential).	cytoplasm
P45142	TRUB_HAEIN		cytoplasm
P45145	Y1297_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45146	Y1298_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45148	CAN_HAEIN		extracellular
P45149	RISB_HAEIN		cytoplasm
P45150	NUSB_HAEIN		cytoplasm
P45153	DAPB_HAEIN	cytoplasm (by similarity).	cytoplasm
P45154	Y1309_HAEIN		cytoplasm
P45157	EX5B_HAEIN		cytoplasm
P45158	EX5A_HAEIN		cytoplasm
P45159	FABA_HAEIN	cytoplasm (by similarity).	cytoplasm
P45161	PBP4_HAEIN	periplasm (potential).	inner membrane
P45162	RRMJ_HAEIN	cytoplasm (by similarity).	cytoplasm
P45164	GLMM_HAEIN		cytoplasm
P45167	UUP2_HAEIN		inner membrane
P45168	POTD1_HAEIN	periplasm (potential).	periplasm
P45169	POTC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45170	POTB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45171	POTA_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P45172	PEPT_HAEIN	cytoplasm (by similarity).	extracellular
P45173	Y1349_HAEIN		cytoplasm
P45174	PUTP_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45176	MALQ_HAEIN	cytoplasm (by similarity).	cytoplasm
P45177	GLGB_HAEIN		extracellular
P45178	GLGX_HAEIN		extracellular

P45179	GLGA_HAEIN		cytoplasm
P45180	PHSG_HAEIN		cytoplasm
P45181	PQQL_HAEIN		extracellular
P45182	Y1369_HAEIN		outer membrane
P45183	MOP_HAEIN		periplasm
P45184	TUSE_HAEIN	cytoplasm (by similarity).	cytoplasm
P45185	MUKF_HAEIN	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
P45188	EX1_HAEIN		cytoplasm
P45189	PHOB_HAEIN	cytoplasm (by similarity).	cytoplasm
P45190	PSTA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45191	PSTC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45192	PSTS_HAEIN	periplasm (potential).	periplasm
P45197	Y1412_HAEIN		cytoplasm
P45198	Y1424_HAEIN		cytoplasm
P45199	FNR_HAEIN	cytoplasm (probable).	cytoplasm
P45200	Y1430_HAEIN		cytoplasm
P45201	USG_HAEIN		cytoplasm
P45202	Y1434_HAEIN		cytoplasm
P45205	DXS_HAEIN		cytoplasm
P45206	SSPB_HAEIN		cytoplasm
P45207	SSPA_HAEIN		cytoplasm
P45208	METF_HAEIN		cytoplasm
P45209	BIOD1_HAEIN		cytoplasm
P45210	MOEA_HAEIN		cytoplasm
P45211	MOEB_HAEIN		cytoplasm
P45213	MSRAB_HAEIN		cytoplasm
P45215	Y1459_HAEIN		cytoplasm
P45217	Y1462_HAEIN		inner membrane
P45219	FTSH2_HAEIN		extracellular
P45220	Y1466_HAEIN		outer membrane
P45221	Y1467_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45239	HTRB_HAEIN	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P45240	GLTS_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45241	RIMK_HAEIN		cytoplasm
P45242	GLRX_HAEIN	cytoplasm (by similarity).	cytoplasm
P45243	SPPA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P45244	Y1542_HAEIN		cytoplasm
P45245	Y1544_HAEIN		cytoplasm
P45246	Y1545_HAEIN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane

P45247	LOLD_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P45248	BIOD2_HAEIN		cytoplasm
P45249	BIOC_HAEIN		cytoplasm
P45250	Y1556_HAEIN		cytoplasm
P45251	KDSA_HAEIN	cytoplasm (by similarity).	cytoplasm
P45252	Y1558_HAEIN		extracellular
P45253	HEMK_HAEIN		cytoplasm
P45255	VPG2_HAEIN		cytoplasm
P45256	DNAB_HAEIN		cytoplasm
P45257	ALR_HAEIN		cytoplasm
P45260	ILVH_HAEIN		cytoplasm
P45261	ILVI_HAEIN		cytoplasm
P45262	Y1590_HAEIN		cytoplasm
P45263	LOLA_HAEIN	periplasm (by similarity).	periplasm
P45264	FTSK_HAEIN		nucleoid
P45265	LRP_HAEIN		cytoplasm
P45266	RADA_HAEIN		cytoplasm
P45267	Y1598_HAEIN		cytoplasm
P45268	Y1604_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45269	CCA_HAEIN		cytoplasm
P45270	LOLB_HAEIN	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P45271	ISPE_HAEIN		cytoplasm
P45272	HMRM_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45273	RISA_HAEIN		cytoplasm
P45274	AMPN_HAEIN	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	extracellular
P45275	Y1618_HAEIN		inner membrane
P45277	ZNTR_HAEIN		cytoplasm
P45279	Y1628_HAEIN		cytoplasm
P45280	Y1629_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45281	RL25_HAEIN		cytoplasm
P45283	PURA_HAEIN	cytoplasm (by similarity).	cytoplasm
P45284	DAPD_HAEIN	cytoplasm (by similarity).	cytoplasm
P45285	SAPA_HAEIN	periplasm (probable).	periplasm
P45286	SAPB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45287	SAPC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45288	SAPD_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane

P45289	SAPF_HAEIN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45290	Y1643_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45291	TRUA_HAEIN		cytoplasm
P45292	F16P_HAEIN	cytoplasm (potential).	cytoplasm
P45293	PDXS_HAEIN		extracellular
P45294	PDXT_HAEIN		cytoplasm
P45295	LDHD_HAEIN	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	inner membrane
P45296	NLPC_HAEIN	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P45297	TLDD_HAEIN		cytoplasm
P45298	Y1654_HAEIN		cytoplasm
P45299	Y1655_HAEIN		inner membrane
P45300	Y1656_HAEIN		inner membrane
P45301	Y1658_HAEIN	periplasm (potential).	periplasm
P45302	ODO2_HAEIN		inner membrane
P45303	ODO1_HAEIN		cytoplasm
P45305	Y1664_HAEIN		extracellular
P45308	MOAE_HAEIN		cytoplasm
P45309	MOAD_HAEIN		cytoplasm
P45310	MOAC_HAEIN		cytoplasm
P45311	MOAA_HAEIN		cytoplasm
P45312	Y1677_HAEIN		cytoplasm
P45313	Y1678_HAEIN		cytoplasm
P45314	KDOP_HAEIN		periplasm
P45315	SOHB_HAEIN		extracellular
P45321	MODC_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P45322	MODB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45323	MODA_HAEIN	periplasm (potential).	periplasm
P45324	MODE_HAEIN	cytoplasm (probable).	cytoplasm
P45331	METE_HAEIN		cytoplasm
P45332	Y1703_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45333	Y1704_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45334	AMPA_HAEIN	cytoplasm (by similarity).	cytoplasm
P45335	Y1706_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P45336	QSEC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45337	QSEB_HAEIN	cytoplasm (potential).	cytoplasm
P45338	PTGA_HAEIN	cytoplasm (by similarity).	cytoplasm

P45339	ENGC_HAEIN		cytoplasm
P45340	ORN_HAEIN	cytoplasm (potential).	cytoplasm
P45341	RFE_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45344	Y1723_HAEIN		inner membrane
P45345	PBPB_HAEIN	bacterial cell inner membrane (by similarity).	inner membrane
P45347	Y1729_HAEIN		inner membrane
P45349	METR_HAEIN	cytoplasm (by similarity).	cytoplasm
P45358	HIS8_ACEPA		cytoplasm
P45365	YPH5_THIVI		cytoplasm
P45366	PHBC_THIVI	cytoplasm (by similarity).	cytoplasm
P45367	YPH2_THIVI		cytoplasm
P45370	PHBC_CHRVI		cytoplasm
P45372	YPH2_CHRVI		cytoplasm
P45373	YPH4_CHRVI		cytoplasm
P45374	YPH5_CHRVI		cytoplasm
P45382	FADH_PARDE		cytoplasm
P45383	RECA_VIBCH	cytoplasm (by similarity).	cytoplasm
P45388	YAEJ_PSEPU		cytoplasm
P45395	KDSD_ECOLI		cytoplasm
P45399	CYCH_BRAJA	bacterial cell inner membrane; multi-pass membrane protein; periplasmic side (potential).	inner membrane
P45400	CYCH_RHIME	bacterial cell inner membrane; multi-pass membrane protein; periplasmic side (potential).	inner membrane
P45401	CCME_BRAJA	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
P45402	CCME_RHIME	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
P45403	CCMF_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45404	CCMF_RHIME	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45405	CCMH_BRAJA	periplasm (probable).	periplasm
P45406	CCMH_RHIME	periplasm (probable).	periplasm
P45408	CCMD_RHILV		inner membrane
P45409	CYCY_RHILV	periplasm (probable).	periplasm
P45410	CITC_KLEPN		cytoplasm
P45414	CITG_KLEPN		cytoplasm
P45416	KDGK_DICD3		cytoplasm
P45417	YHJD_DICD3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45418	YOTL_DICD3		cytoplasm
P45419	PGSA_PSEFL	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane

P45420	YHCD_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P45421	YHCE_ECOLI		extracellular
P45422	YHCF_ECOLI		extracellular
P45423	YHCG_ECOLI		cytoplasm
P45424	YHCH_ECOLI		inner membrane
P45425	NANK_ECOLI		cytoplasm
P45454	DMA_SERMA		cytoplasm
P45455	RPE_SERMA		cytoplasm
P45460	AMPC_YEREN	periplasm (by similarity).	periplasm
P45463	YGIP_ECOLI		cytoplasm
P45464	YRAM_ECOLI		fimbrium
P45465	YRAN_ECOLI		inner membrane
P45468	YRAQ_ECOLI		inner membrane
P45469	YRAR_ECOLI		cytoplasm
P45470	YHBO_ECOLI		extracellular
P45472	YHBQ_ECOLI		cytoplasm
P45475	YHBV_ECOLI		cytoplasm
P45477	PYRD_HAEIN	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P45489	Y990_CAMJE		cytoplasm
P45490	Y987_CAMJE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45491	Y984_CAMJE		cytoplasm
P45492	Y983_CAMJE	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P45493	HIPO_CAMJE		extracellular
P45505	YFAH_ECOLI		inner membrane
P45508	YFAL_ECOLI		extracellular
P45510	DHAK_CITFR		cytoplasm
P45511	GLDA_CITFR		cytoplasm
P45512	DHAR_CITFR		cytoplasm
P45513	DHAT_CITFR		cytoplasm
P45514	DHAB_CITFR		cytoplasm
P45515	YDHW_CITFR		cytoplasm
P45516	YDHX_CITFR		cytoplasm
P45517	YDHY_CITFR		inner membrane
P45524	YHET_ECOLI		inner membrane
P45527	YHBU_ECOLI		periplasm
P45530	TUSB_ECOLI	cytoplasm (probable).	cytoplasm
P45531	TUSC_ECOLI	cytoplasm (probable).	cytoplasm
P45532	TUSD_ECOLI	cytoplasm (probable).	cytoplasm
P45537	YHFK_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45539	FRLA_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45541	FRLC_ECOLI		cytoplasm
P45543	FRLD_ECOLI		cytoplasm
P45544	FRLR_ECOLI		cytoplasm
P45545	YHFS_ECOLI		cytoplasm

P45546	YHFT_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45548	PHP_ECOLI		cytoplasm
P45549	YHFW_ECOLI		cytoplasm
P45550	YHFX_ECOLI		cytoplasm
P45551	YHFY_ECOLI		cytoplasm
P45552	YHFZ_ECOLI		cytoplasm
P45563	XAPA_ECOLI		cytoplasm
P45564	YFEN_ECOLI		outer membrane
P45565	AIS_ECOLI		inner membrane
P45566	YHDT_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45568	DXR_ECOLI		cytoplasm
P45574	DSVA_DESVH		cytoplasm
P45575	DSVB_DESVH		cytoplasm
P45577	PROQ_ECOLI	cytoplasm (potential).	inner membrane
P45578	LUXS_ECOLI		cytoplasm
P45579	YBDH_ECOLI		cytoplasm
P45581	YCFK_ECOLI		extracellular
P45597	PTFAX_XANCP	cytoplasm (potential).	cytoplasm
P45599	FUR_KLEPN	cytoplasm (by similarity).	cytoplasm
P45603	HIS6_KLEOX	cytoplasm (by similarity).	cytoplasm
P45604	PTW3C_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45622	HEM2_BRAJA		cytoplasm
P45632	EXOC_AZOBR		cytoplasm
P45648	RNPA_COXBU		extracellular
P45649	Y1919_COXBU		inner membrane
P45650	OXAA_COXBU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P45670	NTRB_AZOBR		inner membrane
P45671	NTRC_AZOBR		cytoplasm
P45672	DUS_AZOBR		cytoplasm
P45674	YNR3_AZOBR		inner membrane
P45675	NTRY_AZOBR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45676	CHER_CAMJE		cytoplasm
P45677	PEB1C_CAMJE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45679	RSMB_COXBU		cytoplasm
P45680	Y1916_COXBU		cytoplasm
P45681	SURE_HAEIN	cytoplasm (potential).	periplasm
P45682	NLPD_PSEAE	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P45683	PIMT_PSEAE	cytoplasm (by similarity).	cytoplasm
P45684	RPOS_PSEAE		cytoplasm
P45686	RBS_THINE		cytoplasm
P45688	CSOC_THINE		inner membrane
P45689	CSOA_THINE		inner membrane
P45690	CSOB_THINE		inner membrane

P45736	YCJD_ECOLI		cytoplasm
P45748	YRDC_ECOLI		inner membrane
P45750	YRFA_ECOLI		cytoplasm
P45751	YRFB_ECOLI		cytoplasm
P45753	YRFD_ECOLI		cytoplasm
P45755	GSPB_AERHY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45756	GSPA_ECOLI		inner membrane
P45757	GSPC_ECOLI	bacterial cell inner membrane (probable).	inner membrane
P45758	GSPD_ECOLI	bacterial cell outer membrane (probable).	inner membrane
P45759	GSPE_ECOLI	cytoplasm (probable).	inner membrane
P45760	GSPI_ECOLI		inner membrane
P45761	GSPJ_ECOLI		inner membrane
P45762	GSPK_ECOLI	bacterial cell inner membrane (potential).	inner membrane
P45763	GSPL_ECOLI	bacterial cell inner membrane (probable).	inner membrane
P45766	YHDW_ECOLI	periplasm (probable).	periplasm
P45767	YHDX_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P45769	YHDZ_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P45771	YRDD_ECOLI		cytoplasm
P45772	GSPC_AERSA	bacterial cell inner membrane (probable).	inner membrane
P45773	GSPG_VIBCH		inner membrane
P45774	GSPH_VIBCH		inner membrane
P45775	GSPI_VIBCH		inner membrane
P45776	GSPJ_VIBCH		inner membrane
P45777	GSPC_VIBCH		inner membrane
P45778	GSPD_AERSA	bacterial cell outer membrane (probable).	inner membrane
P45779	GSPD_VIBCH	bacterial cell outer membrane; peripheral membrane protein (potential).	inner membrane
P45780	GSPF_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P45781	GSPK_VIBCH	bacterial cell inner membrane (potential).	inner membrane
P45782	GSPL_VIBCH	bacterial cell inner membrane (probable).	inner membrane
P45784	GSPN_VIBCH	bacterial cell inner membrane (probable).	inner membrane
P45785	YEXN_AERSA		cytoplasm
P45789	GSPL_AERHY	bacterial cell inner membrane (probable).	inner membrane
P45790	GSPC_AERHY	bacterial cell inner membrane (probable).	inner membrane
P45791	TAPA_AERHY		inner membrane
P45792	TAPB_AERHY	cytoplasm (potential).	inner membrane

P45793	TAPC_AERHY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P45794	LEP4_AERHY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P45795	YRDB_ECOLI		cytoplasm
P45799	NUDE_ECOLI		cytoplasm
P45804	YHGE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P45805	HYPF1_RALEU		cytoplasm
P45807	YBAM_ECOLI		cytoplasm
P45809	RS12_ERWAM		cytoplasm
P45848	YCIQ_ECOLI		cytoplasm
P45876	DCMR_METDI		cytoplasm
P45955	YBGF_ECOLI		extracellular
P45956	YGBF_ECOLI		cytoplasm
P45992	HIFD1_HAEIN	bacterial cell membrane; lipid-anchor (probable).	extracellular
P45993	HIFD2_HAEIN	bacterial cell membrane; lipid-anchor (probable).	extracellular
P45994	HIFE1_HAEIN		extracellular
P45995	HIFE2_HAEIN		extracellular
P45997	HFC2_HAEIN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P45998	HFC3_HAEIN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P45999	FASC_ECOLI		periplasm
P46001	FASE_ECOLI		periplasm
P46003	ARD1_ECOLI		cytoplasm
P46004	AGGD_ECOLI	periplasm (by similarity).	periplasm
P46005	AGGC_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P46006	AGGB_ECOLI		fimbrium
P46007	AGGA_ECOLI		fimbrium
P46009	FOCD_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P46022	MTGA_ECOLI	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P46028	HLY2_ACTPL		periplasm
P46068	DSDC_ECOLI		cytoplasm
P46072	FRA1_VIBFI		cytoplasm
P46116	AARA_PROST	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P46117	YARA_PROST		cytoplasm
P46118	HEXR_ECOLI		cytoplasm
P46119	YBJC_ECOLI		inner membrane
P46121	YBFK_ECOLI		cytoplasm

P46122	YAJI_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P46126	YFIM_ECOLI		extracellular
P46130	YBHC_ECOLI	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P46142	YGGM_ECOLI		cytoplasm
P46144	YEDJ_ECOLI		cytoplasm
P46145	YAAV_ECOLI		cytoplasm
P46148	ETRA_SHEON		cytoplasm
P46149	Y736_SHEON		cytoplasm
P46154	FADH_PSEPU		cytoplasm
P46173	RL16_BUCAK		cytoplasm
P46174	RL29_BUCAK		cytoplasm
P46175	RS17_BUCAK		cytoplasm
P46176	RL14_BUCAK		cytoplasm
P46177	RL24_BUCAK		cytoplasm
P46178	RL5_BUCAK		cytoplasm
P46179	RS14_BUCAK		cytoplasm
P46180	RS8_BUCAK		cytoplasm
P46181	RL6_BUCAK		cytoplasm
P46182	RL18_BUCAK		cytoplasm
P46183	RS5_BUCAK		cytoplasm
P46184	RL30_BUCAK		cytoplasm
P46185	RL15_BUCAK		cytoplasm
P46206	CATB_PSESY	periplasm (probable).	periplasm
P46231	Y2115_VIBPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P46232	RNT_VIBPA		cytoplasm
P46233	MOTY_VIBPA	bacterial cell membrane; peripheral membrane protein (potential).	inner membrane
P46234	Y2110_VIBPA		cytoplasm
P46235	LGUL_VIBPA		extracellular
P46240	AROE_BUCAP		cytoplasm
P46245	AROH_BUCAP		cytoplasm
P46246	RL20_BUCAP		cytoplasm
P46355	HEM3_YERPE		cytoplasm
P46356	CYAY_YERPE		inner membrane
P46357	DAPF_YERPE	cytoplasm (by similarity).	cytoplasm
P46358	RFAY_XANCP		cytoplasm
P46362	FOLB_HAEIN		cytoplasm
P46363	BIRA_HAEIN		cytoplasm
P46368	DHA2_RALEU		cytoplasm
P46383	TONB_ENTAE	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
P46384	PILG_PSEAE		cytoplasm
P46393	YARQ_ACTPL		cytoplasm
P46399	CH10_ACTAC	cytoplasm (by similarity).	cytoplasm
P46400	RPOD_RHOCA		cytoplasm
P46448	FDXG_HAEIN	cytoplasm (potential).	periplasm
P46449	CSPD_HAEIN	cytoplasm (by similarity).	cytoplasm

P46450	LCFA_HAEIN	membrane; peripheral membrane protein (potential). partially membrane-associated (potential).	cytoplasm
P46451	TRPC_HAEIN		cytoplasm
P46452	GMHB_HAEIN	cytoplasm (by similarity).	cytoplasm
P46453	ENGB_HAEIN		cytoplasm
P46454	LLDD_HAEIN		cytoplasm
P46455	Y974A_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P46456	PURR_HAEIN		cytoplasm
P46457	SFSA_HAEIN		cytoplasm
P46458	CCMH_HAEIN	periplasm (probable).	periplasm
P46474	YHDP_ECOLI		cytoplasm
P46478	AAEX_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P46482	AAEA_ECOLI	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P46490	Y198_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P46491	CRCB_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P46492	METI_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P46493	THD1_HAEIN		cytoplasm
P46494	Y656A_HAEIN		cytoplasm
P46495	Y1572_HAEIN		cytoplasm
P46496	NER_HAEIN		cytoplasm
P46738	NFAE_ECOLI	periplasm (by similarity).	periplasm
P46739	NFAA_ECOLI		fimbrium
P46837	YHGF_ECOLI		cytoplasm
P46846	GNTX_ECOLI		cytoplasm
P46850	RTCB_ECOLI		inner membrane
P46852	YHHW_ECOLI		cytoplasm
P46853	YHHX_ECOLI		cytoplasm
P46854	YHHY_ECOLI		cytoplasm
P46855	YHHZ_ECOLI		cytoplasm
P46856	YRHA_ECOLI		cytoplasm
P46857	YRHB_ECOLI		cytoplasm
P46859	GNTK_ECOLI		cytoplasm
P46879	YQGD_ECOLI		inner membrane
P46887	YECH_ECOLI		cytoplasm
P46888	USPC_ECOLI	cytoplasm (by similarity).	cytoplasm
P46890	YBAE_ECOLI		inner membrane
P46891	COF_ECOLI		cytoplasm
P46924	LEUO_SALTY		cytoplasm
P46928	YKGB_PASHA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P47200	FNRA_PSEST		cytoplasm

P47203	FTSA_PSEAE		inner membrane
P47205	LPXC_PSEAE		cytoplasm
P47206	SCOP_PSEST		periplasm
P47227	BPHB_BURCE		cytoplasm
P47228	BPHC_BURCE		cytoplasm
P47229	BPHD_BURCE		cytoplasm
P47235	GYRA_CAMFE		cytoplasm
P47243	DBFB_PSEPA		cytoplasm
P47734	FADH_METMR	cytoplasm (potential).	cytoplasm
P47737	X19F_ECOLI		cytoplasm
P47765	RPOS_YEREN		cytoplasm
P47846	HEM1_COXBU		cytoplasm
P48027	GACS_PSESY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P48194	RP32_CAUCR		cytoplasm
P48195	GRPE_CAUCR	cytoplasm (probable).	cytoplasm
P48196	RNPH_CAUCR		cytoplasm
P48197	GYRB_CAUCR		cytoplasm
P48198	DPO3B_CAUCR	cytoplasm (by similarity).	cytoplasm
P48204	GRPE_FRATU	cytoplasm (probable).	cytoplasm
P48205	DNAK_FRATU		cytoplasm
P48207	DNAJ_FRATU	cytoplasm (probable).	cytoplasm
P48208	DNAJ_HAEDU	cytoplasm (probable).	cytoplasm
P48209	DNAK_HAEDU		cytoplasm
P48210	CH60_BORPE	cytoplasm (by similarity).	cytoplasm
P48211	CH60_CAUCR	cytoplasm (by similarity).	cytoplasm
P48213	CH60_EHRRW	cytoplasm (by similarity).	cytoplasm
P48214	CH60_EHRRW	cytoplasm (by similarity).	cytoplasm
P48215	CH60_NEIFL	cytoplasm (by similarity).	cytoplasm
P48216	CH60_PSEPU	cytoplasm (by similarity).	cytoplasm
P48219	CH60_YEREN	cytoplasm (by similarity).	cytoplasm
P48220	CH60_ZYMMO	cytoplasm (by similarity).	cytoplasm
P48222	CH10_CAUCR	cytoplasm (by similarity).	cytoplasm
P48224	CH10_EHRRW	cytoplasm (by similarity).	cytoplasm
P48226	CH10_PSEPU	cytoplasm (by similarity).	cytoplasm
P48228	CH10_YEREN	cytoplasm (by similarity).	cytoplasm
P48229	CH10_ZYMMO	cytoplasm (by similarity).	cytoplasm
P48246	HEM4_PSEAE		cytoplasm
P48247	GSA_PSEAE	cytoplasm (potential).	cytoplasm
P48285	ENO_HELPY	cytoplasm (by similarity).	extracellular
P48289	RECA_GLUOX	cytoplasm (by similarity).	cytoplasm
P48291	RECA1_MYXXA	cytoplasm (by similarity).	cytoplasm
P48292	RECA2_MYXXA	cytoplasm (by similarity).	cytoplasm
P48339	HUPG_BRAJA		cytoplasm
P48340	HUPH_BRAJA		cytoplasm
P48341	HUPJ_BRAJA		cytoplasm
P48342	HUPK_BRAJA		cytoplasm
P48343	RUBR_AZOCH		cytoplasm
P48344	RUBL_BRAJA		cytoplasm
P48345	DESR_DESVM		inner membrane
P48364	MDH_MORS5		cytoplasm
P48369	GYRA_AERSA		cytoplasm
P48370	GYRA_HELPY		cytoplasm
P48371	GYRA_NEIGO		cytoplasm
P48372	GYRA_PSEAE		cytoplasm

P48374	PARC_NEIGO	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P48516	IF3_MYXXA	cytoplasm (by similarity).	cytoplasm
P48571	LEU1_BUCRP		cytoplasm
P48573	LEU2_BUCRP		cytoplasm
P48574	LEUD_BUCRP		cytoplasm
P48633	HMWP2_YEREN		cytoplasm
P48636	Y4923_PSEAE		cytoplasm
P48796	FUR_CAMJE	cytoplasm (by similarity).	cytoplasm
P48823	HEXA_ALTSO		cytoplasm
P48839	AGAA_VIBS7		extracellular
P48840	AGAB_VIBS7		extracellular
P48841	GANA_PSEFL		extracellular
P48850	RS10_RICPR		cytoplasm
P48851	RS10_NEIGO		cytoplasm
P48948	RS20_RHIME		cytoplasm
P48952	RL3_RICPR		cytoplasm
P48982	BGAL_XANMN		extracellular
P49007	HEXB_ALTSO		extracellular
P49155	XYLI_PSEPU		cytoplasm
P49156	DMPH_PSEUF		cytoplasm
P49172	4OT_PSEUF		cytoplasm
P49225	RS21_MYXXA		cytoplasm
P49226	RL4_MORMO		cytoplasm
P49239	RL2_YEREN		cytoplasm
P49240	RL35_BUCAP		cytoplasm
P49251	MAOC_KLEAE		cytoplasm
P49304	MOCC_RHIME		cytoplasm
P49305	YMO1_RHIME		cytoplasm
P49306	YMO2_RHIME		cytoplasm
P49307	MOCA_RHIME		cytoplasm
P49308	MOCB_RHIME	periplasm (potential).	periplasm
P49309	MOCR_RHIME		cytoplasm
P49392	RS15_CAMJE		cytoplasm
P49400	RS20_CAUCR		cytoplasm
P49408	ROC2_AGRRH		extracellular
P49409	ROB2_AGRRH		extracellular
P49417	CAT_VIBAN		cytoplasm
P49424	MANA_PSEFL		extracellular
P49617	HSLV_PASHA	cytoplasm (by similarity).	cytoplasm
P49618	LAPT_PASHA	periplasm (by similarity).	periplasm
P49944	BFR_BRUME		cytoplasm
P49988	RP54_PSEAE		cytoplasm
P49989	RP54_RHIET		cytoplasm
P49996	DNAA_VIBHA	cytoplasm (by similarity).	cytoplasm
P49997	RECF_AZOV1	cytoplasm (potential).	cytoplasm
P49998	RECF_CAUCR	cytoplasm (potential).	cytoplasm
P50018	DNAJ_AGRT5	cytoplasm (probable).	cytoplasm
P50019	DNAK_AGRT5		cytoplasm
P50025	DNAJ_LEGPN	cytoplasm (probable).	cytoplasm
P50054	SECE_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P50056	NUSG_RICPR		cytoplasm
P50070	PRIM_MYXXA		cytoplasm

P50173	GLDA_PSEPU		cytoplasm
P50182	MTN4_NEILA		cytoplasm
P50183	T2N4_NEILA		extracellular
P50191	T2H1_HAEPH		extracellular
P50192	MTHA_HAEPH		cytoplasm
P50193	MTHB_HAEPH		cytoplasm
P50194	T2E7_ECOLI		extracellular
P50195	T2E8_ECOLI		extracellular
P50196	MTE8_ECOLI		cytoplasm
P50197	LINC_PSEPA		cytoplasm
P50198	LINX_PSEPA		cytoplasm
P50201	MAS12_AGRRH		cytoplasm
P50202	MAS1_AGRT9		cytoplasm
P50206	BPHB_PSES1		cytoplasm
P50215	IDH_SPHYA		cytoplasm
P50321	G3PC_RALEU		cytoplasm
P50322	G3PP_RALEU		cytoplasm
P50323	NODD1_BRAEL		cytoplasm
P50324	NODD2_BRAEL		cytoplasm
P50327	NODD1_BRASN		cytoplasm
P50328	NODD2_BRASN		cytoplasm
P50330	NOLA_BRASN		cytoplasm
P50331	NODD1_RHIGA		cytoplasm
P50332	NODI_RHIGA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P50333	NODJ_RHIGA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P50335	YHJJ_SALTY	periplasm (potential).	extracellular
P50337	Y4SM_RHISN		cytoplasm
P50350	CHVI_RHIME	cytoplasm (potential).	cytoplasm
P50351	CHVI_RHISN	cytoplasm (potential).	cytoplasm
P50356	NODC_RHIGA	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P50357	NODC_RHISN	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P50358	Y4SN_RHISN		inner membrane
P50359	Y4HO_RHISN		cytoplasm
P50360	Y4HP_RHISN		cytoplasm
P50361	Y4HQ_RHISN		cytoplasm
P50456	MLC_ECOLI		cytoplasm
P50457	PUUE_ECOLI		cytoplasm
P50465	END8_ECOLI		cytoplasm
P50500	IRO_THIFE	periplasm (potential).	periplasm
P50507	RP32_AGRTU		cytoplasm
P50508	RP32_ENTCL		cytoplasm
P50509	RP32_PROMI		cytoplasm
P50510	RP32_SERMA		cytoplasm
P50511	RP32_VIBCH		cytoplasm
P50512	RP32_ZYMMO		cytoplasm
P50513	RLUD_ZYMMO		cytoplasm
P50529	STHA_VIBCH	cytoplasm (by similarity).	cytoplasm
P50587	PYRE_PSEAE		cytoplasm

P50597	RNPH_PSEAE		cytoplasm
P50598	TOLQ_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P50599	TOLR_PSEAE	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P50600	TOLA_PSEAE	bacterial cell inner membrane; single-pass type ii membrane protein (potential).	inner membrane
P50601	TOLB_PSEAE	periplasm (potential).	periplasm
P50612	FLAA_HELMU		flagellar
P50853	RIBD_ACTPL		cytoplasm
P50854	RISA_ACTPL		cytoplasm
P50855	RIBAB_ACTPL		cytoplasm
P50856	RISB_ACTPL		cytoplasm
P50858	AAC6_ENTAE		cytoplasm
P50868	CAT4_ENTAE		cytoplasm
P50869	CAT4_MORMO		cytoplasm
P50872	TRPE_AZOBR		cytoplasm
P50921	TPIS_VIBMA		cytoplasm
P50922	RBL2_RHOCA		cytoplasm
P50932	FMT_RICPR		cytoplasm
P50935	HIS2_RHOS4	cytoplasm (by similarity).	cytoplasm
P50936	HIS4_RHOS4	cytoplasm (by similarity).	cytoplasm
P50937	HIS6_RHOS4	cytoplasm (by similarity).	cytoplasm
P50938	Y612_RICPR		cytoplasm
P50939	NUOL_RHOCA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P50940	NUOK_RHOCA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P50941	FABG_RICPR		cytoplasm
P50973	NUON_RHOCA	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P50974	NUOM_RHOCA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P50975	NUOJ_RHOCA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P50987	ARLY_PSEAE	cytoplasm (probable).	cytoplasm
P51000	DPPC_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P51007	FNRL_RHOS4		cytoplasm
P51008	HEMZ_RHOS4	cytoplasm (potential).	cytoplasm
P51009	G3P_XANFL		cytoplasm
P51010	TKT_XANFL		extracellular
P51012	RPE_RHOCA		cytoplasm
P51013	RPE_RHORU		cytoplasm
P51014	BPHF_PSES1		cytoplasm
P51015	BPHI_BURCE		cytoplasm
P51016	DMPG_PSEUF		cytoplasm

P51017	NAHM_PSEPU		cytoplasm
P51018	TODH_PSEPU		cytoplasm
P51019	XYLK_PSEPU		cytoplasm
P51020	MHPE_ECOLI		cytoplasm
P51024	YAIL_ECOLI		cytoplasm
P51025	YAIM_ECOLI		cytoplasm
P51026	YI2A_ECOLI		cytoplasm
P51033	CISY_BARHE		cytoplasm
P51034	CISY_BARQU		cytoplasm
P51037	CISY1_RHITR		cytoplasm
P51038	CISY2_RHITR		cytoplasm
P51040	CISY_RICFE		cytoplasm
P51042	CISY_RICCN		cytoplasm
P51043	CISY_RICTY		cytoplasm
P51045	CISY_THIFE		cytoplasm
P51053	DHSB_COXBU		cytoplasm
P51054	DHSA_COXBU		inner membrane
P51056	ODO1_COXBU		cytoplasm
P51067	ACEK_SALTY	cytoplasm (by similarity).	cytoplasm
P51185	PTSO_KLEOX	cytoplasm (probable).	cytoplasm
P51560	TETR7_VIBAN		cytoplasm
P51561	TETR8_PASMU		cytoplasm
P51562	TETR8_PASPI		cytoplasm
P51565	TNP9_ECOLI		cytoplasm
P51585	ALGD_AZOVI		cytoplasm
P51594	GCH1_CAMJE		cytoplasm
P51618	GCH2_PHOPO		extracellular
P51691	ARS_PSEAE	cytoplasm (potential).	extracellular
P51696	FOLD_PHOPO		cytoplasm
P51752	MIP_COXBU	secreted protein (potential).	outer membrane
P51758	CYCR_RHOGE	bacterial cell membrane; lipid- anchor (by similarity).	periplasm
P51836	ERA_COXBU		cytoplasm
P51838	RECO_COXBU		cytoplasm
P51852	DCIP_AZOBR		cytoplasm
P51853	BZNB_PSEFL		cytoplasm
P51961	RISA_PHOPO		cytoplasm
P51962	RIBB_PHOPO		cytoplasm
P51963	RISB_PHOPO		cytoplasm
P51967	YGAD_ENTAG		inner membrane
P51973	COMA_NEIGO	bacterial cell inner membrane; multi- pass membrane protein (probable).	inner membrane
P51981	YCJG_ECOLI		cytoplasm
P51995	PNTAB_RICPR	bacterial cell inner membrane; multi- pass membrane protein (by similarity).	inner membrane
P52003	MEXR_PSEAE		cytoplasm
P52005	TORY_ECOLI	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P52006	YFAO_ECOLI		extracellular
P52007	YECM_ECOLI		cytoplasm
P52021	RNH2_VIBCH	cytoplasm (potential).	extracellular
P52022	DPO3A_VIBCH	cytoplasm (by similarity).	cytoplasm
P52024	HOLB_PSEAE		cytoplasm

P52037	YGFF_ECOLI		cytoplasm
P52043	YGFH_ECOLI		cytoplasm
P52044	YGFI_ECOLI		cytoplasm
P52045	MMCD_ECOLI		cytoplasm
P52047	YGGA_AERHY	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P52048	YGGP_ECOLI		cytoplasm
P52052	YGGR_ECOLI		inner membrane
P52053	PROC_VIBAL	cytoplasm (by similarity).	cytoplasm
P52055	YPI1_VIBAL		cytoplasm
P52059	YPI3_VIBAL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P52060	YGGU_ECOLI		inner membrane
P52061	HAM1_ECOLI		extracellular
P52062	YGGW_ECOLI		cytoplasm
P52070	PHAC_METEX	cytoplasm (by similarity).	cytoplasm
P52073	GLCE_ECOLI		cytoplasm
P52074	GLCF_ECOLI		cytoplasm
P52076	QSEB_ECOLI	cytoplasm (potential).	cytoplasm
P52086	COBC_ECOLI		cytoplasm
P52089	YPAH_PSELE		inner membrane
P52095	DCLZ_ECOLI		cytoplasm
P52096	YAER_ECOLI		cytoplasm
P52097	TILS_ECOLI	cytoplasm (probable).	cytoplasm
P52102	YFHL_ECOLI		periplasm
P52106	CSGD_ECOLI		cytoplasm
P52107	CSGC_ECOLI		extracellular
P52108	RSTA_ECOLI	cytoplasm (potential).	cytoplasm
P52112	GLPM_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P52116	SSRP_VIBCH	cytoplasm (potential).	cytoplasm
P52117	SMPA_VIBCH	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P52118	RECN_VIBCH		cytoplasm
P52119	YFJF_ECOLI		inner membrane
P52120	Y850_VIBCH		inner membrane
P52122	Y849_VIBCH		cytoplasm
P52123	YFJH_ECOLI		cytoplasm
P52124	YFJI_ECOLI		cytoplasm
P52125	YFJJ_ECOLI		cytoplasm
P52126	YFJK_ECOLI		cytoplasm
P52127	YFJL_ECOLI		cytoplasm
P52128	YFJM_ECOLI		cytoplasm
P52129	YFJN_ECOLI		cytoplasm
P52130	YFJO_ECOLI		cytoplasm
P52131	YFJP_ECOLI		cytoplasm
P52132	YFJQ_ECOLI		cytoplasm
P52133	YFJR_ECOLI		cytoplasm
P52134	YPJK_ECOLI		inner membrane
P52135	YFJT_ECOLI		extracellular
P52138	YFJW_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

P52139	YFJX_ECOLI		cytoplasm
P52140	YFJY_ECOLI		cytoplasm
P52141	YFJZ_ECOLI		cytoplasm
P52143	YPJA_ECOLI	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P52144	ARSR2_ECOLI		cytoplasm
P52145	ARSA2_ECOLI		inner membrane
P52146	ARB2_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P52147	ARSC2_ECOLI		cytoplasm
P52148	ARD2_ECOLI		cytoplasm
P52151	PURA_THIFE	cytoplasm (by similarity).	cytoplasm
P52152	RHO_CHRVI		inner membrane
P52155	RHO_PSEFL		inner membrane
P52156	RHO_RHOS4		inner membrane
P52218	CCMA_PARDE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P52219	CCMB_PARDE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P52220	CCMC_PARDE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P52223	CYCH_PSEFL	bacterial cell inner membrane; multi-pass membrane protein; periplasmic side (potential).	inner membrane
P52225	CCMF_PSEFL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P52226	CCMH_PSEFL	periplasm (probable).	periplasm
P52233	THIO_THIFE		cytoplasm
P52237	DSBE_PSEFL	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
P52279	PIP_XANCI	cytoplasm (by similarity).	cytoplasm
P52310	AROAHAESO	cytoplasm (probable).	cytoplasm
P52311	MTX2_XANOR		cytoplasm
P52312	AROACAMJE	cytoplasm (probable).	cytoplasm
P52313	URE1_YERPS	cytoplasm (by similarity).	cytoplasm
P52316	URED_YERPS		cytoplasm
P52317	UREE_YERPS	cytoplasm (by similarity).	cytoplasm
P52318	UREF_YERPS		cytoplasm
P52324	RPODCAUOCR		cytoplasm
P52325	RPODNEIGO		cytoplasm
P52326	RPODPSEFL		cytoplasm
P52327	RPODPSEPU		cytoplasm
P52558	PUR6_BRUME		cytoplasm
P52559	PURK_BRUME		cytoplasm
P52595	CBBRRHORU		cytoplasm
P52598	YGBI_ECOLI		cytoplasm

P52599	EMRK_ECOLI	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P52602	KLCA1_ECOLI		cytoplasm
P52603	KLCA2_ECOLI		cytoplasm
P52604	KLCB1_ECOLI		cytoplasm
P52605	KLCB2_ECOLI		cytoplasm
P52606	Y1657_HAEIN		cytoplasm
P52612	FLII_ECOLI	cytoplasm (potential).	inner membrane
P52614	FLIK_ECOLI		flagellar
P52615	FLJB_SALAE		flagellar
P52616	FLJB_SALTY		flagellar
P52617	FLJA_SALAE		cytoplasm
P52618	FLJA_SALAB		cytoplasm
P52619	FLJA_SALTY		cytoplasm
P52621	FLIA_YEREN		cytoplasm
P52625	PATM_VIBHA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P52626	PATH_VIBHA	periplasm (potential).	periplasm
P52627	FLIZ_ECOLI		cytoplasm
P52642	RFBC_SALBO		cytoplasm
P52643	LDHD_ECOLI		cytoplasm
P52644	HSLJ_ECOLI		cytoplasm
P52645	YDBH_ECOLI		cytoplasm
P52647	NIFJ_ECOLI		cytoplasm
P52660	BLAA_PROVU		cytoplasm
P52661	GBPR_AZOBR		cytoplasm
P52662	PECT_DICD3		cytoplasm
P52663	BLAN_ENTCL		extracellular
P52664	BLAB_PROVU		extracellular
P52666	BUDR_KLETE		cytoplasm
P52667	ESTR_ACIAD		cytoplasm
P52669	TTUA4_AGRVI		cytoplasm
P52670	ILVR_CAUCR		cytoplasm
P52671	UDP_KLEPN	cytoplasm (by similarity).	cytoplasm
P52672	CYSH_THIRO	cytoplasm (by similarity).	cytoplasm
P52673	CYSI_THIRO		cytoplasm
P52674	CYSJ_THIRO		cytoplasm
P52676	NMCR_ENTCL		cytoplasm
P52679	PCPR_SPHCR		cytoplasm
P52680	DBHA_SERMA		cytoplasm
P52681	DBHB_SERMA		cytoplasm
P52682	BLAN_SERMA		extracellular
P52683	SMER_SERMA		cytoplasm
P52684	MAUR_KLEPN		cytoplasm
P52685	MAUR_PARDE		cytoplasm
P52686	SDSB_PSES9		cytoplasm
P52687	CITA_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P52688	CITB_KLEPN	cytoplasm (probable).	cytoplasm
P52689	LTRA_KLEPN		cytoplasm
P52690	CBBR_RHOSH		cytoplasm
P52695	HLYU_VIBCH		cytoplasm
P52696	YBHD_ECOLI		cytoplasm

P52697	6PGL_ECOLI		cytoplasm
P52698	PHCA_RALSO		cytoplasm
P52699	BLAB_SERMA		extracellular
P52700	BLA1_XANMA	periplasm (potential).	periplasm
P52831	HYIN_PSESY		extracellular
P52832	GPMI_PSESM		cytoplasm
P52863	RL18_VIBPR		cytoplasm
P52966	SECA_RHOCA	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	inner membrane
P52967	FLAV_RHOCA		cytoplasm
P52973	MURI_HAEIN		cytoplasm
P52975	RNH2_CAUCR	cytoplasm (potential).	extracellular
P52978	NODQ_RHITR		cytoplasm
P52979	GLGB_AGRTU		extracellular
P52988	YENI_YEREN		cytoplasm
P52989	SWRI_SERLI		cytoplasm
P52990	PSYI_PSESZ		cytoplasm
P52993	ODO2_RALEU		inner membrane
P52994	NODH_RHITR		cytoplasm
P52995	NODP_RHITR		cytoplasm
P53000	PROA_CAMJE	cytoplasm (by similarity).	cytoplasm
P53419	NODK_BRAEL		cytoplasm
P53508	F6A1_ECOLI		fimbrium
P53509	F6A2_ECOLI		fimbrium
P53510	F6B1_ECOLI		fimbrium
P53511	F6B2_ECOLI		fimbrium
P53512	CSD1_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P53513	CSD2_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P53514	PMFC_PROMI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P53515	AFAA_ECOLI		cytoplasm
P53516	AFAB_ECOLI	periplasm (potential).	periplasm
P53517	AFAC_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P53518	CSSC1_ECOLI	periplasm (potential).	periplasm
P53519	CSSC2_ECOLI	periplasm (potential).	periplasm
P53520	PMFD_PROMI	periplasm (potential).	periplasm
P53570	ETFB_METME		cytoplasm
P53571	ETFA_METME		cytoplasm
P53572	FIXC_AZOVI		cytoplasm
P53573	ETFA_BRAJA		cytoplasm
P53574	FIXB_AZOVI		cytoplasm
P53575	ETFB_BRAJA		cytoplasm
P53576	FIXA_AZOVI		cytoplasm
P53577	FIXA_BRAJA		cytoplasm
P53591	SUCD_COXBU		cytoplasm
P53592	SUCC_COXBU		cytoplasm
P53593	SUCC_PSEAE		cytoplasm

P53594	MTKA_METEX		cytoplasm
P53595	MTKB_METEX		cytoplasm
P53606	FLA_LEGMI		flagellar
P53607	THD1_BURCE		cytoplasm
P53639	SODF_CAMCO		periplasm
P53640	SODF_CAMJE		periplasm
P53641	SODF_PSEAE		periplasm
P53642	SODM_BORPE		periplasm
P53652	SODM_PSEAE		periplasm
P53655	SODM_YEREN		periplasm
P53656	BIOA_ESCVU		cytoplasm
P53658	FIXX_AZOVI		cytoplasm
P53818	ALF_CAMJE		cytoplasm
P54085	TATC_AZOCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P54203	TRPB_PASMU		cytoplasm
P54220	ARO_A_PASHA	cytoplasm (probable).	cytoplasm
P54222	BETB1_RHIME		cytoplasm
P54291	RHLI_PSEAE		cytoplasm
P54292	RHLR_PSEAE		cytoplasm
P54293	ESAR_ERWST		cytoplasm
P54294	TRAR_AGR5		cytoplasm
P54295	YENR_YEREN		cytoplasm
P54297	LUXL_VIBHA		cytoplasm
P54298	LUXM_VIBHA		cytoplasm
P54299	LUXO_VIBHA		cytoplasm
P54300	LUXP_VIBHA	periplasm (probable).	periplasm
P54301	LUXN_VIBHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P54302	LUXQ_VIBHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P54303	PHZR_PSECL		cytoplasm
P54305	HRC_A_CAUCR		cytoplasm
P54413	CLPP_CAMJE	cytoplasm (by similarity).	cytoplasm
P54414	CLPP_PARDE	cytoplasm (by similarity).	cytoplasm
P54649	HTPG_ACTAC	cytoplasm (by similarity).	cytoplasm
P54656	ESAI_ERWST		extracellular
P54689	ILVE_HAEIN		cytoplasm
P54698	RCSA_SALTY		cytoplasm
P54736	PKN2_MYXXA		inner membrane
P54746	YBGG_ECOLI		cytoplasm
P54794	MOAR_KLEAE		cytoplasm
P54795	MOAE_KLEAE		cytoplasm
P54796	MOAF_KLEAE		cytoplasm
P54901	CSIE_ECOLI		cytoplasm
P54905	CRTB_RHOS4		cytoplasm
P54906	CRTF_RHOS4		cytoplasm
P54907	TRBB_AGR5	cytoplasm (probable).	inner membrane
P54908	TRBC_AGR5	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P54909	TRBD_AGR5		inner membrane
P54910	TRBE_AGR5		cytoplasm

P54911	TRBJ_AGRTU		cytoplasm
P54912	TRBK_AGRTU		cytoplasm
P54913	TRBL_AGRTU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P54914	TRBF_AGRTU		inner membrane
P54915	TRBG_AGRTU		cytoplasm
P54916	TRBH_AGRTU	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P54917	TRBI_AGRTU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P54924	SQHC_BRAJA	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P54925	DEGP_BARHE	periplasm (potential).	periplasm
P54929	NIFA_AZOLI		cytoplasm
P54930	NIFA_ENTAG		cytoplasm
P54931	NIFA_RHIEC		cytoplasm
P54933	SMOK_RHOSH		inner membrane
P54934	BISC_RHOSH		periplasm
P54972	CRTW_PARSN		inner membrane
P54973	CRTZ_PARSN		cytoplasm
P54974	CRTY_PARSN		cytoplasm
P54975	CRTB_PARSN		cytoplasm
P54976	CRTE_RHOS4		cytoplasm
P54978	CRTI_PARSN		cytoplasm
P54979	CRTJ_MYXXA		cytoplasm
P54980	CRTI_RHOS4		cytoplasm
P54988	NTRA_CHEHE		cytoplasm
P54989	NTAA_CHEHE		cytoplasm
P54990	NTAB_CHEHE		cytoplasm
P55044	FPG_NEIMB		cytoplasm
P55110	PROA_LEGLO		extracellular
P55117	LKTA_PASTR	secreted protein (by similarity).	extracellular
P55118	LKA11_PASHA	secreted protein (by similarity).	extracellular
P55120	LKC3_PASHA	cytoplasm (potential).	cytoplasm
P55121	LKCB_PASHA	cytoplasm (potential).	cytoplasm
P55122	LKTB_PASSP	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P55124	LKTC_PASSP	cytoplasm (potential).	cytoplasm
P55125	HLYD_PASSP	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P55132	RT1C_ACTPL	cytoplasm (potential).	cytoplasm
P55133	RELA_VIBSS		cytoplasm
P55135	RUMA_ECOLI		cytoplasm
P55138	YGCE_ECOLI		cytoplasm
P55140	YGCG_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55172	PQQB_PSEFL		cytoplasm
P55174	PQQF_PSEFL		extracellular

P55176	YPQQ_PSEFL		cytoplasm
P55186	ILVD_CAUCR		cytoplasm
P55218	METZ_PSEAE		cytoplasm
P55219	VRP1_SALEN		extracellular
P55220	VRP2_SALEN		extracellular
P55222	VFR_PSEAE		cytoplasm
P55253	RMLA3_ECOLI		cytoplasm
P55254	RMLA_SALAN		cytoplasm
P55255	RMLA_NEIMB		cytoplasm
P55256	RMLA_XANCP		cytoplasm
P55257	RMLA_YEREN		cytoplasm
P55293	RFBB2_ECOLI		cytoplasm
P55294	RFBB_NEIMB		cytoplasm
P55295	RFBB_XANCP		cytoplasm
P55323	MUCR_RHIME		cytoplasm
P55324	ROSA_AGRRD		cytoplasm
P55336	FABG_VIBHA		cytoplasm
P55338	FABF_VIBHA		cytoplasm
P55348	SQHC_RHISN		cytoplasm
P55349	Y4AB_RHISN		cytoplasm
P55350	Y4AC_RHISN		cytoplasm
P55351	Y4AD_RHISN		cytoplasm
P55353	Y4AF_RHISN		inner membrane
P55354	NOEL_RHISN		cytoplasm
P55355	NODZ_RHISN		cytoplasm
P55356	NOEK_RHISN		cytoplasm
P55357	NOEJ_RHISN		cytoplasm
P55358	Y4AK_RHISN		cytoplasm
P55359	NODD1_RHISN		cytoplasm
P55360	Y4AM_RHISN		cytoplasm
P55361	SYRB_RHISN		cytoplasm
P55362	Y4AO_RHISN		cytoplasm
P55363	MUCR_RHISN		cytoplasm
P55364	Y4AQ_RHISN		cytoplasm
P55365	Y4AR_RHISN		cytoplasm
P55366	Y4AS_RHISN		cytoplasm
P55368	Y4BA_RHISN		inner membrane
P55369	Y4BB_RHISN		cytoplasm
P55370	Y4BC_RHISN		cytoplasm
P55371	Y4BD_RHISN		cytoplasm
P55373	Y4BF_RHISN		cytoplasm
P55374	Y4BG_RHISN		cytoplasm
P55375	Y4BH_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55376	Y4BI_RHISN		cytoplasm
P55377	Y4BJ_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	extracellular
P55378	Y4BK_RHISN		cytoplasm
P55379	Y4BL_RHISN		cytoplasm
P55380	Y4BM_RHISN		cytoplasm
P55381	Y4BN_RHISN		cytoplasm
P55382	Y4BO_RHISN		cytoplasm
P55383	Y4CA_RHISN		cytoplasm

P55384	Y4CB_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55385	Y4CC_RHISN		cytoplasm
P55386	Y4CD_RHISN		cytoplasm
P55387	Y4CE_RHISN		cytoplasm
P55388	Y4CF_RHISN		cytoplasm
P55389	Y4CG_RHISN		cytoplasm
P55390	Y4CH_RHISN	cytoplasm (potential).	cytoplasm
P55391	Y4CI_RHISN		nucleoid
P55392	Y4CJ_RHISN		cytoplasm
P55393	Y4CK_RHISN		cytoplasm
P55394	TRAI_RHISN		cytoplasm
P55395	TRBB_RHISN	cytoplasm (probable).	inner membrane
P55396	TRBC_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55397	TRBD_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55398	TRBE1_RHISN		cytoplasm
P55399	TRBE2_RHISN		cytoplasm
P55400	TRBJ_RHISN		cytoplasm
P55401	TRBK_RHISN	periplasm (potential).	periplasm
P55402	TRBL_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55403	TRBF_RHISN		inner membrane
P55404	TRBG_RHISN	periplasm (potential).	periplasm
P55405	TRBH_RHISN	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P55406	TRBI_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55407	TRAR_RHISN		cytoplasm
P55408	TRAM_RHISN		cytoplasm
P55409	Y4DJ_RHISN		cytoplasm
P55410	Y4DK_RHISN		cytoplasm
P55411	Y4DL_RHISN		cytoplasm
P55412	Y4DM_RHISN		cytoplasm
P55413	Y4DN_RHISN		cytoplasm
P55414	Y4DO_RHISN		cytoplasm
P55415	Y4DP_RHISN		cytoplasm
P55416	TRAB_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55417	TRAF_RHISN	periplasm (probable).	periplasm
P55418	TRAA_RHISN		cytoplasm
P55419	TRAC_RHISN		cytoplasm
P55420	TRAD_RHISN		cytoplasm
P55421	TRAG_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55422	Y4DW_RHISN		cytoplasm
P55423	Y4DX_RHISN		cytoplasm
P55424	Y4EA_RHISN		cytoplasm

P55425	Y4EB_RHISN		cytoplasm
P55426	Y4EC_RHISN		cytoplasm
P55427	Y4ED_RHISN		cytoplasm
P55428	Y4EE_RHISN	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P55429	Y4EF_RHISN		cytoplasm
P55431	NOLL_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	cytoplasm
P55432	Y4EI_RHISN		inner membrane
P55434	Y4EK_RHISN		cytoplasm
P55435	Y4EL_RHISN		cytoplasm
P55437	Y4EN_RHISN		cytoplasm
P55438	Y4EO_RHISN		cytoplasm
P55439	Y4FA_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55440	Y4FB_RHISN		cytoplasm
P55441	Y4FC_RHISN		cytoplasm
P55442	Y4FD_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55443	Y4FE_RHISN		cytoplasm
P55444	Y4FF_RHISN		cytoplasm
P55445	Y4FG_RHISN		cytoplasm
P55446	Y4FH_RHISN		cytoplasm
P55448	Y4FJ_RHISN	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P55449	Y4FK_RHISN		cytoplasm
P55450	Y4FL_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55451	Y4FM_RHISN		cytoplasm
P55452	Y4FN_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P55453	Y4FO_RHISN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P55454	Y4FP_RHISN	periplasm (potential).	periplasm
P55455	Y4FQ_RHISN		cytoplasm
P55456	Y4FR_RHISN		cytoplasm
P55457	Y4GA_RHISN		cytoplasm
P55458	Y4GB_RHISN		cytoplasm
P55459	Y4GC_RHISN		cytoplasm
P55460	Y4GD_RHISN		cytoplasm
P55461	Y4GE_RHISN		cytoplasm
P55462	RFBB_RHISN		cytoplasm
P55463	RFBD_RHISN		cytoplasm
P55464	RMLA_RHISN		cytoplasm
P55465	Y4GI_RHISN		cytoplasm
P55466	Y4GJ_RHISN		cytoplasm
P55467	FIXF_RHISN		cytoplasm
P55468	RFBC_RHISN		cytoplasm

P55469	Y4GM_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55470	Y4GN_RHISN		cytoplasm
P55471	Y4HA_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55472	NOEE_RHISN		cytoplasm
P55473	NOEI_RHISN	cytoplasm (potential).	cytoplasm
P55474	NOLO_RHISN	cytoplasm (potential).	inner membrane
P55475	NODJ_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P55476	NODI_RHISN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P55477	Y4HJ_RHISN		cytoplasm
P55478	Y4HK_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55479	Y4HL_RHISN		extracellular
P55480	Y4HM_RHISN		cytoplasm
P55481	Y4HN_RHISN		cytoplasm
P55482	Y4HR_RHISN		cytoplasm
P55485	Y4IR_RHISN		cytoplasm
P55486	Y4IC_RHISN		inner membrane
P55487	Y4ID_RHISN		cytoplasm
P55488	Y4IE_RHISN		cytoplasm
P55489	Y4IF_RHISN		cytoplasm
P55490	Y4IG_RHISN		cytoplasm
P55491	Y4IH_RHISN		cytoplasm
P55492	Y4II_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55493	Y4IJ_RHISN		inner membrane
P55494	Y4IK_RHISN		periplasm
P55495	Y4IL_RHISN		cytoplasm
P55496	Y4IM_RHISN		periplasm
P55497	Y4IN_RHISN		cytoplasm
P55498	Y4IO_RHISN		cytoplasm
P55499	Y4IP_RHISN		inner membrane
P55500	Y4IQ_RHISN		cytoplasm
P55501	Y4JA_RHISN		cytoplasm
P55502	Y4JB_RHISN		cytoplasm
P55503	Y4JC_RHISN		cytoplasm
P55504	Y4JD_RHISN		cytoplasm
P55505	Y4JE_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55507	Y4JG_RHISN		inner membrane
P55508	Y4JH_RHISN		cytoplasm
P55509	Y4JI_RHISN		periplasm
P55510	Y4JJ_RHISN		cytoplasm
P55511	Y4JK_RHISN		cytoplasm
P55512	Y4JL_RHISN		cytoplasm
P55513	Y4JM_RHISN		cytoplasm

P55514	Y4JN_RHISN		cytoplasm
P55515	Y4JO_RHISN		cytoplasm
P55516	Y4JP_RHISN		cytoplasm
P55517	Y4JQ_RHISN		cytoplasm
P55518	Y4JR_RHISN		cytoplasm
P55519	Y4JS_RHISN		cytoplasm
P55520	Y4JT_RHISN		cytoplasm
P55521	Y4KA_RHISN		cytoplasm
P55522	Y4KB_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55524	Y4KD_RHISN		cytoplasm
P55525	Y4KE_RHISN		cytoplasm
P55526	Y4KF_RHISN		cytoplasm
P55527	Y4KG_RHISN		inner membrane
P55528	Y4KH_RHISN		cytoplasm
P55529	Y4KK_RHISN		cytoplasm
P55530	Y4KL_RHISN		cytoplasm
P55531	Y4KM_RHISN		extracellular
P55532	Y4KN_RHISN		cytoplasm
P55533	Y4KO_RHISN		cytoplasm
P55534	Y4KP_RHISN		cytoplasm
P55535	Y4KQ_RHISN		cytoplasm
P55537	Y4KS_RHISN		cytoplasm
P55538	Y4KT_RHISN		cytoplasm
P55539	ISPA_RHISN	cytoplasm (potential).	cytoplasm
P55540	CPXU_RHISN		cytoplasm
P55541	Y4LA_RHISN		cytoplasm
P55542	FE45_RHISN		cytoplasm
P55543	CPXR_RHISN		cytoplasm
P55544	CPXP_RHISN		cytoplasm
P55546	Y4LF_RHISN		inner membrane
P55547	Y4LG_RHISN		cytoplasm
P55548	Y4LH_RHISN		cytoplasm
P55549	Y4LI_RHISN		periplasm
P55550	Y4LJ_RHISN		cytoplasm
P55551	Y4LK_RHISN		cytoplasm
P55552	Y4LL_RHISN		inner membrane
P55554	Y4LN_RHISN		inner membrane
P55555	Y4LO_RHISN		cytoplasm
P55558	Y4LR_RHISN		cytoplasm
P55559	Y4LS_RHISN		cytoplasm
P55560	Y4MA_RHISN		cytoplasm
P55561	Y4MB_RHISN	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P55562	Y4MC_RHISN		cytoplasm
P55563	Y4MD_RHISN		cytoplasm
P55564	Y4ME_RHISN		cytoplasm
P55565	Y4MF_RHISN		cytoplasm
P55566	Y4MG_RHISN		cytoplasm
P55567	Y4MH_RHISN		cytoplasm
P55568	Y4MI_RHISN	periplasm (potential).	periplasm
P55570	Y4MK_RHISN	bacterial cell membrane; peripheral membrane protein (potential).	inner membrane

P55571	Y4ML_RHISN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P55572	Y4MM_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55573	Y4MN_RHISN		cytoplasm
P55574	Y4MO_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55575	Y4MP_RHISN		cytoplasm
P55576	Y4MQ_RHISN		cytoplasm
P55577	Y4NA_RHISN		cytoplasm
P55578	Y4NF_RHISN		cytoplasm
P55579	Y4NG_RHISN		cytoplasm
P55580	Y4NH_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55581	Y4NI_RHISN		inner membrane
P55582	Y4NJ_RHISN		cytoplasm
P55583	Y4NK_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55584	Y4NL_RHISN		cytoplasm
P55585	Y4NM_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55586	Y4OA_RHISN		cytoplasm
P55587	Y4OB_RHISN		cytoplasm
P55588	Y4OC_RHISN		cytoplasm
P55589	Y4OD_RHISN		cytoplasm
P55590	Y4OE_RHISN		cytoplasm
P55597	Y4OL_RHISN		cytoplasm
P55598	Y4OM_RHISN		cytoplasm
P55599	Y4ON_RHISN		cytoplasm
P55601	Y4OP_RHISN	periplasm (potential).	periplasm
P55602	Y4OQ_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P55603	Y4OR_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P55604	Y4OS_RHISN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P55605	Y4OT_RHISN		cytoplasm
P55606	Y4OU_RHISN		inner membrane
P55607	Y4OV_RHISN		cytoplasm
P55608	Y4OW_RHISN		cytoplasm
P55609	Y4OX_RHISN		cytoplasm
P55610	Y4PA_RHISN		cytoplasm
P55611	OTSB_RHISN		cytoplasm
P55612	OTSA_RHISN		cytoplasm
P55613	Y4PD_RHISN		cytoplasm
P55614	Y4PE_RHISN		inner membrane
P55615	Y4PF_RHISN		cytoplasm

P55616	Y4PG_RHISN		inner membrane
P55617	Y4PL_RHISN		cytoplasm
P55618	Y4PM_RHISN		cytoplasm
P55619	SYRM1_RHISN		cytoplasm
P55620	Y4PO_RHISN		cytoplasm
P55623	Y4QB_RHISN		cytoplasm
P55624	Y4QC_RHISN		cytoplasm
P55625	Y4QD_RHISN		cytoplasm
P55626	Y4QE_RHISN		cytoplasm
P55627	Y4QF_RHISN		cytoplasm
P55628	Y4QG_RHISN		cytoplasm
P55629	Y4QH_RHISN		cytoplasm
P55630	Y4QI_RHISN		cytoplasm
P55631	Y4QJ_RHISN		cytoplasm
P55632	Y4QK_RHISN		cytoplasm
P55634	Y4RA_RHISN		cytoplasm
P55635	Y4RB_RHISN		cytoplasm
P55636	Y4RC_RHISN		cytoplasm
P55637	Y4RD_RHISN		cytoplasm
P55638	Y4RE_RHISN		cytoplasm
P55639	Y4RF_RHISN		cytoplasm
P55640	Y4RG_RHISN		cytoplasm
P55641	Y4RH_RHISN		cytoplasm
P55642	Y4RI_RHISN		cytoplasm
P55643	Y4RJ_RHISN		cytoplasm
P55644	Y4RK_RHISN		inner membrane
P55645	Y4RL_RHISN		cytoplasm
P55646	Y4RM_RHISN		cytoplasm
P55647	Y4RN_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55648	Y4RO_RHISN		cytoplasm
P55650	Y4SG_RHISN		cytoplasm
P55651	Y4SH_RHISN		cytoplasm
P55652	Y4SI_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55653	GABD_RHISN		cytoplasm
P55654	Y4SK_RHISN		cytoplasm
P55655	Y4SL_RHISN		cytoplasm
P55656	Y4SO_RHISN		cytoplasm
P55658	Y4TD_RHISN		cytoplasm
P55659	Y4TE_RHISN	periplasm (potential).	periplasm
P55660	Y4TF_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P55661	Y4TG_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P55662	Y4TH_RHISN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P55663	Y4TI_RHISN		extracellular
P55664	Y4TJ_RHISN		cytoplasm
P55665	Y4TK_RHISN		cytoplasm
P55666	Y4TL_RHISN		cytoplasm

P55667	Y4TM_RHISN		cytoplasm
P55668	Y4TN_RHISN		cytoplasm
P55669	Y4TO_RHISN	periplasm (potential).	periplasm
P55670	Y4UD_RHISN		inner membrane
P55671	Y4UG_RHISN		inner membrane
P55673	NIFE_RHISN		cytoplasm
P55674	NIFN_RHISN		periplasm
P55675	NIFX_RHISN		cytoplasm
P55676	Y4VQ_RHISN		cytoplasm
P55677	Y4VR_RHISN		cytoplasm
P55678	FER3_RHISN		periplasm
P55679	Y4WA_RHISN		extracellular
P55680	Y4WB_RHISN		cytoplasm
P55681	Y4WC_RHISN		cytoplasm
P55682	Y4WD_RHISN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P55683	HIS8_RHISN		cytoplasm
P55684	Y4WF_RHISN		cytoplasm
P55685	Y4WG_RHISN		cytoplasm
P55686	Y4WH_RHISN		outer membrane
P55687	Y4WI_RHISN		cytoplasm
P55688	Y4WJ_RHISN		cytoplasm
P55689	NIFW_RHISN		cytoplasm
P55690	NIFS_RHISN		cytoplasm
P55691	Y4WM_RHISN	periplasm (potential).	periplasm
P55693	Y4WO_RHISN		cytoplasm
P55694	Y4WP_RHISN		cytoplasm
P55695	Y4XD_RHISN		cytoplasm
P55696	Y4XE_RHISN		cytoplasm
P55697	Y4XF_RHISN		extracellular
P55698	Y4XQ_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55699	Y4XG_RHISN		inner membrane
P55700	NODD2_RHISN		cytoplasm
P55701	Y4XI_RHISN	cytoplasm (potential).	cytoplasm
P55702	Y4XJ_RHISN		inner membrane
P55703	Y4XK_RHISN	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P55704	Y4XL_RHISN		extracellular
P55705	Y4XM_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P55706	Y4XN_RHISN		outer membrane
P55707	Y4XO_RHISN		cytoplasm
P55708	Y4XP_RHISN		cytoplasm
P55709	Y4YA_RHISN		cytoplasm
P55710	Y4YB_RHISN		cytoplasm
P55711	NOLX_RHISN		inner membrane
P55712	NOLW_RHISN		inner membrane
P55713	NOLB_RHISN		cytoplasm
P55714	NOLT_RHISN	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P55715	NOLU_RHISN		cytoplasm
P55716	NOLV_RHISN		cytoplasm

P55717	Y4YI_RHISN	cytoplasm (potential).	inner membrane
P55718	Y4YJ_RHISN		nucleoid
P55719	Y4YK_RHISN		inner membrane
P55720	Y4YL_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55721	Y4YM_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55722	Y4YN_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55723	Y4YO_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55725	Y4YQ_RHISN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P55726	Y4YR_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P55727	Y4YS_RHISN		inner membrane
P55728	Y4ZA_RHISN		cytoplasm
P55729	Y4ZB_RHISN		cytoplasm
P55730	Y4ZC_RHISN		cytoplasm
P55733	SYRM2_RHISN		cytoplasm
P55746	CGA2_HELPY		cytoplasm
P55757	YOHL_SERMA		nucleoid
P55758	PSIB_RHILP		inner membrane
P55759	YPSI_RHILP		cytoplasm
P55794	YSPU_SHIFL		cytoplasm
P55798	PRP1_ECOLI		cytoplasm
P55799	PRP2_ECOLI		cytoplasm
P55819	SGAA_METEX		cytoplasm
P55827	RS3_ACTAC		cytoplasm
P55829	RS17_ACTAC		cytoplasm
P55834	RL7_HELPY		cytoplasm
P55835	RL2_ACTAC		cytoplasm
P55836	RL4_ACTAC		cytoplasm
P55837	RL16_ACTAC		cytoplasm
P55838	RL22_ACTAC		cytoplasm
P55839	RL23_ACTAC		cytoplasm
P55840	RL29_ACTAC		cytoplasm
P55870	HLY1_AERHY		extracellular
P55880	THIJ_SALTY		cytoplasm
P55881	THIL_SALTY		cytoplasm
P55882	THID_SALTY		cytoplasm
P55883	THIM_SALTY		cytoplasm
P55886	YIGA_SALTY		inner membrane
P55888	XERC_SALTY	cytoplasm (by similarity).	nucleoid
P55912	PRPE_SALTY		cytoplasm
P55914	YJJZ_ECOLI		inner membrane
P55922	RAMA_ENTCL		cytoplasm
P55923	ISTB_BURCE		cytoplasm
P55929	CCPR_NITEU	periplasm (probable).	periplasm

P55969	HPAA_HELPY	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P55970	GRPE_HELPY	cytoplasm (probable).	cytoplasm
P55976	NUSG_HELPY		cytoplasm
P55977	NUSA_HELPY		cytoplasm
P55979	BCP_HELPY		cytoplasm
P55980	CAGA_HELPY		fimbrium
P55982	RIR1_HELPY		cytoplasm
P55983	RIR2_HELPY		cytoplasm
P55984	UPPS_HELPY		inner membrane
P55985	TRUD_HELPY		cytoplasm
P55986	Y1459_HELPY		cytoplasm
P55987	ATPA_HELPY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P55988	ATPB_HELPY		inner membrane
P55990	DHE4_HELPY		cytoplasm
P55991	TOP1_HELPY		cytoplasm
P55992	GYRB_HELPY		cytoplasm
P55993	RPOD_HELPY		cytoplasm
P55994	DNAK_HELPY		cytoplasm
P55996	GCP_HELPY		extracellular
P55997	RNPA_HELPY		extracellular
P56000	SYV_HELPY	cytoplasm (by similarity).	cytoplasm
P56001	RPOA_HELPY		cytoplasm
P56005	SRP54_HELPY		cytoplasm
P56006	SCOA_HELPY		cytoplasm
P56007	SCOB_HELPY		cytoplasm
P56008	RS1_HELPY		cytoplasm
P56009	RS2_HELPY		cytoplasm
P56010	RS3_HELPY		cytoplasm
P56011	RS4_HELPY		cytoplasm
P56013	RS6_HELPY		cytoplasm
P56018	RS11_HELPY		cytoplasm
P56020	RS13_HELPY		cytoplasm
P56021	RS14_HELPY		cytoplasm
P56022	RS15_HELPY		cytoplasm
P56023	RS16_HELPY		cytoplasm
P56026	RS19_HELPY		cytoplasm
P56027	RS20_HELPY		cytoplasm
P56028	RS21_HELPY		cytoplasm
P56029	RL1_HELPY		cytoplasm
P56030	RL2_HELPY		cytoplasm
P56031	RL3_HELPY		cytoplasm
P56032	RL4_HELPY		cytoplasm
P56033	RL5_HELPY		cytoplasm
P56034	RL6_HELPY		cytoplasm
P56035	RL9_HELPY		cytoplasm
P56036	RL10_HELPY		cytoplasm
P56038	RL13_HELPY		cytoplasm
P56039	RL14_HELPY		cytoplasm
P56040	RL15_HELPY		cytoplasm
P56041	RL16_HELPY		cytoplasm
P56042	RL17_HELPY		cytoplasm
P56043	RL18_HELPY		cytoplasm
P56044	RL19_HELPY		cytoplasm

P56045	RL20_HELPY		cytoplasm
P56046	RL21_HELPY		cytoplasm
P56047	RL22_HELPY		cytoplasm
P56049	RL24_HELPY		cytoplasm
P56050	RL27_HELPY		cytoplasm
P56052	RL29_HELPY		cytoplasm
P56059	ERA_HELPY		cytoplasm
P56060	KDSA_HELPY	cytoplasm (by similarity).	cytoplasm
P56061	PANC_HELPY		cytoplasm
P56062	CISY_HELPY		cytoplasm
P56063	IDH_HELPY		cytoplasm
P56064	PRIM_HELPY		cytoplasm
P56065	PAND_HELPY		cytoplasm
P56066	CLPS_HELPY		cytoplasm
P56067	CYSM_HELPY		cytoplasm
P56068	MURI_HELPY		cytoplasm
P56070	PPSA_HELPY		cytoplasm
P56072	SDHL_HELPY		cytoplasm
P56073	AROK_HELPY	cytoplasm (probable).	cytoplasm
P56074	HEM2_HELPY		cytoplasm
P56075	NDK_HELPY	cytoplasm (by similarity).	cytoplasm
P56076	TPIS_HELPY		cytoplasm
P56077	PTH_HELPY	cytoplasm (by similarity).	cytoplasm
P56078	RL25_HELPY		cytoplasm
P56080	Y117_HELPY		cytoplasm
P56081	AROB_HELPY	cytoplasm (probable).	cytoplasm
P56082	ATPG_HELPY		inner membrane
P56084	ATPE_HELPY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P56086	ATPF_HELPY		inner membrane
P56087	ATPL_HELPY		inner membrane
P56088	IMDH_HELPY		cytoplasm
P56096	FTSW_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P56098	RODA_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P56102	AMPM_HELPY		cytoplasm
P56103	KGUA_HELPY	cytoplasm (by similarity).	cytoplasm
P56105	DPO1_HELPY		cytoplasm
P56106	PYRH_HELPY		cytoplasm
P56107	HEMH_HELPY	cytoplasm (by similarity).	extracellular
P56108	TAL_HELPY	cytoplasm (probable).	cytoplasm
P56109	ALF_HELPY		cytoplasm
P56110	G6PD_HELPY		cytoplasm
P56111	EDD_HELPY		cytoplasm
P56112	Y175_HELPY		cytoplasm
P56113	Y169_HELPY		cytoplasm
P56114	GATA_HELPY		cytoplasm
P56115	GSA_HELPY	cytoplasm (potential).	cytoplasm
P56116	HTPG_HELPY	cytoplasm (by similarity).	cytoplasm
P56117	Y190_HELPY	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane

P56118	RNC_HELPY	cytoplasm (by similarity).	cytoplasm
P56119	AROE_HELPY		cytoplasm
P56120	RNH_HELPY	cytoplasm (potential).	extracellular
P56121	RNH2_HELPY	cytoplasm (potential).	extracellular
P56122	AROC_HELPY		cytoplasm
P56123	RNR_HELPY		cytoplasm
P56125	HEM1_HELPY		cytoplasm
P56129	DCDA_HELPY		cytoplasm
P56130	Y285_HELPY		cytoplasm
P56131	Y269_HELPY		cytoplasm
P56132	Y274_HELPY		inner membrane
P56133	PIMT_HELPY	cytoplasm (by similarity).	cytoplasm
P56136	Y258_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P56137	PURA_HELPY	cytoplasm (by similarity).	cytoplasm
P56138	GIDA_HELPY	cytoplasm (by similarity).	cytoplasm
P56139	DXR_HELPY		cytoplasm
P56140	HEM3_HELPY		cytoplasm
P56141	TRPA_HELPY		cytoplasm
P56142	TRPB_HELPY		cytoplasm
P56143	RP54_HELPY		cytoplasm
P56144	TRUA_HELPY		cytoplasm
P56148	RADA_HELPY		cytoplasm
P56149	ASPA_HELPY		cytoplasm
P56153	IPYR_HELPY	cytoplasm (by similarity).	cytoplasm
P56155	PYRF_HELPY		cytoplasm
P56156	CLPP_HELPY	cytoplasm (by similarity).	cytoplasm
P56157	DPO3A_HELPY	cytoplasm (by similarity).	cytoplasm
P56162	PYRE_HELPY		cytoplasm
P56176	Y1363_HELPY		inner membrane
P56184	KPRS_HELPY	cytoplasm (by similarity).	cytoplasm
P56185	Y1430_HELPY		inner membrane
P56186	COAE_AERHY	cytoplasm (by similarity).	cytoplasm
P56188	RPE_HELPY		cytoplasm
P56189	MURA_HELPY	cytoplasm (probable).	inner membrane
P56190	CSTA_HELPY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P56191	DDL_HELPY	cytoplasm (by similarity).	cytoplasm
P56195	DEOB_HELPY	cytoplasm (by similarity).	cytoplasm
P56196	GPMI_HELPY		cytoplasm
P56197	AROA_HELPY	cytoplasm (probable).	cytoplasm
P56200	T2CX_CITFR		extracellular
P56204	Y552_HELPY		cytoplasm
P56214	RECR_HELPY		cytoplasm
P56256	YSAA_ECOLI		periplasm
P56257	YKGL_ECOLI		cytoplasm
P56258	WECF_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P56259	YIFN_ECOLI		cytoplasm
P56262	DLHH_ECOLI		extracellular
P56266	NIFA_KLEOX		cytoplasm
P56267	NIFL_KLEOX		inner membrane
P56372	SELA_DESBA	cytoplasm (by similarity).	cytoplasm

P56397	UNG_HELPY	cytoplasm (by similarity).	extracellular
P56398	RRF_HELPY	cytoplasm (by similarity).	cytoplasm
P56411	KDGL_HELPY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P56414	MOAA_HELPY		cytoplasm
P56415	MOBA_HELPY	cytoplasm (by similarity).	cytoplasm
P56416	FLHB_HELPY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P56417	SYI_HELPY	cytoplasm (by similarity).	cytoplasm
P56418	ACON2_HELPY		cytoplasm
P56419	DEF_HELPY		cytoplasm
P56420	TIG_HELPY		cytoplasm
P56421	MOG_HELPY		cytoplasm
P56422	MOAE_HELPY		cytoplasm
P56428	UVRC_HELPY	cytoplasm (by similarity).	cytoplasm
P56429	DHOM_HELPY		cytoplasm
P56431	TRXB_HELPY	cytoplasm (by similarity).	cytoplasm
P56456	SYI_HELPY	cytoplasm (by similarity).	cytoplasm
P56461	FMT_HELPY		cytoplasm
P56462	GCH1_HELPY		cytoplasm
P56463	DEOD_HELPY		cytoplasm
P56464	ACP_HELPY	cytoplasm (by similarity).	cytoplasm
P56465	PYRC_HELPY		cytoplasm
P56466	RHO_HELPY		inner membrane
P56467	FOLD_HELPY		cytoplasm
P56468	PUR8_HELPY		cytoplasm
P56474	UVRA_HELPY	cytoplasm (by similarity).	cytoplasm
P56507	Y143A_HAEIN		inner membrane
P56585	PYRB_PSEFL		cytoplasm
P56607	YEAC_PSEHA		inner membrane
P56614	YMDF_ECOLI		extracellular
P56621	CNRY_RALME		cytoplasm
P56644	SGAR_HYPME		cytoplasm
P56653	4HBT_PSEUC		cytoplasm
P56737	TRPD_HELPY		cytoplasm
P56744	DAT_ACIBA		cytoplasm
P56834	MRAY_ZYMMO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P56865	LEPA_BORPE		cytoplasm
P56874	UREI_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P56875	RL7_HELPJ		cytoplasm
P56876	TSAA_HELPJ		cytoplasm
P56882	MVIN_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P56883	MUTS_RHIME		cytoplasm
P56884	GLND_RHIME		cytoplasm
P56885	CBBR2_RHIME		cytoplasm
P56886	F16P2_RHIME		cytoplasm
P56887	KPPR2_RHIME		cytoplasm
P56888	ALF2_RHIME		cytoplasm

P56889	RBL12_RHIME		extracellular
P56890	RBS2_RHIME		cytoplasm
P56891	CYSH_RHIME	cytoplasm (by similarity).	cytoplasm
P56892	CYSD_RHIME		cytoplasm
P56893	CYSN_RHIME		cytoplasm
P56894	GREA_RHIME		cytoplasm
P56896	NUOC2_RHIME		inner membrane
P56897	NUOB2_RHIME		cytoplasm
P56898	SSB_RHIME		cytoplasm
P56899	UVRA_RHIME	cytoplasm (by similarity).	cytoplasm
P56900	TKT2_RHIME		extracellular
P56901	LRP_RHIME		cytoplasm
P56902	FABF_RHIME		cytoplasm
P56903	RECF_RHIME	cytoplasm (potential).	cytoplasm
P56904	THID_RHIME		cytoplasm
P56907	NUOD1_RHIME		cytoplasm
P56908	NUOD2_RHIME		cytoplasm
P56909	NUOE1_RHIME		cytoplasm
P56910	NUOE2_RHIME		cytoplasm
P56911	NUON2_RHIME	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P56912	NUOF1_RHIME		periplasm
P56913	NUOF2_RHIME		periplasm
P56914	NUOG2_RHIME		cytoplasm
P56929	TRPB_RHIET		cytoplasm
P56934	LEU2_BUCAI		cytoplasm
P56935	LEUD_BUCAI		cytoplasm
P56936	REPA1_BUCAI		nucleoid
P56938	COXX_RHOSH	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P56939	COXZ_BRAJA		inner membrane
P56940	COXZ_RHOSH		inner membrane
P56949	MDCF_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P56954	MQO_CAMJE		cytoplasm
P56963	FLA1_CAMJE		flagellar
P56964	FLB1_CAMJE		flagellar
P56970	CH10_CAMJE	cytoplasm (by similarity).	cytoplasm
P56977	CYAY_NEIMB		inner membrane
P56978	CYAY_SALTY		inner membrane
P56979	INTM_ECOLI		cytoplasm
P56985	GALE_NEIMB		cytoplasm
P56986	GALE_NEIMC		cytoplasm
P56987	RECA_NEIMA	cytoplasm (by similarity).	cytoplasm
P56988	RECA_NEIMB	cytoplasm (by similarity).	cytoplasm
P56989	FKBP_NEIMA		cytoplasm
P56990	GLYA_NEIMB	cytoplasm (by similarity).	cytoplasm
P56991	AROE_NEIMA		cytoplasm
P56992	AROE_NEIMB		cytoplasm
P56993	CYAY_NEIMA		inner membrane
P56994	HEM1_NEIMB		cytoplasm
P56995	TRPE_NEIMB		cytoplasm
P56996	UVRB_NEIMA	cytoplasm (by similarity).	cytoplasm

P56997	GALE_NEIMA		cytoplasm
P56998	BFRA_NEIMA		cytoplasm
P57001	CYSG_NEIMA		cytoplasm
P57002	PGM_NEIMA		cytoplasm
P57003	TONB_NEIMA	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
P57004	TONB_NEIMB	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
P57005	SODC_NEIMA	periplasm (by similarity).	periplasm
P57006	CH60_NEIMA	cytoplasm (by similarity).	cytoplasm
P57007	SERC_NEIMB	cytoplasm (by similarity).	cytoplasm
P57008	DHAS_NEIMA		cytoplasm
P57009	RPOB_NEIMA		cytoplasm
P57010	FTSY_NEIMA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57011	FTSY_NEIMB	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57012	CTRC_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P57013	CTRD_NEIMA	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P57014	FPG_NEIMA		cytoplasm
P57015	YIFB_SALTY		cytoplasm
P57023	LOLB_NEIMA	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P57024	LOLB_NEIMB	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P57025	H8_NEIMA		periplasm
P57026	H8_NEIMB		periplasm
P57028	PRIM_NEIMA		cytoplasm
P57029	PRIM_NEIMB		cytoplasm
P57030	LOLD_NEIMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P57031	LOLD_NEIMB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P57032	LOLD_XYLFA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P57033	LGTB_NEIMA		cytoplasm
P57034	CTRB_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P57035	PANC_NEIMA		cytoplasm
P57036	PANC_NEIMB		cytoplasm

P57037	LIPM_NEIMA	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (probable).	inner membrane
P57038	LPB1_NEIMA	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (probable).	inner membrane
P57039	FMM1_NEIMA		extracellular
P57040	RMLA_NEIMA		cytoplasm
P57046	TATA_HAEIN	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P57047	TATB_HAEIN	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P57048	TATB_SALTY	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P57051	TATA_VIBCH	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P57061	LOLC_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P57062	LOLC_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P57063	TATB_VIBCH	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P57064	KCY_NEIMA	cytoplasm (by similarity).	cytoplasm
P57065	KCY_NEIMB	cytoplasm (by similarity).	cytoplasm
P57066	LOLD_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P57067	LOLA_NEIMA	periplasm (by similarity).	periplasm
P57068	LOLA_NEIMB	periplasm (by similarity).	periplasm
P57069	LOLA_VIBCH	periplasm (by similarity).	periplasm
P57070	LOLB_VIBCH	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P57072	YHGF_NEIMA		cytoplasm
P57089	NADD_NEIMA		cytoplasm
P57090	NADD_NEIMB		cytoplasm
P57098	GLXK_NEIMA		cytoplasm
P57099	GLXK_NEIMB		cytoplasm
P57109	MAAI_PSEAE		cytoplasm
P57111	RNFH_PSEST		inner membrane
P57112	STHA_PSEAE	cytoplasm (by similarity).	cytoplasm
P57115	HSLV_BUCAI	cytoplasm (by similarity).	cytoplasm
P57116	HSLU_BUCAI	cytoplasm (by similarity).	cytoplasm
P57117	GIDA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57118	ATP6_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57119	ATPL_BUCAI		inner membrane
P57120	ATPF_BUCAI		inner membrane
P57121	ATPD_BUCAI		inner membrane

P57122	ATPA_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57123	ATPG_BUCAI		inner membrane
P57124	ATPB_BUCAI		inner membrane
P57125	ATPE_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57126	GYRB_BUCAI		cytoplasm
P57127	DPO3B_BUCAI	cytoplasm (by similarity).	cytoplasm
P57128	DNAA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57130	RNPA_BUCAI		extracellular
P57131	OXAA_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57132	TRME_BUCAI		cytoplasm
P57133	EFP_BUCAI		cytoplasm
P57134	DNAC_BUCAI		nucleoid
P57135	DNAT_BUCAI		nucleoid
P57136	Y023_BUCAI		cytoplasm
P57137	FTSY_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57138	GLMS_BUCAI	cytoplasm (by similarity).	cytoplasm
P57139	GLMU_BUCAI	cytoplasm (by similarity).	cytoplasm
P57140	Y028_BUCAI		cytoplasm
P57142	METE_BUCAI		cytoplasm
P57143	PUR9_BUCAI		cytoplasm
P57144	DBH_BUCAI		cytoplasm
P57145	RPOC_BUCAI		cytoplasm
P57146	RPOB_BUCAI		cytoplasm
P57147	RL7_BUCAI		cytoplasm
P57148	RL10_BUCAI		cytoplasm
P57149	RL1_BUCAI		cytoplasm
P57150	RL11_BUCAI		cytoplasm
P57151	NUSG_BUCAI		cytoplasm
P57152	SECE_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57153	MURB_BUCAI	cytoplasm (probable).	inner membrane
P57154	METF_BUCAI		cytoplasm
P57155	ARGE_BUCAI	cytoplasm (probable).	extracellular
P57156	ARGC_BUCAI	cytoplasm (probable).	cytoplasm
P57157	ARGB_BUCAI	cytoplasm (probable).	cytoplasm
P57158	ASSY_BUCAI	cytoplasm (probable).	cytoplasm
P57159	ARLY_BUCAI	cytoplasm (probable).	cytoplasm
P57160	Y052_BUCAI		cytoplasm
P57161	SECB_BUCAI	cytoplasm (by similarity).	cytoplasm
P57162	CYSE_BUCAI	cytoplasm (by similarity).	cytoplasm
P57163	RPOD_BUCAI		cytoplasm
P57164	PRIM_BUCAI		cytoplasm
P57165	RS21_BUCAI		cytoplasm
P57166	GCP_BUCAI		extracellular
P57167	RIBB_BUCAI		cytoplasm
P57168	Y060_BUCAI		cytoplasm
P57169	CCA_BUCAI		cytoplasm

P57170	UPPP_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57171	CYSK_BUCAI		cytoplasm
P57172	DNLJ_BUCAI		cytoplasm
P57174	FLIE_BUCAI		flagellar
P57175	FLIF_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57176	FLIG_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57177	FLIH_BUCAI		inner membrane
P57178	FLII_BUCAI	cytoplasm (potential).	inner membrane
P57179	FLIJ_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57180	Y078_BUCAI		cytoplasm
P57181	FLIK_BUCAI		flagellar
P57182	FLIM_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57183	FLIN_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57184	FLIP_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57185	FLIQ_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57186	FLIR_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57187	RL33_BUCAI		cytoplasm
P57188	RL28_BUCAI		cytoplasm
P57189	Y087_BUCAI		inner membrane
P57190	IPYR_BUCAI	cytoplasm (by similarity).	cytoplasm
P57191	PMBA_BUCAI	cytoplasm (potential).	cytoplasm
P57192	Y091_BUCAI		cytoplasm
P57193	FABB_BUCAI	cytoplasm (by similarity).	cytoplasm
P57194	TAL_BUCAI	cytoplasm (probable).	cytoplasm
P57195	TKT_BUCAI		extracellular
P57196	DAPE_BUCAI		cytoplasm
P57197	DAPA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57198	AROC_BUCAI		cytoplasm
P57199	Y098_BUCAI		inner membrane
P57200	HIS1_BUCAI	cytoplasm (by similarity).	cytoplasm
P57201	HISX_BUCAI		cytoplasm
P57202	HIS8_BUCAI		cytoplasm
P57203	HIS7_BUCAI	cytoplasm (by similarity).	cytoplasm
P57204	HIS5_BUCAI	cytoplasm (by similarity).	cytoplasm
P57205	HIS4_BUCAI	cytoplasm (by similarity).	cytoplasm
P57206	HIS6_BUCAI	cytoplasm (by similarity).	cytoplasm
P57207	HIS2_BUCAI	cytoplasm (by similarity).	cytoplasm
P57208	6PGD_BUCAI		cytoplasm

P57209	DCD_BUCAI		extracellular
P57211	TILS_BUCAI	cytoplasm (probable).	cytoplasm
P57212	RISA_BUCAI		cytoplasm
P57213	RNFA_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57215	RNFC_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P57216	RNFD_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57217	RNFG_BUCAI		inner membrane
P57218	RNFE_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57219	END3_BUCAI		cytoplasm
P57220	PRIA_BUCAI		cytoplasm
P57221	SYI_BUCAI	cytoplasm (by similarity).	cytoplasm
P57222	Y122_BUCAI		inner membrane
P57223	Y123_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57224	AROH_BUCAI		cytoplasm
P57227	RL35_BUCAI		cytoplasm
P57228	RL20_BUCAI		cytoplasm
P57231	IHFA_BUCAI		cytoplasm
P57232	QUEA_BUCAI	cytoplasm (potential).	cytoplasm
P57233	TGT_BUCAI		cytoplasm
P57234	Y134_BUCAI		inner membrane
P57237	END4_BUCAI		extracellular
P57238	RL25_BUCAI		cytoplasm
P57239	Y139_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57240	SURA_BUCAI		periplasm
P57241	KSGA_BUCAI		cytoplasm
P57242	APAH_BUCAI		extracellular
P57243	DYR_BUCAI		cytoplasm
P57244	CARB_BUCAI		cytoplasm
P57245	CARA_BUCAI		cytoplasm
P57246	DAPB_BUCAI	cytoplasm (by similarity).	cytoplasm
P57247	ISPH_BUCAI		cytoplasm
P57248	LSPA_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	extracellular
P57249	SYI_BUCAI	cytoplasm (by similarity).	cytoplasm
P57250	RIBF_BUCAI		cytoplasm
P57251	RS20_BUCAI		cytoplasm
P57253	NUOB_BUCAI		cytoplasm
P57254	NUOCD_BUCAI		inner membrane
P57255	NUOE_BUCAI		cytoplasm
P57256	NUOF_BUCAI		periplasm
P57257	NUOG_BUCAI		cytoplasm
P57259	NUOI_BUCAI		cytoplasm

P57260	NUOJ_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57261	NUOK_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57262	NUOL_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57263	NUOM_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57265	FOLC_BUCAI		cytoplasm
P57266	KPRS_BUCAI	cytoplasm (by similarity).	cytoplasm
P57267	ISPE_BUCAI		cytoplasm
P57269	HEMK_BUCAI		cytoplasm
P57270	Y173_BUCAI		extracellular
P57271	NADE_BUCAI		cytoplasm
P57272	ACKA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57273	PTA_BUCAI	cytoplasm (potential).	inner membrane
P57274	Y177_BUCAI		cytoplasm
P57275	RIR2_BUCAI		cytoplasm
P57276	RIR1_BUCAI		cytoplasm
P57277	GYRA_BUCAI		cytoplasm
P57278	Y181_BUCAI		extracellular
P57279	TSAA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57280	UNG_BUCAI	cytoplasm (by similarity).	extracellular
P57281	GRPE2_BUCAI	cytoplasm (probable).	cytoplasm
P57282	PPNK_BUCAI	cytoplasm (by similarity).	cytoplasm
P57283	SMPA_BUCAI	bacterial cell membrane; lipid-anchor (by similarity).	outer membrane
P57284	Y187_BUCAI		cytoplasm
P57285	RNT_BUCAI		cytoplasm
P57286	SODM_BUCAI		periplasm
P57287	PTH_BUCAI	cytoplasm (by similarity).	cytoplasm
P57288	ENGD_BUCAI		cytoplasm
P57289	THRC_BUCAI		cytoplasm
P57290	AKH_BUCAI		cytoplasm
P57291	HPRT_BUCAI	cytoplasm (by similarity).	cytoplasm
P57292	PANC_BUCAI		cytoplasm
P57293	PANB_BUCAI		cytoplasm
P57294	DKSA_BUCAI		cytoplasm
P57295	TRUA_BUCAI		cytoplasm
P57296	PBPB_BUCAI		inner membrane
P57297	SECA_BUCAI	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	inner membrane
P57298	MUTT_BUCAI		cytoplasm
P57299	COAE_BUCAI	cytoplasm (by similarity).	cytoplasm
P57300	GUAC_BUCAI		cytoplasm
P57301	ODP1_BUCAI		cytoplasm
P57302	ODP2_BUCAI		inner membrane
P57303	DLDH_BUCAI	cytoplasm (by similarity).	cytoplasm
P57304	SPED_BUCAI		cytoplasm
P57305	SPEE_BUCAI		cytoplasm

P57306	MTNN_BUCAI		cytoplasm
P57307	Y211_BUCAI		inner membrane
P57309	FTSA_BUCAI		inner membrane
P57310	MURC_BUCAI	cytoplasm (probable).	cytoplasm
P57311	MURG_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57312	FTSW_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57313	MURD_BUCAI	cytoplasm (by similarity).	cytoplasm
P57314	MRAY_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57315	MURF_BUCAI	cytoplasm (probable).	cytoplasm
P57316	MURE_BUCAI	cytoplasm (probable).	cytoplasm
P57317	FTSI_BUCAI	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P57318	FTSL_BUCAI		inner membrane
P57319	MRAW_BUCAI	cytoplasm (by similarity).	cytoplasm
P57320	ILVH_BUCAI		cytoplasm
P57321	ILVI_BUCAI		cytoplasm
P57322	DEGP_BUCAI		cytoplasm
P57323	DAPD_BUCAI	cytoplasm (by similarity).	cytoplasm
P57324	AMPM_BUCAI		cytoplasm
P57325	RS2_BUCAI		cytoplasm
P57327	PYRH_BUCAI		cytoplasm
P57328	RRF_BUCAI	cytoplasm (by similarity).	cytoplasm
P57329	DXR_BUCAI		cytoplasm
P57330	UPPS_BUCAI		inner membrane
P57331	Y237_BUCAI		outer membrane
P57332	DPO3A_BUCAI	cytoplasm (by similarity).	cytoplasm
P57334	FLHB_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57335	FLHA_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57336	GLO2_BUCAI		cytoplasm
P57337	DPO3E_BUCAI		cytoplasm
P57338	GMHA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57339	XGPT_BUCAI	cytoplasm (by similarity).	cytoplasm
P57340	GRPE1_BUCAI	cytoplasm (probable).	cytoplasm
P57341	Y253_BUCAI		inner membrane
P57342	SSRP_BUCAI	cytoplasm (potential).	cytoplasm
P57343	Y255_BUCAI		cytoplasm
P57344	ACPS_BUCAI	cytoplasm (by similarity).	cytoplasm
P57345	ERA_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P57346	RNC_BUCAI	cytoplasm (by similarity).	cytoplasm
P57347	LEP_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57348	LEPA_BUCAI		cytoplasm

P57349	TRMU_BUCAI	cytoplasm (by similarity).	cytoplasm
P57350	Y262_BUCAI		inner membrane
P57351	PUR8_BUCAI		cytoplasm
P57352	MLTE_BUCAI		extracellular
P57353	FABI_BUCAI		cytoplasm
P57354	RNB_BUCAI	cytoplasm (by similarity).	cytoplasm
P57355	Y267_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57356	LIPB_BUCAI	cytoplasm (by similarity).	cytoplasm
P57357	LIPA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57358	PYRF_BUCAI		cytoplasm
P57359	GCH2_BUCAI		extracellular
P57360	HNS_BUCAI		cytoplasm
P57361	CLS_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57362	Y274_BUCAI		cytoplasm
P57363	ISPZ_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57364	Y276_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57365	TRPA_BUCAI		cytoplasm
P57366	TRPC_BUCAI		cytoplasm
P57367	TRPD_BUCAI		cytoplasm
P57368	Y281_BUCAI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P57369	RLUB_BUCAI		cytoplasm
P57370	SOHB_BUCAI		extracellular
P57371	TOP1_BUCAI		cytoplasm
P57372	SUHB_BUCAI		extracellular
P57373	Y286_BUCAI		cytoplasm
P57374	ISPG_BUCAI		cytoplasm
P57377	BIOD_BUCAI		cytoplasm
P57378	BIOB_BUCAI		cytoplasm
P57379	BIOA_BUCAI		cytoplasm
P57380	6PGL_BUCAI		cytoplasm
P57381	MFD_BUCAI		cytoplasm
P57382	LOLC_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57383	LOLD_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57385	FLAV_BUCAI		cytoplasm
P57386	PHR_BUCAI		cytoplasm
P57387	Y301_BUCAI		cytoplasm
P57388	ODO1_BUCAI		cytoplasm
P57389	ODO2_BUCAI		inner membrane
P57390	GPMA_BUCAI		cytoplasm
P57391	K6PF_BUCAI	cytoplasm (by similarity).	cytoplasm

P57392	GLPF_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57393	TPIS_BUCAI		cytoplasm
P57394	IHFB_BUCAI		cytoplasm
P57395	RS1_BUCAI		cytoplasm
P57396	AROA_BUCAI	cytoplasm (probable).	cytoplasm
P57397	SERC_BUCAI	cytoplasm (by similarity).	cytoplasm
P57401	SYD_BUCAI	cytoplasm (by similarity).	cytoplasm
P57402	ZNUB_BUCAI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P57403	ZNUC_BUCAI		inner membrane
P57404	KPYK_BUCAI		cytoplasm
P57405	G6PD_BUCAI		cytoplasm
P57406	HTPX_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	extracellular
P57407	CSPC_BUCAI	cytoplasm (by similarity).	cytoplasm
P57408	Y323_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57409	Y324_BUCAI		cytoplasm
P57410	MINE_BUCAI		inner membrane
P57411	MIND_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57412	MINC_BUCAI		inner membrane
P57413	RSMC_BUCAI		cytoplasm
P57415	MVIN_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57416	PYRC_BUCAI		cytoplasm
P57417	FLGN_BUCAI		flagellar
P57418	FLGA_BUCAI	periplasm (probable).	flagellar
P57419	FLGB_BUCAI		flagellar
P57420	FLGC_BUCAI		flagellar
P57421	FLGD_BUCAI		inner membrane
P57422	FLGE_BUCAI		flagellar
P57423	FLGF_BUCAI		flagellar
P57424	FLGG_BUCAI		flagellar
P57425	FLGH_BUCAI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P57426	FLGI_BUCAI		periplasm
P57427	FLGJ_BUCAI		inner membrane
P57428	FLGK_BUCAI		flagellar
P57429	RNE_BUCAI	cytoplasm (by similarity).	cytoplasm
P57430	RLUC_BUCAI		cytoplasm
P57431	RL32_BUCAI		cytoplasm
P57432	FABG_BUCAI		cytoplasm
P57433	ACP_BUCAI	cytoplasm (by similarity).	cytoplasm
P57434	KTHY_BUCAI		cytoplasm
P57435	HOLB_BUCAI		cytoplasm
P57436	Y355_BUCAI		extracellular

P57437	PTGCB_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57438	YHIT_BUCAI		cytoplasm
P57439	Y358_BUCAI		cytoplasm
P57440	PORL_BUCAI		outer membrane
P57442	PNCB_BUCAI	cytoplasm (by similarity).	cytoplasm
P57443	PYRD_BUCAI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57444	Y363_BUCAI		cytoplasm
P57445	UUP_BUCAI		inner membrane
P57446	Y365_BUCAI		inner membrane
P57447	SYV_BUCAI	cytoplasm (by similarity).	cytoplasm
P57448	AMPA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57449	OTC_BUCAI	cytoplasm (probable).	cytoplasm
P57450	PYRB_BUCAI		cytoplasm
P57451	PYRI_BUCAI		cytoplasm
P57452	Y371_BUCAI		cytoplasm
P57453	DEAD_BUCAI	cytoplasm (probable).	cytoplasm
P57454	PNP_BUCAI	cytoplasm (by similarity).	cytoplasm
P57455	RS15_BUCAI		cytoplasm
P57456	TRUB_BUCAI		cytoplasm
P57457	RBFA_BUCAI	cytoplasm (potential).	cytoplasm
P57459	NUSA_BUCAI		cytoplasm
P57460	SECG_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57461	GLMM_BUCAI		cytoplasm
P57462	FTSH_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57463	RRMJ_BUCAI	cytoplasm (by similarity).	cytoplasm
P57464	GREA_BUCAI		cytoplasm
P57465	Y385_BUCAI		cytoplasm
P57466	MURA_BUCAI	cytoplasm (probable).	inner membrane
P57467	RL21_BUCAI		cytoplasm
P57468	RL27_BUCAI		cytoplasm
P57469	Y389_BUCAI		cytoplasm
P57470	RS9_BUCAI		cytoplasm
P57471	RL13_BUCAI		cytoplasm
P57473	SRP54_BUCAI		cytoplasm
P57474	RS16_BUCAI		cytoplasm
P57475	RIMM_BUCAI	cytoplasm (potential).	cytoplasm
P57476	TRMD_BUCAI	cytoplasm (potential).	cytoplasm
P57477	RL19_BUCAI		cytoplasm
P57478	TLDD_BUCAI		cytoplasm
P57479	AROQ_BUCAI		cytoplasm
P57480	FIS_BUCAI		cytoplasm
P57481	RLUD_BUCAI		cytoplasm
P57482	Y402_BUCAI		cytoplasm
P57484	CSRA_BUCAI		cytoplasm
P57485	GSH1_BUCAI		cytoplasm
P57487	END1_BUCAI		extracellular
P57488	Y410_BUCAI		cytoplasm
P57489	RPIA_BUCAI		cytoplasm

P57491	PYRG_BUCAI		cytoplasm
P57492	ENO_BUCAI	cytoplasm (by similarity).	extracellular
P57493	NLPD_BUCAI		extracellular
P57494	ISPF_BUCAI		extracellular
P57495	ISPD_BUCAI		cytoplasm
P57497	CYSC_BUCAI		cytoplasm
P57498	CYSN_BUCAI		cytoplasm
P57499	CYSD_BUCAI		cytoplasm
P57500	CYSG_BUCAI		cytoplasm
P57501	CYSH_BUCAI	cytoplasm (by similarity).	cytoplasm
P57502	CYSI_BUCAI		cytoplasm
P57503	CYSJ_BUCAI		cytoplasm
P57504	MUTS_BUCAI		cytoplasm
P57505	DSBA_BUCAI		periplasm
P57506	EX53_BUCAI		cytoplasm
P57507	ENGB_BUCAI		cytoplasm
P57508	TYP_A_BUCAI		cytoplasm
P57509	KGUA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57510	Y435_BUCAI		cytoplasm
P57513	DCDA_BUCAI		cytoplasm
P57514	LGT_BUCAI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P57516	Y441_BUCAI		cytoplasm
P57517	Y442_BUCAI		inner membrane
P57518	CORC_BUCAI		inner membrane
P57520	HOLA_BUCAI		cytoplasm
P57521	NADD_BUCAI		cytoplasm
P57522	TUSA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57523	DHAS_BUCAI		cytoplasm
P57524	Y449_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57526	ALF_BUCAI		cytoplasm
P57527	Y452_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57528	EX5C_BUCAI		cytoplasm
P57529	EX5B_BUCAI		cytoplasm
P57530	EX5A_BUCAI		cytoplasm
P57532	THIL_BUCAI		cytoplasm
P57533	RIBD1_BUCAI		cytoplasm
P57534	RIBD2_BUCAI		cytoplasm
P57535	NUSB_BUCAI		cytoplasm
P57536	DXS_BUCAI		cytoplasm
P57537	ISPA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57538	Y466_BUCAI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P57539	TUSE_BUCAI	cytoplasm (by similarity).	cytoplasm
P57540	CYOE_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57541	CYOD_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane

P57542	CYOC_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57543	CYOB_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57544	CYOA_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57545	Y473_BUCAI		cytoplasm
P57546	TIG_BUCAI		cytoplasm
P57547	CLPP_BUCAI	cytoplasm (by similarity).	cytoplasm
P57548	CLPX_BUCAI		cytoplasm
P57550	PPID_BUCAI	bacterial cell membrane; single-pass type ii membrane protein (potential).	outer membrane
P57551	MDLA_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57552	MDLB_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57553	DPO3X_BUCAI		cytoplasm
P57554	Y482_BUCAI		inner membrane
P57555	HTPG_BUCAI	cytoplasm (by similarity).	cytoplasm
P57557	FOLD_BUCAI		cytoplasm
P57559	Y488_BUCAI		inner membrane
P57561	Y494_BUCAI		inner membrane
P57562	SMG_BUCAI		cytoplasm
P57563	DEF_BUCAI		cytoplasm
P57564	FMT_BUCAI		cytoplasm
P57565	RL17_BUCAI		cytoplasm
P57566	RPOA_BUCAI		cytoplasm
P57567	RS4_BUCAI		cytoplasm
P57568	RS11_BUCAI		cytoplasm
P57569	RS13_BUCAI		cytoplasm
P57572	RL15_BUCAI		cytoplasm
P57573	RL30_BUCAI		cytoplasm
P57574	RS5_BUCAI		cytoplasm
P57575	RL18_BUCAI		cytoplasm
P57576	RL6_BUCAI		cytoplasm
P57577	RS8_BUCAI		cytoplasm
P57578	RS14_BUCAI		cytoplasm
P57579	RL5_BUCAI		cytoplasm
P57580	RL24_BUCAI		cytoplasm
P57581	RL14_BUCAI		cytoplasm
P57582	RS17_BUCAI		cytoplasm
P57583	RL29_BUCAI		cytoplasm
P57584	RL16_BUCAI		cytoplasm
P57585	RS3_BUCAI		cytoplasm
P57586	RL22_BUCAI		cytoplasm
P57587	RS19_BUCAI		cytoplasm
P57588	RL2_BUCAI		cytoplasm
P57589	RL23_BUCAI		cytoplasm
P57590	RL4_BUCAI		cytoplasm
P57591	RL3_BUCAI		cytoplasm

P57592	RS10_BUCAI		cytoplasm
P57594	RS7_BUCAI		cytoplasm
P57595	RS12_BUCAI		cytoplasm
P57596	TUSB_BUCAI	cytoplasm (by similarity).	cytoplasm
P57597	TUSC_BUCAI	cytoplasm (by similarity).	cytoplasm
P57598	TUSD_BUCAI	cytoplasm (by similarity).	cytoplasm
P57599	FKBA_BUCAI		cytoplasm
P57600	ARGD_BUCAI	cytoplasm (probable).	cytoplasm
P57601	TSGA_BUCAI	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P57603	RPE_BUCAI		cytoplasm
P57604	AROB_BUCAI	cytoplasm (probable).	cytoplasm
P57605	AROK_BUCAI	cytoplasm (probable).	cytoplasm
P57606	DEOD_BUCAI		cytoplasm
P57607	DEOB_BUCAI	cytoplasm (by similarity).	cytoplasm
P57609	Y544_BUCAI		inner membrane
P57610	SSB_BUCAI		cytoplasm
P57611	DNAB_BUCAI		cytoplasm
P57612	GSHB_BUCAI		cytoplasm
P57613	RUVX_BUCAI	cytoplasm (potential).	cytoplasm
P57614	Y549_BUCAI		cytoplasm
P57615	Y550_BUCAI		cytoplasm
P57616	TRMB_BUCAI		cytoplasm
P57617	MUTY_BUCAI		cytoplasm
P57618	FETP_BUCAI		cytoplasm
P57619	MURI_BUCAI		cytoplasm
P57620	EX1_BUCAI		cytoplasm
P57621	Y556_BUCAI		cytoplasm
P57622	PYRE_BUCAI		cytoplasm
P57623	DUT_BUCAI		extracellular
P57624	CYSQ_BUCAI	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	cytoplasm
P57625	RL9_BUCAI		cytoplasm
P57626	RS18_BUCAI		cytoplasm
P57627	RS6_BUCAI		cytoplasm
P57628	RNR_BUCAI		cytoplasm
P57629	PURA_BUCAI	cytoplasm (by similarity).	cytoplasm
P57630	HFLC_BUCAI		extracellular
P57631	HFLK_BUCAI		inner membrane
P57632	MIAA_BUCAI		cytoplasm
P57633	MUTL_BUCAI		cytoplasm
P57634	MTLD_BUCAI		cytoplasm
P57635	PTM3C_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57636	G6PI_BUCAI	cytoplasm (by similarity).	cytoplasm
P57637	ORN_BUCAI	cytoplasm (potential).	cytoplasm
P57638	AMIB_BUCAI		extracellular
P57639	RL31_BUCAI		cytoplasm
P57640	IBP_BUCAI		cytoplasm
P57641	FENR_BUCAI		cytoplasm
P57642	SYK3_BUCAI		cytoplasm
P57643	COAD_BUCAI	cytoplasm (by similarity).	cytoplasm

P57644	Y584_BUCAI		cytoplasm
P57645	Y585_BUCAI		cytoplasm
P57646	Y586_BUCAI		inner membrane
P57647	PIT_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57648	Y588_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57649	DAPF_BUCAI	cytoplasm (by similarity).	cytoplasm
P57650	CYAY_BUCAI		inner membrane
P57651	HEM3_BUCAI		cytoplasm
P57652	RHO_BUCAI		inner membrane
P57653	THIO_BUCAI		cytoplasm
P57654	REP_BUCAI		cytoplasm
P57655	ILVC_BUCAI		cytoplasm
P57656	ILVD_BUCAI		cytoplasm
P57657	ISCS_BUCAI		cytoplasm
P57658	NIFU_BUCAI		cytoplasm
P57659	HSCB_BUCAI		inner membrane
P57660	HSCA_BUCAI		cytoplasm
P57661	FER_BUCAI		cytoplasm
P57662	ENGA_BUCAI		cytoplasm
P57663	Y608_BUCAI		inner membrane
P57665	ORN_PSEAE	cytoplasm (potential).	cytoplasm
P57668	TPX_PSEAE		cytoplasm
P57683	ATKA_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P57686	ATKC_PSEAE	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
P57696	DHPS_NEIMC		cytoplasm
P57698	ATKB_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P57701	DSBB2_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P57702	CYC1_PSEAE		cytoplasm
P57703	METE_PSEAE		cytoplasm
P57707	ISPD_PSEAE		cytoplasm
P57708	ISPF_PSEAE		extracellular
P57714	METX_PSEAE	cytoplasm (by similarity).	cytoplasm
P57767	PAAD_THAAR		cytoplasm
P57778	HEMH_PSEFL	cytoplasm (by similarity).	extracellular
P57779	HEMH_ZYMMO	cytoplasm (by similarity).	extracellular
P57794	NIFS_ACEDI		cytoplasm
P57798	Y585A_HELPY		cytoplasm
P57800	TATB_PASMU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P57801	GREA_PASMU		cytoplasm
P57802	GREB_PASMU		cytoplasm
P57803	ISCS_PASMU		cytoplasm
P57805	RNC_PASMU	cytoplasm (by similarity).	cytoplasm

P57806	LEPA_PASMU		cytoplasm
P57807	UNG_PASMU	cytoplasm (by similarity).	extracellular
P57808	TYSY_PASMU	cytoplasm (by similarity).	cytoplasm
P57809	NUDH_PASMU		cytoplasm
P57810	RDGC_PASMU	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
P57812	ENGA_PASMU		cytoplasm
P57813	RNH_PASMU	cytoplasm (potential).	extracellular
P57814	GMHA_PASMU	cytoplasm (by similarity).	cytoplasm
P57815	MURE_PASMU	cytoplasm (probable).	cytoplasm
P57816	MRAY_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P57817	MURG_PASMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P57818	MURC_PASMU	cytoplasm (probable).	cytoplasm
P57819	DDL_PASMU	cytoplasm (by similarity).	cytoplasm
P57820	PTH_PASMU	cytoplasm (by similarity).	cytoplasm
P57821	MURA_PASMU	cytoplasm (probable).	inner membrane
P57823	AMPA_PASMU	cytoplasm (by similarity).	cytoplasm
P57824	SYK3_PASMU		cytoplasm
P57825	Y205_PASMU		inner membrane
P57826	RECR_PASMU		cytoplasm
P57827	SSRP_PASMU	cytoplasm (potential).	cytoplasm
P57828	PUR9_PASMU		cytoplasm
P57829	PUR2_PASMU		cytoplasm
P57830	GLYA_PASMU	cytoplasm (by similarity).	cytoplasm
P57831	TGT_PASMU		cytoplasm
P57832	QUEA_PASMU	cytoplasm (potential).	cytoplasm
P57833	ISPE_PASMU		cytoplasm
P57834	LOLB_PASMU	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P57835	LOLA_PASMU	periplasm (by similarity).	periplasm
P57837	KAD_PASMU	cytoplasm (by similarity).	cytoplasm
P57839	ISPZ_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P57840	AROC_PASMU		cytoplasm
P57842	FDHD_PASMU	cytoplasm (potential).	cytoplasm
P57843	METE_PASMU		cytoplasm
P57844	UVRB_PASMU	cytoplasm (by similarity).	cytoplasm
P57845	MINC_PASMU		inner membrane
P57846	HTPX_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P57848	DXS_PASMU		cytoplasm
P57849	THII_PASMU	cytoplasm (potential).	cytoplasm
P57850	ARGR_PASMU	cytoplasm (probable).	cytoplasm
P57853	KDSA_PASMU	cytoplasm (by similarity).	cytoplasm
P57854	TRPA_PASMU		cytoplasm
P57855	TRPC_PASMU		cytoplasm
P57856	TRPD_PASMU		cytoplasm

P57858	PYRD_PASMU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P57861	TRUA_PASMU		cytoplasm
P57862	Y647_PASMU		inner membrane
P57863	GCH2_PASMU		extracellular
P57864	Y681_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57865	GCH1_PASMU		cytoplasm
P57866	ACKA_PASMU	cytoplasm (by similarity).	cytoplasm
P57867	DAPB_PASMU	cytoplasm (by similarity).	cytoplasm
P57868	NUSB_PASMU		cytoplasm
P57869	RISB_PASMU		cytoplasm
P57870	DNAK_PASMU		cytoplasm
P57871	SFSA_PASMU		cytoplasm
P57872	RBFA_PASMU	cytoplasm (potential).	cytoplasm
P57874	HEMH_PASMU	cytoplasm (by similarity).	extracellular
P57875	KCY_PASMU	cytoplasm (by similarity).	cytoplasm
P57876	OTC_PASMU	cytoplasm (probable).	cytoplasm
P57877	ASSY_PASMU	cytoplasm (probable).	cytoplasm
P57878	PUR7_PASMU		cytoplasm
P57880	TPX_PASMU		cytoplasm
P57881	SERC_PASMU	cytoplasm (by similarity).	cytoplasm
P57882	UVRC_PASMU	cytoplasm (by similarity).	cytoplasm
P57883	KDSB_PASMU	cytoplasm (potential).	cytoplasm
P57884	METX_PASMU	cytoplasm (by similarity).	cytoplasm
P57885	ORN_PASMU	cytoplasm (potential).	cytoplasm
P57886	MUTL_PASMU		cytoplasm
P57887	MURI_PASMU		cytoplasm
P57888	KGUA_PASMU	cytoplasm (by similarity).	cytoplasm
P57889	PURA_PASMU	cytoplasm (by similarity).	cytoplasm
P57891	DCD_PASMU		extracellular
P57892	RUVB_PASMU		cytoplasm
P57893	RUVA_PASMU		cytoplasm
P57894	RUVC_PASMU		cytoplasm
P57896	RNT_PASMU		cytoplasm
P57899	GAL1_PASMU	cytoplasm (potential).	cytoplasm
P57900	Y1045_PASMU		inner membrane
P57901	LUXS_PASMU		extracellular
P57902	FIS_PASMU		cytoplasm
P57903	AROQ_PASMU		cytoplasm
P57907	ARGC_PASMU	cytoplasm (probable).	cytoplasm
P57908	ARGB_PASMU	cytoplasm (probable).	cytoplasm
P57909	ARLY_PASMU	cytoplasm (probable).	cytoplasm
P57910	FPG_PASMU		cytoplasm
P57911	Y1146_PASMU		inner membrane
P57912	RL33_PASMU		cytoplasm
P57913	RADC_PASMU		cytoplasm
P57914	DUT_PASMU		extracellular
P57915	RNPA_PASMU		extracellular
P57916	RS18_PASMU		cytoplasm
P57917	LEXA_PASMU		cytoplasm
P57918	IPYR_PASMU	cytoplasm (by similarity).	cytoplasm
P57919	HIS1_PASMU	cytoplasm (by similarity).	cytoplasm
P57920	HIS7_PASMU	cytoplasm (by similarity).	cytoplasm

P57921	HIS5_PASMU	cytoplasm (by similarity).	cytoplasm
P57922	APAH_PASMU		extracellular
P57924	AROB_PASMU	cytoplasm (probable).	cytoplasm
P57925	AROK_PASMU	cytoplasm (probable).	cytoplasm
P57926	KITH_PASMU	cytoplasm (potential).	cytoplasm
P57927	TKT1_PASMU		extracellular
P57928	LYXK_PASMU		cytoplasm
P57929	Y1258_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P57930	THIE_PASMU		cytoplasm
P57931	THIM_PASMU		cytoplasm
P57932	AROE_PASMU		cytoplasm
P57934	CSRA_PASMU		cytoplasm
P57935	RIMM_PASMU	cytoplasm (potential).	cytoplasm
P57936	TPIS_PASMU		cytoplasm
P57937	DEOC_PASMU	cytoplasm (by similarity).	cytoplasm
P57940	RIBB_PASMU		cytoplasm
P57941	RPOA_PASMU		cytoplasm
P57943	CYAY_PASMU		inner membrane
P57944	GLPK_PASMU		cytoplasm
P57945	GIDA_PASMU	cytoplasm (by similarity).	cytoplasm
P57946	GIDB_PASMU		cytoplasm
P57947	ENGB_PASMU		cytoplasm
P57948	DEF_PASMU		cytoplasm
P57949	FMT_PASMU		cytoplasm
P57950	MSCL_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P57951	Y1576_PASMU		cytoplasm
P57952	MURB_PASMU	cytoplasm (probable).	inner membrane
P57953	ISPD_PASMU		cytoplasm
P57954	ISPF_PASMU		extracellular
P57955	SURE_PASMU	cytoplasm (potential).	periplasm
P57957	ILVD_PASMU		cytoplasm
P57958	TKT2_PASMU		extracellular
P57959	LSPA_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P57960	ISPH_PASMU		cytoplasm
P57961	RPIA_PASMU		cytoplasm
P57962	DAPF_PASMU	cytoplasm (by similarity).	cytoplasm
P57963	GLMS_PASMU	cytoplasm (by similarity).	cytoplasm
P57964	DCUP_PASMU	cytoplasm (probable).	cytoplasm
P57965	NUDC_PASMU		cytoplasm
P57967	COAA_PASMU	cytoplasm (probable).	cytoplasm
P57968	HSLU_PASMU	cytoplasm (by similarity).	cytoplasm
P57969	HSLV_PASMU	cytoplasm (by similarity).	cytoplasm
P57970	UBIA_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P57971	MOBA_PASMU	cytoplasm (by similarity).	cytoplasm
P57972	MUTS_PASMU		cytoplasm
P57973	PGK_PASMU	cytoplasm (potential).	cytoplasm
P57974	RECO_PASMU		cytoplasm
P57975	ENO_PASMU	cytoplasm (by similarity).	extracellular

P57976	PLSX_PASMU		cytoplasm
P57977	LIPB_PASMU	cytoplasm (by similarity).	cytoplasm
P57978	LIPA_PASMU	cytoplasm (potential).	cytoplasm
P57979	UVRA_PASMU	cytoplasm (by similarity).	cytoplasm
P57980	TIG_PASMU		cytoplasm
P57981	CLPX_PASMU		cytoplasm
P57982	RS2_PASMU		cytoplasm
P57984	RRF_PASMU	cytoplasm (by similarity).	cytoplasm
P57985	DXR_PASMU		cytoplasm
P57986	RNH2_PASMU	cytoplasm (potential).	extracellular
P57987	ISPG_PASMU		cytoplasm
P57989	CSD_PASMU		cytoplasm
P57998	INSB4_ECOLI		cytoplasm
P58000	TKRA_ENTAG		cytoplasm
P58033	YPJJ_ECOLI		cytoplasm
P58034	YMGF_ECOLI		inner membrane
P58035	SGCB_ECOLI	cytoplasm (probable).	cytoplasm
P58036	YJBS_ECOLI		cytoplasm
P58037	YRHD_ECOLI		cytoplasm
P58040	YF7A_PSEAE		inner membrane
P58041	RZOD_ECOLI	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P58042	RZOR_ECOLI	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P58054	YBEM_ECO57		cytoplasm
P58056	PPNK_CAUCR	cytoplasm (by similarity).	cytoplasm
P58057	PPNK_ECO57	cytoplasm (by similarity).	cytoplasm
P58060	RL10_CAUCR		cytoplasm
P58063	TRUB_CAUCR		cytoplasm
P58065	YNIA_ECO57		cytoplasm
P58066	RPOZ_CAUCR		cytoplasm
P58067	NAGZ_ECO57	cytoplasm (by similarity).	extracellular
P58068	YHDW_ECO57	periplasm (probable).	periplasm
P58070	ERA_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P58071	ERA_CAUCR		cytoplasm
P58072	G3P3_ECO57	cytoplasm (potential).	cytoplasm
P58074	TRMU_CAUCR	cytoplasm (by similarity).	cytoplasm
P58079	HUTI_CAUCR	cytoplasm (potential).	cytoplasm
P58082	HUTH_CAUCR	cytoplasm (potential).	cytoplasm
P58088	TRMB_CAUCR		cytoplasm
P58091	Y874_CAUCR		inner membrane
P58092	Y509_ECO57		inner membrane
P58093	YV6A_VIBCH		inner membrane
P58094	YCIX_ECOLI		inner membrane
P58095	YPJI_ECOLI		cytoplasm
P58096	YSDA_ECOLI		inner membrane
P58097	HSLO_CAUCR	cytoplasm (by similarity).	cytoplasm
P58100	COAE_CAUCR	cytoplasm (by similarity).	cytoplasm
P58103	COAD_CAUCR	cytoplasm (by similarity).	cytoplasm
P58112	Y481_CAUCR		cytoplasm
P58113	Y1473_CAUCR		periplasm
P58114	Y3178_CAUCR		cytoplasm
P58115	YHAK_ECO57		cytoplasm
P58116	YHHW_ECO57		cytoplasm

P58122	RS16_CAUCR		cytoplasm
P58123	RS16_PASMU		cytoplasm
P58127	RTCA_ECO57	cytoplasm (by similarity).	cytoplasm
P58130	PLSB_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P58141	GPDA_CAUCR	cytoplasm (probable).	cytoplasm
P58142	GPDA_RHILO	cytoplasm (probable).	cytoplasm
P58152	MINE_NEIGO		inner membrane
P58153	MINE_RHILO		inner membrane
P58161	CITG_ECO57		cytoplasm
P58162	DSBD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58163	NORM_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58164	MDTK_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58167	RL19_CAUCR		cytoplasm
P58168	RL19_RHILO		cytoplasm
P58184	LPXK_CAUCR		cytoplasm
P58185	LPXK_RHILO		cytoplasm
P58186	LPXK_RICMO		cytoplasm
P58187	LPXK_RICRI		cytoplasm
P58188	LPXK_RICTY		cytoplasm
P58194	UBID_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P58204	FLGI_RHOSH	periplasm (by similarity).	periplasm
P58205	FLGH_RHOSH	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
P58207	DAPA_RHILO	cytoplasm (by similarity).	cytoplasm
P58208	MIOC_ECO57		cytoplasm
P58209	DAPB_ECO57	cytoplasm (by similarity).	cytoplasm
P58210	DAPB1_RHILO	cytoplasm (by similarity).	cytoplasm
P58211	DAPB2_RHILO	cytoplasm (by similarity).	cytoplasm
P58213	YTFI_ECO57		cytoplasm
P58216	YHBX_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58217	DINI2_ECO57		cytoplasm
P58218	DINI3_ECO57		cytoplasm
P58219	YJGN_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58220	TKRA_ECO57	cytoplasm (probable).	cytoplasm
P58221	MOBA_ECO57	cytoplasm (by similarity).	cytoplasm
P58224	YIAT_ECO57	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P58226	SELA_RHIME	cytoplasm (by similarity).	cytoplasm
P58228	DCEA_ECO57		cytoplasm

P58229	GADC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58230	GADX_ECO57		cytoplasm
P58231	FLGJ_ECO57	periplasm (by similarity).	extracellular
P58233	CEST_ECO57	cytoplasm (by similarity).	cytoplasm
P58237	HIS5_ECO57	cytoplasm (by similarity).	cytoplasm
P58244	CLCA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P58245	YJIP_ECO57		cytoplasm
P58256	ILVC_ECO57		cytoplasm
P58259	NRDR_RHIME		cytoplasm
P58263	THIG_ECO57	cytoplasm (by similarity).	cytoplasm
P58264	THIG_RHIME	cytoplasm (by similarity).	cytoplasm
P58277	CLPP1_RHIME	cytoplasm (by similarity).	cytoplasm
P58278	CLPP2_RHIME	cytoplasm (by similarity).	cytoplasm
P58292	YDDE_ECO57		cytoplasm
P58316	YAAW_ECO57		inner membrane
P58318	YFJF_ECO57		inner membrane
P58319	LTAE_ECO57		cytoplasm
P58320	DSBG_ECO57	periplasm (by similarity).	periplasm
P58324	RNFC_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	cytoplasm
P58325	RNFD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P58326	DAPB1_RHIME	cytoplasm (by similarity).	cytoplasm
P58329	ATZD_PSESD		extracellular
P58330	FLAD1_RHIME		flagellar
P58332	CBBR1_RHIME		cytoplasm
P58333	TKT1_RHIME		extracellular
P58336	ALF1_RHIME		cytoplasm
P58337	YY1A_XYLFA	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
P58338	OCD1_RHIME		cytoplasm
P58339	OCD2_RHIME		cytoplasm
P58341	ATCU1_RHIME	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P58342	ATCU2_RHIME	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P58344	RNFE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58345	RNFG_ECO57	bacterial cell inner membrane (potential).	inner membrane
P58346	SYRB3_RHIME		cytoplasm
P58347	KPPR1_RHIME		cytoplasm
P58348	RBL11_RHIME		extracellular
P58349	RBS1_RHIME		cytoplasm
P58350	AATB1_RHIME	cytoplasm (by similarity).	cytoplasm

P58356	TORS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58357	TORR_ECO57	cytoplasm (probable).	cytoplasm
P58358	TORT_ECO57	periplasm (by similarity).	periplasm
P58359	TORC_ECO57	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P58360	TORA_ECO57	periplasm (by similarity).	periplasm
P58361	TORY_ECO57	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P58362	TORZ_ECO57	periplasm (by similarity).	periplasm
P58363	ARCB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58364	TORD_ECO57	cytoplasm (by similarity).	cytoplasm
P58377	LNT1_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58378	HMRR1_RHIME	cytoplasm (probable).	cytoplasm
P58379	HMRR2_RHIME	cytoplasm (probable).	cytoplasm
P58380	FAB11_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P58381	FAB12_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P58382	ACKA1_RHIME	cytoplasm (by similarity).	cytoplasm
P58384	TUS_ECO57	cytoplasm (by similarity).	cytoplasm
P58388	THTM_ECO57	cytoplasm (probable).	cytoplasm
P58393	GLGA1_RHIME		cytoplasm
P58394	GLGA2_RHIME		cytoplasm
P58398	MLRA_SALTY		cytoplasm
P58402	EVGS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P58403	YFDX_ECO57		cytoplasm
P58408	ALLD_ECO57	cytoplasm (probable).	cytoplasm
P58409	YBIC_ECO57	cytoplasm (potential).	cytoplasm
P58411	MACA_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P58422	GPH_ECO57		cytoplasm
P58423	FSAA_ECO57	cytoplasm (by similarity).	cytoplasm
P58424	FSAB_ECO57	cytoplasm (probable).	cytoplasm
P58429	SFSA_AGRT5		cytoplasm
P58431	SFSA_YERPE		cytoplasm
P58470	RL31_RHIME		cytoplasm
P58471	RL31_YERPE		cytoplasm
P58472	RL31B_YERPE		cytoplasm
P58473	PEPB_ECO57	cytoplasm (by similarity).	cytoplasm
P58474	PEPB_HAEIN	cytoplasm (by similarity).	cytoplasm
P58475	PEPB_YERPE	cytoplasm (by similarity).	cytoplasm
P58477	HTPG_RHIME	cytoplasm (by similarity).	cytoplasm
P58478	HTPG_RICCN	cytoplasm (by similarity).	cytoplasm

P58479	HTPG_SALTI	cytoplasm (by similarity).	cytoplasm
P58480	HTPG_SALTY	cytoplasm (by similarity).	cytoplasm
P58482	HTPG_YERPE	cytoplasm (by similarity).	cytoplasm
P58483	Y2857_YERPE	periplasm (potential).	periplasm
P58484	MENC_ECO57		extracellular
P58485	MENC_SALTI		extracellular
P58486	MENC_SALTY		extracellular
P58487	MENC_YERPE		extracellular
P58492	ZIPA_YERPE	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
P58494	PEPE_SALTI	cytoplasm (by similarity).	cytoplasm
P58496	MODD_ECO57		cytoplasm
P58497	MODE_YERPE	cytoplasm (probable).	cytoplasm
P58506	RHAA_ECO57	cytoplasm (by similarity).	cytoplasm
P58507	RHAA_YERPE	cytoplasm (by similarity).	cytoplasm
P58508	ICIA_SALTI		cytoplasm
P58509	ICIA_SALTY		cytoplasm
P58510	ICIA_YERPE		cytoplasm
P58523	MTOX_ECO57		cytoplasm
P58524	MTOX_SALTI		cytoplasm
P58525	MTOX_SALTY		cytoplasm
P58526	MTOX_YERPE		cytoplasm
P58528	GATC_MORCA		cytoplasm
P58529	SOTB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58530	SOTB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58531	SOTB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58532	DTD_SALTI	cytoplasm (probable).	cytoplasm
P58533	DTD_SALTY	cytoplasm (probable).	cytoplasm
P58534	DTD_YERPE	cytoplasm (probable).	cytoplasm
P58535	SGCX_SALTY		extracellular
P58536	SGCX_SALTI		extracellular
P58537	SUHB_SALTY		cytoplasm
P58538	ARAA_ECO57		cytoplasm
P58539	ARAA_SALTI		cytoplasm
P58540	ARAA_YERPE		cytoplasm
P58541	ARAB_ECO57		cytoplasm
P58542	ARAB_SALTI		cytoplasm
P58543	ARAB_YERPE		cytoplasm
P58551	KDKA_PASPI	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
P58578	GSHB_ECO57		cytoplasm
P58579	GSHB_RALSO		cytoplasm
P58580	GSHB_SALTY		cytoplasm
P58581	GSHB_SALTI		cytoplasm
P58582	GSHB_YERPE		cytoplasm
P58584	SOLI1_RALSO		cytoplasm
P58585	YBHK_ECO57		cytoplasm
P58586	YBHK_SALTY		cytoplasm

P58587	YBHK_SALTI		cytoplasm
P58589	Y1158_YERPE		cytoplasm
P58590	SOLR1_RALSO		cytoplasm
P58591	EPSD1_RALSO		cytoplasm
P58592	IPT1_RALSO		cytoplasm
P58594	EPSE1_RALSO	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P58595	EPSF1_RALSO	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P58596	EPSP1_RALSO		extracellular
P58597	EPSA1_RALSO	bacterial cell outer membrane; multi-pass membrane protein (potential).	outer membrane
P58598	PGLR1_RALSO	secreted protein (by similarity).	extracellular
P58599	GUN1_RALSO	bacterial cell membrane; lipid-anchor (probable).	extracellular
P58600	EPSC1_RALSO	cytoplasm (by similarity).	extracellular
P58601	PME1_RALSO		extracellular
P58603	OMPT_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P58607	SKP_YERPE	periplasm (by similarity).	periplasm
P58611	LPXD_YERPE		cytoplasm
P58614	GLGS_SALTY		cytoplasm
P58615	GLGS_SALTI		cytoplasm
P58617	GLK_RALSO	cytoplasm (by similarity).	cytoplasm
P58618	GLK_SALTI	cytoplasm (by similarity).	cytoplasm
P58619	GLK_YERPE	cytoplasm (by similarity).	cytoplasm
P58620	MAMA_ECO57		cytoplasm
P58626	YCEF_ECO57	cytoplasm (potential).	inner membrane
P58627	YCEF_SALTY	cytoplasm (potential).	inner membrane
P58628	YCEF_SALTI	cytoplasm (potential).	inner membrane
P58629	YHDE_ECO57	cytoplasm (potential).	inner membrane
P58630	YHDE_SALTY	cytoplasm (potential).	inner membrane
P58631	YHDE_SALTI	cytoplasm (potential).	inner membrane
P58633	Y1046_RALSO	cytoplasm (potential).	inner membrane
P58634	Y2196_RALSO	cytoplasm (potential).	inner membrane
P58635	Y1593_YERPE	cytoplasm (potential).	inner membrane
P58636	Y3668_YERPE	cytoplasm (potential).	inner membrane
P58638	PYRF_AGRT5		cytoplasm
P58640	PYRF_ECO57		cytoplasm
P58642	PYRF_RALSO		cytoplasm
P58643	PYRF_SALTI		cytoplasm
P58644	PYRF_YERPE		cytoplasm
P58646	ATPE_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P58647	ATPE_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P58650	OADG3_SALTY	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane

P58651	OADG1_SALTY	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
P58652	COBC_SALTI		cytoplasm
P58653	ORGA_SALTY		extracellular
P58654	ORGB_SALTY		extracellular
P58655	PSTA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58662	RCSC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58663	RCSB_SALTY		cytoplasm
P58664	YG EK_ECO57		cytoplasm
P58665	ISPG_AGRT5		cytoplasm
P58669	ISPG_RALSO		cytoplasm
P58670	ISPG_SALTI		cytoplasm
P58671	ISPG_SALTY		cytoplasm
P58672	ISPG_YERPE		cytoplasm
P58673	ISPH_AGRT5		cytoplasm
P58677	ISPH_RALSO		cytoplasm
P58678	ISPH_SALTI		cytoplasm
P58679	ISPH_SALTY		cytoplasm
P58680	ISPH_YERPE		cytoplasm
P58683	APHA_SALTY	periplasm (potential).	periplasm
P58687	AROD_SALTY		cytoplasm
P58688	FOLD_SALTY		cytoplasm
P58699	TRPR_SALTI	cytoplasm (by similarity).	cytoplasm
P58700	TRPR_YERPE	cytoplasm (by similarity).	cytoplasm
P58704	CYNS_ECO57		cytoplasm
P58708	DALD_RALSO		cytoplasm
P58709	DALD_YERPE		cytoplasm
P58713	PDXA_ECO57	cytoplasm (probable).	cytoplasm
P58714	PDXA_RALSO	cytoplasm (probable).	cytoplasm
P58715	PDXA1_SALTI	cytoplasm (probable).	cytoplasm
P58716	PDXA2_SALTI	cytoplasm (probable).	cytoplasm
P58717	PDXA1_SALTY	cytoplasm (probable).	cytoplasm
P58718	PDXA2_SALTY	cytoplasm (probable).	cytoplasm
P58719	PDXA_YERPE	cytoplasm (probable).	cytoplasm
P58720	IGAA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58721	IGAA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58722	IGAA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58723	CH60_BURTH	cytoplasm (by similarity).	cytoplasm
P58726	CSPJ_SALTI	cytoplasm (by similarity).	cytoplasm
P58729	AROC_SALTY		cytoplasm
P58734	DCTA_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58736	ALR1_AGRT5		cytoplasm
P58737	ALR2_AGRT5		cytoplasm

P58739	DADA_AGRT5		cytoplasm
P58740	UPPP1_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58741	UPPP2_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58744	DKGA_SALT1	cytoplasm (by similarity).	cytoplasm
P58745	MRAW_AGRT5		cytoplasm
P58747	MOBA_NEIMB	cytoplasm (by similarity).	cytoplasm
P58758	IPTZ_AGRT5		cytoplasm
P58759	Y2500_AGRT5		cytoplasm
P58760	Y2499_AGRT5		cytoplasm
P58767	TAG1_ECO27	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P58768	MRAZ_AGRT5		inner membrane
P58769	Y1920_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58770	Y1919_AGRT5		cytoplasm
P58777	CAT_KLESP		cytoplasm
P58788	HIS5_AGRT5	cytoplasm (by similarity).	cytoplasm
P58790	HIS4_AGRT5	cytoplasm (by similarity).	cytoplasm
P58792	Y2496_AGRT5		cytoplasm
P58795	WRBA_AGRT5		cytoplasm
P58799	HIS6_AGRT5	cytoplasm (by similarity).	cytoplasm
P58881	HIS7_XANAC	cytoplasm (by similarity).	cytoplasm
P58882	HIS7_XANCP	cytoplasm (by similarity).	cytoplasm
P58891	HIS8_XANAC		cytoplasm
P58892	HIS8_XANCP		cytoplasm
P58895	CARA_XANAC		cytoplasm
P58896	CARA_XANCP		cytoplasm
P58898	LEU1_BUCPS		cytoplasm
P58900	LEU1_XANAC		cytoplasm
P58901	LEU1_XANCP		cytoplasm
P58902	ACPD1_XANAC		extracellular
P58903	ACPD2_XANAC		extracellular
P58904	ACPD_XANCP		extracellular
P58931	BCSA_PSEFL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P58932	BCSA_XANAC	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P58933	BCSB_XANAC	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
P58934	BCSB_PSEFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P58935	GUN_XANAC	secreted protein (by similarity).	extracellular
P58937	BCSC_PSEFL		outer membrane
P58938	BCSC_XANAC		outer membrane
P58942	CARB_XANAC		cytoplasm

P58943	CARB_XANCP		cytoplasm
P58945	LEU2_BUCPS		cytoplasm
P58948	LEU2_XANAC		cytoplasm
P58949	LEU2_XANCP		cytoplasm
P58975	LPXH_XANAC	cytoplasm (by similarity).	cytoplasm
P58976	LPXH_XANCP	cytoplasm (by similarity).	cytoplasm
P58987	HUTU_XANAC	cytoplasm (by similarity).	cytoplasm
P58988	HUTU_XANCP	cytoplasm (by similarity).	cytoplasm
P58994	GSH1_BUCAP		cytoplasm
P58997	BFPA_ECO11		fimbrium
P59001	UPP_XANAC		cytoplasm
P59002	PYRH_BUCAP		cytoplasm
P59005	PYRH_PARZE		cytoplasm
P59008	PYRH_XANAC		cytoplasm
P59009	PYRH_XANCP		cytoplasm
P59019	LEUD_BUCPS		cytoplasm
P59027	LEU3_BUCPS	cytoplasm (by similarity).	cytoplasm
P59030	RS8_BUCAP		cytoplasm
P59039	PYRG_BUCAP		cytoplasm
P59058	END4_BUCAP		extracellular
P59059	RS7_BUCAP		cytoplasm
P59061	RS7_RHOCA		cytoplasm
P59064	RS7_TREPR		cytoplasm
P59085	ARGE_BUCAP	cytoplasm (probable).	extracellular
P59086	ARGD_BUCAP	cytoplasm (probable).	cytoplasm
P59098	ARGB_BUCAP	cytoplasm (probable).	cytoplasm
P59099	ARGA_BUCAP	cytoplasm (probable).	cytoplasm
P59100	OTC_BUCAP	cytoplasm (probable).	cytoplasm
P59107	RNB_SHIFL	cytoplasm (by similarity).	cytoplasm
P59124	RS5_SHEON		cytoplasm
P59128	RS4_BRUSU		cytoplasm
P59131	RS4_SHEON		cytoplasm
P59132	RS4_SHIFL		cytoplasm
P59160	GPMA_BRUSU		cytoplasm
P59166	RS12_SHEON		cytoplasm
P59175	GPMI_SHEON		cytoplasm
P59176	GPMI_SHIFL		cytoplasm
P59180	RS3_BRUSU		cytoplasm
P59183	RS3_SHEON		cytoplasm
P59184	RS3_SHIFL		cytoplasm
P59188	FDHE_SHEON	cytoplasm (by similarity).	cytoplasm
P59189	SSUD_SHIFL		cytoplasm
P59192	Y2497_SHEON		inner membrane
P59193	YFDT_SHIFL		cytoplasm
P59194	YFCJ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59195	YHHS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59196	YFCD_SHIFL		cytoplasm
P59197	MIAA_SHIFL		cytoplasm
P59198	MSBB1_SHIFL	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
P59207	RECX_SHEON	cytoplasm (potential).	cytoplasm

P59236	Y3081_SHEON		inner membrane
P59237	ALR2_ECOL6		cytoplasm
P59238	LDCA_ECOL6	cytoplasm (by similarity).	cytoplasm
P59243	MLTE_ECOL6		extracellular
P59244	TDCD_ECOL6		cytoplasm
P59245	MODD_ECOL6		cytoplasm
P59246	Y3601_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59251	Y1290_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59252	YGIH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59257	Y626_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59269	TSGA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59273	Y2251_SHEON		inner membrane
P59274	Y2072_VIBVU		inner membrane
P59284	ALLA_BRUSU		extracellular
P59285	ALLA_PSEPK		extracellular
P59286	MPGP_ECOL6	cytoplasm (by similarity).	cytoplasm
P59292	ARGA_SHEON	cytoplasm (probable).	cytoplasm
P59293	ARGA_SHIFL	cytoplasm (probable).	cytoplasm
P59294	ARGA_VIBVU	cytoplasm (probable).	cytoplasm
P59296	ARGB_BRUSU	cytoplasm (probable).	cytoplasm
P59298	ARGB_ECOL6	cytoplasm (probable).	cytoplasm
P59300	ARGB_PSEPK	cytoplasm (probable).	cytoplasm
P59301	ARGB_SHEON	cytoplasm (probable).	cytoplasm
P59302	ARGB_SHIFL	cytoplasm (probable).	cytoplasm
P59304	ARGB_VIBVU	cytoplasm (probable).	cytoplasm
P59306	ARGC_ECOL6	cytoplasm (probable).	cytoplasm
P59308	ARGC2_PSEPK	cytoplasm (probable).	cytoplasm
P59309	ARGC_SHEON	cytoplasm (probable).	cytoplasm
P59310	ARGC_SHIFL	cytoplasm (probable).	cytoplasm
P59313	ARGC_VIBVU	cytoplasm (probable).	cytoplasm
P59314	ARGC_BRUSU	cytoplasm (probable).	cytoplasm
P59317	ARGD_ECOL6	cytoplasm (probable).	cytoplasm
P59318	ARGD_MYXXA	cytoplasm (probable).	cytoplasm
P59319	ARGD_PSEPK	cytoplasm (probable).	cytoplasm
P59320	ARGD_SHEON	cytoplasm (probable).	cytoplasm
P59321	ARGD_SHIFL	cytoplasm (probable).	cytoplasm
P59323	ARGD_VIBVU	cytoplasm (probable).	cytoplasm
P59324	ARGD_YERPE	cytoplasm (probable).	cytoplasm
P59329	DCYD_ECOL6		cytoplasm
P59330	DCYD_SHIFL		cytoplasm
P59331	HCHA_ECOL6	cytoplasm (by similarity).	cytoplasm
P59332	HCHA_SHIFL	cytoplasm (by similarity).	cytoplasm
P59333	CAIT_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P59334	CAIT_PROSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59335	CAIT_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59337	DHAA_BRAJA		cytoplasm
P59338	DCUR_ECOL6	cytoplasm (probable).	cytoplasm
P59339	DCUR_SHIFL	cytoplasm (probable).	cytoplasm
P59340	DCUS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59341	DCUS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59342	BARA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59343	DSBB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59344	DSBB1_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59345	DSBB2_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59346	DSBB_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59347	DSBB_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59348	Y6755_BRAJA		inner membrane
P59349	YEAH_ECOL6		inner membrane
P59351	Y396_PSEPK		inner membrane
P59352	Y2883_SHEON		inner membrane
P59353	Y2091_VIBVU		inner membrane
P59361	CHO_SHIFL		cytoplasm
P59362	GLMS_BRAJA	cytoplasm (by similarity).	cytoplasm
P59363	ISPZ_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59364	ISPZ_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59365	ISPZ_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59366	ISPZ_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59369	HDFR_ECOL6		cytoplasm
P59370	RS11_BRAJA		cytoplasm
P59374	RS11_PSEPK		cytoplasm
P59375	RS11_SHEON		cytoplasm

P59380	RS11_WIGBR		cytoplasm
P59385	HYIN1_BRAJA		extracellular
P59392	SELD_PSEPK		cytoplasm
P59393	SELD_SHIFL		cytoplasm
P59394	CAIB_SHIFL	cytoplasm (by similarity).	cytoplasm
P59395	CAID_SHIFL		cytoplasm
P59396	YACL_SHIFL		inner membrane
P59397	HISX_BRAJA		cytoplasm
P59398	HISX_BUCBP		cytoplasm
P59400	HISX_PSEPK		cytoplasm
P59401	HISX_SHIFL		cytoplasm
P59402	NORR_SHIFL		cytoplasm
P59403	NORW_SHIFL	cytoplasm (by similarity).	cytoplasm
P59404	NORV_ECOL6	cytoplasm (probable).	cytoplasm
P59405	NORV_SHIFL	cytoplasm (probable).	cytoplasm
P59408	DUT_BUCBP		extracellular
P59409	TDH_SHIFL	cytoplasm (by similarity).	cytoplasm
P59410	TDH_VIBPA	cytoplasm (by similarity).	cytoplasm
P59411	RBFA_BUCBP	cytoplasm (potential).	cytoplasm
P59412	ASSY_BUCBP	cytoplasm (probable).	cytoplasm
P59413	RNPA_BUCBP		extracellular
P59414	AROE_BUCBP		cytoplasm
P59415	TRPD_BUCBP		cytoplasm
P59416	AROA_BUCBP	cytoplasm (probable).	cytoplasm
P59417	IPYR_BUCBP	cytoplasm (by similarity).	cytoplasm
P59418	MURC_BUCBP	cytoplasm (probable).	cytoplasm
P59419	MURE_BUCBP	cytoplasm (probable).	cytoplasm
P59424	MURG_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P59425	RIMM_BUCBP	cytoplasm (potential).	cytoplasm
P59426	ILVD_BUCBP		cytoplasm
P59427	TYSY_BUCBP	cytoplasm (by similarity).	cytoplasm
P59428	PURA_BUCBP	cytoplasm (by similarity).	cytoplasm
P59429	KAD_BUCBP	cytoplasm (by similarity).	cytoplasm
P59432	GLYA_BUCBP	cytoplasm (by similarity).	cytoplasm
P59434	RNH_BUCBP	cytoplasm (potential).	extracellular
P59435	DDL_BUCBP	cytoplasm (by similarity).	cytoplasm
P59436	MRAY_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P59437	NANK_SHIFL		cytoplasm
P59439	CHO_ECOL6		cytoplasm
P59440	ISPZ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59442	NANE_SHIFL		cytoplasm
P59443	GUAC_BUCBP		cytoplasm
P59444	INSC_SHIFL		cytoplasm
P59445	INSE_SHIFL		cytoplasm
P59447	RS3_BUCBP		cytoplasm
P59448	CARB_BUCBP		cytoplasm
P59449	ACP_BUCBP	cytoplasm (by similarity).	cytoplasm
P59450	MURB_BUCBP	cytoplasm (probable).	inner membrane
P59452	NUSB_BUCBP		cytoplasm
P59453	HIS1_BUCBP	cytoplasm (by similarity).	cytoplasm

P59454	HIS7_BUCBP	cytoplasm (by similarity).	cytoplasm
P59455	RPOA_BUCBP		cytoplasm
P59457	TRPA_BUCBP		cytoplasm
P59458	TRPB_BUCBP		cytoplasm
P59459	TRPC_BUCBP		cytoplasm
P59460	TRMU_BUCBP	cytoplasm (by similarity).	cytoplasm
P59461	PGK_BUCBP	cytoplasm (potential).	cytoplasm
P59462	TPIS_BUCBP		cytoplasm
P59463	CYAY_BUCBP		inner membrane
P59464	DCD_BUCBP		cytoplasm
P59465	UNG_BUCBP	cytoplasm (by similarity).	extracellular
P59471	Y015_BUCBP		inner membrane
P59474	DAPB_BUCBP	cytoplasm (by similarity).	cytoplasm
P59475	ACPS_BUCBP	cytoplasm (by similarity).	cytoplasm
P59476	RNC_BUCBP	cytoplasm (by similarity).	cytoplasm
P59477	DPO4_ECOL6	cytoplasm (probable).	cytoplasm
P59478	Y393_BUCBP		inner membrane
P59479	LEXA2_PSEPK		cytoplasm
P59484	RS11_BUCBP		cytoplasm
P59485	GIDA_BUCBP	cytoplasm (by similarity).	cytoplasm
P59487	AROB_BUCBP	cytoplasm (probable).	cytoplasm
P59488	AROK_BUCBP	cytoplasm (probable).	cytoplasm
P59489	GREA_BUCBP		cytoplasm
P59490	PTH_BUCBP	cytoplasm (by similarity).	cytoplasm
P59491	RS4_BUCBP		cytoplasm
P59492	SERC_BUCBP	cytoplasm (by similarity).	cytoplasm
P59493	DEF_BUCBP		cytoplasm
P59494	LAMB1_VIBPA	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P59495	GSHB_BUCBP		cytoplasm
P59496	END4_BUCBP		extracellular
P59497	RNT_BUCBP		cytoplasm
P59498	CSRA_BUCBP		cytoplasm
P59499	GLMS_BUCBP	cytoplasm (by similarity).	cytoplasm
P59500	KTHY_BUCBP		cytoplasm
P59501	HIS5_BUCBP	cytoplasm (by similarity).	cytoplasm
P59502	RS18_BUCBP		cytoplasm
P59503	MINC_BUCBP		inner membrane
P59507	MIAA_BUCBP		cytoplasm
P59508	TRUA_BUCBP		cytoplasm
P59512	KPRS_BUCBP	cytoplasm (by similarity).	cytoplasm
P59513	RL27_BUCBP		cytoplasm
P59514	RS9_BUCBP		cytoplasm
P59515	LEU3_BUCBP	cytoplasm (by similarity).	cytoplasm
P59516	LEUD_BUCBP		cytoplasm
P59518	TRMD_BUCBP	cytoplasm (potential).	cytoplasm
P59519	LEU2_BUCBP		cytoplasm
P59520	HIS4_BUCBP	cytoplasm (by similarity).	cytoplasm
P59521	HIS6_BUCBP	cytoplasm (by similarity).	cytoplasm
P59522	MRAW_BUCBP	cytoplasm (by similarity).	cytoplasm
P59523	UPPP_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
P59524	KSGA_BUCBP		cytoplasm
P59525	CH10_BUCBP	cytoplasm (by similarity).	cytoplasm

P59526	CH60_BUCBP	cytoplasm (by similarity).	cytoplasm
P59527	THIO_BUCBP		cytoplasm
P59555	GCH2_BUCBP		extracellular
P59556	RIBB_BUCBP		cytoplasm
P59557	FMT_BUCBP		cytoplasm
P59558	RL22_BUCBP		cytoplasm
P59559	HTPX_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	extracellular
P59560	Y1352_PSEPK		inner membrane
P59561	YAJQ_SHIFL		inner membrane
P59562	Y1617_VIBPA		inner membrane
P59563	K6PF_BUCBP	cytoplasm (by similarity).	cytoplasm
P59564	BIOD_BUCBP		cytoplasm
P59565	DNAK_BUCBP		cytoplasm
P59566	ENO_BUCBP	cytoplasm (by similarity).	extracellular
P59567	DNAA_BUCBP	cytoplasm (by similarity).	cytoplasm
P59568	KHSE_BUCBP	cytoplasm (potential).	cytoplasm
P59569	TRME_BUCBP		cytoplasm
P59570	OMP1_VIBPA	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P59571	COPA2_PSESM	periplasm (by similarity).	periplasm
P59572	COPB2_PSESM	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P59574	MURI_BUCBP		cytoplasm
P59575	PYRE_BUCBP		cytoplasm
P59576	CARA_BUCBP		cytoplasm
P59577	PYRG_BUCBP		cytoplasm
P59578	PYRH_BUCBP		cytoplasm
P59579	RS8_BUCBP		cytoplasm
P59580	KCY_BUCBP	cytoplasm (by similarity).	cytoplasm
P59581	HSLV_BUCBP	cytoplasm (by similarity).	cytoplasm
P59582	DAPF_BUCBP	cytoplasm (by similarity).	cytoplasm
P59585	YFEO_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P59586	RS11_XYLFT		cytoplasm
P59588	PLDB_ECOL6	bacterial cell inner membrane (by similarity).	inner membrane
P59590	FIMC_ECOL6	periplasm (by similarity).	periplasm
P59600	ARGE_SHIFL	cytoplasm (probable).	extracellular
P59601	ARGE_VIBPA		extracellular
P59604	ASSY_PSEPK	cytoplasm (probable).	cytoplasm
P59605	ASSY_VIBPA	cytoplasm (probable).	cytoplasm
P59606	ASSY_XYLFT	cytoplasm (probable).	cytoplasm
P59607	ASSY_BRAJA	cytoplasm (probable).	cytoplasm
P59608	ASSY_BURCE	cytoplasm (probable).	cytoplasm
P59609	ASSY_SHIFL	cytoplasm (probable).	cytoplasm
P59610	ARGJ_BRAJA	cytoplasm (probable).	cytoplasm
P59612	ARGJ_PSEPK	cytoplasm (probable).	cytoplasm
P59613	ARLY_BRAJA	cytoplasm (probable).	cytoplasm
P59614	ARLY_BUCBP	cytoplasm (probable).	cytoplasm
P59615	ARLY_BURCE	cytoplasm (probable).	cytoplasm
P59618	ARLY_PSEPK	cytoplasm (probable).	cytoplasm

P59619	ARLY_SHIFL	cytoplasm (probable).	cytoplasm
P59620	ARLY_VIBPA	cytoplasm (probable).	cytoplasm
P59621	ARLY_XYLFT	cytoplasm (probable).	cytoplasm
P59638	CLCB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59639	CLCA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P59640	YEHT_SHIFL		cytoplasm
P59650	DHSL_COXBU		cytoplasm
P59663	MASZ_SHIFL	cytoplasm (by similarity).	cytoplasm
P59673	FIXA_SHIFL		cytoplasm
P59674	FIXB_ECOL6		cytoplasm
P59675	FIXB_SHIFL		cytoplasm
P59688	NAGB_SHIFL		extracellular
P59696	T200_SALTY		cytoplasm
P59697	T200_SALTI		cytoplasm
P59699	NANT_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59714	LEUD_SHIFL		cytoplasm
P59715	YBHO_SHIFL	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P59720	TRMB_NITEU		cytoplasm
P59725	PQQD_ENTIT		cytoplasm
P59727	Y2125_NITEU		cytoplasm
P59728	YBIX_SHIFL		cytoplasm
P59733	Y357_NITEU		inner membrane
P59735	CYSQ_SHIFL	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	cytoplasm
P59736	AROH_SHIFL		cytoplasm
P59737	AROP_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59738	MODC_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P59739	CYAA_SHIFL	cytoplasm (by similarity).	cytoplasm
P59740	CYSD_SHIFL		cytoplasm
P59741	FADL_SHIFL	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	inner membrane
P59746	CHBG_SHIFL		inner membrane
P59748	PQQE_ENTIT		cytoplasm
P59753	RS13_COXBU		cytoplasm
P59755	RS13_NITEU		cytoplasm
P59765	TREA_SALTI	periplasm (by similarity).	periplasm
P59766	LINA1_PSEPA	periplasm (by similarity).	periplasm
P59767	LINA2_PSEPA	periplasm (by similarity).	periplasm
P59783	OXAA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P59784	PTHA_SHIFL	cytoplasm (potential).	cytoplasm
P59785	ALGA_PSEFL		cytoplasm
P59786	ALGL_PSEFL	periplasm (potential).	periplasm
P59788	ALGX_PSEFL	periplasm (potential).	periplasm
P59789	ALGI_PSEFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59791	ALGF_PSEFL	periplasm (by similarity).	periplasm
P59792	ALGJ_PSEFL	bacterial cell inner membrane; peripheral membrane protein; periplasmic side (by similarity).	periplasm
P59793	ALGD_PSESY		cytoplasm
P59794	ATE_HELHP	cytoplasm (potential).	cytoplasm
P59795	ATE_SPHEL	cytoplasm (potential).	cytoplasm
P59807	CABC_PROVU		extracellular
P59810	OXAA_NITEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59813	FABH_HELHP	cytoplasm (probable).	cytoplasm
P59815	RLUB_SHIFL		cytoplasm
P59818	XERC_MYXXA	cytoplasm (by similarity).	nucleoid
P59821	Y839_HELHP		inner membrane
P59828	ALGG_PSEFL	periplasm (probable).	extracellular
P59831	RLUA_HAEDU		cytoplasm
P59835	RLUC_HAEDU		cytoplasm
P59838	RLUD_BLOFL		cytoplasm
P59840	TRUC_HAEDU		cytoplasm
P59841	FRDC_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59842	INSA_HAEDU		cytoplasm
P59843	INSB_HAEDU		cytoplasm
P59844	FRDD_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59874	TRUB_BORPA		cytoplasm
P59876	TRUB_BRUSU		cytoplasm
P59877	TRUB_BLOFL		cytoplasm
P59878	TRUB_HAEDU		cytoplasm
P59885	RRAA_HAEDU		cytoplasm
P59892	TRUD_HELHP		cytoplasm
P59893	TRUD_WOLSU		cytoplasm
P59911	UBIE_HAEDU		cytoplasm
P59917	Y1769_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59927	SSB_RALEU		cytoplasm
P59928	SSB_BORPA		cytoplasm
P59930	SSB_HAEDU		cytoplasm
P59931	SSB_HELHP		cytoplasm
P59933	SSB_WOLSU		cytoplasm
P59988	USPB_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P59992	FLHD_PHOLL	cytoplasm (by similarity).	cytoplasm

P59993	FLHD_SALPU	cytoplasm (by similarity).	cytoplasm
P60001	CLRD_IDEDE	cytoplasm (probable).	cytoplasm
P60004	USPA_PHOLL	cytoplasm (by similarity).	cytoplasm
P60005	USPE_PHOLL	cytoplasm (by similarity).	cytoplasm
P60012	Y3824_CHRVO		inner membrane
P60013	Y3642_PHOLL		inner membrane
P60037	OXAA_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60046	INSJ_SHIFL		inner membrane
P60054	NDPA_PHOLL	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
P60055	NDPA1_VIBVY	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
P60056	NDPA2_VIBVY	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
P60062	ADIC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60063	ADIC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60064	ADIC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60065	ADIC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60066	ADIC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60067	ALX_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60091	PRMA_CHRVO	cytoplasm (by similarity).	cytoplasm
P60092	PRMA_PHOLL	cytoplasm (by similarity).	cytoplasm
P60094	PRMA_VIBVY	cytoplasm (by similarity).	cytoplasm
P60095	PRMA_WOLSU	cytoplasm (by similarity).	cytoplasm
P60100	RL11_CHRVO		cytoplasm
P60103	RL11_PHOLL		cytoplasm
P60104	RL11_PR01		cytoplasm
P60105	RL11_VIBVY		cytoplasm
P60106	RL11_WOLSU		cytoplasm
P60109	HUTG_CHRVO		cytoplasm
P60110	HUTG_PHOLL		cytoplasm
P60111	HUTG_VIBVY		cytoplasm
P60119	REPX_WIGBR		cytoplasm
P60189	SPRT_SALTI	cytoplasm (potential).	extracellular
P60190	SPRT_SALTY	cytoplasm (potential).	extracellular
P60194	RNZ_SALTI		extracellular
P60195	RNZ_SALTY		extracellular
P60216	MTNN_SALTI		cytoplasm
P60217	MTNN_SALTY		cytoplasm

P60240	RAPA_ECOLI		cytoplasm
P60241	RAPA_ECO57		cytoplasm
P60294	MUKF_SHIFL	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
P60317	PROQ_SALTY	cytoplasm (potential).	inner membrane
P60318	PROQ_SALTI	cytoplasm (potential).	inner membrane
P60325	RPOZ_HELPY		cytoplasm
P60326	RPOZ_HELPJ		cytoplasm
P60327	DCAS_AGRSK		cytoplasm
P60340	TRUB_ECOLI		cytoplasm
P60341	TRUB_ECOL6		cytoplasm
P60342	TRUB_ECO57		cytoplasm
P60344	TRUB_GEOSL		cytoplasm
P60345	TRUB_RHOPA		cytoplasm
P60347	TRUD_GEOSL		cytoplasm
P60350	TRUA_GEOSL		cytoplasm
P60352	TRUA_RHOPA		cytoplasm
P60363	HPPA1_RHOPA	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P60364	CH601_RHOPA	cytoplasm (by similarity).	cytoplasm
P60365	CH602_RHOPA	cytoplasm (by similarity).	cytoplasm
P60366	CH101_RHOPA	cytoplasm (by similarity).	cytoplasm
P60367	CH102_RHOPA	cytoplasm (by similarity).	cytoplasm
P60391	MRAW_ECO57	cytoplasm (by similarity).	cytoplasm
P60396	MRAW_GEOSL		cytoplasm
P60398	MRAW_RHOPA		cytoplasm
P60401	RL2_GEOSL		cytoplasm
P60403	RL2_RHOPA		cytoplasm
P60422	RL2_ECOLI		cytoplasm
P60423	RL2_ECOL6		cytoplasm
P60424	RL2_ECO57		cytoplasm
P60427	RL2_SALTI		cytoplasm
P60428	RL2_SALTY		cytoplasm
P60429	RL2_SHIFL		cytoplasm
P60436	RL2_YERPE		cytoplasm
P60437	RL2_YERPS		cytoplasm
P60438	RL3_ECOLI		cytoplasm
P60439	RL3_ECOL6		cytoplasm
P60440	RL3_ECO57		cytoplasm
P60443	RL3_NEIMA		cytoplasm
P60444	RL3_NEIMB		cytoplasm
P60445	RL3_SALTI		cytoplasm
P60446	RL3_SALTY		cytoplasm
P60447	RL3_SHIFL		cytoplasm
P60454	RL3_GEOSL		cytoplasm
P60456	RL3_RHOPA		cytoplasm
P60472	UPPS_ECOLI		inner membrane
P60473	UPPS_ECOL6		inner membrane
P60474	UPPS_ECO57		inner membrane
P60475	UPPS_SHIFL		inner membrane
P60482	UPPS_GEOSL		inner membrane
P60500	GUAA_GEOSL		cytoplasm
P60501	GUAA_RHOPA		cytoplasm
P60511	LEXA2_XANAC		cytoplasm

P60512	LEXA_XANCI		cytoplasm
P60527	HAD_AGRTR		cytoplasm
P60536	HIS2_GEOSL	cytoplasm (by similarity).	cytoplasm
P60538	HIS21_RHOPA	cytoplasm (by similarity).	cytoplasm
P60539	HIS22_RHOPA	cytoplasm (by similarity).	cytoplasm
P60542	HIS3_GEOSL	cytoplasm (by similarity).	cytoplasm
P60544	HIS3_RHOPA	cytoplasm (by similarity).	cytoplasm
P60547	KGUA_ECOL6	cytoplasm (by similarity).	cytoplasm
P60548	KGUA_SHIFL	cytoplasm (by similarity).	cytoplasm
P60549	KGUA_BDEBA	cytoplasm (by similarity).	cytoplasm
P60551	KGUA_GEOSL	cytoplasm (by similarity).	cytoplasm
P60556	KGUA_RHOPA	cytoplasm (by similarity).	cytoplasm
P60560	GUAC_ECOLI		cytoplasm
P60561	GUAC_ECOL6		cytoplasm
P60566	FIXA_ECOLI		cytoplasm
P60567	FIXA_ECO57		cytoplasm
P60581	HIS4_GEOSL	cytoplasm (by similarity).	cytoplasm
P60582	HIS4_RHOPA	cytoplasm (by similarity).	cytoplasm
P60584	CAIA_ECOLI		cytoplasm
P60585	CAIA_ECOL6		cytoplasm
P60586	CAIA_ECO57		cytoplasm
P60587	CAIA_SHIFL		cytoplasm
P60593	SPEE_NEIMA		cytoplasm
P60594	SPEE_NEIMB		cytoplasm
P60596	HIS5_ECOL6	cytoplasm (by similarity).	cytoplasm
P60597	HIS5_BDEBA	cytoplasm (by similarity).	cytoplasm
P60599	HIS5_GEOSL	cytoplasm (by similarity).	cytoplasm
P60601	HIS5_RHOPA	cytoplasm (by similarity).	cytoplasm
P60624	RL24_ECOLI		cytoplasm
P60625	RL24_ECO57		cytoplasm
P60626	RL24_SALTY		cytoplasm
P60631	NANE1_SALTY		cytoplasm
P60633	YOHJ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60634	YOHJ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60635	YOHJ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60636	YOHJ_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60651	SPEB_ECOLI		cytoplasm
P60652	SPEB_ECO57		cytoplasm
P60653	SPEB_NEIMA		cytoplasm
P60654	SPEB_NEIMB		cytoplasm
P60655	SPEB_SALTI		cytoplasm
P60656	SPEB_SALTY		cytoplasm
P60657	SPEB_SHIFL		cytoplasm
P60658	SPEA_SALTI		cytoplasm
P60659	SPEA_SALTY		cytoplasm
P60665	HIS6_ECO57	cytoplasm (by similarity).	cytoplasm
P60667	HIS6_RHOPA	cytoplasm (by similarity).	cytoplasm
P60668	NANE_SALTI		cytoplasm

P60715	HIS6_GEOSL	cytoplasm (by similarity).	cytoplasm
P60717	LIPA_ECOL6	cytoplasm (potential).	cytoplasm
P60718	LIPA_ECO57	cytoplasm (potential).	cytoplasm
P60719	LIPA_SHIFL	cytoplasm (potential).	cytoplasm
P60721	LIPB_ECO57	cytoplasm (potential).	cytoplasm
P60722	LIPB_ECOL6	cytoplasm (potential).	cytoplasm
P60723	RL4_ECOLI		cytoplasm
P60724	RL4_ECOL6		cytoplasm
P60725	RL4_ECO57		cytoplasm
P60726	RL4_SALTY		cytoplasm
P60727	RL4_SALTI		cytoplasm
P60730	RL4_YERPE		cytoplasm
P60731	RL4_YERPS		cytoplasm
P60732	RL24_NEIMA		cytoplasm
P60733	RL24_NEIMB		cytoplasm
P60740	RL24_GEOSL		cytoplasm
P60744	RL24_RHOPA		cytoplasm
P60753	MSBA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60758	HIS1_ECOL6	cytoplasm (by similarity).	cytoplasm
P60776	LPLA_ECO57	cytoplasm (by similarity).	cytoplasm
P60777	SMP_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
P60779	TSGA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60783	YGIH_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60784	YGIH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60786	LEPA_ECOL6		cytoplasm
P60787	LEPA_ECO57		cytoplasm
P60788	LEPA_SHIFL		cytoplasm
P60789	LEPA_GEOSL		cytoplasm
P60793	LEPA_RHOPA		cytoplasm
P60801	PDXB_SALTI		cytoplasm
P60802	PDXB_SALTY		cytoplasm
P60804	HIS11_GEOSL	cytoplasm (by similarity).	cytoplasm
P60806	HIS1_RHOPA	cytoplasm (by similarity).	cytoplasm
P60809	LPLA_BDEBA	cytoplasm (by similarity).	cytoplasm
P60816	YFBU_SALTI		inner membrane
P60817	YFBU_SALTY		inner membrane
P60821	YHBP_SALTI		inner membrane
P60822	YHBP_SALTY		inner membrane
P60828	YJGA_SALTI		cytoplasm
P60829	YJGA_SALTY		cytoplasm
P60836	HIS12_GEOSL	cytoplasm (by similarity).	cytoplasm
P60837	HISZ_GEOSL	cytoplasm (by similarity).	cytoplasm
P60845	AQPZ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60859	HISX_GEOSL		cytoplasm

P60861	HISX_RHOPA		cytoplasm
P60869	YBJL_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P60870	YBJL_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P60871	YBJL_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P60872	YIDE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P60873	YIDE_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P60874	YIDE_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P60885	HIS7_GEOSL	cytoplasm (by similarity).	cytoplasm
P60887	HIS7_RHOPA	cytoplasm (by similarity).	cytoplasm
P60925	AQPZ_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60926	Y507_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60929	Y3119_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60930	LEPA_BDEBA		cytoplasm
P60933	UPPP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60936	UPPP_BDEBA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60938	UPPP_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60940	UPPP_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60956	LGT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60957	LGT_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60958	LGT_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60959	LGT_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P60960	LGT_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60967	LGT_BDEBA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60968	LGT_BUCAP	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P60970	LGT_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60973	LGT_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P60998	HIS8_BDEBA		cytoplasm
P61000	HIS8_GEOSL		cytoplasm
P61002	HIS8_RHOPA		cytoplasm
P61032	LNT_BDEBA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P61033	LNT_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P61034	LNT_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P61035	LNT_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P61036	LNT_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P61056	RL4_NEIMA		cytoplasm
P61057	RL4_NEIMB		cytoplasm
P61061	RL4_BDEBA		cytoplasm
P61063	RL4_GEOSL		cytoplasm
P61068	RL4_RHOPA		cytoplasm
P61071	RL4_WOLPM		cytoplasm
P61090	SLYA_SALCH		cytoplasm
P61091	SLYA_SALTI		cytoplasm
P61115	KAD_BDEBA	cytoplasm (by similarity).	cytoplasm
P61167	Y1785_RHOPA		inner membrane
P61175	RL22_ECOLI		cytoplasm
P61176	RL22_ECOL6		cytoplasm
P61177	RL22_ECO57		cytoplasm
P61178	RL22_SALTI		cytoplasm
P61179	RL22_SALTY		cytoplasm
P61184	HTPG_BDEBA	cytoplasm (by similarity).	cytoplasm
P61185	HTPG_GEOSL	cytoplasm (by similarity).	cytoplasm
P61187	HTPG_RHOPA	cytoplasm (by similarity).	cytoplasm
P61189	HTPG_WOLPM	cytoplasm (by similarity).	cytoplasm
P61196	LIPA_GEOSL	cytoplasm (potential).	cytoplasm
P61198	LIPA_RHOPA	cytoplasm (potential).	cytoplasm
P61200	LIPA_WOLPM	cytoplasm (potential).	cytoplasm
P61302	RRF_BDEBA	cytoplasm (by similarity).	cytoplasm

P61304	RRF_GEOSL	cytoplasm (by similarity).	cytoplasm
P61309	RRF_RHOPA	cytoplasm (by similarity).	cytoplasm
P61311	RRF_WOLPM	cytoplasm (by similarity).	cytoplasm
P61317	LOLA_ECOL6	periplasm (by similarity).	periplasm
P61318	LOLA_ECO57	periplasm (by similarity).	periplasm
P61319	LOLA_SHIFL	periplasm (by similarity).	periplasm
P61321	LOLB_ECO57	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P61322	LOLB_SHIFL	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P61323	HPRK_GEOSL		cytoplasm
P61341	GATB_BDEBA		cytoplasm
P61343	GATB_GEOSL		cytoplasm
P61347	GATB_RHOPA		cytoplasm
P61349	GATB_WOLPM		cytoplasm
P61351	YCEI_SALTY	periplasm (potential).	periplasm
P61352	YCEI_SALTI	periplasm (potential).	periplasm
P61377	CCMA_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P61378	CCMA2_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P61379	GSH1_PSESY		cytoplasm
P61381	SYCN_YERPS	or: cytoplasm (by similarity). or: bacterial cell inner membrane; peripheral membrane protein (by similarity). not exported across the inner membrane (by similarity).	cytoplasm
P61389	CRCB_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P61394	CRCB_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P61396	GSHB_RHOPA		cytoplasm
P61404	Y2873_GEOSL		cytoplasm
P61405	Y679_RHOPA		cytoplasm
P61411	THE2_GEOSL		cytoplasm
P61415	Y307_RHOPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P61422	THIED_GEOSL		cytoplasm
P61425	THIC_GEOSL		cytoplasm
P61427	THIC_RHOPA		cytoplasm
P61429	PHNS2_DESVH	periplasm (by similarity).	periplasm
P61433	MURB_BDEBA	cytoplasm (probable).	inner membrane
P61435	MURB_GEOSL	cytoplasm (probable).	inner membrane
P61450	FABA_RHOPA	cytoplasm (by similarity).	cytoplasm
P61452	FABZ_DESVH	cytoplasm (by similarity).	cytoplasm
P61453	FABZ_GEOSL	cytoplasm (by similarity).	cytoplasm
P61454	FABZ_RHOPA	cytoplasm (by similarity).	cytoplasm
P61455	FABZ_WOLPM	cytoplasm (by similarity).	cytoplasm
P61466	Y1076_DESVH		inner membrane

P61467	YIDD_ECOL6		inner membrane
P61468	YIDD_ECO57		inner membrane
P61469	Y3467_GEOSL		inner membrane
P61471	Y3394_RHOPA		inner membrane
P61472	YIDD_SALTI		inner membrane
P61473	YIDD_SHIFL		inner membrane
P61475	Y4101_YERPE		inner membrane
P61477	HSLV_RHOPA	cytoplasm (by similarity).	cytoplasm
P61479	HSLV_WOLPM	cytoplasm (by similarity).	cytoplasm
P61481	LOLD_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P61482	LOLD_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P61517	CAN_ECOLI		cytoplasm
P61518	CAN_SHIFL		extracellular
P61522	ASSY_DESVH	cytoplasm (probable).	cytoplasm
P61523	ASSY_GEOSL	cytoplasm (probable).	cytoplasm
P61529	RUVB_BDEBA		cytoplasm
P61531	RUVB_DESVH		cytoplasm
P61532	RUVB_GEOSL		cytoplasm
P61536	RUVB_RHOPA		cytoplasm
P61539	RUVB_WOLPM		cytoplasm
P61548	FLBT_RHOPA		cytoplasm
P61605	LEXA_AGRTU		cytoplasm
P61608	LEXA1_GEOSL		cytoplasm
P61609	LEXA2_GEOSL		cytoplasm
P61613	LEXA_RHOCA		cytoplasm
P61614	LEXA_RHOPA		cytoplasm
P61616	MURB_RHOPA	cytoplasm (probable).	inner membrane
P61617	SAHH_GEOSL	cytoplasm (by similarity).	cytoplasm
P61653	KDSA_BDEBA	cytoplasm (by similarity).	cytoplasm
P61654	KDSA_DESVH	cytoplasm (by similarity).	cytoplasm
P61655	KDSA_GEOSL	cytoplasm (by similarity).	cytoplasm
P61657	KDSA_RHOPA	cytoplasm (by similarity).	cytoplasm
P61659	HIS7_DESVH	cytoplasm (by similarity).	cytoplasm
P61662	SFSA_BDEBA		cytoplasm
P61663	SFSA_DESVH		cytoplasm
P61664	SFSA_GEOSL		cytoplasm
P61666	MUTS_DESVH		cytoplasm
P61667	MUTS_GEOSL		cytoplasm
P61670	MUTS_RHOPA		cytoplasm
P61673	MUTS_WOLPM		cytoplasm
P61676	MURC_BDEBA	cytoplasm (probable).	cytoplasm
P61678	MURC_DESVH	cytoplasm (probable).	cytoplasm
P61679	MURC_GEOSL	cytoplasm (probable).	cytoplasm
P61682	MURC_RHOPA	cytoplasm (probable).	cytoplasm
P61683	MURC_WOLPM	cytoplasm (probable).	cytoplasm
P61711	RISB2_BRUAB		cytoplasm
P61712	RISB2_BRUME		cytoplasm
P61713	RISB2_BRUSU		cytoplasm
P61714	RISB_ECOLI		cytoplasm
P61715	RISB_ECOL6		cytoplasm
P61717	RISB_ECO57		cytoplasm
P61718	RISB_SHIFL		cytoplasm

P61720	RISB_BDEBA		cytoplasm
P61723	RISB_GEOSL		cytoplasm
P61726	RISB1_RHOPA		cytoplasm
P61727	RISB2_RHOPA		cytoplasm
P61729	RISB_WOLPM		cytoplasm
P61732	PHS_BDEBA		cytoplasm
P61734	PHS_RHOPA		cytoplasm
P61735	SELA_DESVH	cytoplasm (by similarity).	cytoplasm
P61736	SELA_GEOSL	cytoplasm (by similarity).	cytoplasm
P61739	GPDA_DESVH	cytoplasm (probable).	cytoplasm
P61740	GPDA_GEOSL	cytoplasm (probable).	cytoplasm
P61746	GPDA_RHOPA	cytoplasm (probable).	cytoplasm
P61749	GPDA_WOLPM	cytoplasm (probable).	cytoplasm
P61779	HIS5_DESVH	cytoplasm (by similarity).	cytoplasm
P61786	NUDH_RHOPA		cytoplasm
P61787	NUDH_WOLPM		cytoplasm
P61788	Y1747_RHOPA		inner membrane
P61887	RMLA2_ECOLI		cytoplasm
P61888	RMLA2_SHIFL		cytoplasm
P61889	MDH_ECOLI		cytoplasm
P61890	MDH_ECOL6		cytoplasm
P61891	MDH_ECO57		cytoplasm
P61892	MDH_YERPE		cytoplasm
P61893	MDH_YERPS		cytoplasm
P61906	DUT_BDEBA		extracellular
P61908	DUT_GEOSL		extracellular
P61911	DUT_RHOPA		extracellular
P61913	DUT_WOLPM		extracellular
P61919	ARGA_PSESY	cytoplasm (probable).	cytoplasm
P61940	RISB_DESVH		cytoplasm
P61949	FLAV_ECOLI		cytoplasm
P61950	FLAV_ECOL6		cytoplasm
P61951	FLAV_ECO57		cytoplasm
P61973	MDH_BDEBA		cytoplasm
P61984	CBID_DESVH		cytoplasm
P61985	CBID_GEOSL		cytoplasm
P62033	Y1964_BDEBA		inner membrane
P62035	Y2259_DESVH		inner membrane
P62036	Y1074_GEOSL		inner membrane
P62040	Y1097_RHOPA		inner membrane
P62043	Y484_WOLPM		inner membrane
P62051	LDH_DESVH	cytoplasm (by similarity).	cytoplasm
P62060	ARGJ_DESVH	cytoplasm (probable).	cytoplasm
P62061	ARGJ_GEOSL	cytoplasm (probable).	cytoplasm
P62064	ARGJ_RHOPA	cytoplasm (probable).	cytoplasm
P62066	YCEQ_ECOLI		cytoplasm
P62067	YCEQ_ECO57		cytoplasm
P62088	PRSX_ECOLI	cytoplasm (potential).	cytoplasm
P62089	PRSX_ECOL6	cytoplasm (potential).	cytoplasm
P62209	RECA_AZOLI	cytoplasm (by similarity).	cytoplasm
P62212	RECA_BDEBA	cytoplasm (by similarity).	cytoplasm
P62214	RECA_DESVH	cytoplasm (by similarity).	cytoplasm
P62215	RECA_GEOSL	cytoplasm (by similarity).	cytoplasm
P62220	RECA_SODGL	cytoplasm (by similarity).	cytoplasm
P62222	RECA_WOLPM	cytoplasm (by similarity).	cytoplasm
P62223	GLND_RHOPA		cytoplasm

P62225	MERB_KLEPN		extracellular
P62227	RS16_BDEBA		cytoplasm
P62229	RS16_DESVH		cytoplasm
P62230	RS16_GEOSL		cytoplasm
P62236	RS16_RHOPA		cytoplasm
P62240	RS16_WOLPM		cytoplasm
P62349	HIS2_PHOPR	cytoplasm (by similarity).	cytoplasm
P62353	HIS4_DESVH	cytoplasm (by similarity).	cytoplasm
P62355	HIS4_PHOPR	cytoplasm (by similarity).	cytoplasm
P62361	YCCT_SALTI		inner membrane
P62362	YCCT_SALTY		inner membrane
P62364	HIS1_DESVH	cytoplasm (by similarity).	cytoplasm
P62365	HIS1_PHOPR	cytoplasm (by similarity).	cytoplasm
P62385	Y2789_PHOPR		inner membrane
P62386	HIS3_DESVH	cytoplasm (by similarity).	cytoplasm
P62399	RL5_ECOLI		cytoplasm
P62400	RL5_ECOL6		cytoplasm
P62401	RL5_ECO57		cytoplasm
P62404	RL5_SALTI		cytoplasm
P62405	RL5_SALTY		cytoplasm
P62410	PGK_BDEBA	cytoplasm (potential).	cytoplasm
P62412	PGK_DESVH	cytoplasm (potential).	cytoplasm
P62418	PGK_PHOPR	cytoplasm (potential).	cytoplasm
P62419	PGK_RHOPA	cytoplasm (potential).	cytoplasm
P62422	PGK_WOLPM	cytoplasm (potential).	cytoplasm
P62430	Y966_PHOPR		inner membrane
P62432	RL11_BDEBA		cytoplasm
P62433	RL11_DESVH		cytoplasm
P62434	RL11_GEOSL		cytoplasm
P62440	RL11_PHOPR		cytoplasm
P62441	RL11_RHOPA		cytoplasm
P62444	RL11_WOLPM		cytoplasm
P62448	HIS5_PHOPR	cytoplasm (by similarity).	cytoplasm
P62450	HIS6_DESVH	cytoplasm (by similarity).	cytoplasm
P62452	HIS6_PHOPR	cytoplasm (by similarity).	cytoplasm
P62455	HIS7_PHOPR	cytoplasm (by similarity).	cytoplasm
P62457	HISX_DESVH		cytoplasm
P62459	HISX_PHOPR		cytoplasm
P62464	UPPP_DESVH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P62466	UPPP_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P62469	MRAW_BDEBA		cytoplasm
P62470	MRAW_DESVH		cytoplasm
P62475	MRAW_PHOPR		cytoplasm
P62478	MRAW_WOLPM		cytoplasm
P62479	Y2397_PHOPR		inner membrane
P62518	OPGH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P62519	OPGH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P62533	PAPK_ECOL6	secreted protein (by similarity).	extracellular
P62534	YR7C_ECOLI		inner membrane
P62535	YR7C_ECOL6		cytoplasm
P62536	COPB3_ECOLI		cytoplasm
P62537	COPB_SHIFL		cytoplasm
P62538	REPE2_ECOLI		cytoplasm
P62539	REPE2_SALTI		cytoplasm
P62552	CCDA_ECOLI		cytoplasm
P62553	CCDA_ECO57		cytoplasm
P62554	CCDB_ECOLI		cytoplasm
P62555	CCDB_ECO57		cytoplasm
P62556	SOPA_ECOLI		cytoplasm
P62557	SOPA_ECO57		cytoplasm
P62558	SOPB_ECOLI		cytoplasm
P62559	SOPB_ECO57		cytoplasm
P62572	CSCR_SHIFL		cytoplasm
P62573	HPCR_ECOLI		cytoplasm
P62574	HPCR_SHIFL		cytoplasm
P62577	CAT_ECOLI		cytoplasm
P62578	CAT_ACIAN		cytoplasm
P62579	CAT_ACIBA		cytoplasm
P62580	CAT_SALTI		cytoplasm
P62584	KS7A_ECOLI		cytoplasm
P62585	KS7A_ECOL6		cytoplasm
P62586	KPSE5_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P62587	KPSE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P62590	INT2_ECOLI		cytoplasm
P62591	INT2_PSEAE		cytoplasm
P62592	INT2_SALTI		cytoplasm
P62593	BLAT_ECOLI		extracellular
P62594	BLAT_SALTI		extracellular
P62602	TREF_ECO57	cytoplasm (by similarity).	extracellular
P62604	CSCR_ECOLI		cytoplasm
P62606	FIM1C_ECOL6	fimbrium (by similarity).	extracellular
P62607	FMF2_ECOLI		extracellular
P62608	FMF2_ECOL6		extracellular
P62609	FOCC_ECOLI	periplasm (by similarity).	periplasm
P62610	FOCC_ECOL6	periplasm (by similarity).	periplasm
P62615	ISPE_ECOLI		cytoplasm
P62616	ISPE_ECO57		cytoplasm
P62617	ISPF_ECOLI		cytoplasm
P62618	ISPF_ECO57		extracellular
P62619	ISPF_SHIFL		extracellular
P62620	ISPG_ECOLI		cytoplasm
P62621	ISPG_ECOL6		cytoplasm
P62622	ISPG_ECO57		cytoplasm
P62623	ISPH_ECOLI		cytoplasm
P62624	ISPH_ECOL6		cytoplasm
P62625	ISPH_ECO57		cytoplasm
P62670	FLMA_ECOLI		inner membrane
P62671	FLMA_ECO57		inner membrane

P62672	APAG_ECOLI		inner membrane
P62673	APAG_ECOL6		inner membrane
P62674	APAG_ECO57		inner membrane
P62675	APAG_SHIFL		inner membrane
P62676	SERC_SALGL	cytoplasm (by similarity).	cytoplasm
P62677	SERC_SALTI	cytoplasm (by similarity).	cytoplasm
P62707	GPMA_ECOLI		cytoplasm
P62708	GPMA_ECOL6		cytoplasm
P62709	GPMA_ECO57		cytoplasm
P62710	GPMA_SHIFL		cytoplasm
P62724	YEIH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P62725	YEIH_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P62768	YAEH_ECOLI		inner membrane
P62769	YAEH_ECO57		inner membrane
P62770	YAEH_ECOL6		inner membrane
P62771	YAEH_SHIFL		inner membrane
P62920	GCSP_BRUAB		cytoplasm
P62921	GCSP_BRUME		cytoplasm
P63020	GNTY_ECOLI		inner membrane
P63021	GNTY_ECOL6		inner membrane
P63022	GNTY_ECO57		inner membrane
P63023	GNTY_SHIFL		inner membrane
P63177	RLMB_ECOLI		cytoplasm
P63178	RLMB_ECOL6		cytoplasm
P63179	RLMB_ECO57		cytoplasm
P63180	RLMB_SHIFL		cytoplasm
P63184	KUP_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63187	GRPE_AGRT5	cytoplasm (probable).	cytoplasm
P63188	GRPE_AGRTU	cytoplasm (probable).	cytoplasm
P63194	RS12_BRUSU		cytoplasm
P63195	RS12_PASHA		cytoplasm
P63196	RS12_HAEIN		cytoplasm
P63201	GADW_ECOLI		cytoplasm
P63202	GADW_ECO57		cytoplasm
P63203	GADW_SHIFL		cytoplasm
P63204	GADE_ECOLI		cytoplasm
P63205	GADE_ECOL6		cytoplasm
P63206	GADE_ECO57		cytoplasm
P63207	GADE_SHIFL		cytoplasm
P63222	GMHA_SALTI	cytoplasm (by similarity).	cytoplasm
P63223	GMHA_SALTY	cytoplasm (by similarity).	cytoplasm
P63225	GMHA_ECO57	cytoplasm (by similarity).	cytoplasm
P63226	GMHA_ECOL6	cytoplasm (by similarity).	cytoplasm
P63227	GMHA_SHIFL	cytoplasm (by similarity).	cytoplasm
P63228	GMHB_ECOLI	cytoplasm (by similarity).	cytoplasm
P63229	GMHB_ECO57	cytoplasm (by similarity).	cytoplasm
P63230	GMHB_SHIFL	cytoplasm (by similarity).	cytoplasm
P63236	GADC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P63237	CSPE_BUCAI	cytoplasm (by similarity).	cytoplasm
P63238	CSPE_BUCAP	cytoplasm (by similarity).	cytoplasm
P63262	CBPA_SALTI	nucleoid (by similarity).	inner membrane
P63263	CBPA_SALTY	nucleoid (by similarity).	inner membrane
P63264	CBPM_ECOLI		cytoplasm
P63265	CBPM_ECO57		cytoplasm
P63266	CBPM_SHIFL		cytoplasm
P63285	CLPB_ECO57	cytoplasm (probable).	cytoplasm
P63286	CLPB_ECOL6	cytoplasm (probable).	cytoplasm
P63299	YTFR_ECO57		inner membrane
P63336	6PGL_NEIMA		extracellular
P63337	6PGL_NEIMB		extracellular
P63341	YQEG_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63342	YQEG_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63343	FTSH_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63344	FTSH_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63351	BTUD_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P63352	BTUD_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P63353	CYSA_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P63354	CYSA_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P63355	METN_ECOL6	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P63356	METN_ECO57	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P63359	MSBA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63360	MSBA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63361	PSTB_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P63362	PSTB_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

P63365	PSTB_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P63366	PSTB_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P63386	YRBF_ECOLI		inner membrane
P63387	YRBF_ECOL6		inner membrane
P63388	YRBF_ECO57		inner membrane
P63389	YHES_ECOLI		inner membrane
P63390	YHES_ECO57		inner membrane
P63411	ACKA_SALTY	cytoplasm (by similarity).	cytoplasm
P63412	ACKA_SALTI	cytoplasm (by similarity).	cytoplasm
P63417	YHBS_ECOLI		cytoplasm
P63418	YHBS_ECOL6		cytoplasm
P63419	YHBS_ECO57		cytoplasm
P63420	YPEA_ECOL6		cytoplasm
P63421	YPEA_ECO57		cytoplasm
P63422	YPEA_SHIFL		cytoplasm
P63423	YPEA_SALTY		cytoplasm
P63424	YPEA_SALTI		cytoplasm
P63437	ACP_BRUME	cytoplasm (by similarity).	cytoplasm
P63438	ACP_BRUSU	cytoplasm (by similarity).	cytoplasm
P63441	ACP_NEIMA	cytoplasm (by similarity).	cytoplasm
P63442	ACP_NEIMB	cytoplasm (by similarity).	cytoplasm
P63446	ACP_XANAC	cytoplasm (by similarity).	cytoplasm
P63447	ACP_XANCP	cytoplasm (by similarity).	cytoplasm
P63448	ACP_XYLFA	cytoplasm (by similarity).	cytoplasm
P63449	ACP_XYLFT	cytoplasm (by similarity).	cytoplasm
P63450	ACPXL_BRUME	cytoplasm (by similarity).	cytoplasm
P63451	ACPXL_BRUSU	cytoplasm (by similarity).	cytoplasm
P63462	ACPD_SALTY		extracellular
P63463	ACPD_SALTI		extracellular
P63464	ACPS_BRUME	cytoplasm (by similarity).	cytoplasm
P63465	ACPS_BRUSU	cytoplasm (by similarity).	cytoplasm
P63466	ACPS_SALTY	cytoplasm (by similarity).	cytoplasm
P63467	ACPS_SALTI	cytoplasm (by similarity).	cytoplasm
P63486	ALLA_ECO57		extracellular
P63487	ALLA_SHIFL		extracellular
P63521	LCFA_SALTY	membrane; peripheral membrane protein (potential). partially membrane-associated (potential).	cytoplasm
P63522	LCFA_SALTI	membrane; peripheral membrane protein (potential). partially membrane-associated (potential).	cytoplasm
P63557	ARGB_SALTY	cytoplasm (probable).	cytoplasm
P63558	ARGB_SALTI	cytoplasm (probable).	cytoplasm
P63566	ARGD_BRUME	cytoplasm (probable).	cytoplasm
P63567	ARGD_BRUSU	cytoplasm (probable).	cytoplasm
P63573	ARGJ_NEIMA	cytoplasm (probable).	cytoplasm
P63574	ARGJ_NEIMB	cytoplasm (probable).	cytoplasm
P63599	AROK_NEIMA	cytoplasm (probable).	cytoplasm
P63600	AROK_NEIMB	cytoplasm (probable).	cytoplasm
P63601	AROK_SALTY	cytoplasm (probable).	cytoplasm

P63602	AROK_SALTI	cytoplasm (probable).	cytoplasm
P63603	AROL_SALTY	cytoplasm (probable).	cytoplasm
P63604	AROL_SALTI	cytoplasm (probable).	cytoplasm
P63607	AROC_BRUME		cytoplasm
P63608	AROC_BRUSU		cytoplasm
P63609	AROC_ECOL6		cytoplasm
P63610	AROC_ECO57		cytoplasm
P63615	AROB_BRUME	cytoplasm (probable).	cytoplasm
P63616	AROB_BRUSU	cytoplasm (probable).	cytoplasm
P63621	ARSC_NEIMA		cytoplasm
P63622	ARSC_NEIMB		cytoplasm
P63623	ASNA_ECOL6	cytoplasm (by similarity).	cytoplasm
P63624	ASNA_ECO57	cytoplasm (by similarity).	cytoplasm
P63631	MURI_BRUME		cytoplasm
P63632	MURI_BRUSU		cytoplasm
P63633	MURI_ECO57		cytoplasm
P63634	MURI_SHIFL		cytoplasm
P63646	ATE_BRUME	cytoplasm (potential).	cytoplasm
P63647	ATE_BRUSU	cytoplasm (potential).	cytoplasm
P63660	ATPE_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P63661	ATPE_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P63699	BFRB_NEIMA		cytoplasm
P63700	BFRB_NEIMB		cytoplasm
P63701	Y420_NEIMA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P63702	Y2020_NEIMB	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P63727	C562_SALTY	periplasm (by similarity).	periplasm
P63728	C562_SALTI	periplasm (by similarity).	periplasm
P63737	CARB_ECO57		cytoplasm
P63738	CARB_SHIFL		cytoplasm
P63746	EUTS_ECOLI		cytoplasm
P63747	EUTS_ECOL6		cytoplasm
P63748	EUTS_SHIFL		cytoplasm
P63773	CH10_XYLFA	cytoplasm (by similarity).	cytoplasm
P63774	CH10_XYLFT	cytoplasm (by similarity).	cytoplasm
P63798	CLS_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63799	CLS_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63808	COAA_BRUME	cytoplasm (probable).	cytoplasm
P63809	COAA_BRUSU	cytoplasm (probable).	cytoplasm
P63814	COAD_BRUME	cytoplasm (by similarity).	cytoplasm
P63815	COAD_BRUSU	cytoplasm (by similarity).	cytoplasm
P63816	COAD_NEIMA	cytoplasm (by similarity).	cytoplasm
P63817	COAD_NEIMB	cytoplasm (by similarity).	cytoplasm
P63824	COAE_BRUME	cytoplasm (by similarity).	cytoplasm
P63825	COAE_BRUSU	cytoplasm (by similarity).	cytoplasm

P63828	COAE_SALTY	cytoplasm (by similarity).	cytoplasm
P63829	COAE_SALTI	cytoplasm (by similarity).	cytoplasm
P63850	HEM6_BRUME	cytoplasm (by similarity).	cytoplasm
P63851	HEM6_BRUSU	cytoplasm (by similarity).	cytoplasm
P63860	CRCB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63861	CRCB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63866	CRCB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63867	CRCB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63875	CSRA_PHOLL		cytoplasm
P63876	CSRA_YERPE		cytoplasm
P63877	CSRA_XYLFA		cytoplasm
P63878	CSRA_XYLFT		cytoplasm
P63883	AMIC_ECOLI		extracellular
P63884	AMIC_ECOL6		extracellular
P63889	CYSC_SALTY		cytoplasm
P63890	CYSC_SALTI		cytoplasm
P63901	DCD_ECOL6		extracellular
P63902	DCD_ECO57		extracellular
P63903	DCD_NEIMA		extracellular
P63904	DCD_NEIMB		extracellular
P63905	DCD_SALTY		extracellular
P63906	DCD_SALTI		extracellular
P63913	DEF_BRUME		cytoplasm
P63914	DEF_BRUSU		cytoplasm
P63915	DEF_NEIMA		cytoplasm
P63916	DEF_NEIMB		cytoplasm
P63917	DEF_XYLFA		cytoplasm
P63918	DEF_XYLFT		cytoplasm
P63919	DEFL_BRUME		cytoplasm
P63920	DEFL_BRUSU		cytoplasm
P63923	DEOB_SALTY	cytoplasm (by similarity).	cytoplasm
P63924	DEOB_SALTI	cytoplasm (by similarity).	cytoplasm
P63933	DGTL1_BRUME		cytoplasm
P63934	DGTL1_BRUSU		cytoplasm
P63943	DAPA_ECOL6	cytoplasm (by similarity).	cytoplasm
P63944	DAPA_ECO57	cytoplasm (by similarity).	cytoplasm
P63968	DNAJ_NEIMA	cytoplasm (probable).	cytoplasm
P63969	DNAJ_NEIMB	cytoplasm (probable).	cytoplasm
P63989	DPO4_SALTY	cytoplasm (probable).	cytoplasm
P63990	DPO4_SALTI	cytoplasm (probable).	cytoplasm
P63993	DSBB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P63994	DSBB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64004	DUT_BRUME		extracellular
P64005	DUT_BRUSU		extracellular

P64006	DUT_ECOL6		extracellular
P64007	DUT_ECO57		extracellular
P64008	DUT_SALTY		extracellular
P64009	DUT_SALTI		extracellular
P64046	EFPL_SALTY		cytoplasm
P64047	EFPL_SALTI		cytoplasm
P64066	ENGB_BRUME		cytoplasm
P64067	ENGB_BRUSU		cytoplasm
P64068	ENGB_SALTY		cytoplasm
P64069	ENGB_SALTI		cytoplasm
P64076	ENO_SALTY	cytoplasm (by similarity).	extracellular
P64077	ENO_SALTI	cytoplasm (by similarity).	extracellular
P64095	FIXB_SALTY		cytoplasm
P64096	FIXB_SALTI		cytoplasm
P64101	EXDL2_HELPY	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
P64102	EXDL2_HELPJ	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
P64103	FABA_BRUME	cytoplasm (by similarity).	cytoplasm
P64104	FABA_BRUSU	cytoplasm (by similarity).	cytoplasm
P64105	FABA_SALTY	cytoplasm (by similarity).	cytoplasm
P64106	FABA_SALTI	cytoplasm (by similarity).	cytoplasm
P64127	FIS_VIBCH		cytoplasm
P64128	FIS_VIBPA		cytoplasm
P64129	FIS_VIBVU		cytoplasm
P64130	FISL_NEIMA		cytoplasm
P64131	FISL_NEIMB		cytoplasm
P64132	FMT_BRUME		cytoplasm
P64133	FMT_BRUSU		cytoplasm
P64148	FPG_ECO57		cytoplasm
P64149	FPG_SHIFL		cytoplasm
P64154	FPG_XYLFA		cytoplasm
P64155	FPG_XYLFT		cytoplasm
P64160	FTSB_NEIMA	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
P64161	FTSB_NEIMB	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
P64162	FTSB_SALTY	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
P64163	FTSB_SALTI	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane

P64176	MACA_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P64177	MACA_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P64186	GPDA_BRUME	cytoplasm (probable).	cytoplasm
P64187	GPDA_BRUSU	cytoplasm (probable).	cytoplasm
P64197	GATB_BRUME		cytoplasm
P64198	GATB_BRUSU		cytoplasm
P64203	GATC_BRUME		cytoplasm
P64204	GATC_BRUSU		cytoplasm
P64209	GCH1_SALTY		cytoplasm
P64210	GCH1_SALTI		cytoplasm
P64211	GCSH_BRUME		cytoplasm
P64212	GCSH_BRUSU		cytoplasm
P64222	GCST_SALTY		cytoplasm
P64223	GCST_SALTI		cytoplasm
P64232	GID_BRUME	cytoplasm (by similarity).	cytoplasm
P64233	GID_BRUSU	cytoplasm (by similarity).	cytoplasm
P64237	GIDB_SALTY		cytoplasm
P64238	GIDB_SALTI		cytoplasm
P64243	GLGS_ECOL6		cytoplasm
P64244	GLGS_ECO57		cytoplasm
P64251	GLPE_SALTY	cytoplasm (by similarity).	cytoplasm
P64252	GLPE_SALTI	cytoplasm (by similarity).	cytoplasm
P64253	GLK_NEIMA	cytoplasm (by similarity).	cytoplasm
P64254	GLK_NEIMB	cytoplasm (by similarity).	cytoplasm
P64255	MRAY_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64256	MRAY_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64257	MRAY_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64258	MRAY_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64267	GNS_SALTY		cytoplasm
P64268	GNS_SALTI		cytoplasm
P64271	GREA_BRUME		cytoplasm
P64272	GREA_BRUSU		cytoplasm
P64273	GREB_ECOL6		cytoplasm
P64274	GREB_ECO57		cytoplasm
P64275	GREA_HELPY		cytoplasm
P64276	GREA_HELPJ		cytoplasm
P64281	GREA_SALTY		cytoplasm
P64282	GREA_SALTI		cytoplasm
P64294	GUAA_ECOL6		cytoplasm
P64295	GUAA_ECO57		cytoplasm
P64305	HAM1_BRUME		extracellular
P64306	HAM1_BRUSU		extracellular
P64342	Y405_XYLFA		inner membrane
P64343	Y1667_XYLFT		inner membrane

P64344	HFQ_NEIMA		cytoplasm
P64345	HFQ_NEIMB		cytoplasm
P64346	HIS1_NEIMA	cytoplasm (by similarity).	cytoplasm
P64347	HIS1_NEIMB	cytoplasm (by similarity).	cytoplasm
P64351	HIS2_BRUME	cytoplasm (by similarity).	cytoplasm
P64352	HIS2_BRUSU	cytoplasm (by similarity).	cytoplasm
P64366	HIS7_BRUME	cytoplasm (by similarity).	cytoplasm
P64367	HIS7_BRUSU	cytoplasm (by similarity).	cytoplasm
P64370	HIS7_NEIMA	cytoplasm (by similarity).	cytoplasm
P64371	HIS7_NEIMB	cytoplasm (by similarity).	cytoplasm
P64377	HISZ_BRUME	cytoplasm (by similarity).	cytoplasm
P64378	HISZ_BRUSU	cytoplasm (by similarity).	cytoplasm
P64382	YHIT_HELPY		cytoplasm
P64383	YHIT_HELPJ		cytoplasm
P64388	DBHB_NEIMA		cytoplasm
P64389	DBHB_NEIMB		cytoplasm
P64390	IHFA_BRUME		cytoplasm
P64391	IHFA_BRUSU		cytoplasm
P64392	IHFA_NEIMA		cytoplasm
P64393	IHFA_NEIMB		cytoplasm
P64394	IHFB_SALTY		cytoplasm
P64395	IHFB_SALTI		cytoplasm
P64396	HRCB_BRUME		cytoplasm
P64397	HRCB_BRUSU		cytoplasm
P64413	NIKR_SALTY		cytoplasm
P64414	NIKR_SALTI		cytoplasm
P64419	HYPB_SALTY		cytoplasm
P64420	HYPB_SALTI		cytoplasm
P64421	HYBF_SALTY		cytoplasm
P64422	HYBF_SALTI		cytoplasm
P64424	ZNTB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64425	ZNTB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64426	YDDW_ECOLI	bacterial cell membrane; lipid-anchor (potential).	cytoplasm
P64427	YDDW_ECO57	bacterial cell membrane; lipid-anchor (potential).	cytoplasm
P64428	YDDW_SHIFL	bacterial cell membrane; lipid-anchor (potential).	cytoplasm
P64429	YPFJ_ECOLI		inner membrane
P64430	YPFJ_ECOL6		inner membrane
P64431	YPFJ_ECO57		inner membrane
P64433	YPJD_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64434	YPJD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64435	YBCW_ECOLI		cytoplasm
P64436	YBCW_ECO57		cytoplasm
P64437	YBGS_SALTY		outer membrane
P64438	YBGS_SALTI		outer membrane

P64440	YBJM_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64441	YBJM_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64445	YNAJ_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64446	YNAJ_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64447	YNAJ_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64448	YNBE_ECOLI		cytoplasm
P64449	YNBE_ECOL6		cytoplasm
P64450	YNBE_SHIFL		cytoplasm
P64451	YDCL_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P64452	YDCL_ECO57	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P64453	YDCX_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64454	YDCX_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64455	YDCY_ECOLI		cytoplasm
P64456	YDCY_ECOL6		cytoplasm
P64457	YDCY_ECO57		cytoplasm
P64458	YDCY_SHIFL		cytoplasm
P64459	YNCJ_ECOLI		cytoplasm
P64460	YNCJ_ECO57		cytoplasm
P64461	YNEC_ECOLI		cytoplasm
P64462	YNEC_ECO57		cytoplasm
P64463	YDFZ_ECOLI		cytoplasm
P64464	YDFZ_ECOL6		cytoplasm
P64465	YDFZ_ECO57		cytoplasm
P64466	YDFZ_SHIFL		cytoplasm
P64467	HNSB2_ECOLI		cytoplasm
P64468	HNSB2_ECOL6		cytoplasm
P64469	HNSB2_ECO57		cytoplasm
P64470	HNSB2_SHIFL		cytoplasm
P64471	YDHI_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64472	YDHI_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64473	YDHI_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64474	YDHL_ECOLI		cytoplasm
P64475	YDHL_SHIFL		cytoplasm
P64476	YDIH_ECOLI		cytoplasm

P64477	YDIH_ECO57		cytoplasm
P64478	YDIH_SHIFL		cytoplasm
P64479	YDIZ_ECOLI		cytoplasm
P64480	YDIZ_ECO57		cytoplasm
P64482	YDJM_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64483	YEAQ_ECOLI		cytoplasm
P64484	YEAQ_ECO57		cytoplasm
P64485	YEAQ_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64486	YEAQ_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64487	YEAQ_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64488	YEAR_ECOLI		cytoplasm
P64489	YEAR_ECO57		cytoplasm
P64490	YOAC_ECOLI		cytoplasm
P64491	YOAC_ECO57		cytoplasm
P64492	YOAC_SHIFL		cytoplasm
P64493	YOAF_ECOLI		periplasm
P64494	YOAF_ECOL6		periplasm
P64495	YOAF_SHIFL		periplasm
P64496	YOAG_ECOLI		cytoplasm
P64497	YOAG_ECOL6		cytoplasm
P64498	YOAG_ECO57		cytoplasm
P64499	YEBO_ECOLI	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P64500	YEBO_ECOL6	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P64501	YEBO_SALTY	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P64502	YEBO_SALTI	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P64503	YEBV_ECOLI		cytoplasm
P64504	YEBV_ECOL6		cytoplasm
P64505	YEBV_SHIFL		cytoplasm
P64506	YEBY_ECOLI		cytoplasm
P64507	YEBY_SHIFL		cytoplasm
P64516	YECN_ECOL6	bacterial cell inner membrane (by similarity).	inner membrane
P64517	YODC_ECOLI		cytoplasm
P64518	YODC_ECO57		cytoplasm
P64519	YODD_ECOLI		cytoplasm
P64520	YODD_ECOL6		cytoplasm
P64521	YEET_ECOLI		cytoplasm
P64522	YEET_ECOL6		cytoplasm
P64523	YEET_ECO57		cytoplasm
P64524	YEEV_ECOLI		cytoplasm

P64525	YEEV_ECOL6		cytoplasm
P64526	YEEW_ECOLI		cytoplasm
P64527	YEEW_ECO57		cytoplasm
P64530	YOHL_ECOLI		cytoplasm
P64531	YOHL_ECOL6		cytoplasm
P64532	YOHL_ECO57		cytoplasm
P64533	YOHL_SHIFL		cytoplasm
P64534	YOHN_ECOLI		cytoplasm
P64535	YOHN_ECOL6		cytoplasm
P64536	YEIS_ECOLI		inner membrane
P64537	YEIS_ECOL6		inner membrane
P64538	YEIS_ECO57		inner membrane
P64539	YEIS_SHIFL		inner membrane
P64540	YFCL_ECOLI		cytoplasm
P64541	YFCL_ECO57		cytoplasm
P64542	YPEC_ECOLI		cytoplasm
P64543	YPEC_ECO57		cytoplasm
P64544	YPEC_SHIFL		cytoplasm
P64545	YFGG_ECOLI		inner membrane
P64546	YFGG_ECO57		inner membrane
P64547	YFGG_SHIFL		inner membrane
P64548	YFIR_ECOLI		inner membrane
P64549	YFIR_ECO57		inner membrane
P64550	YGAW_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64551	YGAW_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64552	YGAW_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64553	YGAW_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64554	YGCF_ECOLI		cytoplasm
P64555	YGCF_ECO57		cytoplasm
P64556	YGCF_SHIFL		cytoplasm
P64557	YGFM_ECOLI		cytoplasm
P64558	YGFM_ECO57		cytoplasm
P64559	YGFY_ECOLI		cytoplasm
P64560	YGFY_ECOL6		cytoplasm
P64561	YGFY_ECO57		cytoplasm
P64562	YQFE_ECOLI		cytoplasm
P64563	YQFE_ECOL6		cytoplasm
P64564	YGGT_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64565	YGGT_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64566	YGGT_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64570	YQGC_ECOLI		cytoplasm
P64571	YQGC_ECO57		cytoplasm

P64572	YGHR_ECOLI		cytoplasm
P64573	YGHR_ECOL6		cytoplasm
P64574	YGHW_ECOLI		cytoplasm
P64575	YGHW_ECOL6		cytoplasm
P64576	YGHW_ECO57		cytoplasm
P64577	YGHW_SHIFL		cytoplasm
P64578	YGJN_ECOLI		cytoplasm
P64579	YGJN_ECO57		cytoplasm
P64580	YGJN_SHIFL		cytoplasm
P64581	YQJD_ECOLI		cytoplasm
P64582	YQJD_ECOL6		cytoplasm
P64583	YQJD_ECO57		cytoplasm
P64584	YQJD_SHIFL		cytoplasm
P64586	YQJE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64587	YQJE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64588	YQJI_ECOLI		cytoplasm
P64589	YQJI_SHIFL		cytoplasm
P64591	YHAH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64593	YHAI_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64594	YHAV_ECOLI		cytoplasm
P64595	YHAV_ECO57		cytoplasm
P64596	YRAP_ECOLI	periplasm (potential).	periplasm
P64597	YRAP_ECOL6	periplasm (potential).	periplasm
P64598	YRAP_ECO57	periplasm (potential).	periplasm
P64599	YHBT_ECOLI		cytoplasm
P64600	YHBT_ECOL6		cytoplasm
P64601	YHBT_ECO57		cytoplasm
P64602	YRBB_ECOLI		cytoplasm
P64603	YRBB_SHIFL		cytoplasm
P64604	YRBD_ECOLI		cytoplasm
P64605	YRBD_ECO57		cytoplasm
P64607	YRBE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64608	YRBE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64609	YRBE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P64610	YRBL_ECOLI		cytoplasm
P64611	YRBL_SHIFL		cytoplasm
P64612	YHCM_ECOLI		cytoplasm
P64613	YHCM_ECO57		cytoplasm
P64614	YHCN_ECOLI		outer membrane
P64615	YHCN_ECOL6		outer membrane
P64616	YHCO_ECOLI		cytoplasm
P64617	YHCO_ECOL6		cytoplasm

P64618	YHCO_ECO57		cytoplasm
P64619	YHDU_ECOLI		inner membrane
P64620	YHDU_ECOL6		inner membrane
P64621	YHDU_ECO57		inner membrane
P64622	YHDV_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64623	YHDV_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64624	YHEO_ECOLI		cytoplasm
P64625	YHEO_ECOL6		cytoplasm
P64626	YHEO_SHIFL		cytoplasm
P64627	YHFL_ECOLI		inner membrane
P64628	YHFL_ECOL6		inner membrane
P64629	YHFL_ECO57		inner membrane
P64630	YHFL_SHIFL		inner membrane
P64631	YHFU_ECOLI		cytoplasm
P64632	YHFU_ECOL6		cytoplasm
P64633	YHFU_SHIFL		cytoplasm
P64634	YRFC_ECOLI		cytoplasm
P64635	YRFC_SHIFL		cytoplasm
P64636	YRFG_ECOLI		cytoplasm
P64637	YRFG_SHIFL		cytoplasm
P64638	YHGG_ECOLI		cytoplasm
P64639	YHGG_ECO57		cytoplasm
P64646	YJDO_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64647	YJDO_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64648	YJDO_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P64649	Y031_HELPY		cytoplasm
P64650	Y027_HELPJ		cytoplasm
P64651	Y085_HELPY		inner membrane
P64652	Y078_HELPJ		inner membrane
P64657	Y245_HELPY		cytoplasm
P64658	Y230_HELPJ		cytoplasm
P64659	Y386_HELPY		cytoplasm
P64660	Y995_HELPJ		cytoplasm
P64663	Y664_HELPY		cytoplasm
P64664	Y609_HELPJ		cytoplasm
P64665	Y1531_HELPY		cytoplasm
P64666	Y1419_HELPJ		cytoplasm
P65185	ISPH_HELPY		cytoplasm
P65186	ISPH_HELPJ		cytoplasm
P65191	ISPH_NEIMA		cytoplasm
P65192	ISPH_NEIMB		cytoplasm
P65193	ISPH_XYLFA		cytoplasm
P65194	ISPH_XYLFT		cytoplasm
P65195	ISPZ_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P65196	ISPZ_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65197	ISPZ_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65198	ISPZ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65199	ALCA_BORPE		cytoplasm
P65200	ALCA_BORPA		cytoplasm
P65207	KDGT1_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65208	KDGT1_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65215	KDSA_SALTY	cytoplasm (by similarity).	cytoplasm
P65216	KDSA_SALTI	cytoplasm (by similarity).	cytoplasm
P65217	KGUA_BRUME	cytoplasm (by similarity).	cytoplasm
P65218	KGUA_BRUSU	cytoplasm (by similarity).	cytoplasm
P65226	KHSE_SALTY	cytoplasm (potential).	cytoplasm
P65227	KHSE_SALTI	cytoplasm (potential).	cytoplasm
P65234	KPRS_NEIMA	cytoplasm (by similarity).	cytoplasm
P65235	KPRS_NEIMB	cytoplasm (by similarity).	cytoplasm
P65246	KTHY_SALTY		cytoplasm
P65247	KTHY_SALTI		cytoplasm
P65253	LLDP_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65254	LLDP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65264	LSPA_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P65265	LSPA_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P65275	LEUD_BRUME		cytoplasm
P65276	LEUD_BRUSU		cytoplasm
P65281	LIPA_BRUME	cytoplasm (potential).	cytoplasm
P65282	LIPA_BRUSU	cytoplasm (potential).	cytoplasm
P65290	YFGH_ECOLI	bacterial cell membrane; lipid-anchor (potential).	outer membrane
P65291	YFGH_ECO57	bacterial cell membrane; lipid-anchor (potential).	outer membrane
P65292	YGDI_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P65293	YGDI_ECOL6	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P65294	YGDR_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P65295	YGDR_ECOL6	bacterial cell membrane; lipid-anchor (potential).	inner membrane

P65296	YGDR_ECO57	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P65297	YGDR_SHIFL	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P65298	YQHH_ECOLI	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P65299	YQHH_SHIFL	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P65320	LPXA_BRUME	cytoplasm (by similarity).	cytoplasm
P65321	LPXA_BRUSU	cytoplasm (by similarity).	cytoplasm
P65322	LPXD_ECOL6		cytoplasm
P65323	LPXD_ECO57		cytoplasm
P65324	LPXD_SHIFL		cytoplasm
P65325	LPXK_BRUME		cytoplasm
P65326	LPXK_BRUSU		cytoplasm
P65350	MGSA_SALTY		cytoplasm
P65351	MGSA_SALTI		cytoplasm
P65359	MINC_SALTY		inner membrane
P65360	MINC_SALTI		inner membrane
P65361	MINE_BRUME		inner membrane
P65362	MINE_BRUSU		inner membrane
P65363	MINE_SALTY		inner membrane
P65364	MINE_SALTI		inner membrane
P65365	MINE_XYLFA		inner membrane
P65366	MINE_XYLFT		inner membrane
P65367	YQCA_ECOLI		cytoplasm
P65368	YQCA_ECO57		cytoplasm
P65369	YQCA_SHIFL		cytoplasm
P65382	MOAA_ECOL6		cytoplasm
P65383	MOAA_ECO57		cytoplasm
P65386	MOAA_SALTY		cytoplasm
P65387	MOAA_SALTI		cytoplasm
P65396	MOAE_BRUME		cytoplasm
P65397	MOAE_BRUSU		cytoplasm
P65398	MOAE_SALTY		cytoplasm
P65399	MOAE_SALTI		cytoplasm
P65417	MPGP_SALTY	cytoplasm (by similarity).	cytoplasm
P65418	MPGP_SALTI	cytoplasm (by similarity).	cytoplasm
P65427	MRAW_BRUME		cytoplasm
P65428	MRAW_BRUSU		cytoplasm
P65434	MRAZ_ECOL6		inner membrane
P65435	MRAZ_ECO57		inner membrane
P65447	MSRB_BRUME		cytoplasm
P65448	MSRB_BRUSU		cytoplasm
P65449	MSRB_SALTY		cytoplasm
P65450	MSRB_SALTI		cytoplasm
P65452	MURA_BRUME	cytoplasm (probable).	inner membrane
P65453	MURA_BRUSU	cytoplasm (probable).	inner membrane
P65454	MURA_SALTY	cytoplasm (probable).	inner membrane
P65455	MURA_SALTI	cytoplasm (probable).	inner membrane
P65468	MURC_BRUME	cytoplasm (probable).	cytoplasm
P65469	MURC_BRUSU	cytoplasm (probable).	cytoplasm
P65470	MURC_ECOL6	cytoplasm (probable).	cytoplasm
P65471	MURC_ECO57	cytoplasm (probable).	cytoplasm
P65489	MUTL_BRUME		cytoplasm
P65490	MUTL_BRUSU		cytoplasm

P65510	KEFG_SALTY		cytoplasm
P65511	KEFG_SALTI		cytoplasm
P65524	CVRA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P65525	CVRA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P65532	NDK_NEIMA	cytoplasm (by similarity).	cytoplasm
P65533	NDK_NEIMB	cytoplasm (by similarity).	cytoplasm
P65534	NDK_SALTY	cytoplasm (by similarity).	cytoplasm
P65535	NDK_SALTI	cytoplasm (by similarity).	cytoplasm
P65538	NDK_XANAC	cytoplasm (by similarity).	cytoplasm
P65539	NDK_XANCP	cytoplasm (by similarity).	cytoplasm
P65540	RNFE_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65541	RNFE_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65542	RNFA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65543	RNFA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65544	MNTH_BRUME	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P65545	MNTH_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P65546	NRDI_BRUME		inner membrane
P65547	NRDI_BRUSU		inner membrane
P65552	NUDD_ECO57		cytoplasm
P65553	NUDD_SHIFL		cytoplasm
P65554	NUDH_SALTY		cytoplasm
P65555	NUDH_SALTI		cytoplasm
P65556	YFCD_ECOLI		cytoplasm
P65557	YFCD_ECOL6		cytoplasm
P65558	YFCD_ECO57		cytoplasm
P65559	YFCD_SALTY		cytoplasm
P65560	YFCD_SALTI		cytoplasm
P65591	NUSG_NEIMA		cytoplasm
P65592	NUSG_NEIMB		cytoplasm
P65599	ORN_SALTY	cytoplasm (potential).	cytoplasm
P65600	ORN_SALTI	cytoplasm (potential).	cytoplasm
P65611	PYRB_BRUME		cytoplasm
P65612	PYRB_BRUSU		cytoplasm
P65615	PYRB_NEIMA		cytoplasm
P65616	PYRB_NEIMB		cytoplasm
P65622	OXAA_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P65623	OXAA_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65624	OXAA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65625	OXAA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P65639	BDM_ECOL6		cytoplasm
P65640	BDM_ECO57		cytoplasm
P65641	BDM_SALTY		cytoplasm
P65642	BDM_SALTI		cytoplasm
P65643	EUTT_ECOLI		cytoplasm
P65644	EUTT_ECOL6		cytoplasm
P65654	PANB_BRUME		cytoplasm
P65655	PANB_BRUSU		cytoplasm
P65662	PAND_SALTY		cytoplasm
P65663	PAND_SALTI		cytoplasm
P65672	CYSD_SALTY		cytoplasm
P65673	CYSD_SALTI		cytoplasm
P65680	PDXA_BRUME	cytoplasm (probable).	cytoplasm
P65681	PDXA_BRUSU	cytoplasm (probable).	cytoplasm
P65692	K6PF_SALTY	cytoplasm (by similarity).	cytoplasm
P65693	K6PF_SALTI	cytoplasm (by similarity).	cytoplasm
P65702	PGK_SALTY	cytoplasm (potential).	cytoplasm
P65703	PGK_SALTI	cytoplasm (potential).	cytoplasm
P65722	PHS_BRUME		cytoplasm
P65723	PHS_BRUSU		cytoplasm
P65736	PLSX_ECOL6		cytoplasm
P65737	PLSX_ECO57		cytoplasm
P65744	IPYR_BRUME	cytoplasm (by similarity).	cytoplasm
P65745	IPYR_BRUSU	cytoplasm (by similarity).	cytoplasm
P65748	IPYR_SALTY	cytoplasm (by similarity).	cytoplasm
P65749	IPYR_SALTI	cytoplasm (by similarity).	cytoplasm
P65750	IPYR_XYLFA	cytoplasm (by similarity).	cytoplasm
P65751	IPYR_XYLFT	cytoplasm (by similarity).	cytoplasm
P65764	FKBA_ECOL6	periplasm (by similarity).	periplasm
P65765	FKBA_ECO57	periplasm (by similarity).	periplasm
P65772	PPNK_NEIMA	cytoplasm (by similarity).	cytoplasm
P65773	PPNK_NEIMB	cytoplasm (by similarity).	cytoplasm
P65774	PPNK_SALTY	cytoplasm (by similarity).	cytoplasm
P65775	PPNK_SALTI	cytoplasm (by similarity).	cytoplasm
P65790	PROB_BRUME	cytoplasm (by similarity).	cytoplasm
P65791	PROB_BRUSU	cytoplasm (by similarity).	cytoplasm
P65792	PROB_SALTY	cytoplasm (by similarity).	cytoplasm
P65793	PROB_SALTI	cytoplasm (by similarity).	cytoplasm
P65799	HSLV_XYLFA	cytoplasm (by similarity).	cytoplasm
P65800	HSLV_XYLFT	cytoplasm (by similarity).	cytoplasm
P65803	PEPT_ECOL6	cytoplasm (by similarity).	extracellular
P65804	PEPT_ECO57	cytoplasm (by similarity).	extracellular
P65807	YGEY_ECOLI		cytoplasm
P65808	YGEY_ECOL6		cytoplasm
P65809	YGEY_ECO57		cytoplasm

P65812	HTPX_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P65813	HTPX_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P65814	HTPX_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P65817	HTPX_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P65818	HTPX_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
P65834	RLUD_ECOL6		cytoplasm
P65835	RLUD_SHIFL		cytoplasm
P65836	RLUD_SALTY		cytoplasm
P65837	RLUD_SALTI		cytoplasm
P65840	RSUA_SALTY		cytoplasm
P65841	RSUA_SALTI		cytoplasm
P65844	TRUA_ECOL6		cytoplasm
P65845	TRUA_ECO57		cytoplasm
P65852	TRUB_BORPE		cytoplasm
P65853	TRUB_BORBR		cytoplasm
P65863	PTH_BRUME	cytoplasm (by similarity).	cytoplasm
P65864	PTH_BRUSU	cytoplasm (by similarity).	cytoplasm
P65870	PTPS_ECOLI		extracellular
P65871	PTPS_ECO57		extracellular
P65872	PTPS_SHIFL		extracellular
P65876	PTHP_NEIMA	cytoplasm (by similarity).	cytoplasm
P65877	PTHP_NEIMB	cytoplasm (by similarity).	cytoplasm
P65878	PURA_BRUME	cytoplasm (by similarity).	cytoplasm
P65879	PURA_BRUSU	cytoplasm (by similarity).	cytoplasm
P65882	PURA_SALTY	cytoplasm (by similarity).	cytoplasm
P65883	PURA_SALTI	cytoplasm (by similarity).	cytoplasm
P65889	PUR7_SALTY		cytoplasm
P65890	PUR7_SALTI		cytoplasm
P65910	PYRE_BRUME		cytoplasm
P65911	PYRE_BRUSU		cytoplasm
P65914	PYRE_NEIMA		cytoplasm
P65915	PYRE_NEIMB		cytoplasm
P65921	PYRG_SALTY		cytoplasm
P65922	PYRG_SALTI		cytoplasm
P65931	PYRH_NEIMA		cytoplasm
P65932	PYRH_NEIMB		cytoplasm
P65933	PYRH_SALTY		cytoplasm
P65934	PYRH_SALTI		cytoplasm
P65948	QUEA_BRUME	cytoplasm (potential).	cytoplasm
P65949	QUEA_BRUSU	cytoplasm (potential).	cytoplasm
P65955	RADC_BRUME		cytoplasm
P65956	RADC_BRUSU		cytoplasm
P65957	RADC_ECOL6		cytoplasm
P65958	RADC_ECO57		cytoplasm
P65959	RADC_SHIFL		cytoplasm
P65960	RADC_SALTY		cytoplasm

P65961	RADC_SALTI		cytoplasm
P65973	RDGC_SALTY	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
P65974	RDGC_SALTI	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
P65975	RECA_BRUME	cytoplasm (by similarity).	cytoplasm
P65976	RECA_BRUSU	cytoplasm (by similarity).	cytoplasm
P65977	RECA_SALTY	cytoplasm (by similarity).	cytoplasm
P65978	RECA_SALTI	cytoplasm (by similarity).	cytoplasm
P65992	RECR_SALTY		cytoplasm
P65993	RECR_SALTI		cytoplasm
P65999	RECX_ECOL6	cytoplasm (potential).	inner membrane
P66000	RECX_ECO57	cytoplasm (potential).	inner membrane
P66001	RECX_SHIFL	cytoplasm (potential).	inner membrane
P66008	STHA_SALTY	cytoplasm (by similarity).	cytoplasm
P66009	STHA_SALTI	cytoplasm (by similarity).	cytoplasm
P66030	GCH2_SALTY		cytoplasm
P66031	GCH2_SALTI		cytoplasm
P66032	RIBB_SALTY		cytoplasm
P66033	RIBB_SALTI		cytoplasm
P66036	RISB_NEIMA		cytoplasm
P66037	RISB_NEIMB		cytoplasm
P66038	RISB_SALTY		cytoplasm
P66039	RISB_SALTI		cytoplasm
P66046	RL10_NEIMA		cytoplasm
P66047	RL10_NEIMB		cytoplasm
P66052	RL11_HELPY		cytoplasm
P66053	RL11_HELPJ		cytoplasm
P66067	RL14_HAEIN		cytoplasm
P66068	RL14_PASMU		cytoplasm
P66071	RL15_ECOL6		cytoplasm
P66072	RL15_ECO57		cytoplasm
P66073	RL15_SALTY		cytoplasm
P66074	RL15_SALTI		cytoplasm
P66075	RL15_SHIFL		cytoplasm
P66078	RL19_BRUME		cytoplasm
P66079	RL19_BRUSU		cytoplasm
P66087	RL1_NEIMA		cytoplasm
P66088	RL1_NEIMB		cytoplasm
P66101	RL20_BRUME		cytoplasm
P66102	RL20_BRUSU		cytoplasm
P66119	RL23_HELPY		cytoplasm
P66120	RL23_HELPJ		cytoplasm
P66129	RL27_NEIMA		cytoplasm
P66130	RL27_NEIMB		cytoplasm
P66131	RL27_SALTY		cytoplasm
P66132	RL27_SALTI		cytoplasm
P66140	RL28_BRUME		cytoplasm
P66141	RL28_BRUSU		cytoplasm
P66142	RL28_HELPY		cytoplasm
P66143	RL28_HELPJ		cytoplasm
P66150	RL28_NEIMA		cytoplasm
P66151	RL28_NEIMB		cytoplasm
P66160	RL28_XANAC		cytoplasm

P66161	RL28_XANCP		cytoplasm
P66162	RL28_XYLFA		cytoplasm
P66163	RL28_XYLFT		cytoplasm
P66164	RL29_BRUME		cytoplasm
P66165	RL29_BRUSU		cytoplasm
P66168	RL29_NEIMA		cytoplasm
P66169	RL29_NEIMB		cytoplasm
P66170	RL29_SALTY		cytoplasm
P66171	RL29_SALTI		cytoplasm
P66183	RL31_BRUME		cytoplasm
P66184	RL31_BRUSU		cytoplasm
P66185	RL31_HELPY		cytoplasm
P66186	RL31_HELPJ		cytoplasm
P66189	RL31_NEIMA		cytoplasm
P66190	RL31_NEIMB		cytoplasm
P66191	RL31_SALTY		cytoplasm
P66192	RL31_SALTI		cytoplasm
P66193	RL31B_SALTY		cytoplasm
P66194	RL31B_SALTI		cytoplasm
P66205	RL32_BRUME		cytoplasm
P66206	RL32_BRUSU		cytoplasm
P66215	RL33_BRUME		cytoplasm
P66216	RL33_BRUSU		cytoplasm
P66217	RL33_HELPY		cytoplasm
P66218	RL33_HELPJ		cytoplasm
P66225	RL33_NEIMA		cytoplasm
P66226	RL33_NEIMB		cytoplasm
P66238	RL33_XANAC		cytoplasm
P66239	RL33_XANCP		cytoplasm
P66240	RL33_XYLFA		cytoplasm
P66241	RL33_XYLFT		cytoplasm
P66265	RL35_BRUME		cytoplasm
P66266	RL35_BRUSU		cytoplasm
P66269	RL35_HELPY		cytoplasm
P66270	RL35_HELPJ		cytoplasm
P66273	RL35_NEIMA		cytoplasm
P66274	RL35_NEIMB		cytoplasm
P66282	RL35_XANAC		cytoplasm
P66283	RL35_XANCP		cytoplasm
P66284	RL35_XYLFA		cytoplasm
P66285	RL35_XYLFT		cytoplasm
P66313	RL6_SALTY		cytoplasm
P66314	RL6_SALTI		cytoplasm
P66324	RS10_BRUME		cytoplasm
P66325	RS10_BRUSU		cytoplasm
P66328	RS10_HELPY		cytoplasm
P66329	RS10_HELPJ		cytoplasm
P66332	RS10_NEIMA		cytoplasm
P66333	RS10_NEIMB		cytoplasm
P66346	RS10_VIBPA		cytoplasm
P66347	RS10_VIBVU		cytoplasm
P66348	RS10_XYLFA		cytoplasm
P66349	RS10_XYLFT		cytoplasm
P66350	RS11_BRUME		cytoplasm
P66351	RS11_BRUSU		cytoplasm
P66354	RS11_NEIMA		cytoplasm

P66355	RS11_NEIMB		cytoplasm
P66367	RS11_VIBCH		cytoplasm
P66368	RS11_VIBPA		cytoplasm
P66369	RS11_VIBVU		cytoplasm
P66374	RS12_NEIMA		cytoplasm
P66375	RS12_NEIMB		cytoplasm
P66379	RS12_XYLFA		cytoplasm
P66380	RS12_XYLFT		cytoplasm
P66381	RS13_BRUME		cytoplasm
P66382	RS13_BRUSU		cytoplasm
P66385	RS13_NEIMA		cytoplasm
P66386	RS13_NEIMB		cytoplasm
P66399	RS14_BRUME		cytoplasm
P66400	RS14_BRUSU		cytoplasm
P66407	RS14_NEIMA		cytoplasm
P66408	RS14_NEIMB		cytoplasm
P66409	RS14_SALTY		cytoplasm
P66410	RS14_SALTI		cytoplasm
P66431	RS15_SALTY		cytoplasm
P66432	RS15_SALTI		cytoplasm
P66433	RS16_BRUME		cytoplasm
P66434	RS16_BRUSU		cytoplasm
P66437	RS16_NEIMA		cytoplasm
P66438	RS16_NEIMB		cytoplasm
P66449	RS17_HELPY		cytoplasm
P66450	RS17_HELPJ		cytoplasm
P66451	RS17_SALTY		cytoplasm
P66452	RS17_SALTI		cytoplasm
P66453	RS18_BRUME		cytoplasm
P66454	RS18_BRUSU		cytoplasm
P66457	RS18_HAEIN		cytoplasm
P66458	RS18_HAEDU		cytoplasm
P66459	RS18_HELPY		cytoplasm
P66460	RS18_HELPJ		cytoplasm
P66478	RS18_VIBPA		cytoplasm
P66479	RS18_VIBVU		cytoplasm
P66486	RS19_NEIMA		cytoplasm
P66487	RS19_NEIMB		cytoplasm
P66488	RS19_RHIME		cytoplasm
P66489	RS19_BRUME		cytoplasm
P66490	RS19_BRUSU		cytoplasm
P66491	RS19_SALTY		cytoplasm
P66492	RS19_SALTI		cytoplasm
P66501	RS20_BRUME		cytoplasm
P66502	RS20_BRUSU		cytoplasm
P66507	RS20_NEIMA		cytoplasm
P66508	RS20_NEIMB		cytoplasm
P66512	RS20_XYLFA		cytoplasm
P66513	RS20_XYLFT		cytoplasm
P66514	RS21_BRUME		cytoplasm
P66515	RS21_BRUSU		cytoplasm
P66518	RS21_NEIMA		cytoplasm
P66519	RS21_NEIMB		cytoplasm
P66532	RS21_VIBCH		cytoplasm
P66533	RS21_VIBPA		cytoplasm
P66534	RS21_VIBVU		cytoplasm

P66535	RS21_XANAC		cytoplasm
P66536	RS21_XANCP		cytoplasm
P66539	RS2_NEIMA		cytoplasm
P66540	RS2_NEIMB		cytoplasm
P66541	RS2_SALTY		cytoplasm
P66542	RS2_SALTI		cytoplasm
P66546	RS2_XYLFA		cytoplasm
P66547	RS2_XYLFT		cytoplasm
P66550	RS3_NEIMA		cytoplasm
P66551	RS3_NEIMB		cytoplasm
P66560	RS4_NEIMA		cytoplasm
P66561	RS4_NEIMB		cytoplasm
P66570	RS5_BRUME		cytoplasm
P66571	RS5_BRUSU		cytoplasm
P66572	RS5_HELPY		cytoplasm
P66573	RS5_HELPJ		cytoplasm
P66576	RS5_NEIMA		cytoplasm
P66577	RS5_NEIMB		cytoplasm
P66586	RS5_XANAC		cytoplasm
P66587	RS5_XANCP		cytoplasm
P66593	RS6_SALTY		cytoplasm
P66594	RS6_SALTI		cytoplasm
P66604	RS6_XYLFA		cytoplasm
P66605	RS6_XYLFT		cytoplasm
P66606	RS7_ECOL6		cytoplasm
P66607	RS7_ECO57		cytoplasm
P66608	RS7_SHIFL		cytoplasm
P66609	RS7_HELPY		cytoplasm
P66610	RS7_HELPJ		cytoplasm
P66613	RS7_NEIMA		cytoplasm
P66614	RS7_NEIMB		cytoplasm
P66621	RS8_HELPY		cytoplasm
P66622	RS8_HELPJ		cytoplasm
P66627	RS8_NEIMA		cytoplasm
P66628	RS8_NEIMB		cytoplasm
P66637	RS9_HELPY		cytoplasm
P66638	RS9_HELPJ		cytoplasm
P66641	RS9_NEIMA		cytoplasm
P66642	RS9_NEIMB		cytoplasm
P66643	RS9_SALTY		cytoplasm
P66644	RS9_SALTI		cytoplasm
P66651	RIMM_BRUME	cytoplasm (potential).	cytoplasm
P66652	RIMM_BRUSU	cytoplasm (potential).	cytoplasm
P66664	RNC_BRUME	cytoplasm (by similarity).	cytoplasm
P66665	RNC_BRUSU	cytoplasm (by similarity).	cytoplasm
P66673	RNH_BRUME	cytoplasm (potential).	extracellular
P66674	RNH_BRUSU	cytoplasm (potential).	extracellular
P66679	RNPH_ECOL6		cytoplasm
P66680	RNPH_ECO57		cytoplasm
P66681	RNT_ECOL6		cytoplasm
P66682	RNT_ECO57		cytoplasm
P66683	RNT_SHIFL		cytoplasm
P66684	RNT_SALTY		cytoplasm
P66685	RNT_SALTI		cytoplasm
P66686	RNPA_SALTY		extracellular
P66687	RNPA_SALTI		extracellular

P66692	RPIA_SALTY		cytoplasm
P66693	RPIA_SALTI		cytoplasm
P66703	RPOA_NEIMA		cytoplasm
P66704	RPOA_NEIMB		cytoplasm
P66712	RPOA_XYLFA		cytoplasm
P66713	RPOA_XYLFT		cytoplasm
P66723	RPOZ_NEIMA		cytoplasm
P66724	RPOZ_NEIMB		cytoplasm
P66732	RPOZ_XANAC		cytoplasm
P66733	RPOZ_XANCP		cytoplasm
P66736	RRF_NEIMA	cytoplasm (by similarity).	cytoplasm
P66737	RRF_NEIMB	cytoplasm (by similarity).	cytoplasm
P66738	RRF_SALTY	cytoplasm (by similarity).	cytoplasm
P66739	RRF_SALTI	cytoplasm (by similarity).	cytoplasm
P66742	RUVA_BRUME		cytoplasm
P66743	RUVA_BRUSU		cytoplasm
P66746	RUVA_SALTY		cytoplasm
P66747	RUVA_SALTI		cytoplasm
P66755	RUVB_SALTY		cytoplasm
P66756	RUVB_SALTI		cytoplasm
P66762	RUVC_SALTY		cytoplasm
P66763	RUVC_SALTI		cytoplasm
P66793	SELD_ECOL6		cytoplasm
P66794	SELD_ECO57		cytoplasm
P66795	QSEB_SALTY	cytoplasm (potential).	cytoplasm
P66796	QSEB_SALTI	cytoplasm (potential).	cytoplasm
P66797	UVRY_ECOL6	cytoplasm (probable).	cytoplasm
P66798	UVRY_ECO57	cytoplasm (probable).	cytoplasm
P66817	DIAA_ECOLI		cytoplasm
P66818	DIAA_ECO57		cytoplasm
P66819	DIAA_SHIFL		cytoplasm
P66820	SMG_NEIMA		cytoplasm
P66821	SMG_NEIMB		cytoplasm
P66822	SMG_SALTY		cytoplasm
P66823	SMG_SALTI		cytoplasm
P66826	SODC_BRUME	periplasm (by similarity).	periplasm
P66827	SODC_BRUSU	periplasm (by similarity).	periplasm
P66828	SODM_ECO57		periplasm
P66829	SODM_SHIFL		periplasm
P66833	SPEE_ECOL6		cytoplasm
P66834	SPEE_ECO57		cytoplasm
P66846	SSB_BORPE		cytoplasm
P66847	SSB_BORBR		cytoplasm
P66848	SSB_NEIMA		cytoplasm
P66849	SSB_NEIMB		cytoplasm
P66858	SSRP_BRUME	cytoplasm (potential).	cytoplasm
P66859	SSRP_BRUSU	cytoplasm (potential).	cytoplasm
P66867	SUCC_BRUME		cytoplasm
P66868	SUCC_BRUSU		cytoplasm
P66869	SUCC_SALTY		cytoplasm
P66870	SUCC_SALTI		cytoplasm
P66879	SURE_BRUME	cytoplasm (potential).	periplasm
P66880	SURE_BRUSU	cytoplasm (potential).	periplasm
P66881	SURE_SALTY	cytoplasm (potential).	periplasm
P66882	SURE_SALTI	cytoplasm (potential).	periplasm

P66887	TATA_BRUME	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P66888	TATA_BRUSU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P66891	TATA_NEIMA	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P66892	TATA_NEIMB	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P66899	DPAL_ECOLI		cytoplasm
P66900	DPAL_ECOL6		cytoplasm
P66901	DPAL_ECO57		cytoplasm
P66928	THIO_HELPY		cytoplasm
P66929	THIO_HELPJ		cytoplasm
P66932	TIG_SALTY		cytoplasm
P66933	TIG_SALTI		cytoplasm
P66948	YFGC_ECOLI		inner membrane
P66949	YFGC_SHIFL		inner membrane
P66950	YFGC_SALTY		inner membrane
P66951	YFGC_SALTI		inner membrane
P66955	TALB_SALTY	cytoplasm (probable).	cytoplasm
P66956	TALB_SALTI	cytoplasm (probable).	cytoplasm
P66974	YFHQ_SALTY		cytoplasm
P66975	YFHQ_SALTI		cytoplasm
P66988	TRPC_BRUME		cytoplasm
P66989	TRPC_BRUSU		cytoplasm
P66994	TRPD_NEIMA		cytoplasm
P66995	TRPD_NEIMB		cytoplasm
P67003	TRPF_BRUME		cytoplasm
P67004	TRPF_BRUSU		cytoplasm
P67042	TYSY_BRUME	cytoplasm (by similarity).	cytoplasm
P67043	TYSY_BRUSU	cytoplasm (by similarity).	cytoplasm
P67053	COXX_BRUME	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P67054	COXX_BRUSU	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P67066	WECC_ECO57		cytoplasm
P67067	WECC_SHIFL		cytoplasm
P67073	UNG_SALTY	cytoplasm (by similarity).	extracellular
P67074	UNG_SALTI	cytoplasm (by similarity).	extracellular
P67080	YGGG_ECOLI		cytoplasm
P67081	YGGG_ECOL6		cytoplasm
P67082	YGGG_ECO57		cytoplasm
P67087	YRAL_ECOLI		cytoplasm
P67088	YRAL_ECO57		cytoplasm
P67089	USPD_ECOL6	cytoplasm (by similarity).	cytoplasm
P67090	USPD_ECO57	cytoplasm (by similarity).	cytoplasm
P67091	USPF_SALTY		cytoplasm
P67092	USPF_SALTI		cytoplasm
P67093	USPG_SALTY		cytoplasm
P67094	USPG_SALTI		cytoplasm

P67095	YFCE_ECOLI		cytoplasm
P67096	YFCE_ECOL6		cytoplasm
P67097	YFCE_ECO57		cytoplasm
P67100	TUSA_SALTY	cytoplasm (by similarity).	cytoplasm
P67101	TUSA_SALTI	cytoplasm (by similarity).	cytoplasm
P67102	Y882_NEIMA		cytoplasm
P67103	Y681_NEIMB		cytoplasm
P67128	YGDQ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67129	YGDQ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67132	Y1974_BRUME		inner membrane
P67133	Y2156_BRUSU		inner membrane
P67144	YHGN_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67145	YHGN_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67154	YQFA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67155	YQFA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67156	YQFA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67161	YGIH_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67162	YGIH_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67173	Y321_BRUME		inner membrane
P67174	Y1717_BRUSU		inner membrane
P67175	YEBC_ECOL6		inner membrane
P67176	YEBC_ECO57		inner membrane
P67179	YEBC_SALTY		inner membrane
P67180	YEBC_SALTI		inner membrane
P67193	Y2060_BRUME		cytoplasm
P67194	Y2067_BRUSU		cytoplasm
P67197	YDIA_SALTY		cytoplasm
P67198	YDIA_SALTI		cytoplasm
P67212	Y1968_BRUME		inner membrane
P67213	Y2162_BRUSU		inner membrane
P67216	Y1895_NEIMA		inner membrane
P67217	Y1641_NEIMB		inner membrane
P67218	YHBC_SALTY		inner membrane
P67219	YHBC_SALTI		inner membrane
P67228	Y1769_BRUME		inner membrane
P67229	Y178_BRUSU		inner membrane

P67244	YQHA_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P67245	YQHA_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P67246	YFCN_SALTY		inner membrane
P67247	YFCN_SALTI		inner membrane
P67258	Y1005_NEIMA		inner membrane
P67259	Y796_NEIMB		inner membrane
P67260	Y1909_BRUME		inner membrane
P67261	Y033_BRUSU		inner membrane
P67270	YBGI_SALTY		extracellular
P67271	YBGI_SALTI		extracellular
P67286	YBJQ_SALTY		inner membrane
P67287	YBJQ_SALTI		inner membrane
P67302	Y549_NEIMA		inner membrane
P67303	Y1906_NEIMB		inner membrane
P67311	NRDR_BRUME		cytoplasm
P67312	NRDR_BRUSU		cytoplasm
P67334	YAII_SALTY		inner membrane
P67335	YAII_SALTI		inner membrane
P67338	YOA_H_ECOLI		inner membrane
P67339	YOA_H_ECOL6		inner membrane
P67340	YOA_H_ECO57		inner membrane
P67341	YMDB_SALTY		inner membrane
P67342	YMDB_SALTI		inner membrane
P67347	YEDY_SALTY		cytoplasm
P67348	YEDY_SALTI		cytoplasm
P67349	YHBQ_SALTY		cytoplasm
P67350	YHBQ_SALTI		cytoplasm
P67366	YCFP_SALTY		inner membrane
P67367	YCFP_SALTI		inner membrane
P67384	UPPP_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67385	UPPP_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67386	UPPP_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67387	UPPP_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67388	UPPP_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67389	UPPP_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67408	URK_SALTY	cytoplasm (by similarity).	cytoplasm
P67409	URK_SALTI	cytoplasm (by similarity).	cytoplasm
P67430	YDHM_ECOLI		cytoplasm
P67431	YDHM_ECOL6		cytoplasm

P67445	YGFO_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67446	YGFO_ECO57	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P67454	EX7S_BRUME	cytoplasm (by similarity).	cytoplasm
P67455	EX7S_BRUSU	cytoplasm (by similarity).	cytoplasm
P67458	EX7S_SALTY	cytoplasm (by similarity).	cytoplasm
P67459	EX7S_SALTI	cytoplasm (by similarity).	cytoplasm
P67470	ZUPT_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P67471	ZUPT_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P67479	Y1673_BRUME		cytoplasm
P67480	Y247_BRUSU		cytoplasm
P67481	YACG_SALTY		cytoplasm
P67482	YACG_SALTI		cytoplasm
P67496	RUVX_XYLFA	cytoplasm (potential).	cytoplasm
P67497	RUVX_XYLFT	cytoplasm (potential).	cytoplasm
P67516	Y211_BRUME		inner membrane
P67517	Y1840_BRUSU		inner membrane
P67518	Y418_NEIMA		inner membrane
P67519	Y2022_NEIMB		inner membrane
P67520	YBEA_SALTY		inner membrane
P67521	YBEA_SALTI		inner membrane
P67524	DNAT_SALTY		nucleoid
P67525	DNAT_SALTI		nucleoid
P67535	Y1380_NEIMA		inner membrane
P67536	Y1218_NEIMB		inner membrane
P67537	YBED_SALTY		inner membrane
P67538	YBED_SALTI		inner membrane
P67539	PUR9_BRUME		cytoplasm
P67540	PUR9_BRUSU		cytoplasm
P67551	YAEP_SALTY		inner membrane
P67552	YAEP_SALTI		inner membrane
P67553	YNFC_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P67554	YNFC_SHIFL	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P67555	OPGG_ECOL6	periplasm (by similarity).	periplasm
P67556	OPGG_ECO57	periplasm (by similarity).	periplasm
P67557	OPGG_SALTY	periplasm (by similarity).	periplasm
P67558	OPGG_SALTI	periplasm (by similarity).	periplasm
P67559	OPGC_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P67560	OPGC_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P67567	Y1669_BRUME		inner membrane
P67568	Y251_BRUSU		inner membrane

P67602	YOBD_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67603	YQFB_ECOLI		inner membrane
P67604	YQFB_ECO57		inner membrane
P67605	YEEX_SALTY		inner membrane
P67606	YEEX_SALTI		inner membrane
P67613	LFTR_SALTY	cytoplasm (by similarity).	cytoplasm
P67614	LFTR_SALTI	cytoplasm (by similarity).	cytoplasm
P67615	FETP_NEIMA		cytoplasm
P67616	FETP_NEIMB		cytoplasm
P67617	FETP_SALTY		cytoplasm
P67618	FETP_SALTI		cytoplasm
P67624	YHEU_ECOLI		inner membrane
P67625	YHEU_ECOL6		inner membrane
P67626	YHEU_ECO57		inner membrane
P67627	YHEU_SHIFL		inner membrane
P67638	FRDC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67639	FRDC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67645	FRDD_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67646	FRDD_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67649	CLPS_SALTY		cytoplasm
P67650	CLPS_SALTI		cytoplasm
P67651	RRAA_SALTY		cytoplasm
P67652	RRAA_SALTI		cytoplasm
P67653	KDSC_ECOL6		cytoplasm
P67654	KDSC_SHIFL		cytoplasm
P67660	YHAJ_ECOLI		cytoplasm
P67661	YHAJ_ECO57		cytoplasm
P67662	AAER_ECOLI		cytoplasm
P67663	AAER_ECOL6		cytoplasm
P67664	AAER_ECO57		cytoplasm
P67673	PRIB_BORPE		cytoplasm
P67674	PRIB_BORBR		cytoplasm
P67675	PRIB_BORPA		cytoplasm
P67676	PRIB_ECO57		cytoplasm
P67677	PRIB_SHIFL		cytoplasm
P67678	ENGC_BORPE		cytoplasm
P67679	ENGC_BORBR		cytoplasm
P67680	ENGC_BORPA		cytoplasm
P67686	USPB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67687	USPB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P67688	PRMA_SALTY	cytoplasm (by similarity).	cytoplasm
P67689	PRMA_SALTI	cytoplasm (by similarity).	cytoplasm

P67690	Y3816_BORPE		inner membrane
P67691	Y4433_BORBR		inner membrane
P67692	Y3960_BORPA		inner membrane
P67693	YACF_SALTY		inner membrane
P67694	YACF_SALTI		inner membrane
P67695	Y1620_VIBVU		inner membrane
P67696	Y2783_VIBVY		inner membrane
P67697	YDCQ_ECOLI		cytoplasm
P67698	YDCQ_ECO57		cytoplasm
P67699	YDDM_ECOLI		cytoplasm
P67700	YDDM_ECOL6		cytoplasm
P67701	YGJM_ECOLI		cytoplasm
P67702	YGJM_ECOL6		cytoplasm
P67703	YGJM_SHIFL		cytoplasm
P67706	FLIE_BRUME		flagellar
P67707	FLIE_BRUSU		flagellar
P67708	FLIE_HELPY		flagellar
P67709	FLIE_HELPJ		flagellar
P67710	MNTR_SALTY	cytoplasm (probable).	cytoplasm
P67711	MNTR_SALTI	cytoplasm (probable).	cytoplasm
P67714	NDPA_SALTY	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
P67715	NDPA_SALTI	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
P67729	YFEO_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P67730	YFEO_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P67735	NANR_SALTY		cytoplasm
P67736	NANR_SALTI		cytoplasm
P67762	YHBP_ECOLI		inner membrane
P67763	YHBP_SHIFL		inner membrane
P67825	CUTC_SHIFL	cytoplasm (probable).	periplasm
P67826	CUTC_ECOLI	cytoplasm (probable).	periplasm
P67897	RS19_ACTAC		cytoplasm
P67898	RS19_HAEDU		cytoplasm
P67899	RS19_HAEIN		cytoplasm
P67900	RS19_PASMU		cytoplasm
P67901	RS10_HAEIN		cytoplasm
P67902	RS10_PASMU		cytoplasm
P67903	RS10_PHOLL		cytoplasm
P67904	RS10_SALTY		cytoplasm
P67905	RS10_SALTI		cytoplasm
P67906	RS10_YERPE		cytoplasm
P67910	HLDD_ECOLI		cytoplasm
P67911	HLDD_ECO57		cytoplasm
P67912	HLDD_SALTY		cytoplasm
P67913	HLDD_SALTI		cytoplasm
P67916	RL35_PASMU		cytoplasm
P67917	RL35_HAEIN		cytoplasm
P67918	BLA1_HAEIN		extracellular
P67919	BLA1_PASHA		extracellular

P67920	BLA1_ACTPL		extracellular
P68066	GRCA_ECOLI		cytoplasm
P68067	GRCA_ECO57		cytoplasm
P68188	MALK_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P68206	YJBJ_ECOLI		cytoplasm
P68207	YJBJ_ECO57		cytoplasm
P68208	YJBJ_SHIFL		cytoplasm
P68397	TADA_SHIFL		cytoplasm
P68398	TADA_ECOLI		cytoplasm
P68567	YHIQ_ECOLI		inner membrane
P68568	YHIQ_ECO57		inner membrane
P68587	YSCP_YERPE	cytoplasm (potential).	extracellular
P68588	PSAE_YERPE		cytoplasm
P68589	PSAE_YERPS		cytoplasm
P68590	YSCH_YERPE		extracellular
P68640	YOPN_YERPE	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P68641	ASCD_YERPE		cytoplasm
P68644	FIXC_ECOLI		periplasm
P68645	FIXC_ECOL6		periplasm
P68646	FIXX_ECOLI		periplasm
P68647	FIXX_ECOL6		periplasm
P68648	FIXX_ECO57		periplasm
P68649	HEAD_ECOL6		flagellar
P68651	NINB_ECOL6		cytoplasm
P68653	TERS_ECOL6		cytoplasm
P68655	VG03_ECOL6		cytoplasm
P68657	EA10_ECOL6		cytoplasm
P68659	VHTJ_ECOL6		cytoplasm
P68661	YBCO_ECOLI		cytoplasm
P68663	YQ54_SALTI		cytoplasm
P68665	YQ55_SALTI		cytoplasm
P68667	GTRB_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P68675	VHED_XANCP		cytoplasm
P68679	RS21_ECOLI		cytoplasm
P68680	RS21_ECOL6		cytoplasm
P68681	RS21_ECO57		cytoplasm
P68682	RS21_PHOLL		cytoplasm
P68683	RS21_SALTI		cytoplasm
P68684	RS21_SALTY		cytoplasm
P68685	RS21_SHIFL		cytoplasm
P68686	RS21_YERPE		cytoplasm
P68688	GLRX1_ECOLI		cytoplasm
P68689	GLRX1_SHIFL		cytoplasm
P68700	ATPL_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P68701	ATPL_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P68702	ATPL_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P68703	ATPL_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P68704	ATPL_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P68705	ATPL_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P68706	ATPL_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P68737	NFI_SALTI	cytoplasm (by similarity).	cytoplasm
P68738	NFI_SALTY	cytoplasm (by similarity).	cytoplasm
P68740	NFI_ECOL6	cytoplasm (by similarity).	cytoplasm
P68741	NFI_ECO57	cytoplasm (by similarity).	cytoplasm
P68746	OTC2_PSESF		cytoplasm
P68747	OTC2_PSESH		cytoplasm
P68766	AMPA_ECOL6		cytoplasm
P68767	AMPA_ECOLI		cytoplasm
P68768	AMPA_ECO57		cytoplasm
P68879	IPGE_SHISO	cytoplasm (by similarity).	cytoplasm
P68918	RL25_SHIFL		cytoplasm
P68919	RL25_ECOLI		cytoplasm
P68995	RL30_AGRT5		cytoplasm
P69049	SODC_LEGPN	periplasm (by similarity).	periplasm
P69050	CH60_LEGPN	cytoplasm (by similarity).	cytoplasm
P69053	LLY_LEGPN		extracellular
P69055	DHSC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69064	SICA_SALDU	cytoplasm (probable).	cytoplasm
P69065	SICA_SALTI	cytoplasm (probable).	cytoplasm
P69066	SICA_SALTY	cytoplasm (probable).	cytoplasm
P69211	MDTI_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69213	MDTJ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69214	MDTJ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69228	BAER_ECOLI	cytoplasm (potential).	cytoplasm
P69229	BAER_ECOL6	cytoplasm (potential).	cytoplasm
P69331	CITD_ECO57	cytoplasm (by similarity).	cytoplasm
P69340	MDTB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P69342	INVF_SALTI		cytoplasm
P69343	INVF_SALTY		cytoplasm
P69344	KAD_NEIME	cytoplasm (by similarity).	cytoplasm
P69346	YEFM_ECOLI		cytoplasm

P69347	YEFM_ECOL6		cytoplasm
P69348	YOEB_ECOLI		extracellular
P69349	YOEB_SHIFL		extracellular
P69368	MDTH_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69381	FIEF_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69405	RCSA_ECOLI		cytoplasm
P69406	RCSA_ECO57		cytoplasm
P69407	RCSB_ECOLI		cytoplasm
P69408	RCSB_ECOL6		cytoplasm
P69409	RCSB_ECO57		cytoplasm
P69410	RCSB_SHIFL		cytoplasm
P69412	RCSF_ECOL6	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
P69413	MERR_PSESP		cytoplasm
P69424	TATC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69426	TATB_ECO57	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P69427	TATB_SHIFL	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P69429	TATA_ECOL6	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P69430	TATA_ECO57	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P69431	TATA_SHIFL	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P69433	PGAD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69434	PGAA_ECOLI	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P69435	PGAA_ECO57	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P69442	KAD_ECO57	cytoplasm (by similarity).	cytoplasm
P69451	LCFA_ECOLI	membrane; peripheral membrane protein (by similarity). partially membrane-associated (by similarity).	cytoplasm

P69452	LCFA_ECOL6	membrane; peripheral membrane protein (by similarity). partially membrane-associated (by similarity).	cytoplasm
P69489	CUTA_ECO57	cytoplasm (by similarity).	periplasm
P69490	CCME_ECOLI	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential). stabilized by ccmd in the membrane.	inner membrane
P69491	CCME_ECOL6	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
P69492	CCME_ECO57	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
P69493	CCME_SHIFL	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
P69505	RCMNS_SHIFL	cytoplasm (by similarity).	cytoplasm
P69680	AMTB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69737	RHAD_SALT1	cytoplasm (by similarity).	extracellular
P69740	MBHS_ECOL6	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	periplasm
P69772	PAD1_ECO57		cytoplasm
P69773	PAD1_ECO11		cytoplasm
P69774	PAD1_ECOLI		cytoplasm
P69782	HOSA_ECO27		cytoplasm
P69784	PTGA_ECOL6	cytoplasm (by similarity).	cytoplasm
P69785	PTGA_SHIFL	cytoplasm (by similarity).	cytoplasm
P69787	PTGCB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69788	PTGCB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69789	PTXB_ECOLI		inner membrane
P69790	PTXB_SHIFL		inner membrane
P69792	PTQA_ECOL6	cytoplasm (by similarity).	cytoplasm
P69793	PTQA_ECO57	cytoplasm (by similarity).	cytoplasm
P69794	PTQA_SHIFL	cytoplasm (by similarity).	cytoplasm
P69796	PTQB_ECOL6	cytoplasm (by similarity).	cytoplasm
P69798	PTNAB_ECOL6	cytoplasm (by similarity).	inner membrane
P69799	PTNAB_ECO57	cytoplasm (by similarity).	inner membrane
P69800	PTNAB_SHIFL	cytoplasm (by similarity).	inner membrane
P69802	PTNC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

P69803	PTNC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69804	PTNC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69806	PTND_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69807	PTND_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69808	PTFB1_ECOLI	cytoplasm (potential).	inner membrane
P69809	PTFB1_ECOL6	cytoplasm (potential).	inner membrane
P69810	PTFB1_ECO57	cytoplasm (potential).	inner membrane
P69812	PTFAH_ECO57	cytoplasm (by similarity).	cytoplasm
P69813	PTKA_ECOL6	cytoplasm (potential).	cytoplasm
P69814	PTKA_ECO57	cytoplasm (potential).	cytoplasm
P69815	PTSN_SHIFL	cytoplasm (by similarity).	cytoplasm
P69816	PTFB2_ECOLI	cytoplasm (potential).	inner membrane
P69817	PTFB2_ECOL6	cytoplasm (potential).	inner membrane
P69818	PTFB2_ECO57	cytoplasm (potential).	inner membrane
P69820	ULAC_ECOLI	cytoplasm (probable).	cytoplasm
P69821	ULAC_ECO57	cytoplasm (probable).	cytoplasm
P69822	ULAB_ECOLI	cytoplasm (potential).	cytoplasm
P69823	ULAB_ECO57	cytoplasm (potential).	cytoplasm
P69824	PTMA_ECOLI	cytoplasm (probable).	cytoplasm
P69825	PTMA_ECO57	cytoplasm (probable).	cytoplasm
P69827	PTMCB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69828	PTKA_ECOLI	cytoplasm (potential).	cytoplasm
P69830	PTQB_ECO57	cytoplasm (by similarity).	cytoplasm
P69832	PTKC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69833	LGT_EHRRW	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69854	DMSD_ECO57	under anaerobic conditions: bacterial cell inner membrane; peripheral membrane protein (by similarity). under aerobic conditions; cytoplasm. mainly (by similarity).	cytoplasm
P69855	DMSD_SHIFL		cytoplasm
P69857	NANC_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P69858	NANC_SHIFL	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P69874	POTA_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

P69875	POTA_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P69876	POTA_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P69877	POTA_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P69902	FCTA_ECOLI		cytoplasm
P69903	FCTA_SHIFL		cytoplasm
P69908	DCEA_ECOLI		cytoplasm
P69909	DCEA_ECOL6		cytoplasm
P69911	DCEB_ECO57		cytoplasm
P69912	DCEB_SHIFL		cytoplasm
P69913	CSRA_ECOLI		cytoplasm
P69914	CSRA_ECOL6		cytoplasm
P69915	CSRA_ECO57		cytoplasm
P69916	CSRA_SALTI		cytoplasm
P69917	CSRA_SALTY		cytoplasm
P69918	CSRA_SHIFL		cytoplasm
P69920	CSRA_PSEFL		cytoplasm
P69921	CSRA2_PSESM		cytoplasm
P69923	FUCI_SHIFL	cytoplasm (by similarity).	cytoplasm
P69924	RIR2_ECOLI		cytoplasm
P69925	RIR2_ECO57		cytoplasm
P69931	HDA_ECOLI		cytoplasm
P69932	HDA_ECOL6		nucleoid
P69933	HDA_ECO57		nucleoid
P69934	HDA_SHIFL		nucleoid
P69935	YDFG_SALTI		cytoplasm
P69936	YDFG_SALTY		cytoplasm
P69938	SUGE_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P69953	BIOD2_YERPE		cytoplasm
P69954	BIOD2_YERPS		cytoplasm
P69957	LCRG_YERPE		cytoplasm
P69958	LCRG_YERPS		cytoplasm
P69959	LCRR_YERPE		extracellular
P69960	LCRR_YERPS		extracellular
P69961	LCRS_YERPE		cytoplasm
P69962	LCRS_YERPS		cytoplasm
P69963	RL23_YERPE		cytoplasm
P69964	RL23_YERPS		cytoplasm
P69965	PSAB_YERPE	periplasm (by similarity).	periplasm
P69966	PSAB_YERPS	periplasm (by similarity).	periplasm
P69967	TYEA_YEREN		cytoplasm
P69968	TYEA_YERPE		cytoplasm
P69969	TYEA_YERPS		cytoplasm
P69970	YSCI_YERPE		extracellular
P69971	YSCI_YERPS		extracellular
P69972	YSCJ_YERPE	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
P69974	YSCK_YERPE		extracellular
P69975	YSCK_YERPS		extracellular

P69976	YSCL_YERPE		extracellular
P69977	YSCL_YERPS		extracellular
P69978	YSCM_YERPE		extracellular
P69979	YSCM_YERPS		extracellular
P69980	YSCR_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P69981	YSCR_YERPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P69982	YSCS_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P69983	YSCS_YERPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P69984	YSCT_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P69985	YSCT_YERPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P69986	YSCU_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P69987	YSCU_YERPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P69988	SLYA_YERPE		cytoplasm
P69989	SLYA_YERPS		cytoplasm
P69990	URE2_YERPE	cytoplasm (by similarity).	cytoplasm
P69991	URE2_YERPS	cytoplasm (by similarity).	cytoplasm
P69992	UREG_YERPE		cytoplasm
P69993	UREG_YERPS		cytoplasm
P69994	URE3_YERPE	cytoplasm (by similarity).	cytoplasm
P69995	URE3_YERPS	cytoplasm (by similarity).	cytoplasm
P70714	CYSQ_ACTAC	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	cytoplasm
P70715	6PGL_ACTAC		cytoplasm
P70718	6PGD_ACTAC		cytoplasm
P70720	FABG_ACTAC		cytoplasm
P70727	NIFS_AZOBR		cytoplasm
P70728	NIFV_AZOBR		cytoplasm
P70730	CLPX_AZOBR		cytoplasm
P70736	Y3023_ACIAD		cytoplasm
P70753	BGAL_ACTPL		cytoplasm
P70773	ICIA_AERSA		cytoplasm
P70774	ASAI_AERSA		cytoplasm
P70775	YGGA_AERSA	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P70785	TTUA3_AGRVI		cytoplasm

P70786	TUB3_AGRVI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P70787	TTUC2_AGRVI	cytoplasm (by similarity).	cytoplasm
P70788	TTUD3_AGRVI		cytoplasm
P70789	KPYK1_AGRVI		cytoplasm
P70790	YTUX_AGRVI		cytoplasm
P70791	YTUY_AGRVI	periplasm (potential).	periplasm
P70792	TTUC4_AGRVI	cytoplasm (by similarity).	cytoplasm
P70793	YTZ3_AGRVI		cytoplasm
P70794	YTZ1_AGRVI		inner membrane
P70795	YZ2R_AGRVI		inner membrane
P70796	YTZ4_AGRVI		cytoplasm
P70799	ALGF_AZOVI	periplasm (by similarity).	periplasm
P70805	ALGG_AZOVI	periplasm (potential).	extracellular
P70917	HSPA_BRAJA		cytoplasm
P70918	HSPB_BRAJA		cytoplasm
P70920	ACON_BRAJA		cytoplasm
P70943	RS21_BURPS		cytoplasm
P71128	CYSM_CAMJE		cytoplasm
P71129	CHEY_CAMJE	cytoplasm (potential).	cytoplasm
P71154	COAD_CHRVI	cytoplasm (by similarity).	cytoplasm
P71156	NIFH_DESGI		cytoplasm
P71165	FLAV_DESVM		cytoplasm
P71169	FLAW_ENTAG		cytoplasm
P71198	TRAM5_ECOLI		cytoplasm
P71229	HYFR_ECOLI		cytoplasm
P71237	WCAC_ECOLI		cytoplasm
P71239	WCAE_ECOLI		cytoplasm
P71242	WCAK_ECOLI		cytoplasm
P71243	WCAL_ECOLI		cytoplasm
P71244	WCAM_ECOLI		inner membrane
P71296	YAGM_ECOLI		cytoplasm
P71297	YAGN_ECOLI		extracellular
P71298	INTF_ECOLI		cytoplasm
P71301	YK GK_ECOLI		cytoplasm
P71311	Y AIS_ECOLI		cytoplasm
P71318	OXYR_PECCC		cytoplasm
P71319	METF_PECCC		cytoplasm
P71331	DNAK_ACTAC		cytoplasm
P71333	FUR_HAEDU	cytoplasm (by similarity).	cytoplasm
P71335	Y004_HAEIN		inner membrane
P71336	Y052_HAEIN	periplasm (probable).	periplasm
P71338	FBPB2_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P71339	Y112_HAEIN		cytoplasm
P71340	NANE_HAEIN		cytoplasm
P71342	NQRE_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P71344	T1SI_HAEIN		cytoplasm
P71346	RAIA_HAEIN		cytoplasm
P71348	Y286_HAEIN	cytoplasm (by similarity).	cytoplasm
P71349	SDHL_HAEIN		cytoplasm
P71350	THIE_HAEIN		cytoplasm

P71351	VAPD_HAEIN		extracellular
P71353	Y568_HAEIN		cytoplasm
P71354	TRKA_HAEIN		inner membrane
P71355	Y663_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P71356	Y687_HAEIN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P71357	Y710_HAEIN		cytoplasm
P71358	CYAY_HAEIN		inner membrane
P71359	RECQ_HAEIN		cytoplasm
P71360	Y878_HAEIN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P71362	DDC_HAEIN		cytoplasm
P71363	Y947_HAEIN		cytoplasm
P71364	Y1015_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P71365	Y1050_HAEIN		extracellular
P71366	T3MH_HAEIN		cytoplasm
P71367	Y1064_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P71369	Y1104_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P71373	Y1208_HAEIN		cytoplasm
P71374	GLO2_HAEIN		cytoplasm
P71375	Y1315_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P71376	Y1333_HAEIN		inner membrane
P71378	Y1339_HAEIN		cytoplasm
P71379	Y1343_HAEIN		cytoplasm
P71380	PHOR_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P71381	TRPG_HAEIN		cytoplasm
P71384	FUMC_HAEIN	cytoplasm (by similarity).	cytoplasm
P71385	Y1407_HAEIN		cytoplasm
P71387	VG16_HAEIN		cytoplasm
P71388	VPC_HAEIN		cytoplasm
P71389	VPN_HAEIN		cytoplasm
P71391	Y1525_HAEIN	periplasm (potential).	periplasm
P71392	LICA1_HAEIN		cytoplasm
P71394	Y1627_HAEIN		cytoplasm
P71395	RNFA_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P71397	RNFC_HAEIN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	cytoplasm
P71398	LSG4_HAEIN		cytoplasm

P71399	LSG1_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P71403	CHEY_HELPY	cytoplasm (potential).	cytoplasm
P71404	CLPB_HELPY	cytoplasm (probable).	cytoplasm
P71405	CYSE_HELPY	cytoplasm (by similarity).	cytoplasm
P71422	MDCB_KLEPN		cytoplasm
P71426	MDCG_KLEPN		cytoplasm
P71481	PRIM_LEGPN		cytoplasm
P71505	DLHH_METEX		extracellular
P71512	HPPK_METEX		cytoplasm
P71513	FOLB_METEX		cytoplasm
P71517	PQQE_METEX		cytoplasm
P72074	LST_NEIGO		cytoplasm
P72077	YRSP_NEIGO		inner membrane
P72097	LST_NEIMB		cytoplasm
P72131	PTXR_PSEAE		cytoplasm
P72138	HIS52_PSEAE	cytoplasm (by similarity).	cytoplasm
P72139	HIS62_PSEAE	cytoplasm (by similarity).	cytoplasm
P72151	FLICB_PSEAE		flagellar
P72154	TRUB_PSEAE		cytoplasm
P72156	ATZA_PSESD	cytoplasm (potential).	extracellular
P72157	PUR6_PSEAE		cytoplasm
P72158	PURK_PSEAE		cytoplasm
P72161	PBP5_PSEAE	periplasm (potential).	cytoplasm
P72170	PYRC_PSEAE		cytoplasm
P72171	ORUR_PSEAE		cytoplasm
P72174	UVRB_PSEAE	cytoplasm (by similarity).	cytoplasm
P72190	YCAB_PSEFR		cytoplasm
P72203	EXBD_PASHA	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
P72204	TONB_PASHA	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
P72215	LPXA_PROMI	cytoplasm (by similarity).	cytoplasm
P72220	BPHB_PSEPU		cytoplasm
P72227	YGAD_PSEPU		inner membrane
P72238	CFA2_PSESG		cytoplasm
P72241	ISPG_PROST		cytoplasm
P72255	RADC_RHOCA		cytoplasm
P72273	Y4YJ_RHIFR		nucleoid
P72294	OCCR_RHIME		cytoplasm
P72295	OCCQ_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P72296	OCCM_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P72297	OCCP_RHIME	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P72299	OOXB2_RHIME		cytoplasm
P72300	OOXA2_RHIME		cytoplasm

P72302	CGMA_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P72315	YCOM_RHORU		cytoplasm
P72332	NODG_RHIS3		cytoplasm
P72334	NODC_RHIS3	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
P72335	NODI_RHIS3	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P72336	NODJ_RHIS3	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P72338	NODP_RHIS3		cytoplasm
P72339	NODQ_RHIS3		cytoplasm
P72366	CSPA_STIAU	cytoplasm (by similarity).	cytoplasm
P74839	PRPR_SALTY		cytoplasm
P74840	PRPD_SALTY		cytoplasm
P74841	BLC2_SALTY		extracellular
P74852	SSAJ_SALTY	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
P74853	SSAK_SALTY		inner membrane
P74854	SSAL_SALTY		inner membrane
P74855	SSAM_SALTY		inner membrane
P74856	SSAV_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P74857	SSAN_SALTY	cytoplasm (potential).	inner membrane
P74858	SSAO_SALTY		inner membrane
P74860	SSAQ_SALTY		inner membrane
P74866	GALR_SALTY		cytoplasm
P74867	YBIF_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P74876	DPO3X_SALTY		cytoplasm
P74879	PDUW_SALTY		cytoplasm
P74881	PUR4_SALTY	cytoplasm (by similarity).	cytoplasm
P74883	PUR5_SALTY	cytoplasm (by similarity).	cytoplasm
P74884	FEOB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P74890	YSCR_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P74891	SSAS_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P74945	VANI_VIBAN		cytoplasm
P74946	VANR_VIBAN		cytoplasm
P74956	LON_VIBPA	cytoplasm (by similarity).	cytoplasm
P74963	RPOA_SHESP		cytoplasm
P74965	VENB_VIBVU		cytoplasm
P74984	ARSC_YEREN		cytoplasm

P74985	ARSB_YEREN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
P74996	GRP_ZYMMO		cytoplasm
P74997	LOLD_ZYMMO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
P75012	TERX_SERMA		cytoplasm
P75616	YAAX_ECOLI		periplasm
P75617	YAAW_ECOLI		inner membrane
P75620	YAAY_ECOLI		inner membrane
P75672	YAFS_ECOLI		inner membrane
P75675	YKFJ_ECOLI		inner membrane
P75676	YAFX_ECOLI		cytoplasm
P75677	YKFF_ECOLI		cytoplasm
P75678	YKFA_ECOLI		cytoplasm
P75679	INN1_ECOLI		cytoplasm
P75682	YAGE_ECOLI		cytoplasm
P75683	YAGG_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P75684	YAGP_ECOLI		cytoplasm
P75687	YKGI_ECOLI		extracellular
P75688	Y309_ECOLI		inner membrane
P75691	YAHK_ECOLI		cytoplasm
P75692	YAHM_ECOLI		extracellular
P75693	YAHN_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75694	YAHO_ECOLI		cytoplasm
P75704	YKIA_ECOLI		inner membrane
P75711	YBBV_ECOLI		cytoplasm
P75713	YLBA_ECOLI		cytoplasm
P75715	SFMH_ECOLI		extracellular
P75719	RZPD_ECOLI		extracellular
P75728	UBIF_ECOLI		cytoplasm
P75733	YBFM_ECOLI		outer membrane
P75734	YBFN_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P75736	YBFF_ECOLI		cytoplasm
P75737	YBFP_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P75741	YBFL_ECOLI		cytoplasm
P75745	YBGK_ECOLI		cytoplasm
P75746	YBGL_ECOLI		inner membrane
P75748	YBGO_ECOLI		extracellular
P75749	YBGP_ECOLI	periplasm (by similarity).	periplasm
P75750	YBGQ_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P75764	YBHJ_ECOLI		cytoplasm
P75767	YBHK_ECOLI		cytoplasm
P75769	YBHM_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75777	YBHG_ECOLI	periplasm (potential).	inner membrane

P75779	YBIX_ECOLI		cytoplasm
P75780	YBIL_ECOLI	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P75782	YBIN_ECOLI		cytoplasm
P75783	YBIO_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75791	YBIU_ECOLI		cytoplasm
P75792	SUPH_ECOLI		periplasm
P75793	PFLF_ECOLI	cytoplasm (by similarity).	cytoplasm
P75794	PFLE_ECOLI	cytoplasm (by similarity).	cytoplasm
P75796	YLIA_ECOLI		inner membrane
P75797	YLIB_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P75799	YLID_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P75800	YLIE_ECOLI		inner membrane
P75801	YLIF_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75804	YLII_ECOLI		cytoplasm
P75809	YBJI_ECOLI		periplasm
P75811	YBJK_ECOLI		cytoplasm
P75817	RUMB_ECOLI		cytoplasm
P75818	YBJP_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P75820	YBJR_ECOLI		extracellular
P75821	YBJS_ECOLI		cytoplasm
P75822	YBJT_ECOLI		cytoplasm
P75823	LTAE_ECOLI		cytoplasm
P75824	HCR_ECOLI		cytoplasm
P75826	YBJE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75828	YBJD_ECOLI		cytoplasm
P75829	YBJX_ECOLI		cytoplasm
P75836	YCAN_ECOLI		cytoplasm
P75838	YCAO_ECOLI		cytoplasm
P75839	YCAP_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75843	YCAQ_ECOLI		cytoplasm
P75849	YCBL_ECOLI		cytoplasm
P75853	SSUA_ECOLI	periplasm (potential).	periplasm
P75855	YCBQ_ECOLI		extracellular
P75856	YCBR_ECOLI	periplasm (by similarity).	periplasm
P75857	YCBS_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P75858	YCBT_ECOLI		extracellular
P75859	YCBU_ECOLI		extracellular
P75860	YCBV_ECOLI		extracellular
P75862	YCBW_ECOLI		cytoplasm
P75863	YCBX_ECOLI		cytoplasm

P75864	YCBY_ECOLI		cytoplasm
P75867	LONH_ECOLI		cytoplasm
P75869	YCCR_ECOLI		fimbrium
P75874	YCCU_ECOLI		fimbrium
P75876	YCCW_ECOLI		extracellular
P75882	YMCA_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P75883	YMCB_ECOLI		inner membrane
P75884	YMCC_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P75893	RUTF_ECOLI		cytoplasm
P75894	RUTE_ECOLI		cytoplasm
P75895	RUTD_ECOLI		cytoplasm
P75897	RUTB_ECOLI		cytoplasm
P75898	RUTA_ECOLI		cytoplasm
P75901	YCDN_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75906	PGAB_ECOLI	bacterial cell membrane; lipid-anchor (probable).	cytoplasm
P75910	YCDU_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75913	YCDW_ECOLI		cytoplasm
P75914	YCDX_ECOLI		cytoplasm
P75915	YCDY_ECOLI		cytoplasm
P75917	YMDA_ECOLI		extracellular
P75919	YMDC_ECOLI		cytoplasm
P75920	OPGC_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75931	MVIM_ECOLI		cytoplasm
P75933	FLGA_ECOLI	periplasm (probable).	flagellar
P75936	FLGD_ECOLI		cytoplasm
P75937	FLGE_ECOLI		flagellar
P75938	FLGF_ECOLI		flagellar
P75942	FLGJ_ECOLI	periplasm (by similarity).	extracellular
P75946	YCFL_ECOLI		cytoplasm
P75948	THIK_ECOLI		cytoplasm
P75952	YCFQ_ECOLI		cytoplasm
P75954	YCFS_ECOLI	periplasm (potential).	periplasm
P75959	YCFX_ECOLI		fimbrium
P75960	NPD_ECOLI	cytoplasm (probable).	cytoplasm
P75966	RLUE_ECOLI		cytoplasm
P75967	YMFD_ECOLI		cytoplasm
P75968	YMFE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75969	INTE_ECOLI		cytoplasm
P75970	VXIS_ECOLI		cytoplasm
P75971	Y1142_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75972	YMFI_ECOLI		cytoplasm
P75973	YMFJ_ECOLI		cytoplasm
P75974	YMFK_ECOLI		cytoplasm

P75976	YMFL_ECOLI		cytoplasm
P75977	YMFM_ECOLI		extracellular
P75978	YMFN_ECOLI		cytoplasm
P75979	YMFR_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P75980	YMFO_ECOLI		cytoplasm
P75981	YMFP_ECOLI		cytoplasm
P75982	YMFQ_ECOLI		cytoplasm
P75987	ELBA_ECOLI		cytoplasm
P75988	YCGX_ECOLI		cytoplasm
P75989	YCGE_ECOLI		cytoplasm
P75990	YCGF_ECOLI		inner membrane
P75991	YCGZ_ECOLI		cytoplasm
P75992	YMGA_ECOLI		cytoplasm
P75993	YMGB_ECOLI		cytoplasm
P75994	YMGC_ECOLI		cytoplasm
P75995	YCGG_ECOLI		inner membrane
P76001	YCGJ_ECOLI		cytoplasm
P76002	YCGK_ECOLI		cytoplasm
P76004	YCGM_ECOLI		cytoplasm
P76007	CVRA_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P76009	MLTE_ECOLI		extracellular
P76010	YCGR_ECOLI		cytoplasm
P76011	TAG1_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76012	YCGY_ECOLI		cytoplasm
P76014	DHAL_ECOLI		cytoplasm
P76015	DHAK_ECOLI		cytoplasm
P76016	DHAR_ECOLI		cytoplasm
P76017	YCGV_ECOLI		extracellular
P76034	YCIT_ECOLI		cytoplasm
P76035	YCIW_ECOLI		cytoplasm
P76037	PUUP_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P76038	PUUD_ECOLI		cytoplasm
P76041	SUCP_ECOLI		cytoplasm
P76042	YCJN_ECOLI	periplasm (potential).	periplasm
P76043	YCJQ_ECOLI		cytoplasm
P76044	YCJR_ECOLI		cytoplasm
P76046	YCJX_ECOLI		cytoplasm
P76049	YCJY_ECOLI		cytoplasm
P76052	ABGB_ECOLI		extracellular
P76053	YDAL_ECOLI		cytoplasm
P76055	YDAO_ECOLI		cytoplasm
P76056	INTR_ECOLI		cytoplasm
P76057	YDAQ_ECOLI		cytoplasm
P76062	RACR_ECOLI		cytoplasm
P76063	YDAS_ECOLI		cytoplasm
P76064	YDAT_ECOLI		cytoplasm
P76065	YDAU_ECOLI		cytoplasm
P76066	YDAW_ECOLI		cytoplasm

P76067	Y1364_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76068	YNAK_ECOLI		cytoplasm
P76069	YDAY_ECOLI		fimbrium
P76071	INH5_ECOLI		cytoplasm
P76072	STFR_ECOLI		flagellar
P76073	YNAE_ECOLI		cytoplasm
P76076	YDBL_ECOLI		cytoplasm
P76077	PAAA_ECOLI		cytoplasm
P76078	PAAB_ECOLI		cytoplasm
P76079	PAAC_ECOLI		cytoplasm
P76080	PAAD_ECOLI		cytoplasm
P76081	PAAE_ECOLI		cytoplasm
P76082	PAAF_ECOLI		cytoplasm
P76083	PAAH_ECOLI		cytoplasm
P76084	PAAI_ECOLI		cytoplasm
P76085	PAAK_ECOLI		cytoplasm
P76086	PAAX_ECOLI		cytoplasm
P76091	YNBB_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76092	YNBC_ECOLI		cytoplasm
P76093	YNBD_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76097	YDCJ_ECOLI		cytoplasm
P76100	YDCK_ECOLI		cytoplasm
P76102	YDCM_ECOLI		periplasm
P76104	YDCP_ECOLI		periplasm
P76108	YDCS_ECOLI	periplasm (by similarity).	periplasm
P76112	YNCA_ECOLI		cytoplasm
P76113	YNCB_ECOLI		cytoplasm
P76114	YNCC_ECOLI		cytoplasm
P76115	YNCD_ECOLI	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P76116	YNCE_ECOLI		cytoplasm
P76117	YNCG_ECOLI		cytoplasm
P76118	YNCH_ECOLI		inner membrane
P76121	YDDH_ECOLI		cytoplasm
P76122	YDDJ_ECOLI		cytoplasm
P76123	YDDK_ECOLI		cytoplasm
P76127	BDM_ECOLI		cytoplasm
P76128	YDDS_ECOLI	periplasm (potential).	periplasm
P76129	DOS_ECOLI		cytoplasm
P76134	YDEM_ECOLI		cytoplasm
P76135	YDEO_ECOLI		cytoplasm
P76136	Y1500_ECOLI	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
P76141	YDEW_ECOLI		cytoplasm
P76142	YNEA_ECOLI		cytoplasm
P76143	YNEB_ECOLI		cytoplasm

P76146	YNEE_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76147	YNEF_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76148	YNEG_ECOLI		cytoplasm
P76149	YNEI_ECOLI		cytoplasm
P76150	YNEK_ECOLI		cytoplasm
P76154	YDFK_ECOLI		cytoplasm
P76155	TFAQ_ECOLI		cytoplasm
P76156	YDFO_ECOLI		cytoplasm
P76157	YNFN_ECOLI		cytoplasm
P76158	YDFP_ECOLI		cytoplasm
P76159	LYSQ_ECOLI		extracellular
P76160	YDFR_ECOLI		fimbrium
P76161	REQ2_ECOLI		cytoplasm
P76162	YDFU_ECOLI		inner membrane
P76163	YDFV_ECOLI		cytoplasm
P76165	YDFX_ECOLI		cytoplasm
P76170	YNFB_ECOLI		fimbrium
P76172	YNFD_ECOLI		extracellular
P76173	YNFH_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P76176	YDGD_ECOLI		cytoplasm
P76177	YDGH_ECOLI		cytoplasm
P76182	RNFD_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P76185	YDHJ_ECOLI		inner membrane
P76186	YDHK_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76187	YDHF_ECOLI		cytoplasm
P76190	YDHO_ECOLI		extracellular
P76192	YDHV_ECOLI		cytoplasm
P76193	YNHG_ECOLI	periplasm (potential).	periplasm
P76194	SUFE_ECOLI	cytoplasm (by similarity).	cytoplasm
P76196	YDIL_ECOLI		cytoplasm
P76201	YDIQ_ECOLI		cytoplasm
P76204	YDIV_ECOLI		cytoplasm
P76205	ARPB_ECOLI		cytoplasm
P76206	YDIY_ECOLI		outer membrane
P76208	YNIB_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76210	YDJO_ECOLI		cytoplasm
P76213	CHO_ECOLI		cytoplasm
P76214	YDJR_ECOLI		cytoplasm
P76215	ASTE_ECOLI		periplasm
P76216	ASTB_ECOLI		extracellular
P76217	ASTD_ECOLI		cytoplasm
P76219	YDJX_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

P76220	YDJY_ECOLI		outer membrane
P76222	YNJA_ECOLI		cytoplasm
P76223	YNJB_ECOLI		cytoplasm
P76227	YNJH_ECOLI		inner membrane
P76230	YDJK_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P76231	YEAC_ECOLI		cytoplasm
P76234	YEAE_ECOLI		cytoplasm
P76235	YEAH_ECOLI		inner membrane
P76237	YEAJ_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76241	YEAM_ECOLI		cytoplasm
P76243	YEAO_ECOLI		cytoplasm
P76245	YEAP_ECOLI		inner membrane
P76249	YEAS_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76250	YEAT_ECOLI		cytoplasm
P76251	TTUC_ECOLI	cytoplasm (by similarity).	cytoplasm
P76254	YEAX_ECOLI		cytoplasm
P76256	YEAZ_ECOLI		cytoplasm
P76257	YOAA_ECOLI		cytoplasm
P76261	YOAD_ECOLI		inner membrane
P76268	KDGR_ECOLI		cytoplasm
P76269	YEBQ_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P76270	YEBR_ECOLI		inner membrane
P76272	YEBT_ECOLI		inner membrane
P76273	YEBU_ECOLI		cytoplasm
P76275	YEBW_ECOLI		cytoplasm
P76280	YOBG_ECOLI		cytoplasm
P76290	YECO_ECOLI		cytoplasm
P76291	YECF_ECOLI		cytoplasm
P76296	YECT_ECOLI		cytoplasm
P76297	FLHE_ECOLI		inner membrane
P76308	YECR_ECOLI		cytoplasm
P76316	DCYD_ECOLI		cytoplasm
P76318	YEDK_ECOLI		cytoplasm
P76319	YEDL_ECOLI		cytoplasm
P76321	YEDN_ECOLI		cytoplasm
P76322	YEDM_ECOLI		cytoplasm
P76329	MPGP_ECOLI	cytoplasm (by similarity).	cytoplasm
P76330	YEDQ_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76335	YEDS_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P76340	YEDW_ECOLI	cytoplasm (probable).	cytoplasm
P76341	YEDX_ECOLI	periplasm (potential).	periplasm
P76342	YEDY_ECOLI		cytoplasm
P76344	YODA_ECOLI		cytoplasm
P76346	YEEI_ECOLI		cytoplasm

P76347	YEEJ_ECOLI		extracellular
P76349	YEEL_ECOLI		cytoplasm
P76352	YEEO_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P76359	YEEL_ECOLI		cytoplasm
P76362	YEES_ECOLI		cytoplasm
P76364	YEEU_ECOLI		cytoplasm
P76369	YEEY_ECOLI		cytoplasm
P76373	UDG_ECOLI		cytoplasm
P76389	YEGH_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76393	YEGI_ECOLI		cytoplasm
P76394	YEGJ_ECOLI		cytoplasm
P76395	YEGK_ECOLI		cytoplasm
P76396	Y EGL_ECOLI		cytoplasm
P76397	MDTA_ECOLI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
P76398	MDTB_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P76402	YEGP_ECOLI		extracellular
P76403	YEGQ_ECOLI		cytoplasm
P76406	YEGR_ECOLI		cytoplasm
P76407	YEGS_ECOLI		fimbrium
P76418	YEGU_ECOLI		extracellular
P76419	YEGV_ECOLI		cytoplasm
P76421	YEGX_ECOLI		extracellular
P76422	THID_ECOLI		cytoplasm
P76423	THIM_ECOLI		cytoplasm
P76440	YEIT_ECOLI		inner membrane
P76458	ATOD_ECOLI		cytoplasm
P76459	ATOA_ECOLI		cytoplasm
P76461	ATOB_ECOLI	cytoplasm (potential).	cytoplasm
P76462	YFAP_ECOLI		extracellular
P76463	YFAQ_ECOLI		cytoplasm
P76464	YFAS_ECOLI		cytoplasm
P76466	YFAT_ECOLI		cytoplasm
P76469	YFAU_ECOLI		cytoplasm
P76471	YFAZ_ECOLI		outer membrane
P76472	YFBH_ECOLI		cytoplasm
P76481	YFBK_ECOLI		periplasm
P76482	YFBL_ECOLI		extracellular
P76483	YFBM_ECOLI		cytoplasm
P76484	YFBN_ECOLI		cytoplasm
P76485	YFBO_ECOLI		cytoplasm
P76486	YFBP_ECOLI		extracellular
P76491	YFBR_ECOLI	cytoplasm (potential).	periplasm
P76498	YFCO_ECOLI		extracellular
P76499	YFCP_ECOLI		extracellular
P76500	YFCQ_ECOLI		extracellular
P76501	YFCR_ECOLI		extracellular
P76502	SIXA_ECOLI		extracellular
P76503	FADI_ECOLI	cytoplasm (by similarity).	cytoplasm

P76505	YFDF_ECOLI		cytoplasm
P76506	VACJ_ECOLI	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
P76507	YFDI_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76508	YFDL_ECOLI		cytoplasm
P76509	YFDM_ECOLI		cytoplasm
P76510	YFDN_ECOLI		cytoplasm
P76512	YFDP_ECOLI		cytoplasm
P76513	YFDQ_ECOLI		cytoplasm
P76514	YFDR_ECOLI		cytoplasm
P76515	YFDS_ECOLI		cytoplasm
P76516	YFDT_ECOLI		cytoplasm
P76518	YFDE_ECOLI		cytoplasm
P76520	YFDX_ECOLI		cytoplasm
P76521	YFDY_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P76524	YPDF_ECOLI		cytoplasm
P76535	YFEU_ECOLI		cytoplasm
P76536	YFEX_ECOLI		cytoplasm
P76537	YFEY_ECOLI		extracellular
P76539	YPEA_ECOLI		cytoplasm
P76540	EUTK_ECOLI		cytoplasm
P76541	EUTL_ECOLI		cytoplasm
P76542	INTZ_ECOLI		cytoplasm
P76543	YFFL_ECOLI		cytoplasm
P76544	YFFM_ECOLI		cytoplasm
P76545	YFFN_ECOLI		cytoplasm
P76546	YFFO_ECOLI		cytoplasm
P76547	YFFP_ECOLI		cytoplasm
P76548	YFFQ_ECOLI		cytoplasm
P76549	YFFR_ECOLI		cytoplasm
P76550	YFFS_ECOLI		cytoplasm
P76551	EUTA_ECOLI		inner membrane
P76553	EUTG_ECOLI		cytoplasm
P76555	EUTQ_ECOLI		cytoplasm
P76556	EUTP_ECOLI		cytoplasm
P76558	MAO2_ECOLI		cytoplasm
P76559	YPFG_ECOLI		cytoplasm
P76561	YPFH_ECOLI		cytoplasm
P76562	YPFI_ECOLI		cytoplasm
P76569	YFGD_ECOLI		cytoplasm
P76573	YFGI_ECOLI		inner membrane
P76575	YFGJ_ECOLI		cytoplasm
P76576	YFGM_ECOLI		inner membrane
P76577	PBPC_ECOLI	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
P76578	YFHM_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
P76584	YPHB_ECOLI		cytoplasm
P76585	YPHG_ECOLI		extracellular
P76586	YPHH_ECOLI		cytoplasm
P76594	YFIQ_ECOLI		cytoplasm

P76612	YPJB_ECOLI		cytoplasm
P76613	YPJC_ECOLI		cytoplasm
P76616	YGAQ_ECOLI		cytoplasm
P76620	YQAD_ECOLI		cytoplasm
P76621	CSID_ECOLI		cytoplasm
P76628	YGAY_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P76632	YGCK_ECOLI		cytoplasm
P76633	YGCW_ECOLI		cytoplasm
P76639	YGEH_ECOLI		cytoplasm
P76641	GUAD_ECOLI		cytoplasm
P76655	YQIG_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P76656	YQII_ECOLI		extracellular
P76658	HLDE_ECOLI		cytoplasm
P76909	YNJD_ECOLI		inner membrane
P76938	YFCM_ECOLI		cytoplasm
P77044	MHPC_ECOLI		cytoplasm
P77072	MERB_ECOLI		extracellular
P77087	YLCE_ECOLI		cytoplasm
P77091	HOKE_ECOLI		inner membrane
P77129	YLBE_ECOLI		cytoplasm
P77136	YQEK_ECOLI		cytoplasm
P77147	YDHT_ECOLI		cytoplasm
P77148	YDHS_ECOLI		cytoplasm
P77150	PDXY_ECOLI		cytoplasm
P77154	YCJT_ECOLI		extracellular
P77161	GLXR_ECOLI		cytoplasm
P77162	YKFB_ECOLI		extracellular
P77163	TFAR_ECOLI		cytoplasm
P77165	YAGT_ECOLI		cytoplasm
P77169	YAGJ_ECOLI		cytoplasm
P77170	PINQ_ECOLI		cytoplasm
P77171	YDCI_ECOLI		cytoplasm
P77174	YBDM_ECOLI		cytoplasm
P77180	YKGH_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P77181	PAAY_ECOLI		cytoplasm
P77182	YFCK_ECOLI		cytoplasm
P77183	YAGQ_ECOLI		cytoplasm
P77187	YAHF_ECOLI		cytoplasm
P77188	YAGY_ECOLI		cytoplasm
P77196	YFCU_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P77199	YAIT_ECOLI		extracellular
P77206	YAFZ_ECOLI		cytoplasm
P77211	CUSC_ECOLI	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
P77212	YKGC_ECOLI		cytoplasm
P77213	CAAL_ECOLI		cytoplasm
P77215	YFAW_ECOLI		cytoplasm
P77216	YBDN_ECOLI		cytoplasm

P77218	EUTD_ECOLI		inner membrane
P77219	YAHC_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P77221	YAHG_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P77231	CITG_ECOLI		cytoplasm
P77234	YBEQ_ECOLI		extracellular
P77237	ESSQ_ECOLI		extracellular
P77239	CUSB_ECOLI		inner membrane
P77243	PRPD_ECOLI		cytoplasm
P77245	YFET_ECOLI		cytoplasm
P77247	YNIC_ECOLI		cytoplasm
P77249	SFMC_ECOLI	periplasm (potential).	periplasm
P77252	YKGE_ECOLI		cytoplasm
P77256	YDJG_ECOLI		cytoplasm
P77257	EGO_ECOLI		inner membrane
P77258	NEMA_ECOLI		cytoplasm
P77260	YDFI_ECOLI		cytoplasm
P77263	YAGV_ECOLI		periplasm
P77268	YDDP_ECOLI		inner membrane
P77269	YPHF_ECOLI	periplasm (potential).	periplasm
P77277	EUTJ_ECOLI		cytoplasm
P77279	YBBL_ECOLI		inner membrane
P77280	YDJJ_ECOLI		cytoplasm
P77285	RNFG_ECOLI	bacterial cell inner membrane (potential).	inner membrane
P77286	YDEU_ECOLI		extracellular
P77288	YFCV_ECOLI		extracellular
P77294	YDER_ECOLI		extracellular
P77295	YGAV_ECOLI		cytoplasm
P77296	YBET_ECOLI		extracellular
P77297	YAHE_ECOLI		cytoplasm
P77300	YAGI_ECOLI		cytoplasm
P77301	YBAP_ECOLI		inner membrane
P77302	YDAM_ECOLI		inner membrane
P77309	YNEJ_ECOLI		cytoplasm
P77315	YPHD_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P77316	YBDR_ECOLI		cytoplasm
P77318	YDEN_ECOLI		inner membrane
P77319	HSCC_ECOLI		cytoplasm
P77324	YAGS_ECOLI		cytoplasm
P77326	TFAS_ECOLI		cytoplasm
P77329	HYFG_ECOLI		periplasm
P77330	BORD_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P77333	YCJZ_ECOLI		cytoplasm
P77334	YCIR_ECOLI		cytoplasm
P77337	YDIS_ECOLI		cytoplasm
P77339	YAFT_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P77357	ABGA_ECOLI		extracellular
P77359	YBEV_ECOLI		inner membrane

P77360	YPHC_ECOLI		cytoplasm
P77364	GLXK1_ECOLI		cytoplasm
P77365	YAFY_ECOLI		cytoplasm
P77366	PGMB_ECOLI		cytoplasm
P77374	YNFE_ECOLI	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	periplasm
P77375	YDHX_ECOLI		periplasm
P77376	YDGJ_ECOLI		cytoplasm
P77378	YDIR_ECOLI		cytoplasm
P77379	YKGD_ECOLI		cytoplasm
P77381	YBES_ECOLI		inner membrane
P77390	CITC_ECOLI		cytoplasm
P77393	YAHL_ECOLI		cytoplasm
P77395	YBBN_ECOLI		periplasm
P77396	YPDC_ECOLI		cytoplasm
P77397	MHPA_ECOLI		cytoplasm
P77398	ARNA_ECOLI		cytoplasm
P77399	FADJ_ECOLI	cytoplasm (by similarity).	cytoplasm
P77402	YDIP_ECOLI		cytoplasm
P77409	PHSC_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P77414	WCAA_ECOLI		cytoplasm
P77423	HYFH_ECOLI		periplasm
P77425	ALLC_ECOLI		cytoplasm
P77427	YBEU_ECOLI		cytoplasm
P77432	YDEV_ECOLI		cytoplasm
P77433	YKGG_ECOLI		cytoplasm
P77434	YFDZ_ECOLI	cytoplasm (by similarity).	cytoplasm
P77439	PTFX1_ECOLI	cytoplasm (probable).	cytoplasm
P77444	SUFS_ECOLI	cytoplasm (by similarity).	cytoplasm
P77445	EUTE_ECOLI		cytoplasm
P77453	HYFJ_ECOLI		cytoplasm
P77454	GLSA1_ECOLI		extracellular
P77455	MAOC_ECOLI		cytoplasm
P77460	YBCY_ECOLI		cytoplasm
P77467	PAAG_ECOLI		cytoplasm
P77468	SFMD_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
P77473	YLAB_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P77475	YQAB_ECOLI		cytoplasm
P77481	YCJV_ECOLI		inner membrane
P77488	DXS_ECOLI		cytoplasm
P77489	YAGR_ECOLI		periplasm
P77493	YDJH_ECOLI		cytoplasm
P77495	PRPE_ECOLI		cytoplasm
P77499	SUFC_ECOLI		inner membrane
P77503	YCJS_ECOLI		cytoplasm
P77504	YBBP_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

P77506	YBDJ_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P77509	YPHE_ECOLI		inner membrane
P77515	STFQ_ECOLI		flagellar
P77522	SUFB_ECOLI		fimbrium
P77525	PAAJ_ECOLI		cytoplasm
P77526	YFCG_ECOLI		cytoplasm
P77529	YDJN_ECOLI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P77536	YKGF_ECOLI		cytoplasm
P77538	YFHR_ECOLI		inner membrane
P77539	YDJL_ECOLI		cytoplasm
P77541	PRPB_ECOLI		cytoplasm
P77544	YFCF_ECOLI		cytoplasm
P77546	YDAV_ECOLI		cytoplasm
P77552	YDHQ_ECOLI		cytoplasm
P77554	YAHJ_ECOLI		cytoplasm
P77555	ALLD_ECOLI	cytoplasm (probable).	cytoplasm
P77559	YNFL_ECOLI		cytoplasm
P77561	YDEP_ECOLI		periplasm
P77562	YAIW_ECOLI		cytoplasm
P77564	YDHW_ECOLI		cytoplasm
P77567	NHOA_ECOLI		inner membrane
P77569	MHPR_ECOLI		cytoplasm
P77570	YDHH_ECOLI		inner membrane
P77580	MHPF_ECOLI		cytoplasm
P77581	ARGM_ECOLI		cytoplasm
P77585	YPDE_ECOLI		cytoplasm
P77588	YDEQ_ECOLI		extracellular
P77596	YAGF_ECOLI		cytoplasm
P77598	YBCV_ECOLI		cytoplasm
P77599	YFCS_ECOLI	periplasm (by similarity).	periplasm
P77601	YKGA_ECOLI		cytoplasm
P77607	YAGL_ECOLI		cytoplasm
P77608	MHPD_ECOLI		cytoplasm
P77609	FLXA_ECOLI		cytoplasm
P77611	RNFC_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	cytoplasm
P77615	YCJW_ECOLI		cytoplasm
P77616	YQIH_ECOLI	periplasm (by similarity).	periplasm
P77619	YFEW_ECOLI		extracellular
P77622	YDDO_ECOLI		inner membrane
P77624	ARCM_ECOLI		cytoplasm
P77625	YFBT_ECOLI		cytoplasm
P77626	YDCN_ECOLI		cytoplasm
P77627	YBER_ECOLI		cytoplasm
P77634	YBCM_ECOLI		cytoplasm
P77646	HCAB_ECOLI		cytoplasm
P77649	YDIU_ECOLI		inner membrane
P77650	HCAD_ECOLI		cytoplasm
P77656	YFDK_ECOLI		cytoplasm
P77657	YAGK_ECOLI		cytoplasm
P77658	YNAA_ECOLI		inner membrane

P77667	SUFA_ECOLI		inner membrane
P77668	HYFI_ECOLI		cytoplasm
P77671	ALN_ECOLI		cytoplasm
P77674	YDCW_ECOLI		cytoplasm
P77689	SUFD_ECOLI		fimbrium
P77690	ARNB_ECOLI		cytoplasm
P77692	YKFI_ECOLI		cytoplasm
P77694	YAGW_ECOLI		outer membrane
P77695	GNSB_ECOLI		cytoplasm
P77698	YBCK_ECOLI		cytoplasm
P77699	TFAD_ECOLI		cytoplasm
P77700	YAHB_ECOLI		cytoplasm
P77704	YDJI_ECOLI		cytoplasm
P77712	YBAW_ECOLI		cytoplasm
P77713	YAGH_ECOLI		extracellular
P77714	YDIT_ECOLI		cytoplasm
P77717	YBAY_ECOLI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
P77721	YDJF_ECOLI		cytoplasm
P77730	YDCR_ECOLI		cytoplasm
P77731	ALLA_ECOLI		extracellular
P77732	YFAX_ECOLI		cytoplasm
P77735	YAJO_ECOLI		cytoplasm
P77736	YAHD_ECOLI		cytoplasm
P77739	YNIA_ECOLI		cytoplasm
P77743	PRPR_ECOLI		cytoplasm
P77744	ABGR_ECOLI		cytoplasm
P77746	YBDO_ECOLI		cytoplasm
P77748	YDIJ_ECOLI		inner membrane
P77756	YBAX_ECOLI		cytoplasm
P77766	TRPH_ECOLI		cytoplasm
P77768	YFCI_ECOLI		cytoplasm
P77775	YFCH_ECOLI		cytoplasm
P77779	YBFO_ECOLI		cytoplasm
P77781	YDII_ECOLI		cytoplasm
P77783	YNFF_ECOLI	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	periplasm
P77788	NUDG_ECOLI		cytoplasm
P77789	YDES_ECOLI		extracellular
P77790	DDPX_ECOLI		cytoplasm
P77791	MAA_ECOLI		cytoplasm
P77795	YDCT_ECOLI		inner membrane
P77802	YAGX_ECOLI		cytoplasm
P77806	YBDL_ECOLI	cytoplasm (by similarity).	cytoplasm
P77808	CINA_ECOLI		inner membrane
P77809	G6PD_ACTAC		cytoplasm
P77828	CH101_BRAJA	cytoplasm (by similarity).	cytoplasm
P77829	CH601_BRAJA	cytoplasm (by similarity).	cytoplasm
P77866	DNAJ_ACTAC	cytoplasm (probable).	cytoplasm
P77872	CATA_HELPY	cytoplasm (probable).	periplasm
P77873	NIFH_HERSE		cytoplasm
P77874	NIFD_HERSE		periplasm
P77913	CH10_NEIGO	cytoplasm (by similarity).	cytoplasm
P77914	BFRB_NEIGO		cytoplasm

P77929	SODM_PSEPU		periplasm
P77930	BFR_PSEPU		cytoplasm
P77935	PUR1_RHIET		cytoplasm
P77938	NADC_RHORU		cytoplasm
P77941	RPOB_RICTY		cytoplasm
P77953	SSB_SHEHA		cytoplasm
P77983	KPYK1_SALTY		cytoplasm
P77998	RP54_XANCV		cytoplasm
P78061	PUUA_ECOLI		cytoplasm
P78067	YNJE_ECOLI		cytoplasm
P78218	DYR15_ECOLI		cytoplasm
P78271	YFES_ECOLI		cytoplasm
P78285	LYSD_ECOLI		extracellular
P80011	TRA3_RHIME		cytoplasm
P80064	HPPD_PSEUJ		cytoplasm
P80150	HRC2_XANCV	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P80151	HRPA1_XANCV	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P80152	HRB3_XANCV	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
P80153	HRPB6_XANCV	cytoplasm (potential).	inner membrane
P80176	HIP_THETI		periplasm
P80200	CGA1_HELPY		cytoplasm
P80288	CY550_PARPV		periplasm
P80298	BLAC_PROVU		extracellular
P80306	FER6_RHOCA		cytoplasm
P80357	ASTA_PSEAE		cytoplasm
P80358	ASTG_PSEAE		cytoplasm
P80448	FER2_RHORU		periplasm
P80458	MDH_RHOPA		cytoplasm
P80469	CH10_RICTY	cytoplasm (by similarity).	cytoplasm
P80536	MDH_BURPS		cytoplasm
P80545	BLAC_SERFO		extracellular
P80549	CY551_CHRVI		periplasm
P80563	PGTL_PELAC		periplasm
P80564	PGTS_PELAC		cytoplasm
P80573	CATC_RALEJ		cytoplasm
P80644	SSUE_ECOLI		cytoplasm
P80645	SSUD_ECOLI		cytoplasm
P80649	AZUP_PARDE	periplasm (by similarity).	periplasm
P80668	FEAB_ECOLI		cytoplasm
P80882	HIP_RHOFE		periplasm
P80918	ACP_COMTE	cytoplasm (by similarity).	cytoplasm
P80920	ACP_LEUMU	cytoplasm (by similarity).	cytoplasm
P80921	ACP_MYXXA	cytoplasm (by similarity).	cytoplasm
P80922	ACP_OCELI	cytoplasm (by similarity).	cytoplasm
P80923	ACP_PSESM	cytoplasm (by similarity).	cytoplasm
P81040	CYCX_DESDE		cytoplasm
P81046	PYP_CHRSA		cytoplasm
P81065	GST_OCHAN	cytoplasm (by similarity).	cytoplasm
P81153	CYC21_RHOCE		periplasm
P81154	CYC22_RHOCE		periplasm
P81171	Y174_RICPR		cytoplasm

P81238	SHP_RHOS4		periplasm
P81378	CYB_RHOVI		inner membrane
P81379	CY1_RHOVI		inner membrane
P81432	SYD_BUCAP	cytoplasm (by similarity).	cytoplasm
P81435	SERC_BUCAP	cytoplasm (by similarity).	cytoplasm
P81445	NIR_ALCXX	periplasm (by similarity).	periplasm
P81650	BGAL_PSEHA		extracellular
P81781	BLC6_VIBCH		extracellular
P81891	YFBW_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P81894	CYC7_GEOMG		periplasm
P82177	MDH_SHEON		cytoplasm
P82291	CYB5_ECTVA		cytoplasm
P82802	FER2_AZOVI		cytoplasm
P82973	AMPD_ENTCL	cytoplasm (by similarity).	extracellular
P82974	AMPD_CITFR	cytoplasm (by similarity).	extracellular
P83170	C551_RHOTE		cytoplasm
P83223	FRDA_SHEON	periplasm (by similarity).	periplasm
P83341	HIP1_ECTMO		periplasm
P83342	HIP2_ECTMO		periplasm
P83763	CBNB_RALEU		cytoplasm
P83842	ADPP_ECOL6		periplasm
P83843	ADPP_ECO57		periplasm
P83844	ADPP_SHIFL		periplasm
P84612	SODF_PSEHT		periplasm
P94123	ACP_AZOBR	cytoplasm (by similarity).	cytoplasm
P94127	NOSZ_ACHCY	periplasm (by similarity).	periplasm
P94131	MUCK_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P94132	ETFD_ACIAD		cytoplasm
P94135	TFDF2_RALEJ		cytoplasm
P94136	TFDE2_RALEJ		cytoplasm
P94139	CZCN_RALME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P94147	MTA1_RUEGE		cytoplasm
P94164	DEOD_ACTPL		cytoplasm
P94165	CH10_ACTPL	cytoplasm (by similarity).	cytoplasm
P94166	CH60_ACTPL	cytoplasm (by similarity).	cytoplasm
P94169	GAL1_ACTPL	cytoplasm (potential).	cytoplasm
P94176	CZCB_ALCSC		inner membrane
P94177	CZCA_ALCSC	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P94178	CZCD_ALCSC	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P94185	MERT_ALCSP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P94186	MERP_ALCSP	periplasm (probable).	inner membrane
P94188	MERA_ALCSP		cytoplasm
P94190	RECA_AERSA	cytoplasm (by similarity).	cytoplasm

P94199	ALG8_AZOVI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P94200	ALG44_AZOVI		inner membrane
P94202	ALGK_AZOVI	bacterial cell membrane; lipid-anchor (potential).	outer membrane
P94210	DHLO_AGRVI		cytoplasm
P94281	GYRB_BARBA		cytoplasm
P94287	DSBB_BURCE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P94317	DNAK_BRAJA		cytoplasm
P94319	DNAJ_BRAJA	cytoplasm (probable).	cytoplasm
P94323	NODM_BRAJA	cytoplasm (by similarity).	cytoplasm
P94325	CISY_BRAJA		cytoplasm
P94326	TRPD_BRAJA		cytoplasm
P94327	TRPC_BRAJA		cytoplasm
P94328	MOAC_BRAJA		cytoplasm
P94340	RRF_BRUME	cytoplasm (by similarity).	cytoplasm
P94612	TRME_COXBU		cytoplasm
P94613	GIDA_COXBU	cytoplasm (by similarity).	cytoplasm
P94614	GIDB_COXBU		cytoplasm
P94644	ISPH_CAMJE		cytoplasm
P94700	MERT_ENTAG	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P94702	MERA_ENTAG		cytoplasm
P94777	MODD_ECO27		cytoplasm
P94797	CH10_FRATH	cytoplasm (by similarity).	cytoplasm
P94798	CH60_FRATH	cytoplasm (by similarity).	cytoplasm
P94819	CH10_HOLOB	cytoplasm (by similarity).	cytoplasm
P94820	CH60_HOLOB	cytoplasm (by similarity).	cytoplasm
P94842	Y496_HELPY		cytoplasm
P94844	DAPB_HELPY	cytoplasm (by similarity).	cytoplasm
P94846	UVRB_HELPY	cytoplasm (by similarity).	cytoplasm
P94851	AN36_HELPY		inner membrane
P94958	AMPC_MORMO	periplasm (by similarity).	periplasm
P94966	BIOB_METSK		cytoplasm
P95329	MASY_MYXXA	cytoplasm (by similarity).	cytoplasm
P95331	FUMC_MYXXA	cytoplasm (by similarity).	cytoplasm
P95333	GRPE_MYXXA	cytoplasm (probable).	cytoplasm
P95334	DNAK_MYXXA		cytoplasm
P95337	AROE_NEICI		cytoplasm
P95340	AROE_NEIFL		cytoplasm
P95354	ARSC_NEIGO		cytoplasm
P95357	FTSX_NEIGO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P95361	AROE_NEIGO		cytoplasm
P95364	YORZ_NEIGO		cytoplasm
P95368	AROE_NEILA		cytoplasm
P95370	CYSG_NEIMB		cytoplasm
P95374	TONB_NEIMC	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane

P95377	LPXD_NEIMB		cytoplasm
P95378	FABZ_NEIMB	cytoplasm (by similarity).	cytoplasm
P95379	LPXA_NEIMB	cytoplasm (by similarity).	cytoplasm
P95389	AROE_NEIMU		cytoplasm
P95399	AROE_NEIPO		cytoplasm
P95400	AROE_NEIPH		cytoplasm
P95442	ATZB_PSESD	cytoplasm (potential).	extracellular
P95453	Y4753_PSEAE		inner membrane
P95454	RRMJ_PSEAE	cytoplasm (by similarity).	cytoplasm
P95458	DCUP_PSEAE	cytoplasm (probable).	cytoplasm
P95459	CSPA_PSEAE	cytoplasm (by similarity).	cytoplasm
P95460	DSBA_PSEAE	periplasm (by similarity).	periplasm
P95469	RECA_PARDE	cytoplasm (by similarity).	cytoplasm
P95477	DCHS_PSEFL		cytoplasm
P95508	YHEM_PASHA		inner membrane
P95509	YHEN_PASHA		inner membrane
P95514	KDSA_PASHA	cytoplasm (by similarity).	cytoplasm
P95516	IHFA_PASHA		cytoplasm
P95519	IHFB_PASHA		cytoplasm
P95524	DYR6_PROMI		cytoplasm
P95525	HEM1_PASMU		cytoplasm
P95526	RECA_PASMU	cytoplasm (by similarity).	cytoplasm
P95539	CATE_PSEPU	cytoplasm (probable).	periplasm
P95557	TATA_PSEST	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
P95575	GLMM_PSEU2		cytoplasm
P95576	TPIS_PSEU2		cytoplasm
P95577	SECG_PSEU2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P95578	Y4182_PSEU2		inner membrane
P95582	GACA_PSEVI		cytoplasm
P95596	DLDH_RHOCA	cytoplasm (potential).	cytoplasm
P95629	PUTA_RHIME		cytoplasm
P95645	MOBA_RHOSH	cytoplasm (by similarity).	cytoplasm
P95646	TRPE_RHOS4		cytoplasm
P95647	CH602_RHOSH	cytoplasm (by similarity).	cytoplasm
P95648	CBBX_RHOSH		cytoplasm
P95649	CBBY_RHOSH		cytoplasm
P95650	GPH_RHOSH		cytoplasm
P95651	HEMN_RHOSH	cytoplasm (by similarity).	cytoplasm
P95653	NDK_RHOSU	cytoplasm (by similarity).	cytoplasm
P95656	PUCC_RHOSU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P95677	CH10_RHOCA	cytoplasm (by similarity).	cytoplasm
P95678	CH60_RHOCA	cytoplasm (by similarity).	cytoplasm
P95713	FLIE_SHIBO		flagellar
P95800	CH60_XANMA	cytoplasm (by similarity).	cytoplasm
P95801	CH10_XANMA	cytoplasm (by similarity).	cytoplasm
P95806	LIND_PSEPA		cytoplasm
P96068	SSAT_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

P96069	SSAU_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P96095	HCP_THIFE	cytoplasm (by similarity).	cytoplasm
P96156	CHID_VIBFU	periplasm (probable).	periplasm
P96157	NAGZ_VIBFU	cytoplasm (by similarity).	extracellular
P96166	NAGA_VIBFU		cytoplasm
P96172	OTC_VIBS2	cytoplasm (probable).	cytoplasm
P96174	PYRB_VIBS2		cytoplasm
P96175	PYRI_VIBS2		cytoplasm
P96185	RECA_XENBV	cytoplasm (by similarity).	cytoplasm
P96188	MTX1_XANCR		cytoplasm
P96189	T2X1_XANCR		extracellular
P96190	TPIS_XANFL		cytoplasm
P96192	DUSC_AZOVI		cytoplasm
P96193	IBPB_AZOVI		cytoplasm
P96194	YIBL_AZOVI		cytoplasm
P96195	LEU2_AZOVI		cytoplasm
P96196	LEUD_AZOVI		cytoplasm
P96198	DHAS_AZOVI		cytoplasm
P96199	USG_AZOVI		cytoplasm
P96329	PIFA_ECOLI		cytoplasm
P96335	GLPT_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
P96343	VPB_HAEIN		nucleoid
P96440	EXPG_RHIME		cytoplasm
P96465	BLA2_XANMA		extracellular
P96554	Y319_MYXXA	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
P96747	FLAC_CAMJE		flagellar
P96763	TPIS_FRATH		cytoplasm
P96949	HPUB_NEIMC	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
P96956	ALGK_PSEAE	bacterial cell membrane; lipid-anchor (potential).	outer membrane
P96963	RADA_PSEAE		cytoplasm
P96974	FLGM_PROMI		cytoplasm
P96975	FLGN_PROMI		flagellar
P96977	CHRR_PSEUG		cytoplasm
P96981	GLO2_RHOCA		cytoplasm
P97051	NHAA_PSEPU		extracellular
P97052	NHAB_PSEPU		cytoplasm
P97054	RNFG_RHOCA	bacterial cell inner membrane (potential).	inner membrane
P97055	RNFE_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
P97056	RNFH_RHOCA		inner membrane
P97084	COBD_SALTY		cytoplasm
P97227	CAGS_HELPY		cytoplasm
P97245	CAGT_HELPY	bacterial cell membrane; lipid-anchor (probable).	inner membrane

P98008	NORB_PSEST	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
P98057	COX14_BRAJA		inner membrane
Q00037	TNPA_ECOLI		cytoplasm
Q00042	TNPX_ECOLI		inner membrane
Q00045	FMD1_NEIGO		extracellular
Q00046	FMD3_NEIGO		extracellular
Q00184	TRAG5_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q00186	TRAM4_ECOLI		cytoplasm
Q00187	TRAL5_ECOLI		cytoplasm
Q00188	TRAL4_ECOLI		cytoplasm
Q00189	TRAH5_ECOLI		cytoplasm
Q00190	TRAH4_ECOLI		cytoplasm
Q00191	TRAI6_ECOLI		cytoplasm
Q00192	TRAI5_ECOLI		cytoplasm
Q00267	NHOA_SALTY		inner membrane
Q00329	RFBJ_SALMU		cytoplasm
Q00330	RFBK_SALMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q00456	TMOA_PSEME		cytoplasm
Q00457	TMOB_PSEME		cytoplasm
Q00458	TMOC_PSEME		cytoplasm
Q00459	TMOD_PSEME		cytoplasm
Q00460	TMOE_PSEME		cytoplasm
Q00473	RFBM_SALMU		cytoplasm
Q00474	RFC_SALMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q00499	CY550_PARVE		periplasm
Q00500	CCMF_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q00501	CCMH_RHOCA	periplasm (probable).	periplasm
Q00512	GSPE_PSEAE	cytoplasm (probable).	inner membrane
Q00513	GSPF_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q00514	GSPG_PSEAE		inner membrane
Q00515	GSPH_PSEAE		inner membrane
Q00516	GSPI_PSEAE		inner membrane
Q00517	GSPJ_PSEAE		inner membrane
Q00518	GSPK_PSEAE	bacterial cell inner membrane (potential).	inner membrane
Q00644	YNG2_AZOBR		cytoplasm
Q00678	NOCR_AGRT5		cytoplasm
Q00736	TRAY5_ECOLI		cytoplasm
Q00739	X193_ECOLI		cytoplasm
Q00740	1A1D_PSEUD		cytoplasm
Q00790	NOSR_PSEST	bacterial cell membrane; peripheral membrane protein (potential).	cytoplasm
Q00840	T1106_NEIMB		cytoplasm

Q00879	FHAE_BORPE		extracellular
Q00924	HYUE_PSESN		cytoplasm
Q00932	LCRT_YERPS		cytoplasm
Q00934	PILR_PSEAE	cytoplasm (probable).	cytoplasm
Q00982	BLO5_PSEAE		extracellular
Q00983	BLL1_PSEAE		extracellular
Q01060	PQQE_ESCVU		cytoplasm
Q01095	FLAV_DESGI		cytoplasm
Q01096	FLAW_DESGI		cytoplasm
Q01103	PCAI_PSEPU		cytoplasm
Q01166	BLAC_YEREN		extracellular
Q01179	NIFS_RHOSH		cytoplasm
Q01180	NIFU_RHOSH		cytoplasm
Q01181	NIFV_RHOSH		cytoplasm
Q01182	NIFW_RHOSH		cytoplasm
Q01194	RP54_RHOSH		cytoplasm
Q01195	YNIU_RHOSH		inner membrane
Q01198	LIGD_PSEPA		cytoplasm
Q01234	NFNB_ENTCL		cytoplasm
Q01243	YSCB_YEREN	or: cytoplasm (by similarity). or: bacterial cell inner membrane; peripheral membrane protein (by similarity). not exported across the inner membrane (by similarity).	cytoplasm
Q01244	YSCC_YEREN	bacterial cell outer membrane (probable).	outer membrane
Q01245	YSCD_YEREN	bacterial cell inner membrane; single-pass type ii membrane protein (potential).	inner membrane
Q01246	YSCE_YEREN		extracellular
Q01247	YSCF_YEREN		inner membrane
Q01248	YSCG_YEREN	bacterial cell inner membrane; peripheral membrane protein (potential).	extracellular
Q01249	YSCH_YEREN		extracellular
Q01250	YSCI_YEREN		extracellular
Q01251	YSCJ_YEREN	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q01252	YSCK_YEREN		extracellular
Q01253	YSCL_YEREN		extracellular
Q01254	YSCM_YEREN		extracellular
Q01262	HYUA_PSESN		cytoplasm
Q01263	HYUB_PSESN		cytoplasm
Q01264	HYUC_PSESN		cytoplasm
Q01330	CRTX_ESCVU		cytoplasm
Q01331	CRTY_ESCVU		cytoplasm
Q01332	CRTZ_ESCVU		cytoplasm
Q01333	YCR4_ESCVU		cytoplasm
Q01334	YCR3_ESCVU		inner membrane
Q01335	IDI2_ESCVU	cytoplasm (by similarity).	cytoplasm
Q01398	DEH1_MORSB		cytoplasm
Q01399	DEH2_MORSB		cytoplasm
Q01410	MANC_SALMO		cytoplasm
Q01411	MANB_SALMO		cytoplasm

Q01468	4OT1_PSEPU		cytoplasm
Q01515	AAC3_SERMA		cytoplasm
Q01563	GSPB_DICD3	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q01564	GSPC2_DICD3	bacterial cell inner membrane (probable).	inner membrane
Q01565	GSPD2_DICD3	bacterial cell outer membrane (probable).	inner membrane
Q01567	OUTS_DICD3		inner membrane
Q01578	GNL_ZYMMO	periplasm (probable).	periplasm
Q01602	OPDE_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q01609	Y2218_PSEAE		cytoplasm
Q01610	OPRR_PSEAE		cytoplasm
Q01624	RPSB_STIAU		cytoplasm
Q01670	CRTC_RHOS4		cytoplasm
Q01671	CRTD_RHOS4		cytoplasm
Q01725	LIFO_PSEAE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
Q01856	Y3073_RHOS4		cytoplasm
Q01857	DCTA_RHILE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q01893	TPIS_MORSP		cytoplasm
Q01953	RECA_PSEFL	cytoplasm (by similarity).	cytoplasm
Q01980	AADR_RHOPA		cytoplasm
Q01990	NODU_RHIFR		inner membrane
Q01993	RISA_PHOLE		cytoplasm
Q02006	Y4233_RHOPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q02007	NODS_RHIFR		cytoplasm
Q02008	RIBB_PHOLE		cytoplasm
Q02047	OTC1_PSESH	cytoplasm (probable).	cytoplasm
Q02104	LIP1_PSYIM	bacterial cell outer membrane; lipid-anchor (potential).	extracellular
Q02198	MORA_PSEPU		cytoplasm
Q02219	ANIA_NEIGO	bacterial cell outer membrane; lipid-anchor (probable).	periplasm
Q02322	UVRD_HAEIN		cytoplasm
Q02430	BCHC_RHOS4		cytoplasm
Q02431	BCHX_RHOS4		cytoplasm
Q02432	BCHY_RHOS4	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q02441	NIRQ_PSEST	cytoplasm (potential).	cytoplasm
Q02443	PUCC_RHOS4	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q02458	URER_PROMI		cytoplasm
Q02540	COPR_PSESM	cytoplasm (probable).	cytoplasm

Q02541	COPS_PSESM	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q02584	TRPC_RHOCA		cytoplasm
Q02635	AATA_RHIME	cytoplasm (by similarity).	cytoplasm
Q02636	ATTY_RHIME		cytoplasm
Q02728	EXOF_RHIME	periplasm (probable).	periplasm
Q02730	EXOX_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q02760	CY1_RHOSH		inner membrane
Q02861	CRTI_MYXXA		cytoplasm
Q02869	HIN_SALAE		cytoplasm
Q02876	NODD1_RHITR		cytoplasm
Q02940	PENA_BURCE		cytoplasm
Q02943	URE3_KLEPN	cytoplasm (by similarity).	cytoplasm
Q02944	URED_KLEPN		cytoplasm
Q02987	HYPF_RHOCA		cytoplasm
Q02998	YH19_RHOCA		cytoplasm
Q02999	YH20_RHOCA		cytoplasm
Q03004	HUPD_RHOCA		extracellular
Q03005	HUPF_RHOCA		cytoplasm
Q03006	HUPG_RHOCA		cytoplasm
Q03007	HUPH_RHOCA		cytoplasm
Q03009	HUPJ_RHOCA		cytoplasm
Q03011	MRPA_PROMI		extracellular
Q03025	APRE_PSEAE	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q03030	DCOA_SALTY		inner membrane
Q03055	MTV1_VIBS3		cytoplasm
Q03170	BLP1_PSEAE		extracellular
Q03268	Y2604_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q03282	URE3_ECOLI	cytoplasm (by similarity).	cytoplasm
Q03283	URE2_ECOLI	cytoplasm (by similarity).	cytoplasm
Q03285	URED_ECOLI		cytoplasm
Q03287	UREG_ECOLI		cytoplasm
Q03298	PHHY_ACIAD		cytoplasm
Q03304	TMOF_PSEME		cytoplasm
Q03313	RHIA_RHILV		extracellular
Q03314	RHIB_RHILV		extracellular
Q03316	RHIR_RHILV		cytoplasm
Q03321	ARO_AERSA	cytoplasm (probable).	cytoplasm
Q03324	LUXC1_PHOLE		cytoplasm
Q03381	REGB_PSEAE		extracellular
Q03408	RP54_CAUCR		cytoplasm
Q03417	SCRK_ZYMMO		cytoplasm
Q03421	ARO_HAEIN	cytoplasm (probable).	cytoplasm
Q03432	PURU_HAEIN		cytoplasm
Q03447	KKA1_SALTY		cytoplasm
Q03450	TRAF4_ECOLI	periplasm (probable).	periplasm
Q03456	FUR_PSEAE	cytoplasm (by similarity).	cytoplasm
Q03470	GYRA_CAMJE		cytoplasm
Q03474	LAFS_VIBPA		cytoplasm

Q03476	L AFL_VIBPA		inner membrane
Q03520	PCPC_SPHCR		cytoplasm
Q03583	RFBX_SHIDY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q03584	RFC_SHIDY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q03638	NADE_RHOCA		cytoplasm
Q03708	IMM7_ECOLI		cytoplasm
Q03841	FLAA1_RHIME		flagellar
Q03842	FLAB1_RHIME		flagellar
Q03845	FLHA_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q03974	LIC2A_HAEIN		cytoplasm
Q04152	ROS_AGRTU		cytoplasm
Q04222	TRA1_COMTE		cytoplasm
Q04285	XYLE2_PSEPU		cytoplasm
Q04302	COXZ_EHRRU		inner membrane
Q04314	IPGB_SHIDY		extracellular
Q04474	RT3C_ACTPL	cytoplasm (potential).	cytoplasm
Q04507	AMOA_NITEU	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q04508	AMOB_NITEU	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q04512	HEM1_RHOS4		inner membrane
Q04515	DYR10_ECOLI		cytoplasm
Q04518	ALDC_KLETE		cytoplasm
Q04520	BUDC_KLETE		cytoplasm
Q04524	ILVB_KLETE		cytoplasm
Q04539	RPEP_RALEU		cytoplasm
Q04540	CBXXP_RALEU		cytoplasm
Q04541	CBBYP_RALEU		cytoplasm
Q04551	6B1_AGRVI		cytoplasm
Q04554	VIS_AGRVI		cytoplasm
Q04557	HYIN1_AGRVI		extracellular
Q04564	TR2M_AGRVI		cytoplasm
Q04570	AROA_PASMU	cytoplasm (probable).	cytoplasm
Q04590	IPT1_AGRVI		cytoplasm
Q04596	LEXA_PECCC		cytoplasm
Q04628	Y1545_PSEAE		inner membrane
Q04640	MXIC_SHIFL		outer membrane
Q04761	RECA_BRUAB	cytoplasm (by similarity).	cytoplasm
Q04786	HEX_VIBVU		periplasm
Q04803	PFER_PSEAE	cytoplasm (potential).	cytoplasm
Q04804	PFES_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q04845	MTC1_CITFR		cytoplasm
Q04849	NTRX_AZOCA		cytoplasm
Q04852	T2C1_CITFR		extracellular
Q04855	YNTC_AZOCA		inner membrane
Q04871	YCL2_ECO11		cytoplasm

Q04872	UDG_ECO11		cytoplasm
Q04873	UDG_SALTY		cytoplasm
Q04953	FLBE_CAUCR		inner membrane
Q04954	FLIF_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q04955	FLIG_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q04971	TVIA_SALTI		inner membrane
Q04972	VIPA_SALTI		cytoplasm
Q04973	VIPB_SALTI		cytoplasm
Q04974	TVID_SALTI		extracellular
Q04975	VIPC_SALTI		extracellular
Q04976	VEXA_SALTI	bacterial cell outer membrane; multi-pass membrane protein (potential).	outer membrane
Q04983	TYRC_ZYMMO		cytoplasm
Q05013	LIPM_NEIMB	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (probable).	inner membrane
Q05014	LPB1_NEIMB	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (probable).	inner membrane
Q05026	GALE_NEIGO		cytoplasm
Q05053	PAC1_PSESV		cytoplasm
Q05115	AMDA_BORBR		cytoplasm
Q05119	TRAF5_ECOLI	periplasm (probable).	periplasm
Q05129	EAEB_ECO27		extracellular
Q05181	PHT1_PSEPU	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q05182	PHT2_PSEPU		cytoplasm
Q05183	PHT3_PSEPU		cytoplasm
Q05184	PHT4_PSEPU		cytoplasm
Q05185	PHT5_PSEPU		cytoplasm
Q05197	PMTA_RHOSH		cytoplasm
Q05311	UVRD_SALTY		cytoplasm
Q05338	HEMH_YERPS	cytoplasm (by similarity).	extracellular
Q05347	RFBX_YERPS	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q05353	HPCB_ECOLI		cytoplasm
Q05354	HPCD_ECOLI		cytoplasm
Q05358	RECA_LEGPN	cytoplasm (by similarity).	cytoplasm
Q05395	HRPJ_PSESY		outer membrane
Q05459	VAGC_SALDU		extracellular
Q05527	KDGF_DICD3		cytoplasm
Q05528	KDUD_DICD3		cytoplasm
Q05529	KDUI_DICD3		cytoplasm
Q05559	FIXA_RHILP		cytoplasm
Q05561	FIXX_RHILP		cytoplasm
Q05587	POCR_SALTY		cytoplasm
Q05590	CBIH_SALTY		cytoplasm
Q05591	CBIJ_SALTY		cytoplasm
Q05592	CBIK_SALTY		extracellular

Q05593	CBIL_SALTY		cytoplasm
Q05594	CBIM_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q05596	CBIO_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q05597	CBIP_SALTY		cytoplasm
Q05599	COBU_SALTY		cytoplasm
Q05600	CBIB_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q05601	CBIC_SALTY		cytoplasm
Q05602	COBS_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q05603	COBT_SALTY		cytoplasm
Q05606	EXBD_PSEPU	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
Q05628	CBID_SALTY		cytoplasm
Q05631	CBIG_SALTY		cytoplasm
Q05632	CBIT_SALTY		cytoplasm
Q05755	GLTB_AZOBR		cytoplasm
Q05756	GLTD_AZOBR		cytoplasm
Q05781	FINO4_ECOLI		outer membrane
Q05805	SIR_DESVH		cytoplasm
Q05869	MUTY_SALTY		cytoplasm
Q05938	TOXR_VIBPA		cytoplasm
Q05980	DNAJ_BRUOV	cytoplasm (probable).	cytoplasm
Q05981	DNAK_BRUOV		cytoplasm
Q06064	FLAA_BORBR		flagellar
Q06065	ATOC_ECOLI	cytoplasm (probable).	cytoplasm
Q06081	MXIJ_SHIFL	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q06119	KDGL_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q06126	TNPA_BORPA		cytoplasm
Q06131	YOPD_YERPS		flagellar
Q06171	FLGF_CAUCR		flagellar
Q06172	FLGG_CAUCR		flagellar
Q06191	AATB2_RHIME	cytoplasm (by similarity).	cytoplasm
Q06198	RPSH_PSEAE		cytoplasm
Q06206	UREG_PROMI		cytoplasm
Q06277	P76_HAESO		cytoplasm
Q06365	ALYP_PSESO		extracellular
Q06399	YEDD_SALTY	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q06400	YEDE_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q06447	RHO_NEIGO		inner membrane
Q06457	NASA_KLEOX		periplasm
Q06458	NIRB_KLEOX		cytoplasm
Q06527	ANKH_CHRVI		cytoplasm

Q06552	PRTN_PSEAE		cytoplasm
Q06553	PRTR_PSEAE		cytoplasm
Q06555	AEPA_PECCC		periplasm
Q06578	IMM1_PSEAE		cytoplasm
Q06579	IMM2_PSEAE		cytoplasm
Q06581	PILT_NEIGO	cytoplasm (probable).	inner membrane
Q06583	PYS1_PSEAE		extracellular
Q06584	PYS2_PSEAE		extracellular
Q06607	IHFB_RHOCA		cytoplasm
Q06610	RBCR_THIFE		cytoplasm
Q06741	GSA_XANCH	cytoplasm (potential).	cytoplasm
Q06758	HISJ_NEIGO	bacterial cell membrane; lipid- anchor (probable).	inner membrane
Q06778	BLOB_PSEAE		extracellular
Q06816	HYEP_STIAU		extracellular
Q06823	SP21_STIAU		cytoplasm
Q06877	LUXP_PHOLE		cytoplasm
Q06878	LUXD2_PHOLE		cytoplasm
Q06903	PPCE_AERHY		cytoplasm
Q06909	CARQ_MYXXA		cytoplasm
Q06911	CARS_MYXXA		cytoplasm
Q06929	YCRS_MYXXA		extracellular
Q06951	RFBB_VIBCH	bacterial membrane (probable).	cytoplasm
Q06952	GM4D1_VIBCH		cytoplasm
Q06963	HLDD_VIBCH		cytoplasm
Q06965	HEM0_RHOS4		inner membrane
Q06968	FLIC_SALBE		flagellar
Q06969	FLIC_SALBU		flagellar
Q06970	FLIC_SALDE		flagellar
Q06971	FLIC_SALDU		flagellar
Q06972	FLIC_SALEN		flagellar
Q06973	FLIC_SALMO		flagellar
Q06974	FLIC_SALON		flagellar
Q06981	FLIC_SALMC		flagellar
Q06982	FLIC_SALRO		flagellar
Q06983	FLIC_SALSE		flagellar
Q06988	TBB2_NEIMB	bacterial cell outer membrane; lipid- anchor (probable).	outer membrane
Q06994	RFAB_SALTY		cytoplasm
Q06995	RFAP_SALTY		inner membrane
Q07024	RFBA_VIBCH		cytoplasm
Q07083	RPSC_MYXXA		cytoplasm
Q07087	RBL1B_THIFE		extracellular
Q07088	RBS2_THIFE		cytoplasm
Q07177	NIFS_RHOCA		cytoplasm
Q07178	NIU1_RHOCA		cytoplasm
Q07179	NIFV_RHOCA		cytoplasm
Q07180	NIFW_RHOCA		cytoplasm
Q07184	YNIU_RHOCA		inner membrane
Q07232	ATPB_BUCAP		inner membrane
Q07251	LDH_RALEU	cytoplasm (probable).	cytoplasm
Q07252	MEMP_RALEU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q07267	LEXA_PRORE		cytoplasm

Q07405	ATPA_MYXXA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q07408	MSP4_ANAMA		outer membrane
Q07411	PPK_KLEPN		cytoplasm
Q07447	RECA_PSEPU	cytoplasm (by similarity).	cytoplasm
Q07559	NIFW_AZOBR		cytoplasm
Q07564	ECPC_EIKCO		extracellular
Q07565	ECPD_EIKCO		extracellular
Q07568	IPGF_SHIFL		extracellular
Q07601	CEST_CITFR	cytoplasm (by similarity).	cytoplasm
Q07602	YMSA_RHIME		periplasm
Q07607	MOSA_RHIME		cytoplasm
Q07608	MOSB_RHIME		cytoplasm
Q07609	MOSC_RHIME	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q07698	ABCA_AERSA		inner membrane
Q07736	T1RA_ECOLI		cytoplasm
Q07737	CHVG_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q07755	NODC_AZOCA	bacterial cell membrane; peripheral membrane protein (probable).	inner membrane
Q07756	NODI_AZOCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q07757	NODJ_AZOCA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q07758	NODS_AZOCA		cytoplasm
Q07759	NODU_AZOCA		inner membrane
Q07765	DEVR_MYXXA		cytoplasm
Q07766	DEVS_MYXXA		cytoplasm
Q07783	CHVI_AGRTU	cytoplasm (potential).	cytoplasm
Q07792	ESTE_VIBMI		extracellular
Q07806	PBPA_PSEAE	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q07809	RECA_PSEST	cytoplasm (by similarity).	cytoplasm
Q07910	FLAB_HELMU		flagellar
Q07911	FLAB_HELPY		flagellar
Q07933	ANFD_RHOCA		cytoplasm
Q07934	ANFG_RHOCA		periplasm
Q07935	ANFK_RHOCA		cytoplasm
Q07942	NIFH2_RHOCA		cytoplasm
Q07944	BEDC1_PSEPU		cytoplasm
Q07945	BEDC2_PSEPU		cytoplasm
Q07946	BEDA_PSEPU		cytoplasm
Q07947	BEDB_PSEPU		cytoplasm
Q08015	MIAE_SALTY		cytoplasm
Q08016	OTC_SALTY	cytoplasm (probable).	cytoplasm
Q08021	YJGM_SALTY		cytoplasm

Q08022	YJGN_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q08082	T6501_BRUOV		cytoplasm
Q08111	DUS_RHOCA		cytoplasm
Q08113	ISPDF_RHOCA		cytoplasm
Q08318	DMA_VIBCH		cytoplasm
Q08327	RECA_ACEPO	cytoplasm (by similarity).	cytoplasm
Q08346	PRIM_HAEIN		cytoplasm
Q08370	HRMA_PSESY		extracellular
Q08382	MODB_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q08383	MODA_RHOCA	periplasm (by similarity).	periplasm
Q08384	MODD_RHOCA		cytoplasm
Q08385	MOPA_RHOCA		cytoplasm
Q08386	MOPB_RHOCA		cytoplasm
Q08456	FIMI_SALTI		extracellular
Q08530	BTR_BORPE		cytoplasm
Q08597	NAC_KLEAE		cytoplasm
Q08598	CBL_KLEAE		cytoplasm
Q08633	RTX3D_ACTPL	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q08675	HLY1_AERSA		extracellular
Q08677	HLY4_AERSA		extracellular
Q08678	AVRV_XANCV		extracellular
Q08698	RIR3_SALTY		cytoplasm
Q08715	URE23_HELFE	cytoplasm (by similarity).	cytoplasm
Q08716	URE1_HELFE	cytoplasm (by similarity).	cytoplasm
Q08790	BLC_VIBCH	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q08812	SYRM_RHIET		cytoplasm
Q08815	YSYM_RHIET		cytoplasm
Q08860	FLIC_SHIFL		flagellar
Q08868	PLPA_PASHA	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q08869	PLPB_PASHA	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q08870	PLPC_PASHA	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q08880	DPO3E_BUCAP		cytoplasm
Q08885	RNH_BUCAP	cytoplasm (potential).	extracellular
Q08889	GLO2_BUCAP		cytoplasm
Q09057	TBB1_NEIMB	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q09063	URED_KLEAE		cytoplasm
Q09065	UREF_HELPY		cytoplasm
Q09066	UREG_HELPY		cytoplasm
Q09067	UREH_HELPY		cytoplasm
Q09102	HYIN_AGRRH		extracellular
Q09109	TR2M_AGRRH		cytoplasm
Q10373	NIU2_RHOCA		cytoplasm
Q10725	FSDH_BURPI		cytoplasm
Q27710	CATA_ONCVE		periplasm
Q2A1K3	ACON_FRATH		cytoplasm

Q2G300	RS20_NOVAD		cytoplasm
Q2G348	THIG_NOVAD	cytoplasm (by similarity).	cytoplasm
Q2G3T3	CLPP_NOVAD	cytoplasm (by similarity).	cytoplasm
Q2G480	PPCK_NOVAD	cytoplasm (by similarity).	cytoplasm
Q2G5N7	ATPA_NOVAD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2G630	PURL_NOVAD	cytoplasm (by similarity).	cytoplasm
Q2G646	GLYA_NOVAD	cytoplasm (by similarity).	cytoplasm
Q2G6I4	SYU_NOVAD	cytoplasm (by similarity).	cytoplasm
Q2G733	AROK_NOVAD	cytoplasm (probable).	cytoplasm
Q2G794	RNH2_NOVAD	cytoplasm (potential).	extracellular
Q2G7F1	ISPE_NOVAD		cytoplasm
Q2G8G5	RL9_NOVAD		cytoplasm
Q2G8J0	CCME_NOVAD	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2G8X1	RS17_NOVAD		cytoplasm
Q2G8X7	RL2_NOVAD		cytoplasm
Q2G8Y1	RS10_NOVAD		cytoplasm
Q2G8Y5	RS12_NOVAD		cytoplasm
Q2G998	MRAY_NOVAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2G9Z8	MSCL_NOVAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2GAB9	XYLA_NOVAD	cytoplasm (by similarity).	cytoplasm
Q2GBA2	RUVB_NOVAD		cytoplasm
Q2GC33	RPOZ_NOVAD		cytoplasm
Q2GCD2	RL10_NOVAD		cytoplasm
Q2GCD7	RPOB_NOVAD		cytoplasm
Q2GCU8	CCME_NEOSM	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2GCX9	RS20_NEOSM		cytoplasm
Q2GD19	CLPP_NEOSM	cytoplasm (by similarity).	cytoplasm
Q2GD80	RS12_NEOSM		cytoplasm
Q2GD90	RPOB_NEOSM		cytoplasm
Q2GDR7	HTPG_NEOSM	cytoplasm (by similarity).	cytoplasm
Q2GDS9	THIG_NEOSM	cytoplasm (by similarity).	cytoplasm
Q2GEE0	RS10_NEOSM		cytoplasm
Q2GER2	ISPF_NEOSM		extracellular
Q2GER5	ATPA_NEOSM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2GEY5	RL9_NEOSM		cytoplasm
Q2GFT8	CLPP_EHRCR	cytoplasm (by similarity).	cytoplasm
Q2GFY5	HTPG_EHRCR	cytoplasm (by similarity).	cytoplasm
Q2GG39	RPOZ_EHRCR		cytoplasm
Q2GG46	CCME_EHRCR	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane

Q2GH53	RL2_EHRCR		cytoplasm
Q2GH57	RS10_EHRCR		cytoplasm
Q2GHF2	RL9_EHRCR		cytoplasm
Q2GHV0	ISPF_EHRCR		extracellular
Q2GHX3	ATPA_EHRCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2GI32	RS20_EHRCR		cytoplasm
Q2GIC8	RS20_ANAPZ		cytoplasm
Q2GIG2	ATPA_ANAPZ	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2GIK8	ISPF_ANAPZ		extracellular
Q2GJ68	RPOB_ANAPZ		cytoplasm
Q2GJB4	CLPP_ANAPZ	cytoplasm (by similarity).	cytoplasm
Q2GJI3	HTPG_ANAPZ	cytoplasm (by similarity).	cytoplasm
Q2GKI0	CCME_ANAPZ	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2GKJ2	RPOZ_ANAPZ		cytoplasm
Q2GL56	RL2_ANAPZ		cytoplasm
Q2GL60	RS10_ANAPZ		cytoplasm
Q2GLH4	RL9_ANAPZ		cytoplasm
Q2IEW1	CLPP_ANADE	cytoplasm (by similarity).	cytoplasm
Q2IFJ5	RPOZ_ANADE		cytoplasm
Q2IG26	MRAY_ANADE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2IGV9	PAND_ANADE		cytoplasm
Q2IHC1	PURL_ANADE	cytoplasm (by similarity).	cytoplasm
Q2IHQ9	ATPA_ANADE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2II80	RL10_ANADE		cytoplasm
Q2II86	RPOB_ANADE		cytoplasm
Q2IJ76	RS17_ANADE		cytoplasm
Q2IJ85	RS10_ANADE		cytoplasm
Q2IJ88	RL2_ANADE		cytoplasm
Q2IJ91	RS12_ANADE		cytoplasm
Q2IKW1	THIG_ANADE	cytoplasm (by similarity).	cytoplasm
Q2ILG4	RS20_ANADE		cytoplasm
Q2ILI1	GLYA_ANADE	cytoplasm (by similarity).	cytoplasm
Q2ILU8	RECF_ANADE	cytoplasm (by similarity).	cytoplasm
Q2IM65	RL9_ANADE		cytoplasm
Q2IM67	ISPE_ANADE		cytoplasm
Q2IMM4	RNH2_ANADE	cytoplasm (potential).	extracellular
Q2INZ4	SURE_ANADE	cytoplasm (potential).	periplasm
Q2IPJ5	RUVB_ANADE		cytoplasm
Q2IPT7	SYI_ANADE	cytoplasm (by similarity).	cytoplasm
Q2IQP0	FLGH_ANADE	outer membrane; lipid-anchor (probable).	inner membrane
Q2IQP1	FLGI_ANADE	periplasm (by similarity).	periplasm

Q2JYF0	CCME_RHIEC	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2JZQ2	TPIS2_RHIEC		cytoplasm
Q2K4E6	Y3534_RHIEC		extracellular
Q2K869	TPIS1_RHIEC		cytoplasm
Q2KAE9	RPOZ_RHIEC		cytoplasm
Q2KCH6	CHED_RHIEC		cytoplasm
Q2KCH7	CHEY_RHIEC	cytoplasm (potential).	cytoplasm
Q2KCH9	CHER_RHIEC		cytoplasm
Q2KCI0	CHEW_RHIEC	cytoplasm (by similarity).	cytoplasm
Q2KU34	ATPA_BORA1	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2KUI1	THIG_BORA1	cytoplasm (by similarity).	cytoplasm
Q2KUX6	ISPF_BORA1		extracellular
Q2KV13	SYI_BORA1	cytoplasm (by similarity).	cytoplasm
Q2KV15	GLYA_BORA1	cytoplasm (by similarity).	cytoplasm
Q2KVY2	RUVB_BORA1		cytoplasm
Q2KXM7	NAGZ_BORA1	cytoplasm (by similarity).	extracellular
Q2KY48	PAND_BORA1		cytoplasm
Q2KYY2	HTPG_BORA1	cytoplasm (by similarity).	cytoplasm
Q2KZ23	RL9_BORA1		cytoplasm
Q2KZD7	RS20_BORA1		cytoplasm
Q2L006	SURE_BORA1	cytoplasm (potential).	periplasm
Q2L066	RPOZ_BORA1		cytoplasm
Q2L146	RNH2_BORA1	cytoplasm (potential).	extracellular
Q2L1B1	FLGI_BORA1	periplasm (by similarity).	periplasm
Q2L1Z2	AROK_BORA1	cytoplasm (probable).	cytoplasm
Q2L256	CLPP_BORA1	cytoplasm (by similarity).	cytoplasm
Q2L2B2	RS17_BORA1		cytoplasm
Q2L2F1	RL2_BORA1		cytoplasm
Q2L2G3	RS10_BORA1		cytoplasm
Q2L2H8	RS12_BORA1		cytoplasm
Q2L2M3	RPOB_BORA1		cytoplasm
Q2LQ83	RPOZ_SYNAS		cytoplasm
Q2LQ87	RPOB_SYNAS		cytoplasm
Q2LQ89	RL10_SYNAS		cytoplasm
Q2LQ98	RL2_SYNAS		cytoplasm
Q2LQA2	RS10_SYNAS		cytoplasm
Q2LQB1	RS17_SYNAS		cytoplasm
Q2LQK7	RS20_SYNAS		cytoplasm
Q2LQM6	GLYA_SYNAS	cytoplasm (by similarity).	cytoplasm
Q2LQZ7	ATPA1_SYNAS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2LR51	MRAY_SYNAS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2LR98	SYI_SYNAS	cytoplasm (by similarity).	cytoplasm
Q2LRA8	RUVB_SYNAS		cytoplasm
Q2LSQ4	PAND_SYNAS		cytoplasm
Q2LT20	FLGI_SYNAS	periplasm (by similarity).	periplasm
Q2LT21	FLGH_SYNAS	outer membrane; lipid-anchor (probable).	inner membrane

Q2LTK5	CLPP_SYNAS	cytoplasm (by similarity).	cytoplasm
Q2LUD6	AROK_SYNAS	cytoplasm (probable).	cytoplasm
Q2LUH7	SURE_SYNAS	cytoplasm (potential).	periplasm
Q2LUI9	MSCL_SYNAS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2LUJ5	RL9_SYNAS		cytoplasm
Q2LUJ9	ISPE_SYNAS		cytoplasm
Q2LUS9	ISPD_SYNAS		cytoplasm
Q2LUT1	ISPF_SYNAS		extracellular
Q2LVU5	RNH2_SYNAS	cytoplasm (potential).	extracellular
Q2LY34	ATPA2_SYNAS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2NQ88	ATPA_SODGM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2NQF9	CORA_SODGM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2NQI3	PPCK_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NQI4	HSLO_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NQJ1	AROK_SODGM	cytoplasm (probable).	cytoplasm
Q2NQL1	TUSD_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NQL2	TUSC_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NQL3	TUSB_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NQL4	RS12_SODGM		cytoplasm
Q2NQM1	RS10_SODGM		cytoplasm
Q2NQM5	RL2_SODGM		cytoplasm
Q2NQN1	RS17_SODGM		cytoplasm
Q2NQQ0	MSCL_SODGM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2NQS8	RPOZ_SODGM		cytoplasm
Q2NRM1	RNH2_SODGM	cytoplasm (potential).	extracellular
Q2NRS1	ISPE_SODGM		cytoplasm
Q2NS25	GLYA_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NSC1	ZIPA_SODGM	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q2NT02	SYI_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NTC0	Y1330_SODGM		inner membrane
Q2NTI8	RUVB_SODGM		cytoplasm
Q2NTS8	PGSA_SODGM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2NU63	TUSE_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NV58	HTPG_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NV79	CLPP_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NVB6	AROL_SODGM	cytoplasm (probable).	cytoplasm
Q2NVM2	SURE_SODGM	cytoplasm (potential).	periplasm
Q2NVM3	ISPF_SODGM		extracellular
Q2NVM4	ISPD_SODGM		cytoplasm
Q2NVR4	PAND_SODGM		cytoplasm

Q2NVV4	MRAY_SODGM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2NVY8	RS20_SODGM		cytoplasm
Q2NW52	RL9_SODGM		cytoplasm
Q2NW94	CH60_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NWR6	RPOB_SODGM		cytoplasm
Q2NWR8	RL10_SODGM		cytoplasm
Q2NWX1	TUSA_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NX13	FLGI_SODGM	periplasm (by similarity).	periplasm
Q2NX47	RECF_SODGM	cytoplasm (by similarity).	cytoplasm
Q2NXR2	XYLA_XANOM	cytoplasm (by similarity).	cytoplasm
Q2NYE5	NUDH_XANOM		cytoplasm
Q2NZ83	GLYA_XANOM	cytoplasm (by similarity).	cytoplasm
Q2NZ85	NRDR_XANOM		cytoplasm
Q2NZB0	MRAZ_XANOM		inner membrane
Q2NZB6	MRAY_XANOM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2NZS1	TAL_XANOM	cytoplasm (probable).	cytoplasm
Q2NZW6	ISPE_XANOM		cytoplasm
Q2NZX6	RL10_XANOM		cytoplasm
Q2NZX8	RPOB_XANOM		cytoplasm
Q2NZY0	RS12_XANOM		cytoplasm
Q2NZY4	RS10_XANOM		cytoplasm
Q2NZY8	RL2_XANOM		cytoplasm
Q2NZZ4	RS17_XANOM		cytoplasm
Q2P0I7	MSCL_XANOM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2P0J4	TRMB_XANOM		cytoplasm
Q2P0J5	THIG_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P0W9	Y3053_XANOM		inner membrane
Q2P0X3	TRUB_XANOM		cytoplasm
Q2P1A2	PHS_XANOM		cytoplasm
Q2P1G3	CCME1_XANOM	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2P1L0	ISPD_XANOM		cytoplasm
Q2P1L1	ISPF_XANOM		extracellular
Q2P1L2	TRUD_XANOM		cytoplasm
Q2P1L4	SURE_XANOM	cytoplasm (potential).	periplasm
Q2P2G0	PROB_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P2G1	PROA_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P2N7	FLGI_XANOM	periplasm (by similarity).	periplasm
Q2P2N8	FLGH_XANOM	outer membrane; lipid-anchor (probable).	inner membrane
Q2P2U8	HTPG_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P310	PPNK_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P328	RS6_XANOM		cytoplasm
Q2P330	RL9_XANOM		cytoplasm
Q2P379	PAND_XANOM		cytoplasm
Q2P380	G6PI_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P3E8	LPXK_XANOM		cytoplasm
Q2P3J8	HIS6_XANOM	cytoplasm (by similarity).	cytoplasm

Q2P3J9	HIS4_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P3K0	HIS5_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P3K2	HIS8_XANOM		cytoplasm
Q2P3K3	HISX_XANOM		cytoplasm
Q2P3Q9	GUAA_XANOM		cytoplasm
Q2P3V2	CCME2_XANOM	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2P3Z2	METX_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P497	GLND_XANOM		cytoplasm
Q2P4B6	FABZ_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P4B9	RNH2_XANOM	cytoplasm (potential).	extracellular
Q2P4L0	NAGZ_XANOM	cytoplasm (by similarity).	extracellular
Q2P4L7	RUMA_XANOM		cytoplasm
Q2P575	RUVB_XANOM		cytoplasm
Q2P5B3	RS20_XANOM		cytoplasm
Q2P5R0	GPMA_XANOM		cytoplasm
Q2P6C8	AROK_XANOM	cytoplasm (probable).	cytoplasm
Q2P6M7	RPOZ_XANOM		cytoplasm
Q2P6Z0	CLPP_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P7Q6	ATPA_XANOM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2P892	SYX_XANOM	cytoplasm (by similarity).	cytoplasm
Q2P9F6	GLGA_XANOM		cytoplasm
Q2P9L0	PDXJ_XANOM	cytoplasm (probable).	cytoplasm
Q2RMP9	RS20_RHORT		cytoplasm
Q2RMR8	TRUB_RHORT		cytoplasm
Q2RMS3	Y3778_RHORT		inner membrane
Q2RMS4	TRMB_RHORT		cytoplasm
Q2RMW0	AROB_RHORT	cytoplasm (probable).	cytoplasm
Q2RMW1	AROK_RHORT	cytoplasm (probable).	cytoplasm
Q2RNA5	HIS5_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RNA6	HIS4_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RNA7	HIS6_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RNA8	HIS2_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RNC0	PHS_RHORT		cytoplasm
Q2RNG2	GLND_RHORT		cytoplasm
Q2RP84	METX_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RP86	HIS8_RHORT		cytoplasm
Q2RPE1	RNH2_RHORT	cytoplasm (potential).	extracellular
Q2RPW9	GPH_RHORT		cytoplasm
Q2RQ51	G6PI_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RQ67	MURA_RHORT	cytoplasm (probable).	inner membrane
Q2RQE9	FLGI_RHORT	periplasm (by similarity).	periplasm
Q2RQF3	FLGH_RHORT	outer membrane; lipid-anchor (probable).	inner membrane
Q2RQM6	HIS1_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RQM7	HISX_RHORT		cytoplasm
Q2RQV0	RL1_RHORT		cytoplasm
Q2RQV1	RL10_RHORT		cytoplasm
Q2RQV3	RPOB_RHORT		cytoplasm
Q2RQV5	RS12_RHORT		cytoplasm
Q2RQV9	RS10_RHORT		cytoplasm
Q2RQW3	RL2_RHORT		cytoplasm

Q2RQW6	RS3_RHORT		cytoplasm
Q2RQW9	RS17_RHORT		cytoplasm
Q2RQX4	RS8_RHORT		cytoplasm
Q2RQX7	RS5_RHORT		cytoplasm
Q2RQY2	RS13_RHORT		cytoplasm
Q2RQY3	RS11_RHORT		cytoplasm
Q2RRL3	THIG_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RRZ2	DEOC_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RS50	GLGA_RHORT		cytoplasm
Q2RSB2	PNTAA_RHORT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2RSB3	PNTAB_RHORT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2RSB4	PNTB_RHORT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2RSR3	SYI_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RT88	RPOZ_RHORT		cytoplasm
Q2RT91	PDXJ_RHORT	cytoplasm (probable).	cytoplasm
Q2RTB8	GLYA2_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RTB9	NRDR_RHORT		cytoplasm
Q2RTH6	SURE_RHORT	cytoplasm (potential).	periplasm
Q2RTY3	HIS3_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RTZ9	FABZ_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RU45	CLPP_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RUN7	TAL_RHORT	cytoplasm (probable).	cytoplasm
Q2RV05	PROB_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RV06	PROA_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RV14	NUDH_RHORT		cytoplasm
Q2RV20	ATPA_RHORT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2RVA2	GLYA1_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RVF5	RUVB_RHORT		cytoplasm
Q2RVM4	CYC2_RHORT		periplasm
Q2RVT5	MRAZ_RHORT		inner membrane
Q2RVU1	MRAY_RHORT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2RWI3	PURL_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RWJ0	RS4_RHORT		cytoplasm
Q2RWV9	LPXK_RHORT		cytoplasm
Q2RXC6	RS6_RHORT		cytoplasm
Q2RXC9	RL9_RHORT		cytoplasm
Q2RXI5	GUAA_RHORT		cytoplasm
Q2RXQ9	QUEA_RHORT	cytoplasm (potential).	cytoplasm
Q2RXS7	ISPE_RHORT		cytoplasm
Q2RYB8	HTPG_RHORT	cytoplasm (by similarity).	cytoplasm
Q2RYF4	CCME_RHORT	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane

Q2S6N9	ATPA2_HAHCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2S8R3	RPOZ_HAHCH		cytoplasm
Q2S8Y9	SYU_HAHCH	cytoplasm (by similarity).	cytoplasm
Q2S903	RL10_HAHCH		cytoplasm
Q2S905	RPOB_HAHCH		cytoplasm
Q2S907	RS12_HAHCH		cytoplasm
Q2S911	RS10_HAHCH		cytoplasm
Q2S915	RL2_HAHCH		cytoplasm
Q2S921	RS17_HAHCH		cytoplasm
Q2S9Q5	AROK_HAHCH	cytoplasm (probable).	cytoplasm
Q2S9R4	GLYA2_HAHCH	cytoplasm (by similarity).	cytoplasm
Q2S9T5	RS20_HAHCH		cytoplasm
Q2S9Y9	MRAY_HAHCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2SBR2	RNH2_HAHCH	cytoplasm (potential).	extracellular
Q2SCL3	RUVB_HAHCH		cytoplasm
Q2SDU4	FLGH_HAHCH	outer membrane; lipid-anchor (probable).	inner membrane
Q2SDU5	FLGI2_HAHCH	periplasm (by similarity).	periplasm
Q2SE53	CCME_HAHCH	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2SEX9	FLGI1_HAHCH	periplasm (by similarity).	periplasm
Q2SFI7	GLYA1_HAHCH	cytoplasm (by similarity).	cytoplasm
Q2SK36	CLPP_HAHCH	cytoplasm (by similarity).	cytoplasm
Q2SKD0	HTPG_HAHCH	cytoplasm (by similarity).	cytoplasm
Q2SKW4	SURE_HAHCH	cytoplasm (potential).	periplasm
Q2SKW6	ISPF_HAHCH		extracellular
Q2SKW7	ISPD_HAHCH		cytoplasm
Q2SLA0	ISPE_HAHCH		cytoplasm
Q2SLB3	RL9_HAHCH		cytoplasm
Q2SNG7	ATPA1_HAHCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2SPF8	HSLO_HAHCH	cytoplasm (by similarity).	cytoplasm
Q2SPF9	PPCK_HAHCH	cytoplasm (by similarity).	cytoplasm
Q2SQL5	THIG_HAHCH	cytoplasm (by similarity).	cytoplasm
Q2SQZ4	RECF_HAHCH	cytoplasm (by similarity).	cytoplasm
Q2STE7	ATPA1_BURTA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2SU17	RL10_BURTA		cytoplasm
Q2SU19	RPOB_BURTA		cytoplasm
Q2SU22	RS12_BURTA		cytoplasm
Q2SU26	RS10_BURTA		cytoplasm
Q2SU30	RL2_BURTA		cytoplasm
Q2SU36	RS17_BURTA		cytoplasm
Q2SU71	AROK_BURTA	cytoplasm (probable).	cytoplasm
Q2SU89	THIG_BURTA	cytoplasm (by similarity).	cytoplasm
Q2SW40	XYLA_BURTA	cytoplasm (by similarity).	cytoplasm
Q2SWF5	SURE_BURTA	cytoplasm (potential).	periplasm
Q2SWJ6	RL9_BURTA		cytoplasm

Q2SWQ6	CLPP_BURTA	cytoplasm (by similarity).	cytoplasm
Q2SWT5	ISPF_BURTA		extracellular
Q2SWT6	ISPD_BURTA		cytoplasm
Q2SWU6	M_SCL_BURTA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2SWY4	RNH2_BURTA	cytoplasm (potential).	extracellular
Q2SY71	RPOZ_BURTA		cytoplasm
Q2SYF5	URE3_BURTA	cytoplasm (by similarity).	cytoplasm
Q2SYF6	URE2_BURTA	cytoplasm (by similarity).	cytoplasm
Q2SYS4	GLYA1_BURTA	cytoplasm (by similarity).	cytoplasm
Q2SZ55	RUVB_BURTA		cytoplasm
Q2SZ62	SY_Y_BURTA	cytoplasm (by similarity).	cytoplasm
Q2SZI6	MRAY_BURTA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2SZZ0	HTPG_BURTA	cytoplasm (by similarity).	cytoplasm
Q2T096	PAND_BURTA		cytoplasm
Q2T0K8	RS20_BURTA		cytoplasm
Q2T1B6	ISPE_BURTA		cytoplasm
Q2T437	GLYA2_BURTA	cytoplasm (by similarity).	cytoplasm
Q2T873	ATPA2_BURTA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2T8B0	DNS2_BURTA		extracellular
Q2T8V6	FLGI2_BURTA	periplasm (by similarity).	periplasm
Q2T8V7	FLGH2_BURTA	outer membrane; lipid-anchor (probable).	inner membrane
Q2VYT6	ISPE_MAGMM		cytoplasm
Q2VZ28	HTPG_MAGMM	cytoplasm (by similarity).	cytoplasm
Q2VZH2	CCME_MAGMM	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2VZN0	ATPA_MAGMM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2W095	AROK_MAGMM	cytoplasm (probable).	cytoplasm
Q2W0H6	MRAY_MAGMM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2W0J7	FLGI_MAGMM	periplasm (by similarity).	periplasm
Q2W2A5	RUVB_MAGMM		cytoplasm
Q2W2H9	RL10_MAGMM		cytoplasm
Q2W2I1	RPOB_MAGMM		cytoplasm
Q2W2I6	RS12_MAGMM		cytoplasm
Q2W2J0	RS10_MAGMM		cytoplasm
Q2W2J4	RL2_MAGMM		cytoplasm
Q2W2K0	RS17_MAGMM		cytoplasm
Q2W2T6	SY_Y_MAGMM	cytoplasm (by similarity).	cytoplasm
Q2W3H9	CLPP_MAGMM	cytoplasm (by similarity).	cytoplasm
Q2W3R9	THIG_MAGMM	cytoplasm (by similarity).	cytoplasm
Q2W4A1	SURE_MAGMM	cytoplasm (potential).	periplasm
Q2W4T2	GLYA_MAGMM	cytoplasm (by similarity).	cytoplasm
Q2W519	RPOZ_MAGMM		cytoplasm
Q2W5H3	RL9_MAGMM		cytoplasm

Q2W9N6	RS20_MAGMM		cytoplasm
Q2Y5G5	RUVB_NITMU		cytoplasm
Q2Y5Q1	GLND_NITMU		cytoplasm
Q2Y629	MRAZ_NITMU		inner membrane
Q2Y635	MRAY_NITMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2Y6A2	QUEA_NITMU	cytoplasm (potential).	cytoplasm
Q2Y6B0	PPNK_NITMU	cytoplasm (by similarity).	cytoplasm
Q2Y6G2	GPH_NITMU		cytoplasm
Q2Y6J0	CLPP_NITMU	cytoplasm (by similarity).	cytoplasm
Q2Y6W3	RUMA_NITMU		cytoplasm
Q2Y6Y6	HIS82_NITMU		cytoplasm
Q2Y742	DUT_NITMU		extracellular
Q2Y751	ISPD_NITMU		cytoplasm
Q2Y752	ISPF_NITMU		extracellular
Q2Y7M5	RS6_NITMU		cytoplasm
Q2Y7M8	RL9_NITMU		cytoplasm
Q2Y7W0	Y1866_NITMU		inner membrane
Q2Y7W4	TRUB_NITMU		cytoplasm
Q2Y7Y3	PROB_NITMU	cytoplasm (by similarity).	cytoplasm
Q2Y863	NAGZ_NITMU	cytoplasm (by similarity).	extracellular
Q2Y865	PDXJ_NITMU	cytoplasm (probable).	cytoplasm
Q2Y8G9	ATPA2_NITMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2Y9E0	FLGI_NITMU	periplasm (by similarity).	periplasm
Q2Y9M8	URE2_NITMU	cytoplasm (by similarity).	cytoplasm
Q2Y9M9	URE3_NITMU	cytoplasm (by similarity).	cytoplasm
Q2Y9Q8	CCME_NITMU	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2Y9R9	GUAA_NITMU		cytoplasm
Q2Y9Z7	GPMA2_NITMU		cytoplasm
Q2YA09	HTPG_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YA81	RS20_NITMU		cytoplasm
Q2YAN2	MURA_NITMU	cytoplasm (probable).	inner membrane
Q2YAU4	HIS1_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YAU5	HISX_NITMU		cytoplasm
Q2YAU6	HIS81_NITMU		cytoplasm
Q2YAU8	HIS5_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YAU9	HIS4_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YAV0	HIS6_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YAV1	HIS3_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YAV2	HIS2_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YAX3	RS4_NITMU		cytoplasm
Q2YAX4	RS11_NITMU		cytoplasm
Q2YAX5	RS13_NITMU		cytoplasm
Q2YAY0	RS5_NITMU		cytoplasm
Q2YAY3	RS8_NITMU		cytoplasm
Q2YAY8	RS17_NITMU		cytoplasm
Q2YAZ1	RS3_NITMU		cytoplasm
Q2YAZ4	RL2_NITMU		cytoplasm
Q2YAZ8	RS10_NITMU		cytoplasm
Q2YB02	RS12_NITMU		cytoplasm

Q2YB05	RPOB_NITMU		cytoplasm
Q2YB07	RL10_NITMU		cytoplasm
Q2YB08	RL1_NITMU		cytoplasm
Q2YB48	GLGA_NITMU		cytoplasm
Q2YB52	PHS_NITMU		cytoplasm
Q2YB61	AROB_NITMU	cytoplasm (probable).	cytoplasm
Q2YB96	RNH2_NITMU	cytoplasm (potential).	extracellular
Q2YB97	FABZ_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YBB2	AROK_NITMU	cytoplasm (probable).	cytoplasm
Q2YBH5	ISPE_NITMU		cytoplasm
Q2YBP9	PROA_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YBR1	SYI_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YBR8	SURE_NITMU	cytoplasm (potential).	periplasm
Q2YBU4	G6PI_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YBW4	NUDH_NITMU		cytoplasm
Q2YBZ1	GPMA1_NITMU		cytoplasm
Q2YCA5	ATPA1_NITMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2YCJ2	METX_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YCZ2	THIG_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YCZ3	TRMB_NITMU		cytoplasm
Q2YD11	RPOZ_NITMU		cytoplasm
Q2YD57	NRDR_NITMU		cytoplasm
Q2YD58	GLYA_NITMU	cytoplasm (by similarity).	cytoplasm
Q2YIH4	LPXK_BRUA2		cytoplasm
Q2YIJ8	GLUP_BRUA2	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q2YIJ9	HIS1_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YIQ6	TPIS2_BRUA2		cytoplasm
Q2YJ67	FLGH_BRUA2	outer membrane; lipid-anchor (probable).	inner membrane
Q2YJN6	GPMA_BRUA2		cytoplasm
Q2YJN7	DAPB_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YJP6	UPPP_BRUA2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2YK38	GUAA_BRUA2		cytoplasm
Q2YKX9	GCSP_BRUA2		cytoplasm
Q2YLI5	ATPA_BRUA2	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q2YLI7	PROA_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YLJ4	NUDH_BRUA2		cytoplasm
Q2YLM3	PROB_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YLP6	RL19_BRUA2		cytoplasm
Q2YLZ8	RS12_BRUA2		cytoplasm
Q2YLZ9	RS7_BRUA2		cytoplasm
Q2YM02	RS10_BRUA2		cytoplasm
Q2YM06	RL2_BRUA2		cytoplasm
Q2YM12	RL1_BRUA2		cytoplasm
Q2YM13	RL10_BRUA2		cytoplasm
Q2YM15	RPOB_BRUA2		cytoplasm
Q2YM63	MRAW_BRUA2		cytoplasm

Q2YM68	MRAY_BRUA2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2YMB5	ISPE_BRUA2		cytoplasm
Q2YMC3	RNH2_BRUA2	cytoplasm (potential).	extracellular
Q2YMG9	RS6_BRUA2		cytoplasm
Q2YMH2	RL9_BRUA2		cytoplasm
Q2YMK2	MTB1_BRUA2		cytoplasm
Q2YMQ2	XYLA_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YMU7	CCME_BRUA2	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q2YMX6	DEGP_BRUA2	periplasm (potential).	periplasm
Q2YN02	RNC_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YN04	ACPS_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YN12	RPOZ_BRUA2		cytoplasm
Q2YN35	UVRC_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YN40	NDK_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YN94	NRDR_BRUA2		cytoplasm
Q2YN95	GLYA_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YNG6	PURL_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YNH2	RS4_BRUA2		cytoplasm
Q2YNM0	SECD_BRUA2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2YNM1	YAJC_BRUA2		inner membrane
Q2YNM5	SURE_BRUA2	cytoplasm (potential).	periplasm
Q2YNP1	AROQ_BRUA2		cytoplasm
Q2YNR0	SYU_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YNV0	PHS_BRUA2		cytoplasm
Q2YNZ1	GLND_BRUA2		cytoplasm
Q2YP58	THIG_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YPB6	MURA_BRUA2	cytoplasm (probable).	inner membrane
Q2YPB8	HISX_BRUA2		cytoplasm
Q2YPD5	URE11_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YPD6	URE21_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YPD7	URE31_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YPF3	G6PI_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YPJ2	MSCL_BRUA2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2YPM3	RECF_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YPN6	AROQ_BRUA2	cytoplasm (probable).	cytoplasm
Q2YPW9	HFQ_BRUA2		cytoplasm
Q2YPX1	CLPP_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YPX3	LON_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YPX5	UVRA_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YPX7	SSB_BRUA2		cytoplasm
Q2YPY7	QUEA_BRUA2	cytoplasm (potential).	cytoplasm
Q2YQ02	HIS3_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YQD8	URE12_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YQD9	URE22_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YQE0	URE32_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YQI5	PURA_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YQM7	PDXJ_BRUA2	cytoplasm (probable).	cytoplasm

Q2YQN2	ILVC_BRUA2		cytoplasm
Q2YQP6	RS20_BRUA2		cytoplasm
Q2YQP8	FPG_BRUA2		cytoplasm
Q2YQR5	TRUB_BRUA2		cytoplasm
Q2YQS0	Y2162_BRUA2		inner membrane
Q2YQS1	TRMB_BRUA2		cytoplasm
Q2YQY7	HIS2_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YQY8	HIS6_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YQY9	HIS4_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YQZ0	HIS5_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YR40	XERD_BRUA2	cytoplasm (by similarity).	nucleoid
Q2YR42	AROK_BRUA2	cytoplasm (probable).	cytoplasm
Q2YR43	AROB_BRUA2	cytoplasm (probable).	cytoplasm
Q2YR81	HIS8_BRUA2		cytoplasm
Q2YRA1	RS3_BRUA2		cytoplasm
Q2YRA4	RS17_BRUA2		cytoplasm
Q2YRA9	RS8_BRUA2		cytoplasm
Q2YRD2	RUVB_BRUA2		cytoplasm
Q2YRG4	DUT_BRUA2		extracellular
Q2YRK0	FUR_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YRP9	CDSA_BRUA2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q2YRQ3	LPXD_BRUA2		cytoplasm
Q2YRQ4	FABZ_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YRQ5	LPXA_BRUA2	cytoplasm (by similarity).	cytoplasm
Q2YRR5	TRPD_BRUA2		cytoplasm
Q2YRT3	RS5_BRUA2		cytoplasm
Q2YRT8	RS13_BRUA2		cytoplasm
Q2YRT9	RS11_BRUA2		cytoplasm
Q30NR6	HIS6_THIDN	cytoplasm (by similarity).	cytoplasm
Q30NW3	RS20_THIDN		cytoplasm
Q30NX1	CLPP_THIDN	cytoplasm (by similarity).	cytoplasm
Q30P60	GLYA2_THIDN	cytoplasm (by similarity).	cytoplasm
Q30PV8	PPCK_THIDN	cytoplasm (by similarity).	cytoplasm
Q30PX6	RUVB_THIDN		cytoplasm
Q30PY0	HIS5_THIDN	cytoplasm (by similarity).	cytoplasm
Q30PY1	HIS4_THIDN	cytoplasm (by similarity).	cytoplasm
Q30PZ4	RS6_THIDN		cytoplasm
Q30PZ7	QUEA_THIDN	cytoplasm (potential).	cytoplasm
Q30Q39	MRAY_THIDN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q30QB8	SURE_THIDN	cytoplasm (potential).	periplasm
Q30QJ2	FABZ_THIDN	cytoplasm (by similarity).	cytoplasm
Q30QP9	ATPA_THIDN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q30QS9	RL9_THIDN		cytoplasm
Q30R29	GLYA1_THIDN	cytoplasm (by similarity).	cytoplasm
Q30R39	HTPG_THIDN	cytoplasm (by similarity).	cytoplasm
Q30RL8	PPNK_THIDN	cytoplasm (by similarity).	cytoplasm
Q30RS1	TRUD_THIDN		cytoplasm
Q30RT3	AROB_THIDN	cytoplasm (probable).	cytoplasm
Q30S57	PDXJ_THIDN	cytoplasm (probable).	cytoplasm
Q30S64	HIS1_THIDN	cytoplasm (by similarity).	cytoplasm

Q30SG0	PROA_THIDN	cytoplasm (by similarity).	cytoplasm
Q30SG7	TRMB_THIDN		cytoplasm
Q30SL9	FLGH_THIDN	outer membrane; lipid-anchor (probable).	inner membrane
Q30SN2	SYX_THIDN	cytoplasm (by similarity).	cytoplasm
Q30SS4	Y678_THIDN		inner membrane
Q30T40	FLGI_THIDN	periplasm (by similarity).	periplasm
Q30T46	MURA_THIDN	cytoplasm (probable).	inner membrane
Q30T55	PURL_THIDN	cytoplasm (by similarity).	cytoplasm
Q30TA4	HISX_THIDN		cytoplasm
Q30TC9	HIS8_THIDN		cytoplasm
Q30TG0	ISPE_THIDN		cytoplasm
Q30TG2	TRUB_THIDN		cytoplasm
Q30TH8	GUAA_THIDN		cytoplasm
Q30TP5	RS12_THIDN		cytoplasm
Q30TP7	RPOB_THIDN		cytoplasm
Q30TP9	RL10_THIDN		cytoplasm
Q30TQ0	RL1_THIDN		cytoplasm
Q30TR6	PAND_THIDN		cytoplasm
Q30TS4	RS4_THIDN		cytoplasm
Q30TS5	RS11_THIDN		cytoplasm
Q30TS6	RS13_THIDN		cytoplasm
Q30TV0	RS8_THIDN		cytoplasm
Q30TV5	RS17_THIDN		cytoplasm
Q30TV8	RS3_THIDN		cytoplasm
Q30TW1	RL2_THIDN		cytoplasm
Q30TW5	RS10_THIDN		cytoplasm
Q30U33	THIG_THIDN	cytoplasm (by similarity).	cytoplasm
Q30UH8	UPPP_THIDN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q30VD6	HIS3_DESDG	cytoplasm (by similarity).	cytoplasm
Q30WI7	TRUB_DESDG		cytoplasm
Q30WJ2	Y3160_DESDG		inner membrane
Q30WJ6	FLGH_DESDG	outer membrane; lipid-anchor (probable).	inner membrane
Q30WJ7	FLGI_DESDG	periplasm (by similarity).	periplasm
Q30WZ8	GPMA_DESDG		cytoplasm
Q30X04	RPOC_DESDG		cytoplasm
Q30X05	RPOB_DESDG		cytoplasm
Q30X08	RL10_DESDG		cytoplasm
Q30X09	RL1_DESDG		cytoplasm
Q30XI4	LEPA_DESDG		cytoplasm
Q30XS2	AROK_DESDG	cytoplasm (probable).	cytoplasm
Q30XT0	Y2720_DESDG		inner membrane
Q30XW1	PROB_DESDG	cytoplasm (by similarity).	cytoplasm
Q30Y32	PPNK_DESDG	cytoplasm (by similarity).	cytoplasm
Q30YL7	GLYA_DESDG	cytoplasm (by similarity).	cytoplasm
Q30YV9	SURE_DESDG	cytoplasm (potential).	periplasm
Q30YX5	RUVC_DESDG		cytoplasm
Q30YX6	RUVA_DESDG		cytoplasm
Q30YX7	RUVB_DESDG		cytoplasm
Q30Z13	GLGA_DESDG		cytoplasm
Q30Z36	RS12_DESDG		cytoplasm
Q30Z37	RS7_DESDG		cytoplasm
Q30Z41	RS10_DESDG		cytoplasm

Q30Z45	RL2_DESDG		cytoplasm
Q30Z48	RS3_DESDG		cytoplasm
Q30Z51	RS17_DESDG		cytoplasm
Q30Z56	RS8_DESDG		cytoplasm
Q30Z59	RS5_DESDG		cytoplasm
Q30Z65	RS13_DESDG		cytoplasm
Q30Z66	RS11_DESDG		cytoplasm
Q30Z67	RS4_DESDG		cytoplasm
Q30Z68	RPOA_DESDG		cytoplasm
Q30Z79	CLPP_DESDG	cytoplasm (by similarity).	cytoplasm
Q30ZD3	ILVC_DESDG		cytoplasm
Q30ZH4	ISPE_DESDG		cytoplasm
Q30ZK3	UVRB_DESDG	cytoplasm (by similarity).	cytoplasm
Q30ZK7	DAPB_DESDG	cytoplasm (by similarity).	cytoplasm
Q30ZQ5	ACPS_DESDG	cytoplasm (by similarity).	cytoplasm
Q30ZQ6	PDXJ_DESDG	cytoplasm (probable).	cytoplasm
Q30ZR8	RS20_DESDG		cytoplasm
Q30ZX3	MUTS_DESDG		cytoplasm
Q310A3	UPPP_DESDG	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q310Z1	RNC_DESDG	cytoplasm (by similarity).	cytoplasm
Q311G6	PROA_DESDG	cytoplasm (by similarity).	cytoplasm
Q311P4	THIG_DESDG	cytoplasm (by similarity).	cytoplasm
Q311X7	GUAA_DESDG		cytoplasm
Q311Y2	HIS4_DESDG	cytoplasm (by similarity).	cytoplasm
Q311Z4	HIS8_DESDG		cytoplasm
Q312A8	NDK_DESDG	cytoplasm (by similarity).	cytoplasm
Q312D5	RS6_DESDG		cytoplasm
Q312D7	RL9_DESDG		cytoplasm
Q312H2	FABZ_DESDG	cytoplasm (by similarity).	cytoplasm
Q312I9	SLYX_DESDG		inner membrane
Q313E8	HTPG_DESDG	cytoplasm (by similarity).	cytoplasm
Q313K0	RL19_DESDG		cytoplasm
Q313K1	RNH2_DESDG	cytoplasm (potential).	extracellular
Q313Q1	MURB_DESDG	cytoplasm (probable).	inner membrane
Q313Q3	MURG_DESDG	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q313Q6	MRAY_DESDG	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q313R1	MRAW_DESDG		cytoplasm
Q313R2	MRAZ_DESDG		inner membrane
Q313S2	DNAK_DESDG		cytoplasm
Q313T1	UVRC_DESDG	cytoplasm (by similarity).	cytoplasm
Q313T7	HISX_DESDG		cytoplasm
Q313V8	ATPA_DESDG	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q313W4	GLMU_DESDG	cytoplasm (by similarity).	cytoplasm
Q315P9	MURA_DESDG	cytoplasm (probable).	inner membrane
Q316L3	HIS5_DESDG	cytoplasm (by similarity).	cytoplasm
Q316L4	HIS6_DESDG	cytoplasm (by similarity).	cytoplasm
Q316U8	RPOZ_DESDG		cytoplasm
Q316W7	SYX_DESDG	cytoplasm (by similarity).	cytoplasm

Q317B9	PURA_DESDG	cytoplasm (by similarity).	cytoplasm
Q317H9	ILVD_DESDG		cytoplasm
Q317N3	QUEA_DESDG	cytoplasm (potential).	cytoplasm
Q31DL8	ATPA_THICR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q31DP2	RPOZ_THICR		cytoplasm
Q31DP8	AROK_THICR	cytoplasm (probable).	cytoplasm
Q31DP9	AROB_THICR	cytoplasm (probable).	cytoplasm
Q31E63	HIS2_THICR	cytoplasm (by similarity).	cytoplasm
Q31E64	HIS6_THICR	cytoplasm (by similarity).	cytoplasm
Q31E65	HIS4_THICR	cytoplasm (by similarity).	cytoplasm
Q31E66	HIS5_THICR	cytoplasm (by similarity).	cytoplasm
Q31E84	SYI_THICR	cytoplasm (by similarity).	cytoplasm
Q31EC7	DUT_THICR		extracellular
Q31EJ2	METX_THICR	cytoplasm (by similarity).	cytoplasm
Q31EW3	IDI2_THICR	cytoplasm (by similarity).	cytoplasm
Q31F66	GUAA_THICR		cytoplasm
Q31F72	HTPG_THICR	cytoplasm (by similarity).	cytoplasm
Q31FF9	PAND_THICR		cytoplasm
Q31FL3	FLGH_THICR	outer membrane; lipid-anchor (probable).	inner membrane
Q31FL4	FLGI_THICR	periplasm (by similarity).	periplasm
Q31FS6	GLYA_THICR	cytoplasm (by similarity).	cytoplasm
Q31FS7	NRDR_THICR		cytoplasm
Q31FW4	RS6_THICR		cytoplasm
Q31FW7	RL9_THICR		cytoplasm
Q31FZ4	QUEA_THICR	cytoplasm (potential).	cytoplasm
Q31G32	NAGZ_THICR	cytoplasm (by similarity).	extracellular
Q31G41	GLND_THICR		cytoplasm
Q31G53	FABZ_THICR	cytoplasm (by similarity).	cytoplasm
Q31G56	RNH2_THICR	cytoplasm (potential).	extracellular
Q31G71	SURE_THICR	cytoplasm (potential).	periplasm
Q31GD4	HIS82_THICR		cytoplasm
Q31GF1	CLPP_THICR	cytoplasm (by similarity).	cytoplasm
Q31GK3	TRUB_THICR		cytoplasm
Q31GK7	Y1121_THICR		inner membrane
Q31GX8	NUDH_THICR		cytoplasm
Q31GY9	HISX_THICR		cytoplasm
Q31GZ0	HIS1_THICR	cytoplasm (by similarity).	cytoplasm
Q31GZ1	MURA_THICR	cytoplasm (probable).	inner membrane
Q31H20	LPXK_THICR		cytoplasm
Q31H83	RUVB_THICR		cytoplasm
Q31HB0	PPNK_THICR	cytoplasm (by similarity).	cytoplasm
Q31HP0	PDXJ_THICR	cytoplasm (probable).	cytoplasm
Q31HZ6	PHS_THICR		cytoplasm
Q31I19	G6PI_THICR	cytoplasm (by similarity).	cytoplasm
Q31I36	HIS81_THICR		cytoplasm
Q31I62	MRAY_THICR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q31I69	MRAZ_THICR		inner membrane
Q31IB5	GLGA_THICR		cytoplasm
Q31ID5	RS20_THICR		cytoplasm
Q31IE5	PROA_THICR	cytoplasm (by similarity).	cytoplasm
Q31IT3	PROB_THICR	cytoplasm (by similarity).	cytoplasm

Q31IV9	RS4_THICR		cytoplasm
Q31IW0	RS11_THICR		cytoplasm
Q31IW1	RS13_THICR		cytoplasm
Q31IW5	RS5_THICR		cytoplasm
Q31IW8	RS8_THICR		cytoplasm
Q31IX3	RS17_THICR		cytoplasm
Q31IX6	RS3_THICR		cytoplasm
Q31IX9	RL2_THICR		cytoplasm
Q31IY3	RS10_THICR		cytoplasm
Q31IY7	RS12_THICR		cytoplasm
Q31IY9	RPOB_THICR		cytoplasm
Q31IZ1	RL10_THICR		cytoplasm
Q31IZ2	RL1_THICR		cytoplasm
Q31J60	HIS3_THICR	cytoplasm (by similarity).	cytoplasm
Q31J61	TRUD_THICR		cytoplasm
Q31JD2	THIG_THICR	cytoplasm (by similarity).	cytoplasm
Q31JF8	TRMB_THICR		cytoplasm
Q31SV5	DEOD_SHIBS		cytoplasm
Q31SV7	TYPH_SHIBS		cytoplasm
Q31SX4	DNAT_SHIBS		nucleoid
Q31T20	UXUA_SHIBS		cytoplasm
Q31TA7	PURA_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31TC2	ULAR_SHIBS	cytoplasm (potential).	cytoplasm
Q31TC3	ULAG_SHIBS	cytoplasm (potential).	extracellular
Q31TC4	ULAA_SHIBS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q31TC5	ULAB_SHIBS	cytoplasm (potential).	cytoplasm
Q31TC6	ULAC_SHIBS	cytoplasm (probable).	cytoplasm
Q31TC7	ULAD_SHIBS		cytoplasm
Q31TC8	ULAE_SHIBS		cytoplasm
Q31TD1	RS6_SHIBS		cytoplasm
Q31TD4	RL9_SHIBS		cytoplasm
Q31TK0	SYV_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31TR2	YJDP_SHIBS		nucleoid
Q31TX1	G6PI_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31U01	NUDC_SHIBS		cytoplasm
Q31U06	THIG_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31U09	RPOC_SHIBS		cytoplasm
Q31U10	RPOB_SHIBS		cytoplasm
Q31U12	RL10_SHIBS		cytoplasm
Q31U13	RL1_SHIBS		cytoplasm
Q31U21	MURB_SHIBS	cytoplasm (probable).	inner membrane
Q31UH1	CORA_SHIBS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q31UL0	ILVC_SHIBS		cytoplasm
Q31UL3	ILVD_SHIBS		cytoplasm
Q31UN0	GLMU_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31UN4	ATPA_SHIBS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q31UP5	RAVA_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31UQ7	RPOZ_SHIBS		cytoplasm

Q31UU4	YIDE_SHIBS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q31UV3	RECF_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31UY6	DUT_SHIBS		extracellular
Q31UZ1	FPG_SHIBS		cytoplasm
Q31V09	GPML_SHIBS		cytoplasm
Q31V17	LLDD_SHIBS		cytoplasm
Q31V53	XYLA_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31VF6	TUSA_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31VJ4	GLGA_SHIBS		cytoplasm
Q31VN1	PPCK_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31VN3	HSLO_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31VP4	AROK_SHIBS	cytoplasm (probable).	cytoplasm
Q31VP5	AROB_SHIBS	cytoplasm (probable).	cytoplasm
Q31VP9	GPH_SHIBS		cytoplasm
Q31VU2	SLYX_SHIBS		inner membrane
Q31VU5	TUSD_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31VU6	TUSC_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31VU7	RS12_SHIBS		cytoplasm
Q31VU8	RS7_SHIBS		cytoplasm
Q31VV5	RS10_SHIBS		cytoplasm
Q31VV9	RL2_SHIBS		cytoplasm
Q31VW2	RS3_SHIBS		cytoplasm
Q31VW3	RL16_SHIBS		cytoplasm
Q31VW5	RS17_SHIBS		cytoplasm
Q31VX0	RS8_SHIBS		cytoplasm
Q31VX3	RS5_SHIBS		cytoplasm
Q31VX8	RS13_SHIBS		cytoplasm
Q31VX9	RS11_SHIBS		cytoplasm
Q31VY0	RS4_SHIBS		cytoplasm
Q31VY1	RPOA_SHIBS		cytoplasm
Q31VY6	MSCL_SHIBS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q31W44	RS15_SHIBS		cytoplasm
Q31W45	TRUB_SHIBS		cytoplasm
Q31W49	YHBC_SHIBS		inner membrane
Q31W66	MURA_SHIBS	cytoplasm (probable).	inner membrane
Q31WF1	IDI_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31WM2	TRMB_SHIBS		cytoplasm
Q31WW5	MUG_SHIBS	cytoplasm (potential).	extracellular
Q31WX6	YGIH_SHIBS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q31WX8	UPPP_SHIBS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q31X95	MUTS_SHIBS		cytoplasm
Q31XA6	SURE_SHIBS	cytoplasm (potential).	periplasm
Q31XA7	TRUD_SHIBS		cytoplasm
Q31XA8	ISPF_SHIBS		extracellular
Q31XA9	ISPD_SHIBS		cytoplasm
Q31XD1	PPNK_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31XD9	RL19_SHIBS		cytoplasm
Q31XF7	NUDH_SHIBS		cytoplasm

Q31XK5	RUMA_SHIBS		cytoplasm
Q31XR8	LEPA_SHIBS		cytoplasm
Q31XS0	RNC_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31XS2	RECO_SHIBS		cytoplasm
Q31XS3	PDXJ_SHIBS	cytoplasm (probable).	cytoplasm
Q31XS4	ACPS_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31XT6	GLYA_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31XW7	PEPB_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31XX2	NDK_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31XY3	GUAA_SHIBS		cytoplasm
Q31Y67	ZIPA_SHIBS	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q31YN2	TUSE_SHIBS	cytoplasm (by similarity).	cytoplasm
Q31YT5	LPXK_SHIBS		cytoplasm
Q31YW9	RIHB_SHIBS		extracellular
Q31Z20	CCME_SHIBS	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q31Z29	NAPA_SHIBS	periplasm (by similarity).	periplasm
Q31Z32	ECOT_SHIBS	periplasm (by similarity).	periplasm
Q31Z90	YCDX_SHIBS		cytoplasm
Q31ZB2	YCEI_SHIBS	periplasm (potential).	periplasm
Q31ZD4	FLGH_SHIBS	outer membrane; lipid-anchor (probable).	inner membrane
Q31ZD5	FLGI_SHIBS	periplasm (by similarity).	periplasm
Q31ZE4	RL32_SHIBS		cytoplasm
Q31ZG4	NAGZ_SHIBS	cytoplasm (by similarity).	extracellular
Q31ZQ1	ISPE_SHIBS		cytoplasm
Q320D7	ZNTB_SHIBS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q320T9	Y1556_SHIBS	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q320W6	TUS_SHIBS	cytoplasm (by similarity).	cytoplasm
Q320Z4	SYI_SHIBS	cytoplasm (by similarity).	cytoplasm
Q320Z5	PDXH_SHIBS		cytoplasm
Q321G0	YDIA_SHIBS		cytoplasm
Q321V7	YOA_H_SHIBS		inner membrane
Q322D7	RUVC_SHIBS		cytoplasm
Q322E5	RUVA_SHIBS		cytoplasm
Q322E6	RUVB_SHIBS		cytoplasm
Q322G8	YEBF_SHIBS	secreted protein (by similarity).	inner membrane
Q322L9	PGSA_SHIBS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q322V3	CDD_SHIBS		cytoplasm
Q323I7	HIS6_SHIBS	cytoplasm (by similarity).	cytoplasm
Q323I8	HIS4_SHIBS	cytoplasm (by similarity).	cytoplasm
Q323I9	HIS5_SHIBS	cytoplasm (by similarity).	cytoplasm
Q323J1	HIS8_SHIBS		cytoplasm
Q323J2	HISX_SHIBS		cytoplasm
Q323N6	YBJQ_SHIBS		inner membrane

Q323Q4	YBJL_SHIBS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q324B3	UVRB_SHIBS	cytoplasm (by similarity).	cytoplasm
Q324G4	GPMA_SHIBS		cytoplasm
Q324V2	ENTS_SHIBS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q325C3	HTPG_SHIBS	cytoplasm (by similarity).	cytoplasm
Q325G4	CLPP_SHIBS	cytoplasm (by similarity).	cytoplasm
Q325I8	NRDR_SHIBS		cytoplasm
Q325J6	QUEA_SHIBS	cytoplasm (potential).	cytoplasm
Q325J7	ACPH_SHIBS		cytoplasm
Q325L0	AROL_SHIBS	cytoplasm (probable).	cytoplasm
Q325P3	PROA_SHIBS	cytoplasm (by similarity).	cytoplasm
Q325P4	PROB_SHIBS	cytoplasm (by similarity).	cytoplasm
Q325V5	ACCA_SHIBS	cytoplasm (by similarity).	cytoplasm
Q325V7	RNH2_SHIBS	cytoplasm (potential).	extracellular
Q325W0	FABZ_SHIBS	cytoplasm (by similarity).	cytoplasm
Q325W2	SKP_SHIBS	periplasm (by similarity).	periplasm
Q325X3	GLND_SHIBS		cytoplasm
Q326A6	PAND_SHIBS		cytoplasm
Q326E5	MURG_SHIBS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q326E8	MRAY_SHIBS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q326F3	MRAW_SHIBS	cytoplasm (by similarity).	cytoplasm
Q326F4	MRAZ_SHIBS		inner membrane
Q326G2	LEU3_SHIBS	cytoplasm (by similarity).	cytoplasm
Q326G3	LEU2_SHIBS		cytoplasm
Q326G4	LEUD_SHIBS		cytoplasm
Q326J2	DAPB_SHIBS	cytoplasm (by similarity).	cytoplasm
Q326K0	RS20_SHIBS		cytoplasm
Q326K7	DNAK_SHIBS		cytoplasm
Q326L3	TAL_SHIBS	cytoplasm (probable).	cytoplasm
Q327L2	DEOD_SHIDS		cytoplasm
Q327L4	TYPH_SHIDS		cytoplasm
Q327N4	DNAT_SHIDS		nucleoid
Q328F6	PURA_SHIDS	cytoplasm (by similarity).	cytoplasm
Q328J6	RL9_SHIDS		cytoplasm
Q328J9	RS6_SHIDS		cytoplasm
Q328K0	ULAF_SHIDS		cytoplasm
Q328K1	ULAD_SHIDS		cytoplasm
Q328K2	ULAC_SHIDS	cytoplasm (probable).	cytoplasm
Q328K3	ULAB_SHIDS	cytoplasm (potential).	cytoplasm
Q328K4	ULAA_SHIDS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q328K5	ULAG_SHIDS	cytoplasm (potential).	extracellular
Q328K6	ULAR_SHIDS	cytoplasm (potential).	cytoplasm
Q328S3	SYV_SHIDS	cytoplasm (by similarity).	cytoplasm
Q328X7	G6PI_SHIDS	cytoplasm (by similarity).	cytoplasm
Q329B8	RECF_SHIDS	cytoplasm (by similarity).	cytoplasm

Q329C8	YIDE_SHIDS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q329H0	YJDP_SHIDS		cytoplasm
Q329K7	RPOZ_SHIDS		cytoplasm
Q329L7	DUT_SHIDS		extracellular
Q329M2	FPG_SHIDS		cytoplasm
Q329P2	GPMI_SHIDS		cytoplasm
Q329P9	LLDD_SHIDS		cytoplasm
Q329R9	GLMU_SHIDS	cytoplasm (by similarity).	cytoplasm
Q329S3	ATPA_SHIDS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q329T4	VIAA_SHIDS		nucleoid
Q329T5	RAVA_SHIDS	cytoplasm (by similarity).	cytoplasm
Q329V0	ILVD_SHIDS		cytoplasm
Q329V3	ILVC_SHIDS		cytoplasm
Q329Z2	CORA_SHIDS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q32AE8	MURB_SHIDS	cytoplasm (probable).	inner membrane
Q32AF6	RL1_SHIDS		cytoplasm
Q32AF7	RL10_SHIDS		cytoplasm
Q32AF9	RPOB_SHIDS		cytoplasm
Q32AG0	RPOC_SHIDS		cytoplasm
Q32AG3	THIG_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32AG9	NUDC_SHIDS		cytoplasm
Q32AJ7	GPH_SHIDS		cytoplasm
Q32AK1	AROB_SHIDS	cytoplasm (probable).	cytoplasm
Q32AK2	AROK_SHIDS	cytoplasm (probable).	cytoplasm
Q32AL3	HSLO_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32AL7	PPCK_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32AR4	TUSA_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32AV6	GLGA_SHIDS		cytoplasm
Q32B18	SLYX_SHIDS		inner membrane
Q32B21	TUSD_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32B22	TUSC_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32B23	TUSB_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32B24	RS12_SHIDS		cytoplasm
Q32B25	RS7_SHIDS		cytoplasm
Q32B30	RS10_SHIDS		cytoplasm
Q32B34	RL2_SHIDS		cytoplasm
Q32B37	RS3_SHIDS		cytoplasm
Q32B38	RL16_SHIDS		cytoplasm
Q32B40	RS17_SHIDS		cytoplasm
Q32B45	RS8_SHIDS		cytoplasm
Q32B48	RS5_SHIDS		cytoplasm
Q32B53	RS13_SHIDS		cytoplasm
Q32B54	RS11_SHIDS		cytoplasm
Q32B55	RS4_SHIDS		cytoplasm
Q32B56	RPOA_SHIDS		cytoplasm
Q32BE4	MURA_SHIDS	cytoplasm (probable).	inner membrane
Q32BG3	YHBC_SHIDS		inner membrane
Q32BG7	TRUB_SHIDS		cytoplasm
Q32BG8	RS15_SHIDS		cytoplasm
Q32BP9	MUG_SHIDS	cytoplasm (potential).	extracellular

Q32BQ6	YGIH_SHIDS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q32BV2	IDI_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32C29	TRMB_SHIDS		cytoplasm
Q32C91	NUDH_SHIDS		cytoplasm
Q32CD0	RUMA_SHIDS		cytoplasm
Q32CI3	ISPD_SHIDS		cytoplasm
Q32CI4	ISPF_SHIDS		extracellular
Q32CI5	TRUD_SHIDS		cytoplasm
Q32CI6	SURE_SHIDS	cytoplasm (potential).	periplasm
Q32CJ6	MUTS_SHIDS		cytoplasm
Q32CV2	LEPA_SHIDS		cytoplasm
Q32CV4	RNC_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32CV6	RECO_SHIDS		cytoplasm
Q32CV7	PDXJ_SHIDS	cytoplasm (probable).	cytoplasm
Q32CV8	ACPS_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32CX4	PPNK_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32CY2	RL19_SHIDS		cytoplasm
Q32D21	GLYA_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32D41	PEPB_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32D44	NDK_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32D55	GUAA_SHIDS		cytoplasm
Q32DE1	ZIPA_SHIDS	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q32DY4	YBJQ_SHIDS		inner membrane
Q32E35	LPXK_SHIDS		cytoplasm
Q32E67	YCEI_SHIDS	periplasm (potential).	periplasm
Q32EE9	HISX_SHIDS		cytoplasm
Q32EF0	HIS8_SHIDS		cytoplasm
Q32EF2	HIS5_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32EF3	HIS4_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32EF4	HIS6_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32EM3	CDD_SHIDS		cytoplasm
Q32ER0	RIHB_SHIDS		extracellular
Q32ET9	FLGH_SHIDS	outer membrane; lipid-anchor (probable).	inner membrane
Q32EU0	FLGI_SHIDS	periplasm (by similarity).	periplasm
Q32EU9	RL32_SHIDS		cytoplasm
Q32EW7	NAGZ_SHIDS	cytoplasm (by similarity).	extracellular
Q32FD5	PDXH_SHIDS		cytoplasm
Q32FD6	SYX_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32FG8	TUS_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32FJ7	YDIA_SHIDS		cytoplasm
Q32FR8	YOAH_SHIDS		inner membrane
Q32FX5	Y1659_SHIDS	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q32GI7	ZNTB_SHIDS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q32GZ9	ISPE_SHIDS		cytoplasm
Q32H98	RUVC_SHIDS		cytoplasm
Q32HA0	RUVA_SHIDS		cytoplasm
Q32HA1	RUVB_SHIDS		cytoplasm

Q32HB5	YEBF_SHIDS	secreted protein (by similarity).	inner membrane
Q32HD9	PGSA_SHIDS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q32HN6	YCDX_SHIDS		cytoplasm
Q32HT7	TUSE_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32HZ7	CCME_SHIDS	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q32I06	NAPA_SHIDS	periplasm (by similarity).	periplasm
Q32I48	UVRB_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32IC5	YBJL_SHIDS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q32IH0	GPMA_SHIDS		cytoplasm
Q32IX8	ENTS_SHIDS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q32J26	PROB_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32J27	PROA_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32J53	HTPG_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32JD7	AROL_SHIDS	cytoplasm (probable).	cytoplasm
Q32JF8	ACPH_SHIDS		cytoplasm
Q32JF9	QUEA_SHIDS	cytoplasm (potential).	cytoplasm
Q32JG7	NRDR_SHIDS		cytoplasm
Q32JJ3	CLPP_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32JS4	ACCA_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32JS6	RNH2_SHIDS	cytoplasm (potential).	extracellular
Q32JS9	FABZ_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32JT1	SKP_SHIDS	periplasm (by similarity).	periplasm
Q32JU2	GLND_SHIDS		cytoplasm
Q32K02	MURG_SHIDS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q32K05	MRAY_SHIDS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q32K10	MRAW_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32K11	MRAZ_SHIDS		inner membrane
Q32K21	LEU3_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32K22	LEU2_SHIDS		cytoplasm
Q32K23	LEUD_SHIDS		cytoplasm
Q32K66	DAPB_SHIDS	cytoplasm (by similarity).	cytoplasm
Q32K74	RS20_SHIDS		cytoplasm
Q32K78	PAND_SHIDS		cytoplasm
Q32KA5	DNAK_SHIDS		cytoplasm
Q32KB0	TAL_SHIDS	cytoplasm (probable).	cytoplasm
Q390V3	SURE1_BURS3	cytoplasm (potential).	periplasm
Q391K1	GLYA2_BURS3	cytoplasm (by similarity).	cytoplasm
Q394V3	ILVD1_BURS3		cytoplasm
Q395F2	KPTA_BURS3		cytoplasm
Q396Y9	UXUA_BURS3		cytoplasm
Q39A02	Y595_BURS3		inner membrane
Q39A26	GLYA1_BURS3	cytoplasm (by similarity).	cytoplasm

Q39BX2	METX_BURS3	cytoplasm (by similarity).	cytoplasm
Q39C57	FLGH_BURS3	outer membrane; lipid-anchor (probable).	inner membrane
Q39C92	GLMU_BURS3	cytoplasm (by similarity).	cytoplasm
Q39CN6	GPMA_BURS3		cytoplasm
Q39CT7	HIS82_BURS3		cytoplasm
Q39CU1	ISPE_BURS3		cytoplasm
Q39CU4	FPG_BURS3		cytoplasm
Q39DI6	MURB_BURS3	cytoplasm (probable).	inner membrane
Q39DI9	RS20_BURS3		cytoplasm
Q39DJ6	LPXK_BURS3		cytoplasm
Q39DM5	DUT_BURS3		extracellular
Q39DS9	ILVD2_BURS3		cytoplasm
Q39DU8	PAND_BURS3		cytoplasm
Q39E36	HTPG_BURS3	cytoplasm (by similarity).	cytoplasm
Q39E45	TAL_BURS3	cytoplasm (probable).	cytoplasm
Q39ED2	ILVC_BURS3		cytoplasm
Q39EX8	MUTS_BURS3		cytoplasm
Q39F41	GLND_BURS3		cytoplasm
Q39F54	FABZ_BURS3	cytoplasm (by similarity).	cytoplasm
Q39F57	RNH2_BURS3	cytoplasm (potential).	extracellular
Q39F73	GUAA_BURS3		cytoplasm
Q39FB0	MSCL_BURS3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q39FB8	ISPD_BURS3		cytoplasm
Q39FB9	ISPF_BURS3		extracellular
Q39FE8	CLPP_BURS3	cytoplasm (by similarity).	cytoplasm
Q39FF4	G6PI_BURS3	cytoplasm (by similarity).	cytoplasm
Q39FJ8	RS6_BURS3		cytoplasm
Q39FK1	RL9_BURS3		cytoplasm
Q39FP8	SURE2_BURS3	cytoplasm (potential).	periplasm
Q39FQ6	NDK_BURS3	cytoplasm (by similarity).	cytoplasm
Q39FS0	PURA1_BURS3	cytoplasm (by similarity).	cytoplasm
Q39H28	TRUB_BURS3		cytoplasm
Q39H32	Y4639_BURS3		inner membrane
Q39HK8	UPPP2_BURS3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q39I64	UVRC_BURS3	cytoplasm (by similarity).	cytoplasm
Q39I69	ACPS_BURS3	cytoplasm (by similarity).	cytoplasm
Q39I70	PDXJ_BURS3	cytoplasm (probable).	cytoplasm
Q39I75	LEPA_BURS3		cytoplasm
Q39I88	RL32_BURS3		cytoplasm
Q39ID3	RL19_BURS3		cytoplasm
Q39IL1	RPOZ_BURS3		cytoplasm
Q39IU7	TRMB_BURS3		cytoplasm
Q39IU8	UPPP1_BURS3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q39IW1	URE3_BURS3	cytoplasm (by similarity).	cytoplasm
Q39IW2	URE2_BURS3	cytoplasm (by similarity).	cytoplasm
Q39IW3	URE1_BURS3	cytoplasm (by similarity).	cytoplasm
Q39J54	HGD_BURS3		cytoplasm
Q39J71	NRDR_BURS3		cytoplasm
Q39J72	GLYA3_BURS3	cytoplasm (by similarity).	cytoplasm

Q39JC8	DNAK_BURS3		cytoplasm
Q39JD3	PPNK_BURS3	cytoplasm (by similarity).	cytoplasm
Q39JF4	QUEA_BURS3	cytoplasm (potential).	cytoplasm
Q39JI9	RUVC_BURS3		cytoplasm
Q39JJ0	RUVA_BURS3		cytoplasm
Q39JJ1	RUVB_BURS3		cytoplasm
Q39JJ6	SYU_BURS3	cytoplasm (by similarity).	cytoplasm
Q39JM2	PROA_BURS3	cytoplasm (by similarity).	cytoplasm
Q39JM8	DAPB_BURS3	cytoplasm (by similarity).	cytoplasm
Q39JU4	NUDH_BURS3		cytoplasm
Q39JU6	PROB_BURS3	cytoplasm (by similarity).	cytoplasm
Q39JW0	ARGJ_BURS3	cytoplasm (probable).	cytoplasm
Q39JX0	MURG_BURS3	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q39JX3	MRAY_BURS3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q39JX8	MRAW_BURS3		cytoplasm
Q39JX9	MRAZ_BURS3		inner membrane
Q39JZ6	TRPD_BURS3		cytoplasm
Q39K83	HIS2_BURS3	cytoplasm (by similarity).	cytoplasm
Q39K84	HIS3_BURS3	cytoplasm (by similarity).	cytoplasm
Q39K85	HIS6_BURS3	cytoplasm (by similarity).	cytoplasm
Q39K86	HIS4_BURS3	cytoplasm (by similarity).	cytoplasm
Q39K87	HIS5_BURS3	cytoplasm (by similarity).	cytoplasm
Q39K90	HIS81_BURS3		cytoplasm
Q39K91	HISX_BURS3		cytoplasm
Q39K92	HIS1_BURS3	cytoplasm (by similarity).	cytoplasm
Q39K93	MURA_BURS3	cytoplasm (probable).	inner membrane
Q39KA4	THIG_BURS3	cytoplasm (by similarity).	cytoplasm
Q39KC4	AROB_BURS3	cytoplasm (probable).	cytoplasm
Q39KC5	AROK_BURS3	cytoplasm (probable).	cytoplasm
Q39KE1	RPOA_BURS3		cytoplasm
Q39KE2	RS4_BURS3		cytoplasm
Q39KE3	RS11_BURS3		cytoplasm
Q39KE4	RS13_BURS3		cytoplasm
Q39KF0	RS5_BURS3		cytoplasm
Q39KF3	RS8_BURS3		cytoplasm
Q39KF8	RS17_BURS3		cytoplasm
Q39KG1	RS3_BURS3		cytoplasm
Q39KG4	RL2_BURS3		cytoplasm
Q39KG8	RS10_BURS3		cytoplasm
Q39KH1	RS7_BURS3		cytoplasm
Q39KH2	RS12_BURS3		cytoplasm
Q39KH4	RPOC_BURS3		cytoplasm
Q39KH5	RPOB_BURS3		cytoplasm
Q39KH7	RL10_BURS3		cytoplasm
Q39KH8	RL1_BURS3		cytoplasm
Q39KU1	GCSP_BURS3		cytoplasm
Q39KX8	ATPA_BURS3	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q39L06	PHS_BURS3		cytoplasm
Q39M27	HIS83_BURS3		cytoplasm
Q39NL8	DEOC_BURS3	cytoplasm (by similarity).	cytoplasm

Q39PN9	PURA2_BURS3	cytoplasm (by similarity).	cytoplasm
Q39Q54	ATPA_GEOMG	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q39QR2	PROA_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39QR3	PROB_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39QT6	GLGA2_GEOMG		cytoplasm
Q39QU2	PPCK_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39QX7	UPPP_GEOMG	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q39R75	TRMB_GEOMG		cytoplasm
Q39R76	TRUD_GEOMG		cytoplasm
Q39RH2	THIG_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39RP0	RL19_GEOMG		cytoplasm
Q39RP1	RNH2_GEOMG	cytoplasm (potential).	extracellular
Q39RQ7	ISPE_GEOMG		cytoplasm
Q39RR2	RS6_GEOMG		cytoplasm
Q39RR5	RL9_GEOMG		cytoplasm
Q39RX3	METX_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39S70	NDK_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39SN1	MSCL_GEOMG	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q39SQ3	HTPG_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39SQ6	TRPD_GEOMG		cytoplasm
Q39SU1	G6PI_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39T45	FABZ_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39T52	LPXK_GEOMG		cytoplasm
Q39T73	RPOZ_GEOMG		cytoplasm
Q39T82	RNC_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39TA2	RS20_GEOMG		cytoplasm
Q39TA7	GUAA_GEOMG		cytoplasm
Q39UE8	HIS3_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39UG0	PDXJ_GEOMG	cytoplasm (probable).	cytoplasm
Q39UG1	ACPS_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39UH2	CLPP_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39V40	GPMA_GEOMG		cytoplasm
Q39V70	NRDR_GEOMG		cytoplasm
Q39V87	GLYA_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39V99	DUT_GEOMG		extracellular
Q39VA3	TRUB_GEOMG		cytoplasm
Q39VA9	Y1583_GEOMG		inner membrane
Q39VS1	SURE_GEOMG	cytoplasm (potential).	periplasm
Q39WD1	FPG_GEOMG		cytoplasm
Q39WF4	SYX_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39X04	AROB_GEOMG	cytoplasm (probable).	cytoplasm
Q39X05	AROK_GEOMG	cytoplasm (probable).	cytoplasm
Q39X41	PPNK_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39XD1	QUEA_GEOMG	cytoplasm (potential).	cytoplasm
Q39XN6	RUVB_GEOMG		cytoplasm
Q39XS5	GLGA1_GEOMG		cytoplasm
Q39XY0	RS4_GEOMG		cytoplasm
Q39XY1	RS11_GEOMG		cytoplasm
Q39XY2	RS13_GEOMG		cytoplasm
Q39XY9	RS5_GEOMG		cytoplasm

Q39XZ2	RS8_GEOMG		cytoplasm
Q39XZ7	RS17_GEOMG		cytoplasm
Q39Y00	RS3_GEOMG		cytoplasm
Q39Y03	RL2_GEOMG		cytoplasm
Q39Y07	RS10_GEOMG		cytoplasm
Q39Y10	RS7_GEOMG		cytoplasm
Q39Y11	RS12_GEOMG		cytoplasm
Q39Y13	RPOB_GEOMG		cytoplasm
Q39Y15	RL10_GEOMG		cytoplasm
Q39Y16	RL1_GEOMG		cytoplasm
Q39YJ7	FLGI_GEOMG	periplasm (by similarity).	periplasm
Q39YJ8	FLGH_GEOMG	outer membrane; lipid-anchor (probable).	inner membrane
Q39YM2	MRAY_GEOMG	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q39YM8	MRAZ_GEOMG		inner membrane
Q39YP1	HIS2_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39YP2	HIS6_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39YP3	HIS4_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39YP4	HIS5_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39YP6	HIS8_GEOMG		cytoplasm
Q39YP7	HISX_GEOMG		cytoplasm
Q39YP8	HIS1_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39YP9	MURA_GEOMG	cytoplasm (probable).	inner membrane
Q39Z66	DAPB_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39ZL5	ISPD_GEOMG		cytoplasm
Q39ZL6	ISPF_GEOMG		extracellular
Q39ZN3	HSLO_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39ZS1	RECF_GEOMG	cytoplasm (by similarity).	cytoplasm
Q39ZT9	ATPA1_PELCD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3A076	ATPA2_PELCD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3A0B8	GLGA_PELCD		cytoplasm
Q3A0D8	GLMU_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A131	MURA_PELCD	cytoplasm (probable).	inner membrane
Q3A132	HIS1_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A133	HISX_PELCD		cytoplasm
Q3A135	HIS5_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A136	HIS4_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A137	HIS6_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A1D3	Y2586_PELCD		cytoplasm
Q3A1D9	PROB_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A1E0	PROA_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A1E5	GPMI_PELCD		cytoplasm
Q3A1R3	SYI_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A1T0	Y2438_PELCD		cytoplasm
Q3A1U6	DAPB_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A201	G6PI_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A220	SYI_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A230	RUVB_PELCD		cytoplasm
Q3A231	RUVA_PELCD		cytoplasm
Q3A232	RUVC_PELCD		cytoplasm

Q3A241	PPNK_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A246	ARGJ_PELCD	cytoplasm (probable).	cytoplasm
Q3A247	DEOC_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A253	SYV_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A258	RIMK_PELCD		cytoplasm
Q3A282	PHS_PELCD		cytoplasm
Q3A2E9	RL19_PELCD		cytoplasm
Q3A2F0	RNH2_PELCD	cytoplasm (potential).	extracellular
Q3A2F7	MRAZ_PELCD		inner membrane
Q3A2F8	MRAW_PELCD		cytoplasm
Q3A2G3	MRAY_PELCD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3A2G6	MURG_PELCD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3A2G8	MURB_PELCD	cytoplasm (probable).	inner membrane
Q3A2J4	AROK2_PELCD	cytoplasm (probable).	cytoplasm
Q3A2N6	AROB_PELCD	cytoplasm (probable).	cytoplasm
Q3A2Z2	HIS3_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A311	ISPE_PELCD		cytoplasm
Q3A317	RS6_PELCD		cytoplasm
Q3A320	RL9_PELCD		cytoplasm
Q3A3A5	ILVD_PELCD		cytoplasm
Q3A3B2	LEU3_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A3C1	QUEA_PELCD	cytoplasm (potential).	cytoplasm
Q3A3F2	FPG_PELCD		cytoplasm
Q3A3I8	PAND_PELCD		cytoplasm
Q3A3N8	AROK1_PELCD	cytoplasm (probable).	cytoplasm
Q3A3X5	CLPP_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A445	LEPA_PELCD		cytoplasm
Q3A499	DUT_PELCD		extracellular
Q3A4A2	RS15_PELCD		cytoplasm
Q3A4A3	TRUB_PELCD		cytoplasm
Q3A4B0	Y1552_PELCD		inner membrane
Q3A4F1	MUTS_PELCD		cytoplasm
Q3A4L7	NRDR_PELCD		cytoplasm
Q3A4L9	GLYA_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A4M7	RL32_PELCD		cytoplasm
Q3A4N5	SURE_PELCD	cytoplasm (potential).	periplasm
Q3A4P5	RS20_PELCD		cytoplasm
Q3A4Q8	RNC_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A522	RPOZ_PELCD		cytoplasm
Q3A554	FABZ_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A585	ACCA_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A590	GUAA_PELCD		cytoplasm
Q3A5F4	FLGH_PELCD	outer membrane; lipid-anchor (probable).	inner membrane
Q3A5F5	FLGI_PELCD	periplasm (by similarity).	periplasm
Q3A5V3	ACPS_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A5V4	PDXJ_PELCD	cytoplasm (probable).	cytoplasm
Q3A6I7	TRMB_PELCD		cytoplasm
Q3A6I8	TRUD_PELCD		cytoplasm
Q3A6L6	TRPD_PELCD		cytoplasm
Q3A6M0	RPOA_PELCD		cytoplasm
Q3A6M1	RS4_PELCD		cytoplasm

Q3A6M2	RS11_PELCD		cytoplasm
Q3A6M3	RS13_PELCD		cytoplasm
Q3A6N0	RS5_PELCD		cytoplasm
Q3A6N3	RS8_PELCD		cytoplasm
Q3A6N8	RS17_PELCD		cytoplasm
Q3A6P0	RL16_PELCD		cytoplasm
Q3A6P1	RS3_PELCD		cytoplasm
Q3A6P4	RL2_PELCD		cytoplasm
Q3A6P8	RS10_PELCD		cytoplasm
Q3A6Q1	RS7_PELCD		cytoplasm
Q3A6Q2	RS12_PELCD		cytoplasm
Q3A6Q3	RPOC_PELCD		cytoplasm
Q3A6Q4	RPOB_PELCD		cytoplasm
Q3A6Q6	RL10_PELCD		cytoplasm
Q3A6Q7	RL1_PELCD		cytoplasm
Q3A6U7	Y651_PELCD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3A6V2	UPPP_PELCD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3A6Y9	THIG2_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A734	UVRC_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A7G2	METX_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A7P5	THIG1_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A7R3	HIS8_PELCD		cytoplasm
Q3A7V6	HSLO_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A828	PURA_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A8C2	DNAK_PELCD		cytoplasm
Q3A8C6	ISPD_PELCD		cytoplasm
Q3A8C7	ISPF_PELCD		extracellular
Q3A8D0	UVRB_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A8M6	RECF_PELCD	cytoplasm (by similarity).	cytoplasm
Q3A8N7	RECO_PELCD		cytoplasm
Q3BM92	FPG_XANC5		cytoplasm
Q3BMF2	XYLA_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BNA6	DUT_XANC5		extracellular
Q3BNC2	SYX_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BP13	ATPA_XANC5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3BP20	GLMU_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BP98	SYV_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BPJ6	LEU2_XANC5		cytoplasm
Q3BPJ7	LEUD_XANC5		cytoplasm
Q3BPJ8	LEU3_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BPK3	ILVC_XANC5		cytoplasm
Q3BPS1	RPOZ_XANC5		cytoplasm
Q3BQ08	MSCL_XANC5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3BQ14	TRMB_XANC5		cytoplasm
Q3BQ15	THIG_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BQ65	RIMK_XANC5		cytoplasm
Q3BQF1	RUVC_XANC5		cytoplasm
Q3BQF2	RUVA_XANC5		cytoplasm

Q3BQF5	RUVB_XANC5		cytoplasm
Q3BQS1	AROB_XANC5	cytoplasm (probable).	cytoplasm
Q3BQS2	AROK_XANC5	cytoplasm (probable).	cytoplasm
Q3BQS3	PDXH_XANC5		cytoplasm
Q3BQX2	MURA_XANC5	cytoplasm (probable).	inner membrane
Q3BR53	GPMA_XANC5		cytoplasm
Q3BRL9	GLMM_XANC5		cytoplasm
Q3BRP3	Y2839_XANC5		inner membrane
Q3BRP7	TRUB_XANC5		cytoplasm
Q3BRP8	RS15_XANC5		cytoplasm
Q3BS21	HTPG_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BS36	QUEA_XANC5	cytoplasm (potential).	cytoplasm
Q3BSJ0	PROB_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BSJ1	PROA_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BSK1	METX_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BSK6	CCME2_XANC5	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3BSP7	GUAA_XANC5		cytoplasm
Q3BSV4	PHS_XANC5		cytoplasm
Q3BTC3	UVRC_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BTC7	LPXK_XANC5		cytoplasm
Q3BTH9	Y2203_XANC5		cytoplasm
Q3BTW6	NDK_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BU05	FLGH_XANC5	outer membrane; lipid-anchor (probable).	inner membrane
Q3BU06	FLGI_XANC5	periplasm (by similarity).	periplasm
Q3BUC6	DAPB_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BUF2	HIS6_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BUF3	HIS4_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BUF4	HIS5_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BUF6	HIS8_XANC5		cytoplasm
Q3BUF7	HISX_XANC5		cytoplasm
Q3BUJ8	MURB_XANC5	cytoplasm (probable).	inner membrane
Q3BUL3	G6PI_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BUL4	PAND_XANC5		cytoplasm
Q3BUS4	SURE_XANC5	cytoplasm (potential).	periplasm
Q3BUS6	TRUD_XANC5		cytoplasm
Q3BUS7	ISPF_XANC5		extracellular
Q3BUS8	ISPD_XANC5		cytoplasm
Q3BUW9	CCME1_XANC5	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3BV17	ZIPA_XANC5	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q3BV19	RL9_XANC5		cytoplasm
Q3BV21	RS6_XANC5		cytoplasm
Q3BV46	PPNK_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BV87	SLYX_XANC5		inner membrane
Q3BVB8	DNAK_XANC5		cytoplasm
Q3BVJ6	GLND_XANC5		cytoplasm
Q3BVL5	FABZ_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BVM0	ACCA_XANC5	cytoplasm (by similarity).	cytoplasm

Q3BVU6	NAGZ_XANC5	cytoplasm (by similarity).	extracellular
Q3BVV1	RUMA_XANC5		cytoplasm
Q3BVV3	RECO_XANC5		cytoplasm
Q3BVV6	RNC_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BVV9	LEPA_XANC5		cytoplasm
Q3BVY0	MUTS_XANC5		cytoplasm
Q3BVY7	RL19_XANC5		cytoplasm
Q3BW42	SYI_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BW45	RS20_XANC5		cytoplasm
Q3BWF4	PURA_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BWJ0	RL32_XANC5		cytoplasm
Q3BWQ1	CLPP_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BWW9	RPOA_XANC5		cytoplasm
Q3BWW0	RS4_XANC5		cytoplasm
Q3BWW1	RS11_XANC5		cytoplasm
Q3BWW2	RS13_XANC5		cytoplasm
Q3BWW6	RS5_XANC5		cytoplasm
Q3BWW9	RS8_XANC5		cytoplasm
Q3BWX4	RS17_XANC5		cytoplasm
Q3BWX6	RL16_XANC5		cytoplasm
Q3BWX7	RS3_XANC5		cytoplasm
Q3BWY0	RL2_XANC5		cytoplasm
Q3BWY4	RS10_XANC5		cytoplasm
Q3BWY8	RS7_XANC5		cytoplasm
Q3BWY9	RS12_XANC5		cytoplasm
Q3BWZ0	RPOC_XANC5		cytoplasm
Q3BWZ1	RPOB_XANC5		cytoplasm
Q3BWZ3	RL10_XANC5		cytoplasm
Q3BWZ4	RL1_XANC5		cytoplasm
Q3BX03	ISPE_XANC5		cytoplasm
Q3BX43	TAL_XANC5	cytoplasm (probable).	cytoplasm
Q3BXF2	MURG_XANC5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3BXF4	MRAY_XANC5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3BXF9	MRAW_XANC5		cytoplasm
Q3BXG0	MRAZ_XANC5		inner membrane
Q3BXI6	NRDR_XANC5		cytoplasm
Q3BXI8	GLYA_XANC5	cytoplasm (by similarity).	cytoplasm
Q3BYA7	NUDH_XANC5		cytoplasm
Q3BYB6	TRPD_XANC5		cytoplasm
Q3BYE8	HGD_XANC5		cytoplasm
Q3BYI1	GLGA_XANC5		cytoplasm
Q3BYS5	ILVD_XANC5		cytoplasm
Q3BZ95	UPPP_XANC5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3BZH2	LLDD_XANC5		cytoplasm
Q3BZR9	PDXJ_XANC5	cytoplasm (probable).	cytoplasm
Q3BZS9	RECF_XANC5	cytoplasm (by similarity).	cytoplasm
Q3HS05	NAPA_PSEST	periplasm (by similarity).	periplasm
Q3HY20	HRPA_ERWPY	extracellular and secreted via type iii secretion system (by similarity).	extracellular

Q3IBX5	CDD_PSEHT		cytoplasm
Q3IC08	DNAK_PSEHT		cytoplasm
Q3ICD4	HIS6_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3ICF0	HISX_PSEHT		cytoplasm
Q3ICF4	HIS4_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3ICL0	Y4058_PSEHT		cytoplasm
Q3ICU8	DEOD_PSEHT		cytoplasm
Q3ID04	ILVD2_PSEHT		cytoplasm
Q3ID15	NDK_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IDA4	UPPP2_PSEHT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3IDC8	MUTS_PSEHT		cytoplasm
Q3IDD3	SURE_PSEHT	cytoplasm (potential).	periplasm
Q3IDD5	GPPA_PSEHT		extracellular
Q3IDE8	RECF_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IDK9	PDXJ_PSEHT	cytoplasm (probable).	cytoplasm
Q3IDL2	RNC_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IDL4	LEPA_PSEHT		cytoplasm
Q3IDL9	NUDH_PSEHT		cytoplasm
Q3IDP4	PDXH_PSEHT		cytoplasm
Q3IDQ4	TRUD_PSEHT		cytoplasm
Q3IDQ5	ISPF_PSEHT		extracellular
Q3IDQ6	ISPD_PSEHT		cytoplasm
Q3IDW0	FLGI_PSEHT	periplasm (by similarity).	periplasm
Q3IDW1	FLGH_PSEHT	outer membrane; lipid-anchor (probable).	inner membrane
Q3IE16	CCME_PSEHT	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3IE61	GLMM_PSEHT		cytoplasm
Q3IEA2	SYI_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IEA5	RS20_PSEHT		cytoplasm
Q3IEC6	RL19_PSEHT		cytoplasm
Q3IEY7	PROA_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IEY8	PROB1_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IEZ3	PURA1_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IF18	RL16_PSEHT		cytoplasm
Q3IF19	RS3_PSEHT		cytoplasm
Q3IF22	RL2_PSEHT		cytoplasm
Q3IF26	RS10_PSEHT		cytoplasm
Q3IF46	CLPP_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IFM3	RS17_PSEHT		cytoplasm
Q3IFN6	THIG_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IFW1	GCSP_PSEHT		cytoplasm
Q3IFY0	MURG_PSEHT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3IFZ5	MRAZ_PSEHT		inner membrane
Q3IFZ6	MRAW_PSEHT		cytoplasm
Q3IG01	MRAY_PSEHT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3IG20	MURA_PSEHT	cytoplasm (probable).	inner membrane
Q3IG57	RIMK_PSEHT		cytoplasm

Q3IGX6	LPXK_PSEHT		cytoplasm
Q3IH68	URE1_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IH69	URE2_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IH70	URE3_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IHD7	RL32_PSEHT		cytoplasm
Q3IHJ1	GUAA_PSEHT		cytoplasm
Q3IHK8	PROB2_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IHV1	UPPP1_PSEHT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3IHX2	Y2303_PSEHT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3II23	GLYA_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3II24	NRDR_PSEHT		cytoplasm
Q3II62	SYI1_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3II73	SYV_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3II82	RL9_PSEHT		cytoplasm
Q3IIA2	RS6_PSEHT		cytoplasm
Q3IIE1	SECB_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IIE4	GPMI_PSEHT		cytoplasm
Q3III5	Y1882_PSEHT		inner membrane
Q3IIJ0	RUVC_PSEHT		cytoplasm
Q3IIJ1	RUVA_PSEHT		cytoplasm
Q3IIJ2	RUVB_PSEHT		cytoplasm
Q3IIQ6	PURA2_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IIW9	RNH2_PSEHT	cytoplasm (potential).	extracellular
Q3IIX1	ACCA_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IYY5	FABZ_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IIZ3	PHS_PSEHT		cytoplasm
Q3IIZ7	GLND_PSEHT		cytoplasm
Q3IJ25	TUSA_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IJ55	Y995_PSEHT		inner membrane
Q3IJ74	RS15_PSEHT		cytoplasm
Q3IJ75	TRUB_PSEHT		cytoplasm
Q3IJA2	AROB_PSEHT	cytoplasm (probable).	cytoplasm
Q3IJC5	AROK_PSEHT	cytoplasm (probable).	cytoplasm
Q3IJH1	ILVD1_PSEHT		cytoplasm
Q3IJH7	RPOZ_PSEHT		cytoplasm
Q3IJI6	RPOA_PSEHT		cytoplasm
Q3IJJ5	RS4_PSEHT		cytoplasm
Q3IJJ9	RS8_PSEHT		cytoplasm
Q3IJK2	RS5_PSEHT		cytoplasm
Q3IJK6	RS13_PSEHT		cytoplasm
Q3IJK7	RS11_PSEHT		cytoplasm
Q3IJS3	LEU3_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IJS4	LEU2_PSEHT		cytoplasm
Q3IJS5	LEUD_PSEHT		cytoplasm
Q3IJT8	SLYX_PSEHT		inner membrane
Q3IJU8	MURB_PSEHT	cytoplasm (probable).	inner membrane
Q3IK30	GLMU_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IK48	ATPA_PSEHT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3IK98	ISPE_PSEHT		cytoplasm
Q3IKB2	MTFA_PSEHT		cytoplasm

Q3IKF4	UVRB_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IKH4	G6PI_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IKQ2	HTPG_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IKQ7	DAPB_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IKR4	PPNK_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3IKY4	TRPD_PSEHT		cytoplasm
Q3ILB9	QUEA_PSEHT	cytoplasm (potential).	cytoplasm
Q3ILJ1	TRMB_PSEHT		cytoplasm
Q3ILJ9	SYI2_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3ILP2	HSLO_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3ILP3	PPCK_PSEHT	cytoplasm (by similarity).	cytoplasm
Q3ILP6	RS7_PSEHT		cytoplasm
Q3ILP7	RS12_PSEHT		cytoplasm
Q3ILP8	RPOC_PSEHT		cytoplasm
Q3ILP9	RPOB_PSEHT		cytoplasm
Q3ILQ2	RL10_PSEHT		cytoplasm
Q3ILQ3	RL1_PSEHT		cytoplasm
Q3IW88	TRMB_RHOS4		cytoplasm
Q3IXP4	ILVD_RHOS4		cytoplasm
Q3IXX7	PROA_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3IXX8	PROB_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3IY62	RS20_RHOS4		cytoplasm
Q3IY64	FPG_RHOS4		cytoplasm
Q3IY97	FLGI2_RHOS4	periplasm (by similarity).	periplasm
Q3IYC6	GPH_RHOS4		cytoplasm
Q3IYI5	MUTS_RHOS4		cytoplasm
Q3IYM4	XYLA_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3IYM7	DNAK_RHOS4		cytoplasm
Q3IYN8	Y2778_RHOS4		inner membrane
Q3IYQ0	UPPP_RHOS4	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3IYT8	RS15_RHOS4		cytoplasm
Q3IYU1	TRUB_RHOS4		cytoplasm
Q3IYU4	DAPB_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3IYX6	UVRC_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3IZ06	RL19_RHOS4		cytoplasm
Q3IZC1	NUDH_RHOS4		cytoplasm
Q3IZF4	PHS_RHOS4		cytoplasm
Q3IZN2	GLYA_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3IZY4	MSCL_RHOS4	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3J077	MURA_RHOS4	cytoplasm (probable).	inner membrane
Q3J079	HISX1_RHOS4		cytoplasm
Q3J0B2	DUT_RHOS4		extracellular
Q3J0F5	RUVC_RHOS4		cytoplasm
Q3J0F6	RUVA_RHOS4		cytoplasm
Q3J0F8	RUVB_RHOS4		cytoplasm
Q3J0S2	YK44_RHOS4		inner membrane
Q3J0W6	NRDR_RHOS4		cytoplasm
Q3J0Z3	PURA_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J151	URE2_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J154	URE1_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J156	UREE_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J170	RCEH_RHOS4		inner membrane

Q3J1A0	BCHZ_RHOS4		cytoplasm
Q3J1A1	PUFQ_RHOS4		cytoplasm
Q3J1A5	RCEL_RHOS4	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3J1A6	RCEM_RHOS4	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3J1G6	CLPP_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J1L9	RL9_RHOS4		cytoplasm
Q3J1M1	RS6_RHOS4		cytoplasm
Q3J1T3	FLG11_RHOS4	periplasm (by similarity).	periplasm
Q3J210	GUAA_RHOS4		cytoplasm
Q3J252	QUEA_RHOS4	cytoplasm (potential).	cytoplasm
Q3J273	PURL_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J278	CCME_RHOS4	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3J2D0	NDK_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J2I9	AROK_RHOS4	cytoplasm (probable).	cytoplasm
Q3J2J0	AROB_RHOS4	cytoplasm (probable).	cytoplasm
Q3J2K8	ISPF_RHOS4		extracellular
Q3J2K9	ISPD_RHOS4		cytoplasm
Q3J2U4	G6PI_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J2W6	FABZ_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J363	RL32_RHOS4		cytoplasm
Q3J3D0	SURE_RHOS4	cytoplasm (potential).	periplasm
Q3J3H0	GLMU_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J3I3	SYV_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J433	ATPA_RHOS4	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3J445	HIS8_RHOS4		cytoplasm
Q3J446	RS4_RHOS4		cytoplasm
Q3J4D4	GCSP_RHOS4		cytoplasm
Q3J4H6	HISX2_RHOS4		cytoplasm
Q3J4L9	MURB_RHOS4	cytoplasm (probable).	inner membrane
Q3J4M2	MURG_RHOS4	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3J4M8	MRAY_RHOS4	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3J4N3	MRAW_RHOS4		cytoplasm
Q3J4N4	MRAZ_RHOS4		inner membrane
Q3J4P0	LEPA_RHOS4		cytoplasm
Q3J4Z5	SYV_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J5C0	ILVC_RHOS4		cytoplasm
Q3J5H6	GLND_RHOS4		cytoplasm
Q3J5K7	ISPE_RHOS4		cytoplasm
Q3J5L4	ACCA_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J5P8	RPOA_RHOS4		cytoplasm
Q3J5P9	RS11_RHOS4		cytoplasm
Q3J5Q0	RS13_RHOS4		cytoplasm
Q3J5Q5	RS5_RHOS4		cytoplasm

Q3J5Q8	RS8_RHOS4		cytoplasm
Q3J5R3	RS17_RHOS4		cytoplasm
Q3J5R5	RL16_RHOS4		cytoplasm
Q3J5R6	RS3_RHOS4		cytoplasm
Q3J5R9	RL2_RHOS4		cytoplasm
Q3J5S3	RS10_RHOS4		cytoplasm
Q3J5S6	RS7_RHOS4		cytoplasm
Q3J5S7	RS12_RHOS4		cytoplasm
Q3J5S9	RPOC_RHOS4		cytoplasm
Q3J5T0	RPOB_RHOS4		cytoplasm
Q3J5T2	RL10_RHOS4		cytoplasm
Q3J5T3	RL1_RHOS4		cytoplasm
Q3J5V6	PPCK_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J5V8	RECO_RHOS4		cytoplasm
Q3J5W1	RNC_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J5W3	ACPS_RHOS4	cytoplasm (by similarity).	cytoplasm
Q3J5W4	PDXJ_RHOS4	cytoplasm (probable).	cytoplasm
Q3J5W7	RPOZ_RHOS4		cytoplasm
Q3J6H3	RNH2_RHOS4	cytoplasm (potential).	extracellular
Q3J6I0	LPXK_RHOS4		cytoplasm
Q3J6M9	ATPA_NITOC	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3J6N3	GLMU_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J6P9	HIS2_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J6Q0	HIS3_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J6Q1	HIS6_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J6Q2	HIS4_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J6Q3	HIS5_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J6R8	RS20_NITOC		cytoplasm
Q3J6R9	PROB_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J6W1	DUT_NITOC		extracellular
Q3J716	LEU2_NITOC		cytoplasm
Q3J768	URE3_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J769	URE2_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J770	URE1_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J780	MRAZ_NITOC		inner membrane
Q3J781	MRAW_NITOC		cytoplasm
Q3J786	MRAY_NITOC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3J789	MURG_NITOC	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3J791	MURB_NITOC	cytoplasm (probable).	inner membrane
Q3J7A0	ARGJ_NITOC	cytoplasm (probable).	cytoplasm
Q3J7B5	UPPP_NITOC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3J7D8	DNAK_NITOC		cytoplasm
Q3J7E0	DAPB_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J7G9	MURA_NITOC	cytoplasm (probable).	inner membrane
Q3J7H0	HIS1_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J7H1	HISX_NITOC		cytoplasm
Q3J7H2	HIS82_NITOC		cytoplasm
Q3J7R7	LPXK_NITOC		cytoplasm

Q3J7T1	PROA_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J7U5	FPG_NITOC		cytoplasm
Q3J808	PURA_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J826	GLMM_NITOC		cytoplasm
Q3J875	ILVC_NITOC		cytoplasm
Q3J897	TRPD_NITOC		cytoplasm
Q3J8A0	GPH_NITOC		cytoplasm
Q3J8D2	LEPA_NITOC		cytoplasm
Q3J8D5	RNC_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J8L6	FLGH_NITOC	outer membrane; lipid-anchor (probable).	inner membrane
Q3J8L7	FLGI_NITOC	periplasm (by similarity).	periplasm
Q3J8Q4	RL1_NITOC		cytoplasm
Q3J8Q5	RL10_NITOC		cytoplasm
Q3J8Q7	RPOB_NITOC		cytoplasm
Q3J8Q8	RPOC_NITOC		cytoplasm
Q3J8Q9	RS12_NITOC		cytoplasm
Q3J8R0	RS7_NITOC		cytoplasm
Q3J8R3	RS10_NITOC		cytoplasm
Q3J8R7	RL2_NITOC		cytoplasm
Q3J8S0	RS3_NITOC		cytoplasm
Q3J8S1	RL16_NITOC		cytoplasm
Q3J8S3	RS17_NITOC		cytoplasm
Q3J8S8	RS8_NITOC		cytoplasm
Q3J8T1	RS5_NITOC		cytoplasm
Q3J8T5	RS13_NITOC		cytoplasm
Q3J8T6	RS11_NITOC		cytoplasm
Q3J8T7	RS4_NITOC		cytoplasm
Q3J8T8	RPOA_NITOC		cytoplasm
Q3J8Y1	RL19_NITOC		cytoplasm
Q3J910	PHS_NITOC		cytoplasm
Q3J953	NAGZ_NITOC	cytoplasm (by similarity).	extracellular
Q3J9B4	Y2122_NITOC		inner membrane
Q3J9B8	TRUB_NITOC		cytoplasm
Q3J9B9	RS15_NITOC		cytoplasm
Q3J9G2	HSLO_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J9K8	GLYA_NITOC	cytoplasm (by similarity).	cytoplasm
Q3J9L3	RNH2_NITOC	cytoplasm (potential).	extracellular
Q3J9L7	NUDH_NITOC		cytoplasm
Q3J9N3	ILVD_NITOC		cytoplasm
Q3JAF6	HTPG_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JAJ8	CLPP_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JAK5	RL32_NITOC		cytoplasm
Q3JAM9	QUEA_NITOC	cytoplasm (potential).	cytoplasm
Q3JAN0	UVRC_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JAP1	THIG_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JAW3	RIMK_NITOC		cytoplasm
Q3JBA8	GPMA_NITOC		cytoplasm
Q3JBT2	RPOZ_NITOC		cytoplasm
Q3JBV4	PPNK_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JC03	UVRB_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JC50	SYV_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JCB2	TRMB_NITOC		cytoplasm
Q3JCC4	LEU3_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JCC5	LEUD_NITOC		cytoplasm

Q3JCJ0	CCME_NITOC	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3JCL5	MUTS_NITOC		cytoplasm
Q3JCM3	SYU_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JCN1	G6PI_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JCN5	NDK_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JCP7	PAND_NITOC		cytoplasm
Q3JCS7	TRUD_NITOC		cytoplasm
Q3JCS8	ISPF_NITOC		extracellular
Q3JCS9	ISPD_NITOC		cytoplasm
Q3JCW5	FABZ_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JCX7	GLND_NITOC		cytoplasm
Q3JCY2	RUMA_NITOC		cytoplasm
Q3JCY7	ACPS_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JCY8	PDXJ_NITOC	cytoplasm (probable).	cytoplasm
Q3JCY9	RECO_NITOC		cytoplasm
Q3JCZ4	SURE_NITOC	cytoplasm (potential).	periplasm
Q3JD12	GLGA2_NITOC		cytoplasm
Q3JDG3	GUAA_NITOC		cytoplasm
Q3JDR0	ISPE_NITOC		cytoplasm
Q3JEG3	AROB_NITOC	cytoplasm (probable).	cytoplasm
Q3JEG4	AROK_NITOC	cytoplasm (probable).	cytoplasm
Q3JEJ4	RL9_NITOC		cytoplasm
Q3JEJ6	RS6_NITOC		cytoplasm
Q3JEN8	HIS81_NITOC		cytoplasm
Q3JES3	RUVB_NITOC		cytoplasm
Q3JES4	RUVA_NITOC		cytoplasm
Q3JES5	RUVC_NITOC		cytoplasm
Q3JEW3	GLGA1_NITOC		cytoplasm
Q3JF14	Y040_NITOC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3JF23	GPMI_NITOC		cytoplasm
Q3JF36	RECF_NITOC	cytoplasm (by similarity).	cytoplasm
Q3JGP5	GLYA2_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JJK4	DNS2_BURP1		extracellular
Q3JJP7	ATPA2_BURP1	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3JMQ0	RL1_BURP1		cytoplasm
Q3JMQ1	RL10_BURP1		cytoplasm
Q3JMQ3	RPOB_BURP1		cytoplasm
Q3JMQ4	RPOC_BURP1		cytoplasm
Q3JMQ7	RS12_BURP1		cytoplasm
Q3JMQ8	RS7_BURP1		cytoplasm
Q3JMR2	RS10_BURP1		cytoplasm
Q3JMR6	RL2_BURP1		cytoplasm
Q3JMR9	RS3_BURP1		cytoplasm
Q3JMS2	RS17_BURP1		cytoplasm
Q3JMS7	RS8_BURP1		cytoplasm
Q3JMT0	RS5_BURP1		cytoplasm
Q3JMT6	RS13_BURP1		cytoplasm
Q3JMT7	RS11_BURP1		cytoplasm
Q3JMT8	RS4_BURP1		cytoplasm

Q3JMT9	RPOA_BURP1		cytoplasm
Q3JMW0	AROK_BURP1	cytoplasm (probable).	cytoplasm
Q3JMW1	AROB_BURP1	cytoplasm (probable).	cytoplasm
Q3JMY2	THIG_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JMZ4	MURA_BURP1	cytoplasm (probable).	inner membrane
Q3JMZ5	HIS1_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JMZ6	HISX_BURP1		cytoplasm
Q3JMZ7	HIS82_BURP1		cytoplasm
Q3JN00	HIS5_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JN01	HIS4_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JN02	HIS6_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JN03	HIS3_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JN04	HIS2_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JNB1	TRPD_BURP1		cytoplasm
Q3JND0	MRAW_BURP1		cytoplasm
Q3JND5	MRAY_BURP1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3JND8	MURG_BURP1	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3JNE9	ARGJ_BURP1	cytoplasm (probable).	cytoplasm
Q3JNG1	PROB_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JNG3	NUDH_BURP1		cytoplasm
Q3JNM7	DAPB_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JNN5	PROA_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JNS0	SYJ_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JNS5	RUVB_BURP1		cytoplasm
Q3JNS6	RUVA_BURP1		cytoplasm
Q3JNS7	RUVC_BURP1		cytoplasm
Q3JNW2	QUEA_BURP1	cytoplasm (potential).	cytoplasm
Q3JP04	PPNK_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JP10	DNAK_BURP1		cytoplasm
Q3JP81	GLYA1_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JP82	NRDR_BURP1		cytoplasm
Q3JPA3	HGD_BURP1		cytoplasm
Q3JPJ6	URE1_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JPJ7	URE2_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JPJ8	URE3_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JPL1	UPPP2_BURP1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3JPL2	TRMB_BURP1		cytoplasm
Q3JPT0	RPOZ_BURP1		cytoplasm
Q3JQ09	RL19_BURP1		cytoplasm
Q3JQ63	RL32_BURP1		cytoplasm
Q3JQ75	LEPA_BURP1		cytoplasm
Q3JQ80	PDXJ_BURP1	cytoplasm (probable).	cytoplasm
Q3JQS6	MUTS_BURP1		cytoplasm
Q3JR26	GLND_BURP1		cytoplasm
Q3JR40	FABZ_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JR43	RNH2_BURP1	cytoplasm (potential).	extracellular
Q3JR65	GUAA_BURP1		cytoplasm
Q3JR90	MSCL_BURP1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q3JR99	ISPD_BURP1		cytoplasm
Q3JRA0	ISPF_BURP1		extracellular
Q3JRC9	CLPP_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JRE2	G6PI_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JRJ2	RS6_BURP1		cytoplasm
Q3JRJ5	RL9_BURP1		cytoplasm
Q3JRP0	SURE_BURP1	cytoplasm (potential).	periplasm
Q3JRP8	RUMA_BURP1		cytoplasm
Q3JRQ0	NDK_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JRR4	PURA_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JSY7	TRUB_BURP1		cytoplasm
Q3JSZ1	Y1913_BURP1		inner membrane
Q3JT80	Y1822_BURP1		inner membrane
Q3JT18	UPPP1_BURP1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3JUC2	ILVC_BURP1		cytoplasm
Q3JUL7	TAL_BURP1	cytoplasm (probable).	cytoplasm
Q3JUM5	HTPG_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JUY9	PAND_BURP1		cytoplasm
Q3JV12	ILVD_BURP1		cytoplasm
Q3JV72	DUT_BURP1		extracellular
Q3JVA8	LPXK_BURP1		cytoplasm
Q3JVB6	RS20_BURP1		cytoplasm
Q3JVB9	MURB_BURP1	cytoplasm (probable).	inner membrane
Q3JW82	FPG_BURP1		cytoplasm
Q3JW85	ISPE_BURP1		cytoplasm
Q3JW89	HIS81_BURP1		cytoplasm
Q3JWH7	GPMA_BURP1		cytoplasm
Q3JWX1	GLMU_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JX21	FLGH_BURP1	outer membrane; lipid-anchor (probable).	inner membrane
Q3JXB2	METX_BURP1	cytoplasm (by similarity).	cytoplasm
Q3JXT0	PHS_BURP1		cytoplasm
Q3JXV6	ATPA1_BURP1	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3JY08	GCSP_BURP1		cytoplasm
Q3K439	ATPA_PSEPF	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3K443	GLMU_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K4L7	RPOZ_PSEPF		cytoplasm
Q3K4M6	DUT_PSEPF		extracellular
Q3K4Z1	GCSP2_PSEPF		cytoplasm
Q3K539	NUDH_PSEPF		cytoplasm
Q3K559	ILVD_PSEPF		cytoplasm
Q3K566	FPG_PSEPF		cytoplasm
Q3K587	THIG_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K588	TRMB_PSEPF		cytoplasm
Q3K594	METX_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K5K9	GLYA3_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K5U8	GPH_PSEPF		cytoplasm
Q3K5V3	TRPD_PSEPF		cytoplasm
Q3K5W9	SYI_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K5X8	RL1_PSEPF		cytoplasm

Q3K5X9	RL10_PSEPF		cytoplasm
Q3K5Y1	RPOB_PSEPF		cytoplasm
Q3K5Y2	RPOC_PSEPF		cytoplasm
Q3K5Y3	RS12_PSEPF		cytoplasm
Q3K5Y4	RS7_PSEPF		cytoplasm
Q3K5Y7	RS10_PSEPF		cytoplasm
Q3K5Z1	RL2_PSEPF		cytoplasm
Q3K5Z4	RS3_PSEPF		cytoplasm
Q3K5Z7	RS17_PSEPF		cytoplasm
Q3K602	RS8_PSEPF		cytoplasm
Q3K605	RS5_PSEPF		cytoplasm
Q3K609	RS13_PSEPF		cytoplasm
Q3K610	RS11_PSEPF		cytoplasm
Q3K611	RS4_PSEPF		cytoplasm
Q3K612	RPOA_PSEPF		cytoplasm
Q3K647	NRDR_PSEPF		cytoplasm
Q3K693	PROA_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K6I0	MSCL_PSEPF	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3K6J0	GLYA2_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K6L0	PROB_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K6L2	RS20_PSEPF		cytoplasm
Q3K6Q4	G6PI_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K6T1	ILVC_PSEPF		cytoplasm
Q3K6W5	ISPE_PSEPF		cytoplasm
Q3K735	MRAZ_PSEPF		inner membrane
Q3K736	MRAW_PSEPF		cytoplasm
Q3K741	MRAY_PSEPF	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3K744	MURG_PSEPF	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3K795	QUEA_PSEPF	cytoplasm (potential).	cytoplasm
Q3K7B2	NDK_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K7D0	GUAA_PSEPF		cytoplasm
Q3K7U0	ARGJ_PSEPF	cytoplasm (probable).	cytoplasm
Q3K7V1	SLYX_PSEPF		inner membrane
Q3K7V8	RUVC_PSEPF		cytoplasm
Q3K7V9	RUVA_PSEPF		cytoplasm
Q3K7W0	RUVB_PSEPF		cytoplasm
Q3K7X5	GCSP1_PSEPF		cytoplasm
Q3K8E9	RUMA_PSEPF		cytoplasm
Q3K8J2	LPXK_PSEPF		cytoplasm
Q3K8J6	MURB_PSEPF	cytoplasm (probable).	inner membrane
Q3K8K7	RL32_PSEPF		cytoplasm
Q3K8U2	HIS82_PSEPF		cytoplasm
Q3K9C9	TUSA_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3K9H0	TAL_PSEPF	cytoplasm (probable).	cytoplasm
Q3K9J0	UPPP2_PSEPF	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3K9W9	CLPP_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KBQ0	UVRC_PSEPF	cytoplasm (by similarity).	cytoplasm

Q3KC07	CCME2_PSEPF	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3KD12	UPPP1_PSEPF	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3KD65	GLGA_PSEPF		cytoplasm
Q3KDV1	GLYA1_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KDW0	XYLA_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KE68	PPNK_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KF38	UVRB_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KFB3	ZIPA_PSEPF	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q3KFG8	Y1745_PSEPF		inner membrane
Q3KFT8	HTPG_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KFY2	CCME1_PSEPF	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3KG59	FLGI_PSEPF	periplasm (by similarity).	periplasm
Q3KG60	FLGH_PSEPF	outer membrane; lipid-anchor (probable).	inner membrane
Q3KG65	PHS_PSEPF		cytoplasm
Q3KH79	MUTS_PSEPF		cytoplasm
Q3KH84	SURE_PSEPF	cytoplasm (potential).	periplasm
Q3KH85	TRUD_PSEPF		cytoplasm
Q3KH86	ISPF_PSEPF		extracellular
Q3KH90	ISPD_PSEPF		cytoplasm
Q3KH98	RNH2_PSEPF	cytoplasm (potential).	extracellular
Q3KHA1	FABZ_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KHB4	GLND_PSEPF		cytoplasm
Q3KHJ1	RL19_PSEPF		cytoplasm
Q3KHL6	PDXJ_PSEPF	cytoplasm (probable).	cytoplasm
Q3KHL7	RECO_PSEPF		cytoplasm
Q3KHL9	RNC_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KHM1	LEPA_PSEPF		cytoplasm
Q3KHV3	HGD_PSEPF		cytoplasm
Q3KHZ1	HIS81_PSEPF		cytoplasm
Q3KHZ2	HISX_PSEPF		cytoplasm
Q3KHZ3	HIS1_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KHZ4	MURA_PSEPF	cytoplasm (probable).	inner membrane
Q3KI82	TRUB_PSEPF		cytoplasm
Q3KI86	Y777_PSEPF		inner membrane
Q3KI98	DAPB_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KIA0	DNAK_PSEPF		cytoplasm
Q3KIS8	URE3_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KIT1	URE2_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KIT2	URE1_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KIX5	RL9_PSEPF		cytoplasm
Q3KIX8	RS6_PSEPF		cytoplasm
Q3KIY4	PURA_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KJA1	AROB_PSEPF	cytoplasm (probable).	cytoplasm
Q3KJA2	AROK_PSEPF	cytoplasm (probable).	cytoplasm
Q3KJC8	HIS3_PSEPF	cytoplasm (by similarity).	cytoplasm

Q3KJC9	HIS2_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KJI5	HIS6_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KJI6	HIS4_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KJI8	HIS5_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KJP4	PPCK_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KJP5	HSLO_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3KKF9	RECF_PSEPF	cytoplasm (by similarity).	cytoplasm
Q3SEU6	HIS5_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SEU7	HIS4_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SEU8	HIS6_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SEU9	HIS3_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SF64	ATPA_THIDA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3SF69	GLMU_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SFG5	UPPP_THIDA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3SFR7	DUT_THIDA		extracellular
Q3SG48	MSCL_THIDA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3SG61	PROA_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SG76	SLYX_THIDA		inner membrane
Q3SGA3	GPML_THIDA		cytoplasm
Q3SGB5	Y2382_THIDA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3SGC1	RL19_THIDA		cytoplasm
Q3SGR5	GPH_THIDA		cytoplasm
Q3SGS1	TRPD_THIDA		cytoplasm
Q3SGT0	RUVC_THIDA		cytoplasm
Q3SGT1	RUVA_THIDA		cytoplasm
Q3SGT3	RUVB_THIDA		cytoplasm
Q3SGX4	NRDR_THIDA		cytoplasm
Q3SGX5	GLYA_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SH10	THIG_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SH11	TRMB_THIDA		cytoplasm
Q3SH26	NUDH_THIDA		cytoplasm
Q3SH31	RS6_THIDA		cytoplasm
Q3SH34	RL9_THIDA		cytoplasm
Q3SH47	LEPA_THIDA		cytoplasm
Q3SH50	RNC_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SH52	RECO_THIDA		cytoplasm
Q3SH53	PDXJ_THIDA	cytoplasm (probable).	cytoplasm
Q3SH62	QUEA_THIDA	cytoplasm (potential).	cytoplasm
Q3SH73	G6PI2_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SH79	GLGA_THIDA		cytoplasm
Q3SH81	PAND_THIDA		cytoplasm
Q3SHA5	PPNK_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SHE4	ILVC_THIDA		cytoplasm
Q3SHL1	LEU2_THIDA		cytoplasm
Q3SHL2	LEUD_THIDA		cytoplasm
Q3SHL3	LEU3_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SHP2	MURA_THIDA	cytoplasm (probable).	inner membrane
Q3SHP3	HIS1_THIDA	cytoplasm (by similarity).	cytoplasm

Q3SHS3	SYI_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SHS6	RS20_THIDA		cytoplasm
Q3SHT9	PHS_THIDA		cytoplasm
Q3SI29	GUAA_THIDA		cytoplasm
Q3SI67	HISX_THIDA		cytoplasm
Q3SI68	HIS82_THIDA		cytoplasm
Q3SI69	HIS2_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SI98	CLPP_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SIE4	FLGH_THIDA	outer membrane; lipid-anchor (probable).	inner membrane
Q3SIE5	FLGI_THIDA	periplasm (by similarity).	periplasm
Q3SIK7	UVRB_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SIM2	RL32_THIDA		cytoplasm
Q3SIN4	DNAK_THIDA		cytoplasm
Q3SIR4	LPXK_THIDA		cytoplasm
Q3SJP0	MUTS_THIDA		cytoplasm
Q3SJS2	DAPB_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SJS7	PROB_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SJW8	HTPG_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SK37	ISPF_THIDA		extracellular
Q3SK38	ISPD_THIDA		cytoplasm
Q3SK65	G6PI1_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SK67	RUMA_THIDA		cytoplasm
Q3SK85	HIS81_THIDA		cytoplasm
Q3SKJ3	SURE_THIDA	cytoplasm (potential).	periplasm
Q3SKM7	RNH2_THIDA	cytoplasm (potential).	extracellular
Q3SKN0	FABZ_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SKP1	GLND_THIDA		cytoplasm
Q3SKU1	ACPS_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SKU2	NAGZ_THIDA	cytoplasm (by similarity).	extracellular
Q3SKW9	Y699_THIDA		inner membrane
Q3SKX3	TRUB_THIDA		cytoplasm
Q3SKX4	RS15_THIDA		cytoplasm
Q3SL47	Y617_THIDA		cytoplasm
Q3SL57	PURA_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SL74	NDK_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SL76	ACCA_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SL86	SYV_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SLI5	RPOZ_THIDA		cytoplasm
Q3SLJ8	SYI_THIDA	cytoplasm (by similarity).	cytoplasm
Q3SLM4	RPOA_THIDA		cytoplasm
Q3SLM5	RS4_THIDA		cytoplasm
Q3SLM6	RS11_THIDA		cytoplasm
Q3SLM7	RS13_THIDA		cytoplasm
Q3SLN2	RS5_THIDA		cytoplasm
Q3SLN5	RS8_THIDA		cytoplasm
Q3SLP0	RS17_THIDA		cytoplasm
Q3SLP2	RL16_THIDA		cytoplasm
Q3SLP3	RS3_THIDA		cytoplasm
Q3SLP6	RL2_THIDA		cytoplasm
Q3SLQ0	RS10_THIDA		cytoplasm
Q3SLQ3	RS7_THIDA		cytoplasm
Q3SLQ4	RS12_THIDA		cytoplasm
Q3SLQ5	RPOC_THIDA		cytoplasm
Q3SLQ6	RPOB_THIDA		cytoplasm
Q3SLQ8	RL10_THIDA		cytoplasm

Q3SLQ9	RL1_THIDA		cytoplasm
Q3SLR6	ISPE_THIDA		cytoplasm
Q3SLR9	FPG_THIDA		cytoplasm
Q3SLY4	AROB_THIDA	cytoplasm (probable).	cytoplasm
Q3SM89	AROK_THIDA	cytoplasm (probable).	cytoplasm
Q3SMH1	MURB_THIDA	cytoplasm (probable).	inner membrane
Q3SMH3	MURG_THIDA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3SMH6	MRAY_THIDA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3SMI1	MRAW_THIDA		cytoplasm
Q3SMI2	MRAZ_THIDA		inner membrane
Q3SMQ9	ARGJ_THIDA	cytoplasm (probable).	cytoplasm
Q3SMV1	RS20_NITWN		cytoplasm
Q3SN92	SLYX_NITWN		inner membrane
Q3SNB8	GLMM_NITWN		cytoplasm
Q3SNC6	PURA_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SNS6	HTPG_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SNS9	PHS_NITWN		cytoplasm
Q3SNU3	LEU3_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SNU7	LEUD_NITWN		cytoplasm
Q3SNV3	LEU2_NITWN		cytoplasm
Q3SNV5	RL19_NITWN		cytoplasm
Q3SP12	RUVC_NITWN		cytoplasm
Q3SP13	RUVA_NITWN		cytoplasm
Q3SP14	RUVB_NITWN		cytoplasm
Q3SPE0	RNH2_NITWN	cytoplasm (potential).	extracellular
Q3SPE5	ISPE_NITWN		cytoplasm
Q3SPI3	LPXK_NITWN		cytoplasm
Q3SPJ6	SYI_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SPS2	THIG_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SPZ6	HIS52_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SQ46	ILVC_NITWN		cytoplasm
Q3SQJ9	XGPT_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SQN7	RRMJ_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SQP4	GUAA_NITWN		cytoplasm
Q3SR23	Y2009_NITWN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3SRA3	MSCL_NITWN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3SRA8	RPOZ_NITWN		cytoplasm
Q3SRB0	PDXJ_NITWN	cytoplasm (probable).	cytoplasm
Q3SRB1	ACPS_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SRB3	RNC_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SRD2	CLPP_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SRI2	FABZ_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SRJ4	TRPD_NITWN		cytoplasm
Q3SRN4	QUEA_NITWN	cytoplasm (potential).	cytoplasm
Q3SRP6	GLMU_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SRQ5	SURE_NITWN	cytoplasm (potential).	periplasm
Q3SRV3	GLYA_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SRV4	NRDR_NITWN		cytoplasm

Q3SRY6	RL9_NITWN		cytoplasm
Q3SRY9	RS6_NITWN		cytoplasm
Q3SS12	SYV_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SS79	NDK_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SSP4	SYV_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SSU2	RPOA_NITWN		cytoplasm
Q3SSU3	RS11_NITWN		cytoplasm
Q3SSU4	RS13_NITWN		cytoplasm
Q3SSU9	RS5_NITWN		cytoplasm
Q3SSV2	RS8_NITWN		cytoplasm
Q3SSV7	RS17_NITWN		cytoplasm
Q3SSV9	RL16_NITWN		cytoplasm
Q3SSW0	RS3_NITWN		cytoplasm
Q3SSW3	RL2_NITWN		cytoplasm
Q3SSW7	RS10_NITWN		cytoplasm
Q3SSX0	RS7_NITWN		cytoplasm
Q3SSX1	RS12_NITWN		cytoplasm
Q3SSX9	RPOC_NITWN		cytoplasm
Q3SSY0	RPOB_NITWN		cytoplasm
Q3SSY2	RL10_NITWN		cytoplasm
Q3SSY3	RL1_NITWN		cytoplasm
Q3ST00	HIS3_NITWN	cytoplasm (by similarity).	cytoplasm
Q3ST14	RS4_NITWN		cytoplasm
Q3ST22	PURL_NITWN	cytoplasm (by similarity).	cytoplasm
Q3ST46	GCSP_NITWN		cytoplasm
Q3STD7	CCME_NITWN	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3STL0	FLGH_NITWN	outer membrane; lipid-anchor (probable).	inner membrane
Q3STL8	FLGI_NITWN	periplasm (by similarity).	periplasm
Q3STS5	MURB_NITWN	cytoplasm (probable).	inner membrane
Q3STS8	MURG_NITWN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3STT1	MRAY_NITWN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3STT6	MRAW_NITWN		cytoplasm
Q3SV02	PDXH_NITWN		cytoplasm
Q3SV39	METX_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SV41	HIS8_NITWN		cytoplasm
Q3SVC7	RL32_NITWN		cytoplasm
Q3SVD4	MUTS_NITWN		cytoplasm
Q3SVH3	NUDH_NITWN		cytoplasm
Q3SVI0	PROA_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SVI1	PROB_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SVJ4	ATPA_NITWN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3SVM8	AROB_NITWN	cytoplasm (probable).	cytoplasm
Q3SVM9	AROK_NITWN	cytoplasm (probable).	cytoplasm
Q3SVN2	ACCA_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SVN6	ARGJ_NITWN	cytoplasm (probable).	cytoplasm
Q3SVS4	PPCK_NITWN	cytoplasm (by similarity).	cytoplasm

Q3SVT1	LEPA_NITWN		cytoplasm
Q3SW26	HISX_NITWN		cytoplasm
Q3SW27	MURA_NITWN	cytoplasm (probable).	inner membrane
Q3SW60	ILVD_NITWN		cytoplasm
Q3SW71	GPMA_NITWN		cytoplasm
Q3SW72	DAPB_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SW76	DNAK_NITWN		cytoplasm
Q3SWA8	UPPP_NITWN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3SWE0	GLND_NITWN		cytoplasm
Q3SWE9	HIS51_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SWF0	HIS4_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SWF1	HIS6_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SWF2	HIS2_NITWN	cytoplasm (by similarity).	cytoplasm
Q3SWH1	Y102_NITWN		cytoplasm
Q3SWM9	DUT_NITWN		extracellular
Q3SWN3	FPG_NITWN		cytoplasm
Q3SWP6	RS15_NITWN		cytoplasm
Q3SWP7	TRUB_NITWN		cytoplasm
Q3SWQ2	Y021_NITWN		inner membrane
Q3SWQ3	TRMB_NITWN		cytoplasm
Q3V7F6	PDXJ_LEGPL	cytoplasm (probable).	cytoplasm
Q3V7G1	PDXJ_LEGPA	cytoplasm (probable).	cytoplasm
Q3V7G6	PDXJ_ACIAD	cytoplasm (probable).	cytoplasm
Q3V7I6	PDXJ_PHOPR	cytoplasm (probable).	cytoplasm
Q3V7J3	PDXJ_SILPO	cytoplasm (probable).	cytoplasm
Q3V7J9	PDXJ_ANAMM	cytoplasm (probable).	cytoplasm
Q3V7K1	PDXJ_SALPA	cytoplasm (probable).	cytoplasm
Q3V7M2	PDXJ_BURMA	cytoplasm (probable).	cytoplasm
Q3V7N3	PDXJ_ERWCT	cytoplasm (probable).	cytoplasm
Q3V7P0	PDXJ_YERPS	cytoplasm (probable).	cytoplasm
Q3V7R4	PDXJ_BDEBA	cytoplasm (probable).	cytoplasm
Q3V7S0	PDXJ_RHOPA	cytoplasm (probable).	cytoplasm
Q3V7S9	PDXJ_BURPS	cytoplasm (probable).	cytoplasm
Q3V870	PDXJ_LEGPH	cytoplasm (probable).	cytoplasm
Q3V891	PDXJ_DESVH	cytoplasm (probable).	cytoplasm
Q3V8A2	PDXJ_METCA	cytoplasm (probable).	cytoplasm
Q3V8B3	PDXJ_WOLPM	cytoplasm (probable).	cytoplasm
Q3V8C9	PDXJ_GEOSL	cytoplasm (probable).	cytoplasm
Q3YQP2	RL19_EHRCJ		cytoplasm
Q3YQT1	NDK_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YQV9	FABZ_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YQX7	RNC_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YR47	SECB_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YRC8	RS6_EHRCJ		cytoplasm
Q3YRD1	GLYA_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YRD9	RUVB_EHRCJ		cytoplasm
Q3YRE0	RUVA_EHRCJ		cytoplasm
Q3YRK8	RS10_EHRCJ		cytoplasm
Q3YRL2	RL2_EHRCJ		cytoplasm
Q3YRL5	RS3_EHRCJ		cytoplasm
Q3YRL6	RL16_EHRCJ		cytoplasm
Q3YRL8	RS17_EHRCJ		cytoplasm
Q3YRM6	RS5_EHRCJ		cytoplasm
Q3YRN0	RS13_EHRCJ		cytoplasm

Q3YRN2	RPOA_EHRCJ		cytoplasm
Q3YRP5	RL32_EHRCJ		cytoplasm
Q3YRQ6	PURA_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YRR6	DNAK_EHRCJ		cytoplasm
Q3YRU2	DUT_EHRCJ		extracellular
Q3YRU6	GPMI_EHRCJ		cytoplasm
Q3YRW8	RRMJ_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YRX3	SYI_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YRX7	MRAW_EHRCJ		cytoplasm
Q3YSC2	LEPA_EHRCJ		cytoplasm
Q3YSC5	RS15_EHRCJ		cytoplasm
Q3YSC6	TRUB_EHRCJ		cytoplasm
Q3YSD5	ACPS_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YSE4	ISPE_EHRCJ		cytoplasm
Q3YSH2	CCME_EHRCJ	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3YSH3	SURE_EHRCJ	cytoplasm (potential).	periplasm
Q3YSH8	RPOZ_EHRCJ		cytoplasm
Q3YSI5	PDXJ_EHRCJ	cytoplasm (probable).	cytoplasm
Q3YSJ8	MUTS_EHRCJ		cytoplasm
Q3YSL9	HTPG_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YSP8	Y207_EHRCJ		cytoplasm
Q3YSQ3	CLPP_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YSS1	PDXH_EHRCJ		cytoplasm
Q3YST1	RNH2_EHRCJ	cytoplasm (potential).	extracellular
Q3YST4	RPOC_EHRCJ		cytoplasm
Q3YST5	RPOB_EHRCJ		cytoplasm
Q3YSU4	RS7_EHRCJ		cytoplasm
Q3YSX9	NRDR_EHRCJ		cytoplasm
Q3YT02	ISPF_EHRCJ		extracellular
Q3YT16	SYV_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YT21	ATPA_EHRCJ	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3YT30	GUAA_EHRCJ		cytoplasm
Q3YT49	SYI_EHRCJ	cytoplasm (by similarity).	cytoplasm
Q3YT63	RS20_EHRCJ		cytoplasm
Q3YT96	RUVC_EHRCJ		cytoplasm
Q3YT97	Y007_EHRCJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3YU09	DEOD_SHISS		cytoplasm
Q3YU11	TYPH_SHISS		cytoplasm
Q3YU31	DNAT_SHISS		nucleoid
Q3YU59	UXUA_SHISS		cytoplasm
Q3YU91	SYV_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YUE4	RL9_SHISS		cytoplasm
Q3YUE7	RS6_SHISS		cytoplasm
Q3YUF0	ULAF_SHISS		cytoplasm
Q3YUF1	ULAE_SHISS		cytoplasm
Q3YUF2	ULAD_SHISS		cytoplasm
Q3YUF3	ULAC_SHISS	cytoplasm (probable).	cytoplasm
Q3YUF4	ULAB_SHISS	cytoplasm (potential).	cytoplasm
Q3YUF5	ULAG_SHISS	cytoplasm (potential).	extracellular

Q3YUF6	ULAR_SHISS	cytoplasm (potential).	cytoplasm
Q3YUG6	PURA_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YUQ0	YJDP_SHISS		nucleoid
Q3YUU9	LAMB_SHISS	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q3YUW0	G6PI_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YUY8	NUDC_SHISS		cytoplasm
Q3YUZ4	THIG_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YUZ6	RPOC_SHISS		cytoplasm
Q3YUZ7	RPOB_SHISS		cytoplasm
Q3YUZ9	RL10_SHISS		cytoplasm
Q3YV00	RL1_SHISS		cytoplasm
Q3YV08	MURB_SHISS	cytoplasm (probable).	inner membrane
Q3YVC1	FADB_SHISS		cytoplasm
Q3YVC2	FADA_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YVE9	CORA_SHISS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3YVI5	GPPA_SHISS		extracellular
Q3YVJ0	ILVC_SHISS		cytoplasm
Q3YVJ3	ILVD_SHISS		cytoplasm
Q3YVN4	GLMU_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YVN8	ATPA_SHISS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3YVP9	VIAA_SHISS		nucleoid
Q3YVQ0	RAVA_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YVV0	XYLA_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YVX0	LLDD_SHISS		cytoplasm
Q3YVX7	GPMI_SHISS		cytoplasm
Q3YVZ7	FPG_SHISS		cytoplasm
Q3YW02	DUT_SHISS		extracellular
Q3YW12	RPOZ_SHISS		cytoplasm
Q3YW58	TUSA_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YW96	GLGA_SHISS		cytoplasm
Q3YWB4	RECF_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YWC7	YIDE_SHISS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q3YWM2	PPCK_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YWM3	HSLO_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YWN3	AROK_SHISS	cytoplasm (probable).	cytoplasm
Q3YWN4	AROB_SHISS	cytoplasm (probable).	cytoplasm
Q3YWN8	GPH_SHISS		cytoplasm
Q3YWS4	SLYX_SHISS		inner membrane
Q3YWS7	TUSD_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YWS8	TUSC_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YWS9	TUSB_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YWT0	RS12_SHISS		cytoplasm
Q3YWT1	RS7_SHISS		cytoplasm
Q3YWT8	RS10_SHISS		cytoplasm
Q3YWU2	RL2_SHISS		cytoplasm
Q3YWU5	RS3_SHISS		cytoplasm
Q3YWU6	RL16_SHISS		cytoplasm
Q3YWU8	RS17_SHISS		cytoplasm

Q3YWV3	RS8_SHISS		cytoplasm
Q3YWV6	RS5_SHISS		cytoplasm
Q3YWW1	RS13_SHISS		cytoplasm
Q3YWW2	RS11_SHISS		cytoplasm
Q3YWW3	RS4_SHISS		cytoplasm
Q3YWW4	RPOA_SHISS		cytoplasm
Q3YWW9	MSCL_SHISS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3YX52	MURA_SHISS	cytoplasm (probable).	inner membrane
Q3YX65	GLMM_SHISS		cytoplasm
Q3YX71	YHBC_SHISS		inner membrane
Q3YX75	TRUB_SHISS		cytoplasm
Q3YX76	RS15_SHISS		cytoplasm
Q3YXE7	TRMB_SHISS		cytoplasm
Q3YXH4	MUG_SHISS	cytoplasm (potential).	extracellular
Q3YXI3	YGIH_SHISS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3YXI5	UPPP_SHISS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3YXY0	IDI_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YY30	NUDH_SHISS		cytoplasm
Q3YY51	MTFA_SHISS		cytoplasm
Q3YY71	RUMA_SHISS		cytoplasm
Q3YYB5	ISPD_SHISS		cytoplasm
Q3YYB6	ISPF_SHISS		extracellular
Q3YYB7	TRUD_SHISS		cytoplasm
Q3YYB8	SURE_SHISS	cytoplasm (potential).	periplasm
Q3YYC9	MUTS_SHISS		cytoplasm
Q3YYM4	PPNK_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YYN2	RL19_SHISS		cytoplasm
Q3YYU7	LEPA_SHISS		cytoplasm
Q3YYU9	RNC_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YYV1	RECO_SHISS		cytoplasm
Q3YYV2	PDXJ_SHISS	cytoplasm (probable).	cytoplasm
Q3YYV3	ACPS_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YZ04	GLYA_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YZ29	PEPB_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YZ34	NDK_SHISS	cytoplasm (by similarity).	cytoplasm
Q3YZ45	GUAA_SHISS		cytoplasm
Q3YZ89	TAL2_SHISS	cytoplasm (probable).	cytoplasm
Q3YZD0	ZIPA_SHISS	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q3YZN3	MEPA_SHISS	periplasm (by similarity).	extracellular
Q3YZZ8	ECOT_SHISS	periplasm (by similarity).	periplasm
Q3Z001	NAPA_SHISS	periplasm (by similarity).	periplasm
Q3Z010	CCME_SHISS	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q3Z044	RIHB_SHISS		extracellular
Q3Z062	CDD_SHISS		cytoplasm
Q3Z0G0	HIS6_SHISS	cytoplasm (by similarity).	cytoplasm

Q3Z0G1	HIS4_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z0G2	HIS5_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z0G4	HIS8_SHISS		cytoplasm
Q3Z0G5	HISX_SHISS		cytoplasm
Q3Z0S6	ISPE_SHISS		cytoplasm
Q3Z192	ZNTB_SHISS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3Z1P4	Y1623_SHISS	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q3Z1W3	TUS_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z1Z3	SYX_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z1Z4	PDXH_SHISS		cytoplasm
Q3Z250	YDIA_SHISS		cytoplasm
Q3Z2F2	YOA_H_SHISS		inner membrane
Q3Z2J8	YEBF_SHISS	secreted protein (by similarity).	inner membrane
Q3Z2L8	RUVB_SHISS		cytoplasm
Q3Z2L9	RUVA_SHISS		cytoplasm
Q3Z2M1	RUVC_SHISS		cytoplasm
Q3Z2T6	PGSA_SHISS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3Z310	NAGZ_SHISS	cytoplasm (by similarity).	extracellular
Q3Z328	RL32_SHISS		cytoplasm
Q3Z337	FLGI_SHISS	periplasm (by similarity).	periplasm
Q3Z338	FLGH_SHISS	outer membrane; lipid-anchor (probable).	inner membrane
Q3Z360	YCEI_SHISS	periplasm (potential).	periplasm
Q3Z392	YCDX_SHISS		cytoplasm
Q3Z3F2	TUSE_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z3K6	LPXK_SHISS		cytoplasm
Q3Z3R7	YBJQ_SHISS		inner membrane
Q3Z3T2	RIMK_SHISS		cytoplasm
Q3Z3T7	YBJL_SHISS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q3Z405	UVRB_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z455	GPMA_SHISS		cytoplasm
Q3Z4E1	RIHA_SHISS		extracellular
Q3Z4K2	ENTS_SHISS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3Z4Q0	SELU_SHISS		cytoplasm
Q3Z4S6	HTPG_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z4W6	CLPP_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z4Z6	NRDR_SHISS		cytoplasm
Q3Z504	QUEA_SHISS	cytoplasm (potential).	cytoplasm
Q3Z505	ACPH_SHISS		cytoplasm
Q3Z522	AROL_SHISS	cytoplasm (probable).	cytoplasm
Q3Z594	PROA_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z595	PROB_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z599	XGPT_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z5H3	ACCA_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z5H5	RNH2_SHISS	cytoplasm (potential).	extracellular

Q3Z5H8	FABZ_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z5I0	SKP_SHISS	periplasm (by similarity).	periplasm
Q3Z5I1	YAET_SHISS	bacterial cell outer membrane (by similarity).	outer membrane
Q3Z5J1	GLND_SHISS		cytoplasm
Q3Z5M9	PAND_SHISS		cytoplasm
Q3Z5R9	MURG_SHISS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q3Z5S2	MRAY_SHISS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q3Z5S7	MRAW_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z5S8	MRAZ_SHISS		inner membrane
Q3Z5T7	LEU3_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z5T8	LEU2_SHISS		cytoplasm
Q3Z5T9	LEUD_SHISS		cytoplasm
Q3Z5X9	DAPB_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z5Y0	RIHC_SHISS		extracellular
Q3Z5Y4	SYI_SHISS	cytoplasm (by similarity).	cytoplasm
Q3Z5Y7	RS20_SHISS		cytoplasm
Q3Z601	DNAK_SHISS		cytoplasm
Q3Z606	TAL1_SHISS	cytoplasm (probable).	cytoplasm
Q43895	YNIU_AZOBR		inner membrane
Q43908	DDC_ACIBA		cytoplasm
Q43909	NIFU_AZOBR		cytoplasm
Q43914	DLHH_AZOBR		extracellular
Q43922	QUIC_ACIAD		cytoplasm
Q43923	QUIX_ACIAD		outer membrane
Q43925	HIS3_AZOCH	cytoplasm (by similarity).	cytoplasm
Q43926	HIS2_AZOCH	cytoplasm (by similarity).	cytoplasm
Q43931	CATB_ACIAD		cytoplasm
Q43932	CATC_ACIAD		cytoplasm
Q43935	CATF_ACIAD		cytoplasm
Q43948	HYPB_AZOCH		cytoplasm
Q43949	HYPB_AZOCH		cytoplasm
Q43950	HYPF_AZOCH		cytoplasm
Q43951	HYPC_AZOCH		cytoplasm
Q43954	HUPM_AZOCH		extracellular
Q43955	HUPN_AZOCH		cytoplasm
Q43957	HUPQ_AZOCH		cytoplasm
Q43958	HUPT_AZOCH		cytoplasm
Q43959	HUPV_AZOCH		cytoplasm
Q43963	TRMD_ACIAD	cytoplasm (potential).	cytoplasm
Q43966	NODZ_AZOCA		cytoplasm
Q43967	NOEC_AZOCA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q43971	HFQ_AZOCA		cytoplasm
Q43973	PCAI_ACIAD		cytoplasm
Q43974	PCAF_ACIAD		cytoplasm
Q43975	PCAK_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q43984	CATA_ACIGB		cytoplasm
Q43989	COBQ_ACIAD		cytoplasm

Q43992	POBR_ACIAD		cytoplasm
Q44002	DHET_ACEEU	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	periplasm
Q44004	CYSM_RALME		cytoplasm
Q44006	CZCR_RALME	cytoplasm (probable).	cytoplasm
Q44007	CZCS_RALME	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q44009	CZCI_RALME		cytoplasm
Q44015	YGB4_RALEU		cytoplasm
Q44018	YGB7_RALEU	periplasm (potential).	periplasm
Q44020	YGB2_RALEU		inner membrane
Q44022	LEUD_RALEU		cytoplasm
Q44044	NIFH_ALCFA		cytoplasm
Q44045	NIFD_ALCFA		periplasm
Q44056	BLA1_AERHY		extracellular
Q44058	AHYI_AERHY		cytoplasm
Q44066	HLYA_AERHY		extracellular
Q44069	LEXA_AERHY		cytoplasm
Q44123	FBPB_ACTPL	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q44184	HYDA_AGRTU		extracellular
Q44185	DCAS_AGRTU		cytoplasm
Q44256	CBAA_COMTE		cytoplasm
Q44257	CBAB_COMTE		cytoplasm
Q44258	CBAC_COMTE		cytoplasm
Q44261	CRTW_PARS1		inner membrane
Q44262	CRTZ_PARS1		cytoplasm
Q44264	HSLR_AERSA		cytoplasm
Q44287	LAMB_AERSA	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q44333	PUTR_AGRTU		cytoplasm
Q44335	FLGB_AGRT5		flagellar
Q44336	FLGC_AGRT5		flagellar
Q44337	FLIE_AGRT5		flagellar
Q44338	FLGG_AGRT5		flagellar
Q44339	FLGA_AGRT5	periplasm (probable).	flagellar
Q44340	FLGI_AGRT5	periplasm (by similarity).	periplasm
Q44342	FLGH_AGRT5	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q44344	FLIP_AGRT5	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q44346	TRAG_AGRT5	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q44347	TRAD_AGRT5		cytoplasm
Q44348	TRAC_AGRT5		cytoplasm
Q44349	TRAA_AGRT5		cytoplasm
Q44350	TRAF_AGRT5	periplasm (probable).	periplasm
Q44351	TRAB_AGRT5	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q44360	TRAG_AGRTU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q44361	TRAD_AGRTU		cytoplasm
Q44362	TRAC_AGRTU		cytoplasm
Q44363	TRAA_AGRTU		cytoplasm
Q44364	TRAF_AGRTU	periplasm (probable).	periplasm
Q44433	YPI2_AGRTU		cytoplasm
Q44452	TRAM_AGRT5		outer membrane
Q44458	FLIG_AGRT5	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q44470	TUB4_AGRVI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q44471	TTUC1_AGRVI	cytoplasm (by similarity).	cytoplasm
Q44472	TTUD4_AGRVI		cytoplasm
Q44473	KPYK2_AGRVI		cytoplasm
Q44504	DSBA_AZOVI	periplasm (by similarity).	periplasm
Q44524	DHNO_AGRVI		cytoplasm
Q44532	FENR_AZOVI		cytoplasm
Q44538	MODC2_AZOVI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q44540	YNIU_AZOVI		inner membrane
Q44581	NCCY_ALCXX		cytoplasm
Q44582	NCCX_ALCXX		cytoplasm
Q44583	NCCH_ALCXX		cytoplasm
Q44584	NCCC_ALCXX		inner membrane
Q44585	NCCB_ALCXX		inner membrane
Q44586	NCCA_ALCXX	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q44587	NCCN_ALCXX	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q44598	TRPE_BUCSC		cytoplasm
Q44601	YTR1_BUCSC	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q44602	TRPD_BUCSC		cytoplasm
Q44603	TRPC_BUCSC		cytoplasm
Q44604	TRPA_BUCSC		cytoplasm
Q44607	AROE_BUCAI		cytoplasm
Q44613	LOLD_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q44633	TRME_BUCAP		cytoplasm
Q44634	Y020_BUCAP		cytoplasm
Q44653	RS1_BUCAP		cytoplasm
Q44654	IHFB_BUCAP		cytoplasm
Q44685	TRPB_BUCAI		cytoplasm
Q44687	TRPB_BUCMH		cytoplasm
Q44689	TRPE_BUCRM		cytoplasm
Q44691	TRPE_BUCRP		cytoplasm
Q44695	TRPE_BUCAI		cytoplasm

Q44696	TRPG_BUCAI		cytoplasm
Q44697	TRPE_BUCDN		cytoplasm
Q44730	FLHD_BORBR	cytoplasm (by similarity).	cytoplasm
Q44740	ALCA_BORBR		cytoplasm
Q44879	CTPA_BARBA	secreted protein (potential).	cytoplasm
Q45072	TFTE_BURCE		cytoplasm
Q45218	FE45_BRAJA		cytoplasm
Q45219	Y2146_BRAJA		cytoplasm
Q45220	ISPA_BRAJA	cytoplasm (potential).	cytoplasm
Q45221	Y2149_BRAJA		cytoplasm
Q45222	Y2150_BRAJA		cytoplasm
Q45223	HBD_BRAJA		cytoplasm
Q45247	HUPN_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q45251	HUPD_BRAJA		extracellular
Q45252	HUPF_BRAJA		cytoplasm
Q45256	HYP A2_BRAJA		cytoplasm
Q45257	HYPB_BRAJA		cytoplasm
Q45263	NOD21_BRAJA		cytoplasm
Q45264	NOD11_BRAJA		cytoplasm
Q45269	NOLO_BRAJA		inner membrane
Q45271	NODZ_BRAJA		cytoplasm
Q45338	BAF_BORPE		cytoplasm
Q45388	TEX_BORPE		cytoplasm
Q45389	BUGT_BORPE	periplasm (potential).	periplasm
Q45407	EPSA2_RALSO	bacterial cell outer membrane; multi-pass membrane protein (potential).	outer membrane
Q45408	EPSP2_RALSO		extracellular
Q45410	EPSD2_RALSO		cytoplasm
Q45411	EPSE2_RALSO	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q45412	EPSF2_RALSO	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q45423	TFDA_BURSR		cytoplasm
Q45918	PYRE_COXBU		cytoplasm
Q45920	DUT_COXBU		extracellular
Q45968	TRA1_COXBU		cytoplasm
Q45971	MTC1_CAUCR		cytoplasm
Q45972	FER1_CAUCR		periplasm
Q45973	Y1074_CAUCR		cytoplasm
Q45974	FLIQ_CAUCR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q45975	FLIR_CAUCR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q45977	HFAD_CAUCR	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
Q45978	HFAC_CAUCR		inner membrane

Q45979	Y952_CAUCR	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q45980	FLIP_CAUCR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q45982	PTMB_CAMCO		inner membrane
Q45983	PTMA_CAMCO		cytoplasm
Q45994	CTRA_CAUCR		cytoplasm
Q46087	TRAT_CHEHE		cytoplasm
Q46097	DSBI_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q46100	CDTA_CAMJE	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q46104	ARLY_CAMJE	cytoplasm (probable).	cytoplasm
Q46105	CBF2_CAMJE		cytoplasm
Q46108	TIG_CAMJE		cytoplasm
Q46113	FLA3_CAMJE		flagellar
Q46114	FLB3_CAMJE		flagellar
Q46121	DBH_CAMJE		cytoplasm
Q46122	MAPA_CAMJE	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q46124	RPOB_CAMJE		cytoplasm
Q46125	HISJ_CAMJE	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q46372	BPHA_COMTE		cytoplasm
Q46373	BPHE_COMTE		cytoplasm
Q46381	BPHB_COMTE		cytoplasm
Q46444	QHED_COMTE	periplasm (potential).	periplasm
Q46463	FUR_CAMUP	cytoplasm (by similarity).	cytoplasm
Q46476	DSBE_CHRVI	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q46481	LPXA_CHRVI	cytoplasm (by similarity).	cytoplasm
Q46495	DESR_DESBR		inner membrane
Q46509	MOP_DESGI		cytoplasm
Q46525	LEP4_BACNO	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q46526	COAE_BACNO	cytoplasm (by similarity).	cytoplasm
Q46550	AROA_BACNO	cytoplasm (probable).	cytoplasm
Q46558	VAPB_BACNO		extracellular
Q46560	VAPI_BACNO		extracellular
Q46561	VAPZ_BACNO		extracellular
Q46565	VAPD_BACNO		extracellular
Q46582	DSVD_DESVH		cytoplasm
Q46618	HRPA_ERWAM	extracellular and secreted via type iii secretion system (by similarity).	extracellular
Q46629	AMSH_ERWAM	bacterial cell outer membrane; multi-pass membrane protein (potential).	outer membrane
Q46630	AMSI_ERWAM		extracellular

Q46631	AMSA_ERWAM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q46632	AMSB_ERWAM		cytoplasm
Q46633	AMSC_ERWAM	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q46634	AMSD_ERWAM		cytoplasm
Q46635	AMSE_ERWAM		cytoplasm
Q46636	AMSF_ERWAM	periplasm (potential).	extracellular
Q46637	AMSJ_ERWAM		cytoplasm
Q46638	AMSK_ERWAM		cytoplasm
Q46646	HRCR_ERWAM	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q46668	CDTA_ECOLI	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q46670	CDTC_ECOLI	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q46717	HLYB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q46751	CARR_PECCE		cytoplasm
Q46769	RFBD2_ECOLI		cytoplasm
Q46786	YGEF_ECOLI		cytoplasm
Q46787	YGEG_ECOLI		cytoplasm
Q46789	YGEI_ECOLI		extracellular
Q46790	PBL_ECOLI	periplasm (potential).	periplasm
Q46791	YGEK_ECOLI		cytoplasm
Q46796	YGEP_ECOLI		cytoplasm
Q46797	YGEQ_ECOLI		cytoplasm
Q46798	YGER_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q46799	XDHA_ECOLI		periplasm
Q46800	XDHB_ECOLI		cytoplasm
Q46801	XDHC_ECOLI		cytoplasm
Q46802	YGEV_ECOLI		cytoplasm
Q46803	YGEW_ECOLI		cytoplasm
Q46806	YGEZ_ECOLI		extracellular
Q46807	ARCL_ECOLI		cytoplasm
Q46808	YQEB_ECOLI		cytoplasm
Q46809	YQEC_ECOLI		cytoplasm
Q46810	YGFJ_ECOLI		cytoplasm
Q46811	YGFK_ECOLI		cytoplasm
Q46812	SSNA_ECOLI		extracellular
Q46814	XDHD_ECOLI		cytoplasm
Q46817	YGFQ_ECOLI		inner membrane
Q46819	YGFS_ECOLI		periplasm
Q46820	YGFT_ECOLI		cytoplasm
Q46824	YGFY_ECOLI		inner membrane
Q46829	BGLA_ECOLI		cytoplasm
Q46831	YQGA_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q46832	YGHD_ECOLI	bacterial cell inner membrane (probable).	inner membrane

Q46833	YGHE_ECOLI	bacterial cell inner membrane (probable).	inner membrane
Q46834	YGHF_ECOLI	periplasm (potential).	inner membrane
Q46835	YGHG_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q46837	ACFD_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q46843	YGHS_ECOLI		inner membrane
Q46844	YGHT_ECOLI		cytoplasm
Q46845	YGHU_ECOLI		cytoplasm
Q46851	YGHZ_ECOLI		inner membrane
Q46855	YQHC_ECOLI		cytoplasm
Q46856	YQHD_ECOLI		cytoplasm
Q46857	DKGA_ECOLI	cytoplasm (by similarity).	cytoplasm
Q46858	YQHG_ECOLI		cytoplasm
Q46861	YGIQ_ECOLI		cytoplasm
Q46863	YGIS_ECOLI	periplasm (potential).	periplasm
Q46864	YGIT_ECOLI		cytoplasm
Q46865	YGIU_ECOLI		cytoplasm
Q46866	YGIV_ECOLI		cytoplasm
Q46868	YQIC_ECOLI		cytoplasm
Q46871	YQJH_ECOLI	cytoplasm (potential).	cytoplasm
Q46888	YGBJ_ECOLI		cytoplasm
Q46889	YGBK_ECOLI		cytoplasm
Q46890	YGBL_ECOLI		cytoplasm
Q46891	YGBM_ECOLI		extracellular
Q46893	ISPD_ECOLI		cytoplasm
Q46896	YGBT_ECOLI		cytoplasm
Q46897	YGCH_ECOLI		cytoplasm
Q46898	YGCI_ECOLI		cytoplasm
Q46899	YGCJ_ECOLI		cytoplasm
Q46901	YGCL_ECOLI		cytoplasm
Q46904	YGCN_ECOLI		cytoplasm
Q46905	YGCO_ECOLI		cytoplasm
Q46906	YGCP_ECOLI		cytoplasm
Q46907	YGCQ_ECOLI		cytoplasm
Q46908	YGCR_ECOLI		cytoplasm
Q46911	YGCU_ECOLI		cytoplasm
Q46915	GUDX_ECOLI		cytoplasm
Q46919	YQCC_ECOLI		cytoplasm
Q46920	QUEF_ECOLI	cytoplasm (probable).	cytoplasm
Q46925	CSDA_ECOLI		cytoplasm
Q46927	YGDL_ECOLI		inner membrane
Q46938	KDUI_ECOLI		cytoplasm
Q46939	YQEF_ECOLI	cytoplasm (potential).	cytoplasm
Q46941	YQEH_ECOLI		cytoplasm
Q46942	YQEI_ECOLI		cytoplasm
Q46943	YQEJ_ECOLI		inner membrane
Q46948	THIJ_ECOLI		cytoplasm
Q46953	YPJF_ECOLI		inner membrane
Q46967	ECHR_ERWCH		cytoplasm
Q46968	ECHI_ERWCH		cytoplasm
Q46970	IM24_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q46971	MTFS_ECOLI		extracellular

Q46978	HPAG_ECOLI		cytoplasm
Q46995	CCDA2_ECOLI		cytoplasm
Q46996	CCDB2_ECOLI		cytoplasm
Q46999	TRAK2_ECOLI		outer membrane
Q46N53	HISX2_RALEJ		cytoplasm
Q46PF7	FLGI_RALEJ	periplasm (by similarity).	periplasm
Q46PF8	FLGH_RALEJ	outer membrane; lipid-anchor (probable).	inner membrane
Q46QJ0	SLYX_RALEJ		inner membrane
Q46QQ8	PURA2_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46RR4	GLYA2_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46RW7	CCME_RALEJ	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q46RX3	NAPA_RALEJ	periplasm (by similarity).	periplasm
Q46UA4	HGD_RALEJ		cytoplasm
Q46VT8	PHS2_RALEJ		cytoplasm
Q46VX8	ATPA_RALEJ	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q46VX9	ATPG_RALEJ		inner membrane
Q46VZ5	GCSP_RALEJ		cytoplasm
Q46WD1	RL1_RALEJ		cytoplasm
Q46WD2	RL10_RALEJ		cytoplasm
Q46WD4	RPOB_RALEJ		cytoplasm
Q46WD5	RPOC_RALEJ		cytoplasm
Q46WD8	RS12_RALEJ		cytoplasm
Q46WD9	RS7_RALEJ		cytoplasm
Q46WE2	RS10_RALEJ		cytoplasm
Q46WE7	RL2_RALEJ		cytoplasm
Q46WF0	RS3_RALEJ		cytoplasm
Q46WF1	RL16_RALEJ		cytoplasm
Q46WF3	RS17_RALEJ		cytoplasm
Q46WF8	RS8_RALEJ		cytoplasm
Q46WG1	RS5_RALEJ		cytoplasm
Q46WG3	RL15_RALEJ		cytoplasm
Q46WG6	RS13_RALEJ		cytoplasm
Q46WG7	RS11_RALEJ		cytoplasm
Q46WG8	RS4_RALEJ		cytoplasm
Q46WG9	RPOA_RALEJ		cytoplasm
Q46WJ3	AROK_RALEJ	cytoplasm (probable).	cytoplasm
Q46WJ4	AROB_RALEJ	cytoplasm (probable).	cytoplasm
Q46WL0	MURA_RALEJ	cytoplasm (probable).	inner membrane
Q46WL1	HIS1_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46WL2	HISX1_RALEJ		cytoplasm
Q46WL3	HIS82_RALEJ		cytoplasm
Q46WL6	HIS5_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46WL7	HIS4_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46WL8	HIS6_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46WL9	HIS3_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46WM0	HIS2_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46WM9	MSCL_RALEJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q46WU8	TRPD_RALEJ		cytoplasm

Q46WY5	MRAZ_RALEJ		inner membrane
Q46WY6	MRAW_RALEJ		cytoplasm
Q46WZ1	MRAY_RALEJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q46WZ4	MURG_RALEJ	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q46X04	ARGJ_RALEJ	cytoplasm (probable).	cytoplasm
Q46X18	PROB_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46X20	NUDH_RALEJ		cytoplasm
Q46XD7	DAPB_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46XE1	PROA_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46XG5	QUEA_RALEJ	cytoplasm (potential).	cytoplasm
Q46XI7	DNAK_RALEJ		cytoplasm
Q46XK7	RS20_RALEJ		cytoplasm
Q46XL1	MURB_RALEJ	cytoplasm (probable).	inner membrane
Q46XM7	SYI_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46Y48	HIS81_RALEJ		cytoplasm
Q46Y85	RL19_RALEJ		cytoplasm
Q46YE9	RPOZ_RALEJ		cytoplasm
Q46YI9	ILVD2_RALEJ		cytoplasm
Q46YS9	HTPG_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46YU4	MDH_RALEJ		cytoplasm
Q46YV7	LEU2_RALEJ		cytoplasm
Q46YV9	LEUD_RALEJ		cytoplasm
Q46YW0	LEU3_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46Z03	RL32_RALEJ		cytoplasm
Q46Z15	LEPA_RALEJ		cytoplasm
Q46Z18	RNC_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46Z21	PDXJ_RALEJ	cytoplasm (probable).	cytoplasm
Q46Z22	ACPS_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46Z26	UVRC_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46Z98	RRMJ_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46ZA1	GLMM_RALEJ		cytoplasm
Q46ZH1	SURE_RALEJ	cytoplasm (potential).	periplasm
Q46ZH7	RUMA_RALEJ		cytoplasm
Q46ZH9	NDK_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46ZJ3	PURA1_RALEJ	cytoplasm (by similarity).	cytoplasm
Q46ZK0	TAL_RALEJ	cytoplasm (probable).	cytoplasm
Q46ZN9	Y2030_RALEJ		inner membrane
Q46ZP3	TRUB_RALEJ		cytoplasm
Q46ZR3	RS6_RALEJ		cytoplasm
Q46ZR6	RL9_RALEJ		cytoplasm
Q47005	NAC_ECOLI		cytoplasm
Q47013	ELAD_ECOLI		inner membrane
Q47015	CEST_ECOLI	cytoplasm (by similarity).	cytoplasm
Q47036	INTL_ECOLI		cytoplasm
Q47037	AFAF_ECOLI		cytoplasm
Q47038	AFAD_ECOLI		fimbrium
Q47066	BLT1_ECOLI		extracellular
Q47068	BFPB_ECO27	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q47077	BGAL_ENTCL		cytoplasm
Q47083	CBL_ECOLI		cytoplasm

Q47085	CBRB_DICD3	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q47086	CBRC_DICD3	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q47098	HPAI_ECOLI		cytoplasm
Q470D5	GLND_RALEJ		cytoplasm
Q470E8	FABZ_RALEJ	cytoplasm (by similarity).	cytoplasm
Q470F1	RNH2_RALEJ	cytoplasm (potential).	extracellular
Q470F3	Y1865_RALEJ		cytoplasm
Q470G4	GUAA_RALEJ		cytoplasm
Q47112	CEA7_ECOLI		extracellular
Q47125	CE10_ECOLI		extracellular
Q47129	FEAR_ECOLI		cytoplasm
Q47132	DAAF_ECOLI		cytoplasm
Q47133	DAAA_ECOLI		cytoplasm
Q47140	HCAF_ECOLI		cytoplasm
Q47141	HCAR_ECOLI		cytoplasm
Q47142	HCAT_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q47146	FADE_ECOLI		cytoplasm
Q47147	YAFJ_ECOLI		cytoplasm
Q47149	YAFQ_ECOLI		inner membrane
Q47150	DINJ_ECOLI		cytoplasm
Q47151	YAFL_ECOLI	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q47152	YAFM_ECOLI		cytoplasm
Q47153	FHIA_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q47154	MBHA_ECOLI		outer membrane
Q47155	DPO4_ECOLI	cytoplasm (probable).	cytoplasm
Q47156	YAFN_ECOLI		cytoplasm
Q47157	YAFO_ECOLI		cytoplasm
Q47158	YAFP_ECOLI		cytoplasm
Q47163	T1MP_ECOLI		cytoplasm
Q47187	EXPI_DICD3		extracellular
Q47188	EXPR_DICD3		cytoplasm
Q47189	EXPR_PECCC		cytoplasm
Q471B8	G6PI2_RALEJ	cytoplasm (by similarity).	cytoplasm
Q47208	FDRA_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q47258	HLYB3_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q47269	YBCN_ECOLI		cytoplasm
Q47270	NINE_ECOLI		inner membrane
Q47274	REQ1_ECOLI		cytoplasm
Q47281	T1RE_ECOLI		cytoplasm
Q47282	T1ME_ECOLI		cytoplasm
Q472C7	G6PI1_RALEJ	cytoplasm (by similarity).	cytoplasm
Q472D3	CLPP_RALEJ	cytoplasm (by similarity).	cytoplasm
Q472F1	ISPF_RALEJ		extracellular

Q472F2	ISPD_RALEJ		cytoplasm
Q47302	INSB2_ECOLI		cytoplasm
Q47309	YI9A_ECOLI		inner membrane
Q47316	IUCA_ECOLI		outer membrane
Q47317	IUCB_ECOLI		cytoplasm
Q47318	IUCC_ECOLI		outer membrane
Q47319	YFIP_ECOLI		cytoplasm
Q47329	UDG5_ECOLI		cytoplasm
Q47334	KSF5_ECOLI		cytoplasm
Q473D1	ACCA_RALEJ	cytoplasm (by similarity).	cytoplasm
Q473E4	MUTS_RALEJ		cytoplasm
Q473K3	UVRB_RALEJ	cytoplasm (by similarity).	cytoplasm
Q473L9	PPNK_RALEJ	cytoplasm (by similarity).	cytoplasm
Q473Q8	UREE_RALEJ	cytoplasm (by similarity).	cytoplasm
Q473Q9	URE1_RALEJ	cytoplasm (by similarity).	cytoplasm
Q473R0	URE2_RALEJ	cytoplasm (by similarity).	cytoplasm
Q473R2	URE3_RALEJ	cytoplasm (by similarity).	cytoplasm
Q473U8	RS15_RALEJ		cytoplasm
Q473V5	ILVC_RALEJ		cytoplasm
Q47400	NEUC_ECOLI		inner membrane
Q47417	TRUC_PECCC		cytoplasm
Q47418	YQCA_PECCC		cytoplasm
Q47422	GSPM_ERWCH	bacterial cell inner membrane (probable).	inner membrane
Q47427	TFAB_ECOLI		cytoplasm
Q47452	PCOA_ECOLI	periplasm (potential).	periplasm
Q47453	PCOB_ECOLI	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
Q47454	PCOC_ECOLI	periplasm (by similarity).	periplasm
Q47455	PCOD_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q47456	PCOR_ECOLI	cytoplasm (probable).	cytoplasm
Q47457	PCOS_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q474E1	SYV_RALEJ	cytoplasm (by similarity).	cytoplasm
Q474I9	PDXH_RALEJ		cytoplasm
Q474L3	GLYA1_RALEJ	cytoplasm (by similarity).	cytoplasm
Q474L4	NRDR_RALEJ		cytoplasm
Q474P5	TRMB_RALEJ		cytoplasm
Q474P6	UPPP_RALEJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q47500	CE05_ECOLI		extracellular
Q47502	CEAK_ECOLI		extracellular
Q47503	IMMK_ECOLI		inner membrane
Q47511	MCCF_ECOLI		extracellular
Q47534	YAIO_ECOLI		outer membrane
Q47536	YAIP_ECOLI		inner membrane
Q47537	TAUA_ECOLI	periplasm (potential).	periplasm
Q47538	TAUB_ECOLI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q47587	RDGA_PECCC		cytoplasm

Q47588	RDGB_PECCE		cytoplasm
Q475B2	ILVD1_RALEJ		cytoplasm
Q475F8	LPXK_RALEJ		cytoplasm
Q475H6	Y575_RALEJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q475R5	RUVC_RALEJ		cytoplasm
Q475R7	RUVA_RALEJ		cytoplasm
Q475R8	RUVB_RALEJ		cytoplasm
Q475S5	SYX_RALEJ	cytoplasm (by similarity).	cytoplasm
Q475Y4	PHS1_RALEJ		cytoplasm
Q47620	CSRA_PECCE		cytoplasm
Q47622	SAPA_ECOLI	periplasm (probable).	periplasm
Q47679	YAFV_ECOLI		cytoplasm
Q47684	YAFW_ECOLI		cytoplasm
Q47685	YKFG_ECOLI		cytoplasm
Q47688	YKFC_ECOLI		cytoplasm
Q47689	MMUP_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q47690	MMUM_ECOLI		cytoplasm
Q476F5	FPG_RALEJ		cytoplasm
Q476F8	ISPE_RALEJ		cytoplasm
Q476J4	SECB_RALEJ	cytoplasm (by similarity).	cytoplasm
Q476J7	GPMA_RALEJ		cytoplasm
Q476K0	SELU_RALEJ		cytoplasm
Q476S2	GLMU_RALEJ	cytoplasm (by similarity).	cytoplasm
Q476U3	THIG_RALEJ	cytoplasm (by similarity).	cytoplasm
Q476W5	METX_RALEJ	cytoplasm (by similarity).	cytoplasm
Q47702	YFEK_ECOLI		cytoplasm
Q47706	UIDC_ECOLI		outer membrane
Q47710	YQJK_ECOLI		cytoplasm
Q47718	INSM_ECOLI		cytoplasm
Q47719	Y6C6_ECOLI		cytoplasm
Q477Z2	ATPG_DECAR		inner membrane
Q477Z3	ATPA_DECAR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q47829	BIOF_ENTAG		cytoplasm
Q47851	IPT_PANAY		cytoplasm
Q47856	HRCR_PANAY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q47860	HYIN_PANAY		extracellular
Q47861	TR2M_PANAY		cytoplasm
Q47862	BIOB_ESCVU		cytoplasm
Q47877	YHLA_EDWTA		inner membrane
Q478E2	RUVC_DECAR		cytoplasm
Q478E3	RUVA_DECAR		cytoplasm
Q478E5	RUVB_DECAR		cytoplasm
Q478K9	CCME_DECAR	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane

Q478U2	UPPP_DECAR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q478Z2	THIG_DECAR	cytoplasm (by similarity).	cytoplasm
Q478Z3	TRMB_DECAR		cytoplasm
Q47908	MSBA_FRANO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q47909	LPXK_FRANO		cytoplasm
Q47945	CYCA_GLUOX	bacterial cell membrane; peripheral membrane protein; periplasmic side (potential).	periplasm
Q47953	FTPA_HAEDU		cytoplasm
Q47956	ISPF_HAEDU		extracellular
Q479A9	RPOZ_DECAR		cytoplasm
Q479H7	LEU32_DECAR	cytoplasm (by similarity).	cytoplasm
Q479M3	ISPE_DECAR		cytoplasm
Q479M6	FPG_DECAR		cytoplasm
Q47A87	NAPA_DECAR	periplasm (by similarity).	periplasm
Q47A95	MRAZ_DECAR		inner membrane
Q47A96	MRAW_DECAR		cytoplasm
Q47AA1	MRAY_DECAR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q47AA4	MURG_DECAR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q47AB5	ARGJ_DECAR	cytoplasm (probable).	cytoplasm
Q47AC6	TRPD_DECAR		cytoplasm
Q47AC9	PROB_DECAR	cytoplasm (by similarity).	cytoplasm
Q47AK9	MURA_DECAR	cytoplasm (probable).	inner membrane
Q47AL4	HIS1_DECAR	cytoplasm (by similarity).	cytoplasm
Q47AL5	HISX_DECAR		cytoplasm
Q47AL9	HIS82_DECAR		cytoplasm
Q47AM1	HIS5_DECAR	cytoplasm (by similarity).	cytoplasm
Q47AM2	HIS4_DECAR	cytoplasm (by similarity).	cytoplasm
Q47AM3	HIS6_DECAR	cytoplasm (by similarity).	cytoplasm
Q47AM4	HIS3_DECAR	cytoplasm (by similarity).	cytoplasm
Q47AM5	HIS2_DECAR	cytoplasm (by similarity).	cytoplasm
Q47AR9	MSCL_DECAR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q47AX4	QUEA_DECAR	cytoplasm (potential).	cytoplasm
Q47B43	LPXK_DECAR		cytoplasm
Q47B64	PAND_DECAR		cytoplasm
Q47B76	SYI_DECAR	cytoplasm (by similarity).	cytoplasm
Q47BB1	DUT_DECAR		extracellular
Q47BG6	SYV_DECAR	cytoplasm (by similarity).	cytoplasm
Q47BH8	ILVC_DECAR		cytoplasm
Q47BI8	RL19_DECAR		cytoplasm
Q47BK5	SYI_DECAR	cytoplasm (by similarity).	cytoplasm
Q47BL6	RS20_DECAR		cytoplasm
Q47BR2	NDK_DECAR	cytoplasm (by similarity).	cytoplasm
Q47BS7	PURA_DECAR	cytoplasm (by similarity).	cytoplasm
Q47C34	MDH_DECAR		cytoplasm
Q47D24	SURE_DECAR	cytoplasm (potential).	periplasm

Q47D81	GCSP_DECAR		cytoplasm
Q47D92	Y2454_DECAR		inner membrane
Q47D96	TRUB_DECAR		cytoplasm
Q47D97	RS15_DECAR		cytoplasm
Q47DJ8	MUTS_DECAR		cytoplasm
Q47DK8	GUAA_DECAR		cytoplasm
Q47EE8	UVRC_DECAR	cytoplasm (by similarity).	cytoplasm
Q47EF3	ACPS_DECAR	cytoplasm (by similarity).	cytoplasm
Q47EF4	PDXJ_DECAR	cytoplasm (probable).	cytoplasm
Q47EF5	RECO_DECAR		cytoplasm
Q47EF7	RNC_DECAR	cytoplasm (by similarity).	cytoplasm
Q47EG0	LEPA_DECAR		cytoplasm
Q47EH3	RL32_DECAR		cytoplasm
Q47EI2	UVRB_DECAR	cytoplasm (by similarity).	cytoplasm
Q47EL1	ISPF_DECAR		extracellular
Q47EL2	ISPD_DECAR		cytoplasm
Q47F76	Y1758_DECAR		cytoplasm
Q47F78	RNH2_DECAR	cytoplasm (potential).	extracellular
Q47F81	FABZ_DECAR	cytoplasm (by similarity).	cytoplasm
Q47F86	DXR_DECAR		cytoplasm
Q47F95	GLND_DECAR		cytoplasm
Q47FB6	CLPP_DECAR	cytoplasm (by similarity).	cytoplasm
Q47G53	URE3_DECAR	cytoplasm (by similarity).	cytoplasm
Q47G54	URE2_DECAR	cytoplasm (by similarity).	cytoplasm
Q47G55	URE1_DECAR	cytoplasm (by similarity).	cytoplasm
Q47G56	UREE_DECAR	cytoplasm (by similarity).	cytoplasm
Q47GP2	HIS81_DECAR		cytoplasm
Q47GQ6	RL9_DECAR		cytoplasm
Q47GQ9	RS6_DECAR		cytoplasm
Q47GZ8	HTPG_DECAR	cytoplasm (by similarity).	cytoplasm
Q47HH9	GLMM_DECAR		cytoplasm
Q47HI2	RRMJ_DECAR	cytoplasm (by similarity).	cytoplasm
Q47HI9	DAPB_DECAR	cytoplasm (by similarity).	cytoplasm
Q47HJ4	PPNK_DECAR	cytoplasm (by similarity).	cytoplasm
Q47HK2	DNAK_DECAR		cytoplasm
Q47HR1	LEU31_DECAR	cytoplasm (by similarity).	cytoplasm
Q47HR2	LEUD_DECAR		cytoplasm
Q47HR4	LEU2_DECAR		cytoplasm
Q47HS4	RUMA_DECAR		cytoplasm
Q47HT1	ACCA_DECAR	cytoplasm (by similarity).	cytoplasm
Q47I19	FLGI_DECAR	periplasm (by similarity).	periplasm
Q47I20	FLGH_DECAR	outer membrane; lipid-anchor (probable).	inner membrane
Q47IC9	NUDH_DECAR		cytoplasm
Q47IG1	SECB_DECAR	cytoplasm (by similarity).	cytoplasm
Q47IH0	NRDR_DECAR		cytoplasm
Q47IH1	GLYA_DECAR	cytoplasm (by similarity).	cytoplasm
Q47IJ3	G6PI_DECAR	cytoplasm (by similarity).	cytoplasm
Q47IJ4	GLGA_DECAR		cytoplasm
Q47IN4	PROA_DECAR	cytoplasm (by similarity).	cytoplasm
Q47IP1	Y533_DECAR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q47IX1	PHS_DECAR		cytoplasm
Q47J77	RPOA_DECAR		cytoplasm
Q47J78	RS4_DECAR		cytoplasm

Q47J79	RS11_DECAR		cytoplasm
Q47J80	RS13_DECAR		cytoplasm
Q47J84	RL15_DECAR		cytoplasm
Q47J86	RS5_DECAR		cytoplasm
Q47J89	RS8_DECAR		cytoplasm
Q47J94	RS17_DECAR		cytoplasm
Q47J96	RL16_DECAR		cytoplasm
Q47J97	RS3_DECAR		cytoplasm
Q47JA0	RL2_DECAR		cytoplasm
Q47JA4	RS10_DECAR		cytoplasm
Q47JA7	RS7_DECAR		cytoplasm
Q47JA8	RS12_DECAR		cytoplasm
Q47JA9	RPOC_DECAR		cytoplasm
Q47JB0	RPOB_DECAR		cytoplasm
Q47JB2	RL10_DECAR		cytoplasm
Q47JB3	RL1_DECAR		cytoplasm
Q47JC0	ILVD_DECAR		cytoplasm
Q47JK2	GLMU_DECAR	cytoplasm (by similarity).	cytoplasm
Q47JK6	AROB_DECAR	cytoplasm (probable).	cytoplasm
Q47JK7	AROK_DECAR	cytoplasm (probable).	cytoplasm
Q47JU2	METX_DECAR	cytoplasm (by similarity).	cytoplasm
Q47K28	MURB_DECAR	cytoplasm (probable).	inner membrane
Q47U38	GIDA_COLP3	cytoplasm (by similarity).	cytoplasm
Q47U39	GIDB_COLP3		cytoplasm
Q47UB2	RPOZ_COLP3		cytoplasm
Q47UE0	GLMU_COLP3	cytoplasm (by similarity).	cytoplasm
Q47UN7	ILVD_COLP3		cytoplasm
Q47UQ0	PROA_COLP3	cytoplasm (by similarity).	cytoplasm
Q47UV6	RL1_COLP3		cytoplasm
Q47UV7	RL10_COLP3		cytoplasm
Q47UV9	RPOB_COLP3		cytoplasm
Q47UW0	RPOC_COLP3		cytoplasm
Q47UW1	RS12_COLP3		cytoplasm
Q47UW2	RS7_COLP3		cytoplasm
Q47UY2	MURB_COLP3	cytoplasm (probable).	inner membrane
Q47V01	SYU_COLP3	cytoplasm (by similarity).	cytoplasm
Q47VD0	PPCK_COLP3	cytoplasm (by similarity).	cytoplasm
Q47VD1	HSLO_COLP3	cytoplasm (by similarity).	cytoplasm
Q47VJ4	MURA_COLP3	cytoplasm (probable).	inner membrane
Q47VL0	MDH_COLP3		cytoplasm
Q47VQ0	MRAZ_COLP3		inner membrane
Q47VQ1	MRAW_COLP3		cytoplasm
Q47VQ6	MRAY_COLP3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q47VQ7	MURD_COLP3	cytoplasm (by similarity).	cytoplasm
Q47VQ9	MURG_COLP3	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q47VY2	GPMI_COLP3		cytoplasm
Q47VY5	SECB_COLP3	cytoplasm (by similarity).	cytoplasm
Q47W31	UPPP_COLP3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q47W34	Y4339_COLP3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q47W59	PANC_COLP3		cytoplasm
Q47W65	GLUQ_COLP3		cytoplasm
Q47WB0	MAO1_COLP3		cytoplasm
Q47WB6	NDK_COLP3	cytoplasm (by similarity).	cytoplasm
Q47WD1	GUAA_COLP3		cytoplasm
Q47WG1	LEUD_COLP3		cytoplasm
Q47WG2	LEU2_COLP3		cytoplasm
Q47WG3	LEU3_COLP3	cytoplasm (by similarity).	cytoplasm
Q47WM2	TRMB_COLP3		cytoplasm
Q47WN0	MUTS_COLP3		cytoplasm
Q47WN6	Y4131_COLP3		cytoplasm
Q47WP3	LEPA_COLP3		cytoplasm
Q47WP5	RNC_COLP3	cytoplasm (by similarity).	cytoplasm
Q47WP8	PDXJ_COLP3	cytoplasm (probable).	cytoplasm
Q47WR3	TAL_COLP3	cytoplasm (probable).	cytoplasm
Q47WV1	RL19_COLP3		cytoplasm
Q47WY2	GLYA4_COLP3	cytoplasm (by similarity).	cytoplasm
Q47XA7	HTPG_COLP3	cytoplasm (by similarity).	cytoplasm
Q47XB3	HIS6_COLP3	cytoplasm (by similarity).	cytoplasm
Q47XB4	HIS4_COLP3	cytoplasm (by similarity).	cytoplasm
Q47XB5	HIS5_COLP3	cytoplasm (by similarity).	cytoplasm
Q47XB7	HIS8_COLP3		cytoplasm
Q47XB8	HISX1_COLP3		cytoplasm
Q47XG2	GCSP2_COLP3		cytoplasm
Q47XG4	GLYA3_COLP3	cytoplasm (by similarity).	cytoplasm
Q47XI3	PPNK_COLP3	cytoplasm (by similarity).	cytoplasm
Q47XI6	DNAK2_COLP3		cytoplasm
Q47XL8	CLPP_COLP3	cytoplasm (by similarity).	cytoplasm
Q47XN8	PHS_COLP3		cytoplasm
Q47Y27	NUDH_COLP3		cytoplasm
Q47Y90	ISPE_COLP3		cytoplasm
Q47YA5	MTFA_COLP3		cytoplasm
Q47YB3	QUEF_COLP3	cytoplasm (probable).	cytoplasm
Q47YC2	TRPD_COLP3		cytoplasm
Q47YI6	DAPB_COLP3	cytoplasm (by similarity).	cytoplasm
Q47YJ3	RRMJ_COLP3	cytoplasm (by similarity).	cytoplasm
Q47YJ7	GLMM_COLP3		cytoplasm
Q47Z18	UVRC_COLP3	cytoplasm (by similarity).	cytoplasm
Q47ZB6	FADI_COLP3	cytoplasm (by similarity).	cytoplasm
Q47ZQ7	Y3013_COLP3		cytoplasm
Q48082	RS1_HAEIN		cytoplasm
Q480C7	UVRB_COLP3	cytoplasm (by similarity).	cytoplasm
Q481E4	Y2611_COLP3		inner membrane
Q481S6	GLYA2_COLP3	cytoplasm (by similarity).	cytoplasm
Q481U4	Y2458_COLP3		inner membrane
Q48211	LST_HAEIN		cytoplasm
Q48214	Y1696_HAEIN		cytoplasm
Q48215	Y1695_HAEIN		cytoplasm
Q48248	CDH_HELPY	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q48252	CAGE_HELPY		cytoplasm

Q48254	HPAA2_HELPY	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q48255	AROQ_HELPY		cytoplasm
Q48260	MOAC_HELPY		cytoplasm
Q48261	HPAA3_HELPY	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q48262	NIXA_HELPY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q48264	HPAA1_HELPY	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q48271	COPP_HELPY		inner membrane
Q482K3	RL32_COLP3		cytoplasm
Q482T6	RS15_COLP3		cytoplasm
Q482T7	TRUB_COLP3		cytoplasm
Q482U1	Y2201_COLP3		inner membrane
Q48303	DHDM_HYPSX		cytoplasm
Q483B5	LPXK_COLP3		cytoplasm
Q483C4	RUVB_COLP3		cytoplasm
Q483C5	RUVA_COLP3		cytoplasm
Q483C6	RUVC_COLP3		cytoplasm
Q483D3	G6PI2_COLP3	cytoplasm (by similarity).	cytoplasm
Q483H8	HISX2_COLP3		cytoplasm
Q483Q8	DEOD_COLP3		cytoplasm
Q483R6	TYPH_COLP3		cytoplasm
Q48409	CASB_KLEOX		cytoplasm
Q48413	RAMA_KLEPN		cytoplasm
Q48414	YBDJ_KLEPN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q48436	BUDC_KLEPN		cytoplasm
Q48440	HPAB_KLEOX		cytoplasm
Q48441	HPAC_KLEOX		cytoplasm
Q48446	YC18_KLEPN		inner membrane
Q48447	GALF_KLEPN		cytoplasm
Q48448	YC02_KLEPN		inner membrane
Q48449	YC03_KLEPN		outer membrane
Q48450	YC04_KLEPN	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q48451	YOR5_KLEPN		extracellular
Q48452	YC06_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q48453	YC07_KLEPN		cytoplasm
Q48454	YC08_KLEPN		cytoplasm
Q48455	YC09_KLEPN		cytoplasm
Q48456	YC10_KLEPN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q48457	YCP11_KLEPN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q48458	YC12_KLEPN		cytoplasm
Q48459	YC13_KLEPN		cytoplasm

Q48460	YC14_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q48462	MANC_KLEPN		cytoplasm
Q48465	MTGA_KLEOX	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q48468	NASR_KLEOX		cytoplasm
Q48475	RFBA1_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q48476	RFBB1_KLEPN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q48478	RFBA2_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q48479	RFBB2_KLEPN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q48481	GLF8_KLEPN		cytoplasm
Q48485	GLF1_KLEPN		cytoplasm
Q485F2	ACCA_COLP3	cytoplasm (by similarity).	cytoplasm
Q485F4	RNH2_COLP3	cytoplasm (potential).	extracellular
Q485F9	FABZ_COLP3	cytoplasm (by similarity).	cytoplasm
Q485G4	DXR_COLP3		cytoplasm
Q485H4	GLND_COLP3		cytoplasm
Q485H5	DAPD_COLP3	cytoplasm (by similarity).	cytoplasm
Q485N9	FLGI_COLP3	periplasm (by similarity).	periplasm
Q485P0	FLGH_COLP3	outer membrane; lipid-anchor (probable).	inner membrane
Q486F9	DNAK1_COLP3		cytoplasm
Q486G0	ARGJ_COLP3	cytoplasm (probable).	cytoplasm
Q486J6	GCSP1_COLP3		cytoplasm
Q486U0	SYI_COLP3	cytoplasm (by similarity).	cytoplasm
Q486U3	RS20_COLP3		cytoplasm
Q48739	IPNS_LYSLA		cytoplasm
Q48743	AMPC_LYSLA	periplasm (by similarity).	periplasm
Q487A3	QUEA_COLP3	cytoplasm (potential).	cytoplasm
Q487E6	SURE_COLP3	cytoplasm (potential).	periplasm
Q487E7	TRUD_COLP3		cytoplasm
Q487E8	ISPF_COLP3		extracellular
Q487E9	ISPD_COLP3		cytoplasm
Q487G4	NAPA_COLP3	periplasm (by similarity).	periplasm
Q487G8	PDXH_COLP3		cytoplasm
Q487I6	CCME_COLP3	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q487J6	G6PI1_COLP3	cytoplasm (by similarity).	cytoplasm
Q487Z2	RS10_COLP3		cytoplasm
Q487Z6	RL2_COLP3		cytoplasm
Q487Z9	RS3_COLP3		cytoplasm
Q48806	DLPA_LEGPH		cytoplasm

Q48815	HELA_LEGPN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q48833	GLRXA_LEGPH		cytoplasm
Q48835	FUR_LEGPN	cytoplasm (by similarity).	cytoplasm
Q488A0	RL16_COLP3		cytoplasm
Q488A2	RS17_COLP3		cytoplasm
Q488J6	SELU_COLP3		cytoplasm
Q488M6	SYV_COLP3	cytoplasm (by similarity).	cytoplasm
Q488N5	NRDR_COLP3		cytoplasm
Q488N6	GLYA1_COLP3	cytoplasm (by similarity).	cytoplasm
Q488Y8	RPOA_COLP3		cytoplasm
Q488Y9	RS4_COLP3		cytoplasm
Q488Z0	RS11_COLP3		cytoplasm
Q488Z1	RS13_COLP3		cytoplasm
Q488Z4	RL15_COLP3		cytoplasm
Q488Z6	RS5_COLP3		cytoplasm
Q488Z9	RS8_COLP3		cytoplasm
Q48907	HPS_METAM		cytoplasm
Q48935	APHA_MYCRA		cytoplasm
Q48952	RBL_MOBAS		extracellular
Q489A5	RBN_COLP3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q489N3	AROB_COLP3	cytoplasm (probable).	cytoplasm
Q489N4	AROK_COLP3	cytoplasm (probable).	cytoplasm
Q489T8	RL9_COLP3		cytoplasm
Q489U1	RS6_COLP3		cytoplasm
Q489W3	FADB_COLP3		cytoplasm
Q489W4	FADA_COLP3	cytoplasm (by similarity).	cytoplasm
Q48A17	PURA_COLP3	cytoplasm (by similarity).	cytoplasm
Q48A85	GPH_COLP3		cytoplasm
Q48AD4	FPG_COLP3		cytoplasm
Q48AT8	TUSA_COLP3	cytoplasm (by similarity).	cytoplasm
Q48AW1	ATPG_COLP3		inner membrane
Q48AW2	ATPA_COLP3	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q48BG3	ATPA_PSE14	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q48BG4	ATPG_PSE14		inner membrane
Q48BG7	GLMU_PSE14	cytoplasm (by similarity).	cytoplasm
Q48BY7	PPCK_PSE14	cytoplasm (by similarity).	cytoplasm
Q48BY8	HSLO_PSE14	cytoplasm (by similarity).	cytoplasm
Q48BZ2	RIMK_PSE14		cytoplasm
Q48C78	HIS5_PSE14	cytoplasm (by similarity).	cytoplasm
Q48C80	HIS4_PSE14	cytoplasm (by similarity).	cytoplasm
Q48C81	HIS6_PSE14	cytoplasm (by similarity).	cytoplasm
Q48C87	GPMI_PSE14		cytoplasm
Q48C90	SECB_PSE14	cytoplasm (by similarity).	cytoplasm
Q48CC6	NUDH_PSE14		cytoplasm
Q48CK1	FPG_PSE14		cytoplasm
Q48CL9	THIG_PSE14	cytoplasm (by similarity).	cytoplasm
Q48CM0	TRMB_PSE14		cytoplasm
Q48CM6	BETB_PSE14		cytoplasm

Q48CP3	GLYA2_PSE14	cytoplasm (by similarity).	cytoplasm
Q48CT3	PQQE_PSE14		cytoplasm
Q48D26	RL1_PSE14		cytoplasm
Q48D27	RL10_PSE14		cytoplasm
Q48D29	RPOB_PSE14		cytoplasm
Q48D30	RPOC_PSE14		cytoplasm
Q48D31	RS12_PSE14		cytoplasm
Q48D32	RS7_PSE14		cytoplasm
Q48D35	RS10_PSE14		cytoplasm
Q48D39	RL2_PSE14		cytoplasm
Q48D42	RS3_PSE14		cytoplasm
Q48D43	RL16_PSE14		cytoplasm
Q48D45	RS17_PSE14		cytoplasm
Q48D50	RS8_PSE14		cytoplasm
Q48D53	RS5_PSE14		cytoplasm
Q48D55	RL15_PSE14		cytoplasm
Q48D58	RS13_PSE14		cytoplasm
Q48D59	RS11_PSE14		cytoplasm
Q48D60	RS4_PSE14		cytoplasm
Q48D61	RPOA_PSE14		cytoplasm
Q48DB7	NRDR_PSE14		cytoplasm
Q48DC9	UREE_PSE14	cytoplasm (by similarity).	cytoplasm
Q48DE6	URE1_PSE14	cytoplasm (by similarity).	cytoplasm
Q48DE7	URE2_PSE14	cytoplasm (by similarity).	cytoplasm
Q48DF0	URE3_PSE14	cytoplasm (by similarity).	cytoplasm
Q48DL4	PROA_PSE14	cytoplasm (by similarity).	cytoplasm
Q48DQ3	KPTA_PSE14		cytoplasm
Q48DU1	MSCL_PSE14	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q48DU7	GLYA1_PSE14	cytoplasm (by similarity).	cytoplasm
Q48E62	DNAK_PSE14		cytoplasm
Q48E64	DAPB_PSE14	cytoplasm (by similarity).	cytoplasm
Q48E69	RRMJ_PSE14	cytoplasm (by similarity).	cytoplasm
Q48E72	GLMM_PSE14		cytoplasm
Q48E75	Y4191_PSE14		inner membrane
Q48E79	TRUB_PSE14		cytoplasm
Q48E80	RS15_PSE14		cytoplasm
Q48EC7	MURA_PSE14	cytoplasm (probable).	inner membrane
Q48EC8	HIS1_PSE14	cytoplasm (by similarity).	cytoplasm
Q48EC9	HISX_PSE14		cytoplasm
Q48ED0	HIS8_PSE14		cytoplasm
Q48EE9	MRAZ_PSE14		inner membrane
Q48EF0	MRAW_PSE14		cytoplasm
Q48EF5	MRAY_PSE14	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q48EF8	MURG_PSE14	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q48EG7	ARGJ_PSE14	cytoplasm (probable).	cytoplasm
Q48EU3	Y3958_PSE14		cytoplasm
Q48EV0	LEPA_PSE14		cytoplasm
Q48EV3	RNC_PSE14	cytoplasm (by similarity).	cytoplasm
Q48EV5	RECO_PSE14		cytoplasm
Q48EV6	PDXJ_PSE14	cytoplasm (probable).	cytoplasm

Q48F57	GLND_PSE14		cytoplasm
Q48F65	DXR_PSE14		cytoplasm
Q48F70	FABZ_PSE14	cytoplasm (by similarity).	cytoplasm
Q48F73	RNH2_PSE14	cytoplasm (potential).	extracellular
Q48F75	ACCA_PSE14	cytoplasm (by similarity).	cytoplasm
Q48F81	ISPD_PSE14		cytoplasm
Q48F85	ISPF_PSE14		extracellular
Q48F86	TRUD_PSE14		cytoplasm
Q48F87	SURE_PSE14	cytoplasm (potential).	periplasm
Q48F92	MUTS_PSE14		cytoplasm
Q48FB7	SLYX_PSE14		inner membrane
Q48FC3	RUVC_PSE14		cytoplasm
Q48FC4	RUVA_PSE14		cytoplasm
Q48FC5	RUVB_PSE14		cytoplasm
Q48FH7	RUMA_PSE14		cytoplasm
Q48FT7	PPNK_PSE14	cytoplasm (by similarity).	cytoplasm
Q48G05	PHS_PSE14		cytoplasm
Q48GC5	FLGH_PSE14	outer membrane; lipid-anchor (probable).	inner membrane
Q48GC6	FLGI_PSE14	periplasm (by similarity).	periplasm
Q48GK0	MTFA_PSE14		cytoplasm
Q48GL4	CCME_PSE14	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q48GS7	HGD_PSE14		cytoplasm
Q48GW3	FADB_PSE14		cytoplasm
Q48GW4	FADA_PSE14	cytoplasm (by similarity).	cytoplasm
Q48HZ1	ARNA_PSE14		cytoplasm
Q48IH8	NUDC_PSE14		cytoplasm
Q48J73	XYLA_PSE14	cytoplasm (by similarity).	cytoplasm
Q48J96	UVRC_PSE14	cytoplasm (by similarity).	cytoplasm
Q48JG9	GLGA_PSE14		cytoplasm
Q48JI5	UPPP_PSE14	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q48JZ6	Y2060_PSE14		cytoplasm
Q48K64	HTPG_PSE14	cytoplasm (by similarity).	cytoplasm
Q48K97	LEU3_PSE14	cytoplasm (by similarity).	cytoplasm
Q48K98	LEUD_PSE14		cytoplasm
Q48K99	LEU2_PSE14		cytoplasm
Q48KA6	UVRB_PSE14	cytoplasm (by similarity).	cytoplasm
Q48KH5	TAL_PSE14	cytoplasm (probable).	cytoplasm
Q48KI2	QUEF_PSE14	cytoplasm (probable).	cytoplasm
Q48KQ4	TUSA_PSE14	cytoplasm (by similarity).	cytoplasm
Q48KR3	ZIPA_PSE14	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q48KZ0	CLPP_PSE14	cytoplasm (by similarity).	cytoplasm
Q48L44	RL32_PSE14		cytoplasm
Q48L51	MURB_PSE14	cytoplasm (probable).	inner membrane
Q48L54	LPXK_PSE14		cytoplasm
Q48LC8	MAO1_PSE14		cytoplasm
Q48LQ7	PDXH_PSE14		cytoplasm
Q48LV7	RL19_PSE14		cytoplasm
Q48LY0	GUAA_PSE14		cytoplasm

Q48LZ8	NDK_PSE14	cytoplasm (by similarity).	cytoplasm
Q48M15	QUEA_PSE14	cytoplasm (potential).	cytoplasm
Q48ME3	GCSP_PSE14		cytoplasm
Q48MF2	SYV_PSE14	cytoplasm (by similarity).	cytoplasm
Q48MV8	ISPE_PSE14		cytoplasm
Q48N66	ILVC_PSE14		cytoplasm
Q48N88	G6PI_PSE14	cytoplasm (by similarity).	cytoplasm
Q48NK6	SYI_PSE14	cytoplasm (by similarity).	cytoplasm
Q48NK9	RS20_PSE14		cytoplasm
Q48NL1	PROB_PSE14	cytoplasm (by similarity).	cytoplasm
Q48NL9	SELU_PSE14		cytoplasm
Q48NN3	SYI_PSE14	cytoplasm (by similarity).	cytoplasm
Q48NP8	TRPD_PSE14		cytoplasm
Q48NS2	GPH_PSE14		cytoplasm
Q48NU7	Y623_PSE14	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q48NZ0	RL9_PSE14		cytoplasm
Q48NZ3	RS6_PSE14		cytoplasm
Q48NZ7	PURA_PSE14	cytoplasm (by similarity).	cytoplasm
Q48P74	CVRA_PSE14	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q48PA1	METX_PSE14	cytoplasm (by similarity).	cytoplasm
Q48PA6	ILVD_PSE14		cytoplasm
Q48PB8	Y448_PSE14	periplasm (potential).	periplasm
Q48PH0	AROB_PSE14	cytoplasm (probable).	cytoplasm
Q48PH1	AROK_PSE14	cytoplasm (probable).	cytoplasm
Q48PJ7	HIS3_PSE14	cytoplasm (by similarity).	cytoplasm
Q48PJ8	HIS2_PSE14	cytoplasm (by similarity).	cytoplasm
Q48Q06	DUT_PSE14		extracellular
Q48Q16	RPOZ_PSE14		cytoplasm
Q48QJ8	RECF_PSE14	cytoplasm (by similarity).	cytoplasm
Q49091	GATA_MORCA		cytoplasm
Q49092	GATB_MORCA		cytoplasm
Q49115	MEAA_METEX		cytoplasm
Q49116	YMEB_METEX		cytoplasm
Q49118	FTR_METEX	cytoplasm (by similarity).	cytoplasm
Q49123	MAUF_METEX	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q49125	MAUE_METEX	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q49126	MAUD_METEX		inner membrane
Q49127	MAUJ_METEX		cytoplasm
Q49129	MAUL_METEX		periplasm
Q49130	MAUM_METEX		periplasm
Q49131	MAUN_METEX		periplasm
Q49149	PQQB_METEX		cytoplasm
Q49150	PQQCD_METEX		cytoplasm
Q491V0	G6PI_BLOPB	cytoplasm (by similarity).	cytoplasm
Q491W4	RPOZ_BLOPB		cytoplasm
Q491W8	DUT_BLOPB		extracellular
Q491X9	SECB_BLOPB	cytoplasm (by similarity).	cytoplasm
Q491Z0	ILVD_BLOPB		cytoplasm

Q491Z9	CORA_BLOPB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q492A4	AROK_BLOPB	cytoplasm (probable).	cytoplasm
Q492A5	AROB_BLOPB	cytoplasm (probable).	cytoplasm
Q492A9	RS12_BLOPB		cytoplasm
Q492B0	RS7_BLOPB		cytoplasm
Q492B6	RL1_BLOPB		cytoplasm
Q492B9	RPOB_BLOPB		cytoplasm
Q492C0	RPOC_BLOPB		cytoplasm
Q492C6	PPNK_BLOPB	cytoplasm (by similarity).	cytoplasm
Q492C9	LEPA_BLOPB		cytoplasm
Q492D1	RNC_BLOPB	cytoplasm (by similarity).	cytoplasm
Q492D2	PDXJ_BLOPB	cytoplasm (probable).	cytoplasm
Q492D3	ACPS_BLOPB	cytoplasm (by similarity).	cytoplasm
Q492D5	GLYA_BLOPB	cytoplasm (by similarity).	cytoplasm
Q492E5	GUAA_BLOPB		cytoplasm
Q492E7	URE3_BLOPB	cytoplasm (by similarity).	cytoplasm
Q492E8	URE2_BLOPB	cytoplasm (by similarity).	cytoplasm
Q492E9	URE1_BLOPB	cytoplasm (by similarity).	cytoplasm
Q492K0	HIS5_BLOPB	cytoplasm (by similarity).	cytoplasm
Q492K2	HIS8_BLOPB		cytoplasm
Q492K3	HISX_BLOPB		cytoplasm
Q492N9	GCH2_BLOPB		extracellular
Q492P7	PGSA_BLOPB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q492Q2	RL32_BLOPB		cytoplasm
Q492T0	LPXK_BLOPB		cytoplasm
Q492T7	SYU_BLOPB	cytoplasm (by similarity).	cytoplasm
Q492T8	PDXH_BLOPB		cytoplasm
Q492W0	ISPE_BLOPB		cytoplasm
Q492W5	GPMA_BLOPB		cytoplasm
Q493B6	ACCA_BLOPB	cytoplasm (by similarity).	cytoplasm
Q493B8	RNH2_BLOPB	cytoplasm (potential).	extracellular
Q493C1	FABZ_BLOPB	cytoplasm (by similarity).	cytoplasm
Q493C3	SKP_BLOPB	periplasm (by similarity).	periplasm
Q493D4	DAPD_BLOPB	cytoplasm (by similarity).	cytoplasm
Q493D9	NUDH_BLOPB		cytoplasm
Q493F8	CLPP_BLOPB	cytoplasm (by similarity).	cytoplasm
Q493I4	RPOA_BLOPB		cytoplasm
Q493I5	RS4_BLOPB		cytoplasm
Q493I6	RS11_BLOPB		cytoplasm
Q493I7	RS13_BLOPB		cytoplasm
Q493J0	RL15_BLOPB		cytoplasm
Q493J1	RS5_BLOPB		cytoplasm
Q493J4	RS8_BLOPB		cytoplasm
Q493J9	RS17_BLOPB		cytoplasm
Q493K1	RL16_BLOPB		cytoplasm
Q493K2	RS3_BLOPB		cytoplasm
Q493K5	RL2_BLOPB		cytoplasm
Q493K9	RS10_BLOPB		cytoplasm
Q493L3	MURB_BLOPB	cytoplasm (probable).	inner membrane
Q493M0	RL19_BLOPB		cytoplasm
Q493M8	ISPF_BLOPB		extracellular

Q493Q1	MURG_BLOPB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q493Q4	MRAY_BLOPB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q493Q9	MRAW_BLOPB	cytoplasm (by similarity).	cytoplasm
Q493R1	LEU3_BLOPB	cytoplasm (by similarity).	cytoplasm
Q493R2	LEU2_BLOPB		cytoplasm
Q493R3	LEUD_BLOPB		cytoplasm
Q493S0	DAPB_BLOPB	cytoplasm (by similarity).	cytoplasm
Q493S3	SYI_BLOPB	cytoplasm (by similarity).	cytoplasm
Q493S5	RS20_BLOPB		cytoplasm
Q493S7	DNAK_BLOPB		cytoplasm
Q493T4	RS15_BLOPB		cytoplasm
Q493T5	TRUB_BLOPB		cytoplasm
Q493U0	GLMM_BLOPB		cytoplasm
Q493U3	RRMJ_BLOPB	cytoplasm (by similarity).	cytoplasm
Q493V3	RL9_BLOPB		cytoplasm
Q493V5	RS6_BLOPB		cytoplasm
Q493X5	UPPP_BLOPB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q493X7	Y061_BLOPB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q493Y9	MURA_BLOPB	cytoplasm (probable).	inner membrane
Q493Z9	SYV_BLOPB	cytoplasm (by similarity).	cytoplasm
Q494C1	GLMU_BLOPB	cytoplasm (by similarity).	cytoplasm
Q494C4	ATPG_BLOPB		inner membrane
Q494C5	ATPA_BLOPB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q494D6	HIS6_BLOPB	cytoplasm (by similarity).	cytoplasm
Q494D7	HIS4_BLOPB	cytoplasm (by similarity).	cytoplasm
Q494E8	ISPD_BLOPB		cytoplasm
Q494F3	PURA_BLOPB	cytoplasm (by similarity).	cytoplasm
Q4AAX7	TRUB_COXBU		cytoplasm
Q4FL49	FTHS_PELUB		cytoplasm
Q4FLD7	RPOZ_PELUB		cytoplasm
Q4FLJ0	RS4_PELUB		cytoplasm
Q4FLJ4	PURL_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FLL0	RL10_PELUB		cytoplasm
Q4FLL2	RPOB_PELUB		cytoplasm
Q4FLL3	RPOC_PELUB		cytoplasm
Q4FLL4	RS12_PELUB		cytoplasm
Q4FLL5	RS7_PELUB		cytoplasm
Q4FLL7	RS10_PELUB		cytoplasm
Q4FLM4	RS3_PELUB		cytoplasm
Q4FLM5	RL16_PELUB		cytoplasm
Q4FLM7	RS17_PELUB		cytoplasm
Q4FLN2	RS8_PELUB		cytoplasm
Q4FLN5	RS5_PELUB		cytoplasm
Q4FLN6	RL15_PELUB		cytoplasm
Q4FLN9	RS13_PELUB		cytoplasm
Q4FLP0	RS11_PELUB		cytoplasm

Q4FLP1	RPOA_PELUB		cytoplasm
Q4FLQ2	Y1082_PELUB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4FLS6	PDXJ_PELUB	cytoplasm (probable).	cytoplasm
Q4FLS7	ACPS_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FLS9	RNC_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FLT4	GLYA_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FLU5	RL32_PELUB		cytoplasm
Q4FM19	ILVD_PELUB		cytoplasm
Q4FM20	SYV_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FM31	ISPDF_PELUB		cytoplasm
Q4FM64	DXR_PELUB		cytoplasm
Q4FM94	CLPP_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FMH0	HISX2_PELUB		cytoplasm
Q4FMN5	SYV_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FMP0	THIG_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FMR5	RS6_PELUB		cytoplasm
Q4FMR7	RL9_PELUB		cytoplasm
Q4FMV1	GCSP_PELUB		cytoplasm
Q4FMW8	GUAA_PELUB		cytoplasm
Q4FMX1	RRMJ_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FN05	PDXH_PELUB		cytoplasm
Q4FN13	PURA_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FN15	GLMM_PELUB		cytoplasm
Q4FNA6	ACCA_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FNE2	HISX1_PELUB		cytoplasm
Q4FNE4	MURA_PELUB	cytoplasm (probable).	inner membrane
Q4FNF4	DAPD_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FNG5	AROB_PELUB	cytoplasm (probable).	cytoplasm
Q4FNH3	LEPA_PELUB		cytoplasm
Q4FNI5	UPPP_PELUB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4FNK9	RS20_PELUB		cytoplasm
Q4FNM7	RS15_PELUB		cytoplasm
Q4FNP9	DNAK_PELUB		cytoplasm
Q4FNQ1	DAPB_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FNR6	GIDA_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FNS3	Y343_PELUB		cytoplasm
Q4FNS8	SECB_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FNT5	HIS5_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FNT6	HIS4_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FNT7	HIS6_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FNY3	CCME_PELUB	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q4FP13	TRMD_PELUB	cytoplasm (potential).	cytoplasm
Q4FP14	RL19_PELUB		cytoplasm
Q4FP15	LEU2_PELUB		cytoplasm
Q4FP16	LEUD_PELUB		cytoplasm
Q4FP17	LEU3_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FP36	ATPA_PELUB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q4FP37	ATPG_PELUB		inner membrane
Q4FP40	NUDH_PELUB		cytoplasm
Q4FP45	PROB_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FP51	METX_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FP52	HIS8_PELUB		cytoplasm
Q4FP74	GPMA_PELUB		cytoplasm
Q4FPA9	SYI_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FPF7	RNH2_PELUB	cytoplasm (potential).	extracellular
Q4FPG0	ISPE_PELUB		cytoplasm
Q4FPI1	UVRC_PELUB	cytoplasm (by similarity).	cytoplasm
Q4FPK2	MRAY_PELUB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4FPK7	MURB_PELUB	cytoplasm (probable).	inner membrane
Q4FPN3	MRAZ_PELUB		inner membrane
Q4FPN5	MRAW_PELUB		cytoplasm
Q4FPQ6	ILVC_PELUB		cytoplasm
Q4FPR9	NRDR_PSYAR		cytoplasm
Q4FPS9	DNAK_PSYAR		cytoplasm
Q4FPT7	GPH_PSYAR		cytoplasm
Q4FPU1	HIS5_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FPV8	HIS4_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FPY8	GLMU_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FPZ3	HIS6_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FQ05	MRAW_PSYAR		cytoplasm
Q4FQ10	MRAY_PSYAR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4FQ23	GLUQ_PSYAR		cytoplasm
Q4FQ35	ATPA_PSYAR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4FQ36	ATPG_PSYAR		inner membrane
Q4FQ45	RL19_PSYAR		cytoplasm
Q4FQ48	UVRC_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FQ54	NUDH_PSYAR		cytoplasm
Q4FQ91	SYI_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FQB9	CLPP_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FQC6	FADB_PSYAR		cytoplasm
Q4FQC7	FADA_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FQF6	MURA_PSYAR	cytoplasm (probable).	inner membrane
Q4FQF7	HIS1_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FQF8	HISX_PSYAR		cytoplasm
Q4FQF9	HIS82_PSYAR		cytoplasm
Q4FQG3	RS12_PSYAR		cytoplasm
Q4FQG4	RS7_PSYAR		cytoplasm
Q4FQH0	RL1_PSYAR		cytoplasm
Q4FQH1	RL10_PSYAR		cytoplasm
Q4FQH3	RPOB_PSYAR		cytoplasm
Q4FQH4	RPOC_PSYAR		cytoplasm
Q4FQI2	AROK_PSYAR	cytoplasm (probable).	cytoplasm
Q4FQI3	AROB_PSYAR	cytoplasm (probable).	cytoplasm
Q4FQK7	Y1853_PSYAR		inner membrane
Q4FQL0	SYV_PSYAR	cytoplasm (by similarity).	cytoplasm

Q4FQN5	CCME_PSYAR	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q4FQU7	MDH_PSYAR		cytoplasm
Q4FQV9	MURG_PSYAR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4FQX1	RRMJ_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FQX4	Y1736_PSYAR		cytoplasm
Q4FQX9	TRMB_PSYAR		cytoplasm
Q4FQY6	RPOZ_PSYAR		cytoplasm
Q4FQZ1	HTPG_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FR05	UPPP_PSYAR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4FR14	TRUD_PSYAR		cytoplasm
Q4FR76	ISPD_PSYAR		cytoplasm
Q4FRF4	SECB_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FRH9	DXR_PSYAR		cytoplasm
Q4FRI3	FABZ_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FRK7	PROB_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FRM3	RS20_PSYAR		cytoplasm
Q4FRP5	PPNK_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FRU7	LEU2_PSYAR		cytoplasm
Q4FRU8	LEUD_PSYAR		cytoplasm
Q4FRV0	LEU3_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FRV6	QUEA_PSYAR	cytoplasm (potential).	cytoplasm
Q4FRX3	MAO1_PSYAR		cytoplasm
Q4FS01	GLMM_PSYAR		cytoplasm
Q4FS02	PDXH_PSYAR		cytoplasm
Q4FS19	RL9_PSYAR		cytoplasm
Q4FS21	RS6_PSYAR		cytoplasm
Q4FS24	Y1334_PSYAR		cytoplasm
Q4FS43	LPXK_PSYAR		cytoplasm
Q4FS54	ILVD_PSYAR		cytoplasm
Q4FSB5	ISPF_PSYAR		extracellular
Q4FSF9	ARGJ_PSYAR	cytoplasm (probable).	cytoplasm
Q4FSH2	HIS81_PSYAR		cytoplasm
Q4FT57	MURB_PSYAR	cytoplasm (probable).	inner membrane
Q4FT93	RNH2_PSYAR	cytoplasm (potential).	extracellular
Q4FTK9	GCSP_PSYAR		cytoplasm
Q4FTR7	DAPD_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FTT9	RUVB_PSYAR		cytoplasm
Q4FTU0	RUVA_PSYAR		cytoplasm
Q4FTU8	GPMI_PSYAR		cytoplasm
Q4FTV0	RUVC_PSYAR		cytoplasm
Q4FTX1	NDK_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FTX2	PURA_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FU67	DUT_PSYAR		extracellular
Q4FUB6	ILVC_PSYAR		cytoplasm
Q4FUC6	RL32_PSYAR		cytoplasm
Q4FUD1	RPOA_PSYAR		cytoplasm
Q4FUD2	RS4_PSYAR		cytoplasm
Q4FUD3	RS11_PSYAR		cytoplasm
Q4FUD4	RS13_PSYAR		cytoplasm

Q4FUD7	RL15_PSYAR		cytoplasm
Q4FUD9	RS5_PSYAR		cytoplasm
Q4FUE2	RS8_PSYAR		cytoplasm
Q4FUE7	RS17_PSYAR		cytoplasm
Q4FUE9	RL16_PSYAR		cytoplasm
Q4FUF0	RS3_PSYAR		cytoplasm
Q4FUF3	RL2_PSYAR		cytoplasm
Q4FUF7	RS10_PSYAR		cytoplasm
Q4FUL7	QUEF_PSYAR	cytoplasm (probable).	cytoplasm
Q4FUP5	GLND_PSYAR		cytoplasm
Q4FUR4	METX_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FUS5	SYI_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FUU7	FPG_PSYAR		cytoplasm
Q4FUU9	MSCL_PSYAR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4FUV3	PDXJ_PSYAR	cytoplasm (probable).	cytoplasm
Q4FUV6	RNC_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FUV9	LEPA_PSYAR		cytoplasm
Q4FUZ5	PROA_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FUZ8	GLYA_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FV41	MUTS_PSYAR		cytoplasm
Q4FV61	THIG_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FVB2	ISPE_PSYAR		cytoplasm
Q4FVB6	PAND_PSYAR		cytoplasm
Q4FVF3	PHS_PSYAR		cytoplasm
Q4FVH5	G6PI_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FVL1	RS15_PSYAR		cytoplasm
Q4FVL3	TRUB_PSYAR		cytoplasm
Q4FVL7	Y067_PSYAR		inner membrane
Q4FVQ6	DAPB_PSYAR	cytoplasm (by similarity).	cytoplasm
Q4FVS5	GUAA_PSYAR		cytoplasm
Q4K399	GIDA_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K3A0	GIDB_PSEF5		cytoplasm
Q4K3A7	ATPA_PSEF5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4K3A8	ATPG_PSEF5		inner membrane
Q4K3B1	GLMU_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K3E4	HIS32_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K3R3	RPOZ_PSEF5		cytoplasm
Q4K3S2	DUT_PSEF5		extracellular
Q4K3W5	CYAY_PSEF5		inner membrane
Q4K3Y6	Y5989_PSEF5		inner membrane
Q4K416	GCSP2_PSEF5		cytoplasm
Q4K475	NUDH_PSEF5		cytoplasm
Q4K498	ILVD_PSEF5		cytoplasm
Q4K4A5	FPG_PSEF5		cytoplasm
Q4K4C5	THIG_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K4C6	TRMB_PSEF5		cytoplasm
Q4K4C7	KPTA_PSEF5		cytoplasm
Q4K4D3	METX_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K4H3	Y5802_PSEF5	periplasm (potential).	periplasm
Q4K4K8	BETB_PSEF5		cytoplasm
Q4K4P6	GLYA2_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K4U8	PQQE_PSEF5		cytoplasm

Q4K4W5	Y5659_PSEF5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4K4Z4	GPH_PSEF5		cytoplasm
Q4K502	TRPD_PSEF5		cytoplasm
Q4K517	SYI_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K523	RL1_PSEF5		cytoplasm
Q4K524	RL10_PSEF5		cytoplasm
Q4K526	RPOB_PSEF5		cytoplasm
Q4K527	RPOC_PSEF5		cytoplasm
Q4K528	RS12_PSEF5		cytoplasm
Q4K529	RS7_PSEF5		cytoplasm
Q4K532	RS10_PSEF5		cytoplasm
Q4K536	RL2_PSEF5		cytoplasm
Q4K539	RS3_PSEF5		cytoplasm
Q4K540	RL16_PSEF5		cytoplasm
Q4K542	RS17_PSEF5		cytoplasm
Q4K547	RS8_PSEF5		cytoplasm
Q4K550	RS5_PSEF5		cytoplasm
Q4K552	RL15_PSEF5		cytoplasm
Q4K554	RS13_PSEF5		cytoplasm
Q4K555	RS11_PSEF5		cytoplasm
Q4K556	RS4_PSEF5		cytoplasm
Q4K557	RPOA_PSEF5		cytoplasm
Q4K592	NRDR_PSEF5		cytoplasm
Q4K5F9	PROA_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K5G2	Y5453_PSEF5		inner membrane
Q4K5Q6	MSCL_PSEF5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4K5R9	GLYA1_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K5T9	PROB_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K5U1	RS20_PSEF5		cytoplasm
Q4K5U4	SYI_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K5Y1	G6PI_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K5Y3	PANC_PSEF5		cytoplasm
Q4K5Z0	GLUQ_PSEF5		cytoplasm
Q4K608	ILVC_PSEF5		cytoplasm
Q4K691	ISPE_PSEF5		cytoplasm
Q4K6I4	MRAZ_PSEF5		inner membrane
Q4K6I5	MRAW_PSEF5		cytoplasm
Q4K6J0	MRAY_PSEF5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4K6J1	MURD_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K6J3	MURG_PSEF5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4K6S8	QUEA_PSEF5	cytoplasm (potential).	cytoplasm
Q4K6U5	NDK_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K6W4	GUAA_PSEF5		cytoplasm
Q4K7C2	ARGJ_PSEF5	cytoplasm (probable).	cytoplasm
Q4K7D1	SLYX_PSEF5		inner membrane
Q4K7D7	RUVC_PSEF5		cytoplasm
Q4K7D8	RUVA_PSEF5		cytoplasm
Q4K7D9	RUVB_PSEF5		cytoplasm

Q4K7Q8	GCSP1_PSEF5		cytoplasm
Q4K898	RUMA_PSEF5		cytoplasm
Q4K8H3	ALLC_PSEF5		extracellular
Q4K8N0	HIS82_PSEF5		cytoplasm
Q4K8U2	TPMT_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4K9J6	CLPP_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KAD4	DNAE2_PSEF5	cytoplasm (probable).	cytoplasm
Q4KAR9	UVRC_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KB60	Y3418_PSEF5		inner membrane
Q4KBJ2	PPNK_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KBL2	NUDC_PSEF5		cytoplasm
Q4KC13	UPPP_PSEF5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4KC31	ECOT_PSEF5	periplasm (by similarity).	periplasm
Q4KC82	ARNA_PSEF5		cytoplasm
Q4KC84	ARNB_PSEF5		cytoplasm
Q4KCP0	GLGA_PSEF5		cytoplasm
Q4KEE7	SELU_PSEF5		cytoplasm
Q4KF05	LEU3_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KF07	LEUD_PSEF5		cytoplasm
Q4KF08	LEU2_PSEF5		cytoplasm
Q4KF19	UVRB_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KF90	TAL_PSEF5	cytoplasm (probable).	cytoplasm
Q4KF97	QUEF_PSEF5	cytoplasm (probable).	cytoplasm
Q4KFC3	FADA_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KFC4	FADB_PSEF5		cytoplasm
Q4KFD3	TUSA_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KFD4	MTFA_PSEF5		cytoplasm
Q4KFH2	ZIPA_PSEF5	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q4KFJ6	Y1868_PSEF5		cytoplasm
Q4KFS0	RL32_PSEF5		cytoplasm
Q4KFT1	MURB_PSEF5	cytoplasm (probable).	inner membrane
Q4KFT5	LPXK_PSEF5		cytoplasm
Q4KFX8	HTPG_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KG23	CCME_PSEF5	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q4KG91	FLGI_PSEF5	periplasm (by similarity).	periplasm
Q4KG92	FLGH_PSEF5	outer membrane; lipid-anchor (probable).	inner membrane
Q4KG97	PHS_PSEF5		cytoplasm
Q4KGJ3	Y1513_PSEF5		inner membrane
Q4KGR0	Y1445_PSEF5		nucleoid
Q4KH71	PDXH_PSEF5		cytoplasm
Q4KHA4	KUP_PSEF5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4KHC1	RECA_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KHE3	MUTS_PSEF5		cytoplasm
Q4KHE8	SURE_PSEF5	cytoplasm (potential).	periplasm
Q4KHE9	TRUD_PSEF5		cytoplasm
Q4KHF0	ISPF_PSEF5		extracellular

Q4KHF4	ISPD_PSEF5		cytoplasm
Q4KHG0	ACCA_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KHG2	RNH2_PSEF5	cytoplasm (potential).	extracellular
Q4KHG5	FABZ_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KHH0	DXR_PSEF5		cytoplasm
Q4KHH3	RRF_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KHH8	GLND_PSEF5		cytoplasm
Q4KHJ7	Y1155_PSEF5		inner membrane
Q4KHQ5	RL19_PSEF5		cytoplasm
Q4KHQ6	TRMD_PSEF5	cytoplasm (potential).	cytoplasm
Q4KHS8	PDXJ_PSEF5	cytoplasm (probable).	cytoplasm
Q4KHS9	RECO_PSEF5		cytoplasm
Q4KHT1	RNC_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KHT3	LEPA_PSEF5		cytoplasm
Q4KHT9	SYV_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KI35	HGD_PSEF5		cytoplasm
Q4KI58	CYSZ_PSEF5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4KI72	HIS81_PSEF5		cytoplasm
Q4KI73	HISX_PSEF5		cytoplasm
Q4KI74	HIS1_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KI75	MURA_PSEF5	cytoplasm (probable).	inner membrane
Q4KIF3	RS15_PSEF5		cytoplasm
Q4KIF4	TRUB_PSEF5		cytoplasm
Q4KIF8	Y842_PSEF5		inner membrane
Q4KIG0	GLMM_PSEF5		cytoplasm
Q4KIG3	RRMJ_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KIG9	DAPB_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KIH1	DNAK_PSEF5		cytoplasm
Q4KJ06	URE3_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJ09	URE2_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJ10	URE1_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJ32	UREE_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJ59	RL9_PSEF5		cytoplasm
Q4KJ62	RS6_PSEF5		cytoplasm
Q4KJ68	PURA_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJ81	ORN_PSEF5	cytoplasm (potential).	cytoplasm
Q4KJF1	CVRA_PSEF5	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q4KJI8	AROB_PSEF5	cytoplasm (probable).	cytoplasm
Q4KJI9	AROK_PSEF5	cytoplasm (probable).	cytoplasm
Q4KJL5	HIS31_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJL6	HIS2_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJR5	SECB_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJR8	GPMI_PSEF5		cytoplasm
Q4KJS4	HIS6_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJS5	HIS4_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KJS7	HIS5_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KK03	PPCK_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KK04	HSLO_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4KK09	RIMK_PSEF5		cytoplasm
Q4KKS8	RECF_PSEF5	cytoplasm (by similarity).	cytoplasm
Q4QJK4	RPOZ_HAEI8		cytoplasm
Q4QJK8	LLDD_HAEI8		cytoplasm

Q4QJL8	Y2036_HAEI8		inner membrane
Q4QJM6	GLND_HAEI8		cytoplasm
Q4QJM8	ORN_HAEI8	cytoplasm (potential).	cytoplasm
Q4QJT7	Y1958_HAEI8		inner membrane
Q4QJW4	DNAK_HAEI8		cytoplasm
Q4QJW6	PROA_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QJX4	UVRB_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QJY8	METX_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QK35	Y1846_HAEI8		inner membrane
Q4QK42	RBFA_HAEI8	cytoplasm (potential).	cytoplasm
Q4QK43	TRUB_HAEI8		cytoplasm
Q4QK45	QUEF_HAEI8	cytoplasm (probable).	cytoplasm
Q4QK58	PEPT_HAEI8	cytoplasm (by similarity).	extracellular
Q4QK60	CDD_HAEI8		cytoplasm
Q4QK70	GLGA_HAEI8		cytoplasm
Q4QKA1	TRPD_HAEI8		cytoplasm
Q4QKA4	SYV_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QKG0	Y1697_HAEI8		inner membrane
Q4QKG3	THII_HAEI8	cytoplasm (potential).	cytoplasm
Q4QKI9	GLMM_HAEI8		cytoplasm
Q4QKJ2	RRMJ_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QKJ8	RS15_HAEI8		cytoplasm
Q4QKM7	DAPB_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QKP3	RIMK_HAEI8		cytoplasm
Q4QL07	G6PI_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QL43	ISPE_HAEI8		cytoplasm
Q4QL45	SYI_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QL68	PURA_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QL69	DAPD_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QL89	MDH_HAEI8		cytoplasm
Q4QL95	ACKA_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QLA4	MTFA_HAEI8		cytoplasm
Q4QLD1	HIS82_HAEI8		cytoplasm
Q4QLF6	DDL_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QLF7	MURC_HAEI8	cytoplasm (probable).	cytoplasm
Q4QLF8	MURG_HAEI8	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4QLG0	MURD_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QLG1	MRAY_HAEI8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4QLG6	MRAW_HAEI8		cytoplasm
Q4QLG7	MRAZ_HAEI8		inner membrane
Q4QLG9	TAL_HAEI8	cytoplasm (probable).	cytoplasm
Q4QLH8	DEOC_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QLI2	XYLA_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QLI8	CYSZ_HAEI8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4QLI9	ZIPA_HAEI8	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane

Q4QLJ5	CCME_HAEI8	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q4QLK7	MURA_HAEI8	cytoplasm (probable).	inner membrane
Q4QLM4	FABZ_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QLM7	RNH2_HAEI8	cytoplasm (potential).	extracellular
Q4QLP6	CORA_HAEI8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4QLR9	RECF_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QLS1	LEUD_HAEI8		cytoplasm
Q4QLS2	LEU2_HAEI8		cytoplasm
Q4QLS3	LEU3_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QLS9	SSRP_HAEI8	cytoplasm (potential).	cytoplasm
Q4QLU2	RS20_HAEI8		cytoplasm
Q4QLU5	SYI_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QLU8	NAGZ_HAEI8	cytoplasm (by similarity).	extracellular
Q4QLV3	DUT_HAEI8		extracellular
Q4QLV5	RADC_HAEI8		cytoplasm
Q4QLW2	FPG_HAEI8		cytoplasm
Q4QLW5	NRDR_HAEI8		cytoplasm
Q4QM04	TYSY_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QM07	NUDH_HAEI8		cytoplasm
Q4QM08	PROB_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QM19	GLYA_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QM32	NDK_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QM33	PEPB_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QM38	PDXH_HAEI8		cytoplasm
Q4QM52	Y1015_HAEI8		inner membrane
Q4QM60	Y1007_HAEI8		inner membrane
Q4QM89	ARLY_HAEI8	cytoplasm (probable).	cytoplasm
Q4QM90	HSLO_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QM91	PPCK_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QM92	RRF_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QM93	DXR_HAEI8		cytoplasm
Q4QM97	RPOA_HAEI8		cytoplasm
Q4QM98	RS4_HAEI8		cytoplasm
Q4QM99	RS11_HAEI8		cytoplasm
Q4QMA0	RS13_HAEI8		cytoplasm
Q4QMA2	RL15_HAEI8		cytoplasm
Q4QMA4	RS5_HAEI8		cytoplasm
Q4QMA7	RS8_HAEI8		cytoplasm
Q4QMB3	RS17_HAEI8		cytoplasm
Q4QMB5	RL16_HAEI8		cytoplasm
Q4QMB6	RS3_HAEI8		cytoplasm
Q4QMB9	RL2_HAEI8		cytoplasm
Q4QMC3	RS10_HAEI8		cytoplasm
Q4QME2	GPMA_HAEI8		cytoplasm
Q4QME9	LEXA_HAEI8		cytoplasm
Q4QMF4	SECB_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QMF8	ILVD_HAEI8		cytoplasm
Q4QMG7	CYAY_HAEI8		inner membrane
Q4QMJ5	PSIE_HAEI8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q4QMJ8	TUSA_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QMK5	CLPP_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QML2	MUTS_HAEI8		cytoplasm
Q4QML8	SURE_HAEI8	cytoplasm (potential).	periplasm
Q4QML9	TRUD_HAEI8		cytoplasm
Q4QMM6	XGPT2_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QMN4	ILVC_HAEI8		cytoplasm
Q4QMP2	XGPT1_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QMP4	ISPD_HAEI8		cytoplasm
Q4QMP5	ISPF_HAEI8		extracellular
Q4QMS5	GLMU_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QMS7	RL10_HAEI8		cytoplasm
Q4QMT7	RS7_HAEI8		cytoplasm
Q4QMT8	RS12_HAEI8		cytoplasm
Q4QMT9	GIDA_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QMV2	RECA_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QMW0	MSCL_HAEI8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4QMW1	Y720_HAEI8		cytoplasm
Q4QMX2	SLYX_HAEI8		inner membrane
Q4QMY0	GPH_HAEI8		cytoplasm
Q4QN01	RS6_HAEI8		cytoplasm
Q4QN04	RL9_HAEI8		cytoplasm
Q4QN06	CH10_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QN07	URE3_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QN08	URE2_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QN09	URE1_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QN10	UREE_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QN30	DEOD_HAEI8		cytoplasm
Q4QN32	RL1_HAEI8		cytoplasm
Q4QN33	RPOB_HAEI8		cytoplasm
Q4QN34	RPOC_HAEI8		cytoplasm
Q4QN56	GIDB_HAEI8		cytoplasm
Q4QN62	ATPA_HAEI8	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4QN63	ATPG_HAEI8		inner membrane
Q4QN69	HIS6_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QN70	HIS4_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QN71	HIS5_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QN73	HIS81_HAEI8		cytoplasm
Q4QN74	HISX_HAEI8		cytoplasm
Q4QNA2	RECR_HAEI8		cytoplasm
Q4QNB3	NUDC_HAEI8		cytoplasm
Q4QND7	ACCA_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QNJ6	NAPA_HAEI8	periplasm (by similarity).	periplasm
Q4QNK0	TRMB_HAEI8		cytoplasm
Q4QNK7	RUMA_HAEI8		cytoplasm
Q4QNK8	RECO_HAEI8		cytoplasm
Q4QNM4	RUVC_HAEI8		cytoplasm
Q4QNM5	RUVA_HAEI8		cytoplasm
Q4QNM6	RUVB_HAEI8		cytoplasm
Q4QNR3	RBN_HAEI8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q4QNS0	MURB_HAEI8	cytoplasm (probable).	inner membrane
Q4QNS2	Y373_HAEI8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4QNS5	HXUB_HAEI8	bacterial cell outer membrane (by similarity).	outer membrane
Q4QNS6	HXUC_HAEI8	bacterial cell outer membrane (by similarity).	outer membrane
Q4QNU2	QUEA_HAEI8	cytoplasm (potential).	cytoplasm
Q4QNW4	GUAA_HAEI8		cytoplasm
Q4QNX8	GCH2_HAEI8		extracellular
Q4QNY2	AROB_HAEI8	cytoplasm (probable).	cytoplasm
Q4QNY3	AROK_HAEI8	cytoplasm (probable).	cytoplasm
Q4QNY7	RIMM_HAEI8	cytoplasm (potential).	cytoplasm
Q4QNY8	TRMD_HAEI8	cytoplasm (potential).	cytoplasm
Q4QNY9	RL19_HAEI8		cytoplasm
Q4QP29	RL32_HAEI8		cytoplasm
Q4QP40	SIAT_HAEI8	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q4QP41	SIAP_HAEI8	periplasm (by similarity).	periplasm
Q4QP49	RNH_HAEI8	cytoplasm (potential).	extracellular
Q4QP67	TDHA_HAEI8	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
Q4QP81	HTPG_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QPH3	PPNK_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QPI5	LPXK_HAEI8		cytoplasm
Q4QPI7	UVRC_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QPI9	UXUA_HAEI8		cytoplasm
Q4QPK9	Y043_HAEI8	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q4QPL1	Y041_HAEI8		inner membrane
Q4QPL7	LIPB_HAEI8	cytoplasm (potential).	cytoplasm
Q4QPM8	LEPA_HAEI8		cytoplasm
Q4QPN0	RNC_HAEI8	cytoplasm (by similarity).	cytoplasm
Q4QPN9	FDHD_HAEI8	cytoplasm (potential).	cytoplasm
Q4QPP0	Y004_HAEI8		inner membrane
Q4TVS0	SIAT2_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q4UJK6	DNAJ_RICFE	cytoplasm (probable).	cytoplasm
Q4UJK7	DNAK_RICFE		cytoplasm
Q4UJU4	HEMK_RICFE		cytoplasm
Q4UJV5	HTPG_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UK16	ATPA_RICFE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4UK17	ATPG_RICFE		inner membrane
Q4UK48	RL32_RICFE		cytoplasm
Q4UK63	RECA_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UK96	GLYA_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UKA8	RNC_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UKB1	RUVC_RICFE		cytoplasm
Q4UKC5	RS12_RICFE		cytoplasm

Q4UKC6	RS7_RICFE		cytoplasm
Q4UKD1	RL1_RICFE		cytoplasm
Q4UKD2	RL10_RICFE		cytoplasm
Q4UKD4	RPOB_RICFE		cytoplasm
Q4UKD5	RPOC_RICFE		cytoplasm
Q4UKE7	DAPB_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UKF3	RRF_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UKG2	RRMJ_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UKK3	DAPD_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UKL2	FER2_RICFE		cytoplasm
Q4UKL4	HSCB_RICFE		inner membrane
Q4UKL5	RNH2_RICFE	cytoplasm (potential).	extracellular
Q4UKL6	UVRB_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UKM5	NUDH_RICFE		cytoplasm
Q4UKN0	PSD_RICFE		cytoplasm
Q4UKP0	MURB_RICFE	cytoplasm (probable).	inner membrane
Q4UKP8	LPXC_RICFE		cytoplasm
Q4UKS2	LEPA_RICFE		cytoplasm
Q4UKS5	PHS_RICFE		cytoplasm
Q4UKW1	CCME_RICFE	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q4UKX7	MURA_RICFE	cytoplasm (probable).	inner membrane
Q4UKX8	RPOZ_RICFE		cytoplasm
Q4UKX9	ACPS_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UL10	UVRC_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UL17	MRAZ_RICFE		inner membrane
Q4UL18	MRAW_RICFE		cytoplasm
Q4UL45	SYI_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UL49	Y873_RICFE		inner membrane
Q4ULD7	IDI2_RICFE	cytoplasm (by similarity).	cytoplasm
Q4ULF0	CLPP_RICFE	cytoplasm (by similarity).	cytoplasm
Q4ULJ9	TRUB_RICFE		cytoplasm
Q4ULK0	RS15_RICFE		cytoplasm
Q4ULQ0	RECR_RICFE		cytoplasm
Q4ULQ7	SUCC_RICFE		cytoplasm
Q4ULR2	DAPA_RICFE	cytoplasm (by similarity).	cytoplasm
Q4ULR6	UPPS_RICFE		cytoplasm
Q4ULT6	MURG_RICFE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4ULT8	MURD_RICFE	cytoplasm (by similarity).	cytoplasm
Q4ULV2	DUT_RICFE		extracellular
Q4ULW6	RUVB_RICFE		cytoplasm
Q4ULW8	RUVA_RICFE		cytoplasm
Q4ULY2	MDH_RICFE		cytoplasm
Q4UM23	RS4_RICFE		cytoplasm
Q4UM50	CYAY_RICFE		inner membrane
Q4UM86	MUTS_RICFE		cytoplasm
Q4UMD8	SYI_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UMI7	MRAY_RICFE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4UMM7	QUEA_RICFE	cytoplasm (potential).	cytoplasm
Q4UMQ3	RS20_RICFE		cytoplasm

Q4UMQ5	RPOA_RICFE		cytoplasm
Q4UMQ6	RS11_RICFE		cytoplasm
Q4UMQ7	RS13_RICFE		cytoplasm
Q4UMR0	RL15_RICFE		cytoplasm
Q4UMR2	RS5_RICFE		cytoplasm
Q4UMR3	RL18_RICFE		cytoplasm
Q4UMR5	RS8_RICFE		cytoplasm
Q4UMS0	RS17_RICFE		cytoplasm
Q4UMS1	RL29_RICFE		cytoplasm
Q4UMS2	RL16_RICFE		cytoplasm
Q4UMS3	RS3_RICFE		cytoplasm
Q4UMS6	RL2_RICFE		cytoplasm
Q4UMT0	RS10_RICFE		cytoplasm
Q4UMW5	FPG_RICFE		cytoplasm
Q4UMX8	SYV_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UN11	LPXK_RICFE		cytoplasm
Q4UN27	RNH_RICFE	cytoplasm (potential).	extracellular
Q4UN64	RS6_RICFE		cytoplasm
Q4UN66	RL9_RICFE		cytoplasm
Q4UN87	NDK_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UN88	GIDA_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UN89	GIDB_RICFE		cytoplasm
Q4UNA7	RL19_RICFE		cytoplasm
Q4UNA8	TRMD_RICFE	cytoplasm (potential).	cytoplasm
Q4UNF0	QUEF_RICFE	cytoplasm (probable).	cytoplasm
Q4UNF2	SECB_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UNG8	RECF_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UNK0	FABZ_RICFE	cytoplasm (by similarity).	cytoplasm
Q4UNK5	Y001_RICFE		cytoplasm
Q4UNU6	FPG_XANC8		cytoplasm
Q4UNZ4	XYLA2_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UP54	GIDB_XANC8		cytoplasm
Q4UPP2	DUT_XANC8		extracellular
Q4UPQ9	SYV_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UPX6	QUEF_XANC8	cytoplasm (probable).	cytoplasm
Q4UQF2	ATPA_XANC8	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4UQF3	ATPG_XANC8		inner membrane
Q4UQF8	GLMU_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UQP3	SYV_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UQT6	GLYA_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UQT7	NRDR_XANC8		cytoplasm
Q4UQW2	MRAZ_XANC8		inner membrane
Q4UQW3	MRAW_XANC8		cytoplasm
Q4UQW8	MRAY_XANC8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4UQX0	MURG_XANC8	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4UR65	MTFA_XANC8		cytoplasm
Q4UR79	TAL_XANC8	cytoplasm (probable).	cytoplasm
Q4URA9	RBN_XANC8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q4URC0	ISPE_XANC8		cytoplasm
Q4URC9	RL1_XANC8		cytoplasm
Q4URD0	RL10_XANC8		cytoplasm
Q4URD2	RPOB_XANC8		cytoplasm
Q4URD3	RPOC_XANC8		cytoplasm
Q4URD4	RS12_XANC8		cytoplasm
Q4URD5	RS7_XANC8		cytoplasm
Q4URD8	RS10_XANC8		cytoplasm
Q4URE2	RL2_XANC8		cytoplasm
Q4URE5	RS3_XANC8		cytoplasm
Q4URE6	RL16_XANC8		cytoplasm
Q4URE8	RS17_XANC8		cytoplasm
Q4URF3	RS8_XANC8		cytoplasm
Q4URF6	RS5_XANC8		cytoplasm
Q4URF8	RL15_XANC8		cytoplasm
Q4URG0	RS13_XANC8		cytoplasm
Q4URG1	RS11_XANC8		cytoplasm
Q4URG2	RS4_XANC8		cytoplasm
Q4URG3	RPOA_XANC8		cytoplasm
Q4URH2	MDH_XANC8		cytoplasm
Q4URL4	CLPP_XANC8	cytoplasm (by similarity).	cytoplasm
Q4URP9	RL32_XANC8		cytoplasm
Q4URT6	PURA_XANC8	cytoplasm (by similarity).	cytoplasm
Q4URY1	DNAE2_XANC8	cytoplasm (probable).	cytoplasm
Q4US37	RS20_XANC8		cytoplasm
Q4US40	SYI_XANC8	cytoplasm (by similarity).	cytoplasm
Q4US84	TRMD_XANC8	cytoplasm (potential).	cytoplasm
Q4US85	RL19_XANC8		cytoplasm
Q4US90	MUTS_XANC8		cytoplasm
Q4USF4	LEPA_XANC8		cytoplasm
Q4USF7	RNC_XANC8	cytoplasm (by similarity).	cytoplasm
Q4USF9	RECO_XANC8		cytoplasm
Q4USG1	RUMA_XANC8		cytoplasm
Q4USG7	NAGZ_XANC8	cytoplasm (by similarity).	extracellular
Q4USP4	ACCA_XANC8	cytoplasm (by similarity).	cytoplasm
Q4USP6	RNH2_XANC8	cytoplasm (potential).	extracellular
Q4USP9	FABZ_XANC8	cytoplasm (by similarity).	cytoplasm
Q4USQ4	DXR_XANC8		cytoplasm
Q4USS0	GLND_XANC8		cytoplasm
Q4UT11	DNAK_XANC8		cytoplasm
Q4UT42	SLYX_XANC8		inner membrane
Q4UT86	PPNK_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UTA3	RS6_XANC8		cytoplasm
Q4UTA5	RL9_XANC8		cytoplasm
Q4UTA8	ZIPA_XANC8	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q4UTK2	CCME1_XANC8	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q4UTP4	ISPD_XANC8		cytoplasm
Q4UTP5	ISPF_XANC8		extracellular
Q4UTP6	TRUD_XANC8		cytoplasm
Q4UTP8	SURE_XANC8	cytoplasm (potential).	periplasm
Q4UTQ4	RRMJ_XANC8	cytoplasm (by similarity).	cytoplasm

Q4UTR4	RECA_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UTU6	XYLA1_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UTV6	PANC_XANC8		cytoplasm
Q4UTV7	PAND_XANC8		cytoplasm
Q4UTV8	G6PI_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UTX4	MURB_XANC8	cytoplasm (probable).	inner membrane
Q4UU40	HISX_XANC8		cytoplasm
Q4UU41	HIS8_XANC8		cytoplasm
Q4UU43	HIS5_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UU44	HIS4_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UU45	HIS6_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UU71	DAPB_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UU85	RPFQ_XANC8		cytoplasm
Q4UUH7	FLGI_XANC8	periplasm (by similarity).	periplasm
Q4UUH8	FLGH_XANC8	outer membrane; lipid-anchor (probable).	inner membrane
Q4UUL4	NDK_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UUV5	PHS_XANC8		cytoplasm
Q4UV60	UVRC_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UV64	LPXK_XANC8		cytoplasm
Q4UV74	Y1986_XANC8		cytoplasm
Q4UVA7	Y1953_XANC8		cytoplasm
Q4UVC5	GUAA_XANC8		cytoplasm
Q4UVG6	CCME2_XANC8	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q4UVH1	METX_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UVI0	PROA_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UVI1	PROB_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UVL3	GPH_XANC8		cytoplasm
Q4UVN3	Y1827_XANC8		inner membrane
Q4UVN8	GLUQ_XANC8		cytoplasm
Q4UVX3	QUEA_XANC8	cytoplasm (potential).	cytoplasm
Q4UVY7	HTPG_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UW79	UVRB_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UW99	RS15_XANC8		cytoplasm
Q4UWA0	TRUB_XANC8		cytoplasm
Q4UWA4	Y1603_XANC8		inner membrane
Q4UWC8	GLMM_XANC8		cytoplasm
Q4UWV1	GPMA_XANC8		cytoplasm
Q4UWX9	MURD_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UX36	MURA_XANC8	cytoplasm (probable).	inner membrane
Q4UX84	PDXH_XANC8		cytoplasm
Q4UX85	AROK_XANC8	cytoplasm (probable).	cytoplasm
Q4UX86	AROB_XANC8	cytoplasm (probable).	cytoplasm
Q4UXI4	PQQE_XANC8		cytoplasm
Q4UXL7	RUVB_XANC8		cytoplasm
Q4UXM0	RUVA_XANC8		cytoplasm
Q4UXM1	RUVC_XANC8		cytoplasm
Q4UXV5	RIMK_XANC8		cytoplasm
Q4UXY4	THIG_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UXY5	TRMB_XANC8		cytoplasm
Q4UXZ0	MSCL_XANC8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q4UY42	RPOZ_XANC8		cytoplasm
Q4UYF7	ILVC_XANC8		cytoplasm
Q4UYG2	LEU3_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UYG4	LEUD_XANC8		cytoplasm
Q4UYG5	LEU2_XANC8		cytoplasm
Q4UYN4	BETB_XANC8		cytoplasm
Q4UZF0	NUDH_XANC8		cytoplasm
Q4UZF9	TRPD_XANC8		cytoplasm
Q4UZI9	HGD_XANC8		cytoplasm
Q4UZL8	GLGA_XANC8		cytoplasm
Q4UZP9	GIDA_XANC8	cytoplasm (by similarity).	cytoplasm
Q4UZT2	ILVD_XANC8		cytoplasm
Q4V073	SECB_XANC8	cytoplasm (by similarity).	cytoplasm
Q4V093	UPPP_XANC8	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4V0H2	LLDD_XANC8		cytoplasm
Q4V0R7	PDXJ_XANC8	cytoplasm (probable).	cytoplasm
Q4V0S6	RECF_XANC8	cytoplasm (by similarity).	cytoplasm
Q4W557	KHSE_NEIMB		cytoplasm
Q4W568	TILS_NEIMB	cytoplasm (probable).	cytoplasm
Q4W580	COAE_NEIMB	cytoplasm (by similarity).	cytoplasm
Q4ZL14	GIDB_PSEU2		cytoplasm
Q4ZL22	ATPA_PSEU2	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4ZL23	ATPG_PSEU2		inner membrane
Q4ZL26	GLMU_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZLP9	HIS5_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZLQ1	HIS4_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZLQ2	HIS6_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZLR0	GPMI_PSEU2		cytoplasm
Q4ZLR3	SECB_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZLV2	NUDH_PSEU2		cytoplasm
Q4ZM34	FPG_PSEU2		cytoplasm
Q4ZM55	THIG_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZM56	TRMB_PSEU2		cytoplasm
Q4ZM62	BETB_PSEU2		cytoplasm
Q4ZM83	GLYA2_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZMC1	PQQE_PSEU2		cytoplasm
Q4ZMF5	Y4637_PSEU2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4ZMI2	GPH_PSEU2		cytoplasm
Q4ZML2	TRPD_PSEU2		cytoplasm
Q4ZMM7	SYU_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZMN4	RL1_PSEU2		cytoplasm
Q4ZMN5	RL10_PSEU2		cytoplasm
Q4ZMN7	RPOB_PSEU2		cytoplasm
Q4ZMN8	RPOC_PSEU2		cytoplasm
Q4ZMN9	RS12_PSEU2		cytoplasm
Q4ZMP0	RS7_PSEU2		cytoplasm
Q4ZMP3	RS10_PSEU2		cytoplasm
Q4ZMP7	RL2_PSEU2		cytoplasm
Q4ZMQ0	RS3_PSEU2		cytoplasm
Q4ZMQ1	RL16_PSEU2		cytoplasm

Q4ZMQ3	RS17_PSEU2		cytoplasm
Q4ZMQ8	RS8_PSEU2		cytoplasm
Q4ZMR1	RS5_PSEU2		cytoplasm
Q4ZMR3	RL15_PSEU2		cytoplasm
Q4ZMR5	RS13_PSEU2		cytoplasm
Q4ZMR6	RS11_PSEU2		cytoplasm
Q4ZMR7	RS4_PSEU2		cytoplasm
Q4ZMR8	RPOA_PSEU2		cytoplasm
Q4ZMV1	DEOC_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZMX9	NRDR_PSEU2		cytoplasm
Q4ZMZ1	UREE2_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZN06	URE12_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZN07	URE2_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZN10	URE3_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZN73	PROA_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZNG6	MSCL_PSEU2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4ZNH2	GLYA1_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZNP7	DNAK_PSEU2		cytoplasm
Q4ZNP9	DAPB_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZNP4	RRMJ_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZNR4	TRUB_PSEU2		cytoplasm
Q4ZNR5	RS15_PSEU2		cytoplasm
Q4ZNV7	MURA_PSEU2	cytoplasm (probable).	inner membrane
Q4ZNV8	HIS1_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZNV9	HISX_PSEU2		cytoplasm
Q4ZNW0	HIS8_PSEU2		cytoplasm
Q4ZNY1	MRAZ_PSEU2		inner membrane
Q4ZNY2	MRAW_PSEU2		cytoplasm
Q4ZNY7	MRAY_PSEU2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4ZNY8	MURD_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZNZ0	MURG_PSEU2	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q4ZNZ9	ARGJ_PSEU2	cytoplasm (probable).	cytoplasm
Q4ZPD1	Y3961_PSEU2		cytoplasm
Q4ZPD8	LEPA_PSEU2		cytoplasm
Q4ZPE1	RNC_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZPE3	RECO_PSEU2		cytoplasm
Q4ZPE4	PDXJ_PSEU2	cytoplasm (probable).	cytoplasm
Q4ZPN9	PDXH_PSEU2		cytoplasm
Q4ZQ44	RUMA_PSEU2		cytoplasm
Q4ZQG4	PHS_PSEU2		cytoplasm
Q4ZQR6	FLGH_PSEU2	outer membrane; lipid-anchor (probable).	inner membrane
Q4ZQR7	FLGI_PSEU2	periplasm (by similarity).	periplasm
Q4ZQY7	MTFA_PSEU2		cytoplasm
Q4ZR01	CCME_PSEU2	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q4ZRA0	FADB_PSEU2		cytoplasm
Q4ZRA1	FADA_PSEU2	cytoplasm (by similarity).	cytoplasm

Q4ZS30	UPPP_PSEU2	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q4ZS46	GLGA_PSEU2		cytoplasm
Q4ZSF5	XYLA_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZSZ2	ARNA_PSEU2		cytoplasm
Q4ZSZ4	ARNB_PSEU2		cytoplasm
Q4ZTG0	DNAE2_PSEU2	cytoplasm (probable).	cytoplasm
Q4ZTN0	NUDC_PSEU2		cytoplasm
Q4ZUC9	UREE1_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZUD1	URE23_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZUD2	URE11_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZUP0	Y2089_PSEU2		cytoplasm
Q4ZUW2	HTPG_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZUZ4	LEU3_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZUZ5	LEUD_PSEU2		cytoplasm
Q4ZUZ6	LEU2_PSEU2		cytoplasm
Q4ZV05	UVRB_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZV64	TAL_PSEU2	cytoplasm (probable).	cytoplasm
Q4ZV71	QUEF_PSEU2	cytoplasm (probable).	cytoplasm
Q4ZVE8	TUSA_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZVF7	ZIPA_PSEU2	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q4ZVM7	CLPP_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZVT9	PPNK_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZVY0	RL32_PSEU2		cytoplasm
Q4ZVY7	MURB_PSEU2	cytoplasm (probable).	inner membrane
Q4ZVY9	LPXK_PSEU2		cytoplasm
Q4ZW63	MAO1_PSEU2		cytoplasm
Q4ZWL1	RUVB_PSEU2		cytoplasm
Q4ZWL2	RUVA_PSEU2		cytoplasm
Q4ZWL3	RUVC_PSEU2		cytoplasm
Q4ZWL9	SLYX_PSEU2		inner membrane
Q4ZWP3	RECA_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZWP5	MUTS_PSEU2		cytoplasm
Q4ZWQ0	SURE_PSEU2	cytoplasm (potential).	periplasm
Q4ZWQ1	TRUD_PSEU2		cytoplasm
Q4ZWQ2	ISPF_PSEU2		extracellular
Q4ZWQ6	ISPD_PSEU2		cytoplasm
Q4ZWR2	ACCA_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZWR4	RNH2_PSEU2	cytoplasm (potential).	extracellular
Q4ZWR7	FABZ_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZWS2	DXR_PSEU2		cytoplasm
Q4ZWT0	GLND_PSEU2		cytoplasm
Q4ZWY6	RL19_PSEU2		cytoplasm
Q4ZWY7	TRMD_PSEU2	cytoplasm (potential).	cytoplasm
Q4ZX09	GUAA_PSEU2		cytoplasm
Q4ZX27	NDK_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZXH2	GCSP_PSEU2		cytoplasm
Q4ZXI0	SYV_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZXX1	ISPE_PSEU2		cytoplasm
Q4ZY06	LLDD_PSEU2		cytoplasm
Q4ZY66	ILVC_PSEU2		cytoplasm
Q4ZY80	GLUQ_PSEU2		cytoplasm
Q4ZY87	PANC_PSEU2		cytoplasm

Q4ZY88	G6PI_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZYJ4	SYI_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZYJ7	RS20_PSEU2		cytoplasm
Q4ZYK0	PROB_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZYK5	SELU_PSEU2		cytoplasm
Q4ZYW8	RL9_PSEU2		cytoplasm
Q4ZYX1	RS6_PSEU2		cytoplasm
Q4ZYX5	PURA_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZZ50	CVRA_PSEU2	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q4ZZ78	METX_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZZ83	ILVD_PSEU2		cytoplasm
Q4ZZ95	Y457_PSEU2	periplasm (potential).	periplasm
Q4ZZE3	AROB_PSEU2	cytoplasm (probable).	cytoplasm
Q4ZZE4	AROK_PSEU2	cytoplasm (probable).	cytoplasm
Q4ZZG6	HIS3_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZZG7	HIS2_PSEU2	cytoplasm (by similarity).	cytoplasm
Q4ZZX9	DUT_PSEU2		extracellular
Q4ZZY9	RPOZ_PSEU2		cytoplasm
Q500C6	RIMK_PSEU2		cytoplasm
Q500D0	HSLO_PSEU2	cytoplasm (by similarity).	cytoplasm
Q500D1	PPCK_PSEU2	cytoplasm (by similarity).	cytoplasm
Q500U5	RECF_PSEU2	cytoplasm (by similarity).	cytoplasm
Q50228	FMDA_METME		cytoplasm
Q50229	FMDB_METME	cytoplasm (potential).	periplasm
Q50230	MAUF_METME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q50231	MAUE_METME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q50232	MAUD_METME		inner membrane
Q50234	MAUL_METME		periplasm
Q50235	MAUM_METME		periplasm
Q50400	AZUR_METFL	periplasm (potential).	periplasm
Q50412	RNH2_MAGMM	cytoplasm (potential).	extracellular
Q50414	MAUE_METFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q50422	MAUL_METFL		cytoplasm
Q50423	MAUM_METFL		periplasm
Q50437	PQQB_METFL		cytoplasm
Q50862	RFBA_MYXXA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q50863	RFBB_MYXXA	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q50864	RFBC_MYXXA		inner membrane
Q50926	CYCX_NITEU	periplasm (potential).	periplasm
Q50939	UVRB_NEIGO	cytoplasm (by similarity).	cytoplasm
Q50940	CAH_NEIGO	periplasm (potential).	periplasm
Q50947	LGTB_NEIGO		cytoplasm
Q50950	LGTE_NEIGO		cytoplasm
Q50961	YPIL_NEIGO		cytoplasm

Q50962	COAE_NEIGO	cytoplasm (by similarity).	cytoplasm
Q50966	FBPC_NEIGO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q50967	PSD_NEIGO		cytoplasm
Q50968	UVRA_NEIGO	cytoplasm (by similarity).	cytoplasm
Q50973	T2B1_NEIGO		extracellular
Q50976	T2F7_NEIGO		extracellular
Q50983	CARA_NEIGO		cytoplasm
Q50986	GLMU_NEIGO	cytoplasm (by similarity).	cytoplasm
Q51003	VAPD_NEIGO		extracellular
Q51005	MTGA_NEIGO	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q51008	FUR_NEIGO	cytoplasm (by similarity).	cytoplasm
Q51061	HLDD_NEIGO		cytoplasm
Q51083	T2N3_NEILA		extracellular
Q51116	LGTB_NEIMB		cytoplasm
Q51117	LGTE_NEIMB		cytoplasm
Q51151	LCBA_NEIMB		cytoplasm
Q51152	YHGF_NEIMB		cytoplasm
Q51161	DHPS_NEIMB		cytoplasm
Q51238	TETM_NEIME		cytoplasm
Q51342	PUR1_PSEAE		cytoplasm
Q51344	DHAS_PSEAE		cytoplasm
Q51355	BLC4_PSEAE		extracellular
Q51363	NADB_PSEAE	cytoplasm (by similarity).	cytoplasm
Q51366	GM4D_PSEAE		cytoplasm
Q51368	TONB_PSEAE	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
Q51371	ALGG_PSEAE	periplasm (probable).	extracellular
Q51372	ALGX_PSEAE	periplasm (potential).	periplasm
Q51373	GACA_PSEAE		cytoplasm
Q51382	HSCA_PSEAE		cytoplasm
Q51383	FER_PSEAE		cytoplasm
Q51384	Y3808_PSEAE		cytoplasm
Q51385	Y3806_PSEAE		cytoplasm
Q51389	GLPF_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q51390	GLPK1_PSEAE		cytoplasm
Q51391	GLPR_PSEAE		cytoplasm
Q51397	OPRJ_PSEAE	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q51404	FUMC1_PSEAE	cytoplasm (by similarity).	cytoplasm
Q51416	AMIB_PSEAE		cytoplasm
Q51417	AMIS_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q51422	SYD_PSEAE	cytoplasm (by similarity).	cytoplasm
Q51423	Y964_PSEAE		inner membrane
Q51424	RUVC_PSEAE		cytoplasm
Q51425	RUVA_PSEAE		cytoplasm
Q51426	RUVB_PSEAE		cytoplasm

Q51429	BLO3_PSEAE		extracellular
Q51455	CHEY_PSEAE	cytoplasm (potential).	cytoplasm
Q51462	FLIE_PSEAE		flagellar
Q51463	FLIF_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q51464	FLIG_PSEAE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q51465	FLIM_PSEAE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q51466	FLIN_PSEAE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q51468	FLIP_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q51470	Y3980_PSEAE		cytoplasm
Q51472	IHFA_PSEAE		cytoplasm
Q51473	IHFB_PSEAE		cytoplasm
Q51481	NIRQ_PSEAE	cytoplasm (potential).	cytoplasm
Q51483	Y522_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q51484	Y525_PSEAE	cytoplasm (probable).	cytoplasm
Q51493	NDOA_PSEAE		cytoplasm
Q51494	NDOB_PSEAE		cytoplasm
Q51495	NDOC_PSEAE		cytoplasm
Q51506	SOXR_PSEAE		cytoplasm
Q51507	PCHB_PSEAE		cytoplasm
Q51508	PCHA_PSEAE		cytoplasm
Q51517	PHZC_PSECL		extracellular
Q51518	PHZA_PSECL		cytoplasm
Q51519	PHZB_PSECL		cytoplasm
Q51520	PHZF_PSECL		extracellular
Q51521	PHZD_PSECL		extracellular
Q51522	PHZI_PSECL		cytoplasm
Q51547	PHOU_PSEAE	cytoplasm (by similarity).	cytoplasm
Q51548	PVDA_PSEAE		cytoplasm
Q51551	PYRX_PSEAE		cytoplasm
Q51553	PHAG_PSEAE		cytoplasm
Q51559	RHLA_PSEAE		cytoplasm
Q51561	RPOB_PSEAE		cytoplasm
Q51564	DAPF_PSEAE	cytoplasm (by similarity).	cytoplasm
Q51566	XERC_PSEAE	cytoplasm (by similarity).	nucleoid
Q51567	SUCD_PSEAE		cytoplasm
Q51574	BLOF_PSEAE		extracellular
Q51576	Y3106_PSEAE		cytoplasm
Q51601	CBDA_BURCE		cytoplasm
Q51602	CBDB_BURCE		cytoplasm
Q51603	CBDC_BURCE		cytoplasm
Q51645	HAD4_BURCE		cytoplasm
Q51647	YISX_BURCE		cytoplasm
Q51649	T408_BURCE		cytoplasm
Q51651	REP9_ECOLI		cytoplasm

Q51659	MAUM_PARDE		periplasm
Q51660	MAUN_PARDE		periplasm
Q51662	NORC_PARDE	cell membrane; single-pass membrane protein (by similarity). may be attached to the membrane by a signal- anchor (by similarity).	inner membrane
Q51664	NORQ_PARDE		cytoplasm
Q51669	GFA_PARDE		extracellular
Q51676	HEMN_PARDE	cytoplasm (by similarity).	cytoplasm
Q51687	HIS8_PARDE		cytoplasm
Q51697	IORA_BREDI		cytoplasm
Q51698	IORB_BREDI		cytoplasm
Q51700	NIRS_PARDE	periplasm (by similarity).	periplasm
Q51701	NIRE_PARDE		cytoplasm
Q51703	NIRD_PARDE		cytoplasm
Q51752	CCME_PSEFC	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q51758	EST1_PSEFL		cytoplasm
Q51761	TRA2_PSEFL		cytoplasm
Q51762	ISTB_PSEFL		cytoplasm
Q51769	MERT_PSEFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q51770	MERP_PSEFL	periplasm (probable).	inner membrane
Q51772	MERA_PSEFL		cytoplasm
Q51785	PHZI_PSEFL		cytoplasm
Q51786	PHZR_PSEFL		cytoplasm
Q51787	PHZA_PSEFL		extracellular
Q51788	PHZB_PSEFL		extracellular
Q51789	PHZC_PSEFL		extracellular
Q51790	PHZD_PSEFL		cytoplasm
Q51791	PHZE_PSEFL		cytoplasm
Q51792	PHZF_PSEFL		extracellular
Q51793	PHZG_PSEFL		extracellular
Q51813	1A1D_PSEFL		cytoplasm
Q51856	RBL_PSEHY		extracellular
Q51857	RBS_PSEHY		cytoplasm
Q51858	CBBQ_PSEHY		cytoplasm
Q51872	LUMQ_PHOLE		cytoplasm
Q51886	PAL_PASMU	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q51892	PDEXY_PROMI		cytoplasm
Q51947	NAHE1_PSEPU		cytoplasm
Q51948	NAHD_PSEPU		periplasm
Q51955	PCAK_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q51956	PCAF_PSEPU		cytoplasm
Q51977	CMTB_PSEPU		cytoplasm
Q51983	CMTG_PSEPU		cytoplasm
Q52028	BPHA_PSEPS		cytoplasm
Q52042	CCDB3_ECOLI		cytoplasm
Q52043	CCDB4_ECOLI		cytoplasm

Q52069	NIFS_ENTAG		cytoplasm
Q52070	NIFV_ENTAG		cytoplasm
Q52071	NIFW_ENTAG		cytoplasm
Q52081	FLGH_PSEPU	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q52082	FLGI_PSEPU	periplasm (by similarity).	periplasm
Q52083	FUR_PSEPU	cytoplasm (by similarity).	cytoplasm
Q52086	HADD_PSEPU		extracellular
Q52087	HADL_PSEPU		cytoplasm
Q52095	SYCN_YEREN	or: cytoplasm (by similarity). or: bacterial cell inner membrane; peripheral membrane protein (by similarity). not exported across the inner membrane (by similarity).	cytoplasm
Q52106	MERT_ACICA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q52107	MERP_ACICA	periplasm (probable).	inner membrane
Q52109	MERA_ACICA		cytoplasm
Q52110	MERD_ACICA		cytoplasm
Q52118	YMO3_ERWST		extracellular
Q52126	NDOR_PSEPU		cytoplasm
Q52154	PCAR_PSEPU		cytoplasm
Q52185	POBA_PSEPS		cytoplasm
Q52186	POBB_PSEPS		cytoplasm
Q52219	REP10_ECOLI		cytoplasm
Q52221	REP7_ECOLI		cytoplasm
Q52243	VAPD_ACTAC		extracellular
Q52277	KORC1_ECOLI		cytoplasm
Q52278	KLEA1_ECOLI		cytoplasm
Q52279	KLEB1_ECOLI		cytoplasm
Q52280	KLEE1_ECOLI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52281	KLEF1_ECOLI		cytoplasm
Q52282	KLEG1_ECOLI		cytoplasm
Q52309	AGP_PRORE	periplasm (by similarity).	periplasm
Q52312	INCC1_ECOLI		cytoplasm
Q52313	KORB1_ECOLI		cytoplasm
Q52328	KLAB_ECOLI		cytoplasm
Q52329	KLAC_ECOLI		cytoplasm
Q52331	KORC2_ECOLI		cytoplasm
Q52346	REP8_ECOLI		nucleoid
Q52347	REP11_ECOLI		cytoplasm
Q52353	TERZ_SERMA		cytoplasm
Q52357	TERD_SERMA		cytoplasm
Q52358	TERE_SERMA		cytoplasm
Q52371	HRCN_PSESY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q52374	HRCQA_PSESY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	extracellular
Q52376	GACA_PSEU2		cytoplasm

Q52377	UVRC_PSEU2	cytoplasm (by similarity).	cytoplasm
Q52402	AARD_PROST	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q52419	DAPB_PSESZ	cytoplasm (by similarity).	cytoplasm
Q52420	HRPA_PSESY	extracellular and secreted via type iii secretion system (by similarity).	extracellular
Q52424	AAC2_PROST		cytoplasm
Q52438	BPHA1_PSES1		cytoplasm
Q52439	BPHA2_PSES1		cytoplasm
Q52440	BPHA3_PSES1		cytoplasm
Q52462	NAHD_PSEU8		periplasm
Q52463	ALG8_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52480	HRPA_PSESG	extracellular and secreted via type iii secretion system (by similarity).	extracellular
Q52488	HRCR_RALSO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52498	HRPA_RALSO	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
Q52522	NIRD_PSEST		cytoplasm
Q52523	NIRL_PSEST		inner membrane
Q52524	NIRG_PSEST		cytoplasm
Q52525	NIRH_PSEST		cytoplasm
Q52561	STBC_PSESM		cytoplasm
Q52562	STBB_PSESM		cytoplasm
Q52563	STBA_PSESM		cytoplasm
Q52581	OMPL_PHOPR	bacterial cell outer membrane; multi-pass membrane protein (potential).	outer membrane
Q52620	PQRA_PROVU		cytoplasm
Q52658	SCA4_RICCN	cytoplasm (probable).	cytoplasm
Q52663	BZTA_RHOCA	periplasm (probable).	periplasm
Q52664	BZTB_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q52665	BZTC_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q52666	BZTD_RHOCA	bacterial cell membrane; peripheral membrane protein (potential).	inner membrane
Q52673	RUVX_RHOCA	cytoplasm (potential).	cytoplasm
Q52679	COBT_RHOCA		cytoplasm
Q52683	COBD_RHOCA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52685	BLUB_RHOCA		cytoplasm
Q52686	COBQ_RHOCA		cytoplasm
Q52698	RNC_RHOCA	cytoplasm (by similarity).	cytoplasm
Q52701	DNAK_RHOCA		cytoplasm

Q52702	DNAJ_RHOCA	cytoplasm (probable).	cytoplasm
Q52710	PUTR_RHOCA		cytoplasm
Q52715	RNFD_RHOCA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q52716	RNFC_RHOCA	bacterial cell inner membrane; peripheral membrane protein (probable).	cytoplasm
Q52718	RNFF_RHOCA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52720	SENC_RHOCA		cytoplasm
Q52723	TKT_RHOCA		extracellular
Q52728	PHBC_RHIET	cytoplasm (by similarity).	cytoplasm
Q52733	CCMA_RHIET	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q52741	ROSR_RHIET		cytoplasm
Q52742	YROS_RHIET		cytoplasm
Q52764	17KD_RICJA	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q52772	KLEF2_ECOLI		cytoplasm
Q52778	NOLL_RHILO	bacterial cell membrane; multi-pass membrane protein (potential).	cytoplasm
Q52779	NODD3_RHILO		cytoplasm
Q52813	AAPQ_RHILV	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q52814	AAPM_RHILV	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q52815	AAPP_RHILV		inner membrane
Q52819	CCME_RHILE	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q52822	EXOR_RHILV		cytoplasm
Q52825	AZUP_RHILV	periplasm (by similarity).	periplasm
Q52828	GSTA_RHILE		cytoplasm
Q52838	NODD1_RHILO		cytoplasm
Q52871	YTF3_RHILT		cytoplasm
Q52873	TRA5_RHIME		cytoplasm
Q52877	MCPE_RHIME		cytoplasm
Q52880	CHEA_RHIME	cytoplasm (potential).	inner membrane
Q52881	CHEW_RHIME	cytoplasm (by similarity).	cytoplasm
Q52892	NOEA_RHIME		cytoplasm
Q52893	NOEB_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52894	DEGP_RHIME	periplasm (potential).	periplasm
Q52905	GLNB_RHIME		cytoplasm
Q52910	LNT2_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q52911	Y410_RHIME		cytoplasm

Q52913	ACTR_RHIME		cytoplasm
Q52915	CYA2_RHIME		cytoplasm
Q52923	EXOD_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52926	EXOR_RHIME		cytoplasm
Q52928	EXOI_RHIME		cytoplasm
Q52938	RKPI_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52942	FLAD2_RHIME		flagellar
Q52943	FLAC_RHIME		flagellar
Q52944	FLGC_RHIME		flagellar
Q52945	FLIE_RHIME		flagellar
Q52946	FLGG_RHIME		flagellar
Q52947	FLGA_RHIME	periplasm (probable).	flagellar
Q52948	FLGI_RHIME	periplasm (by similarity).	periplasm
Q52950	FLGH_RHIME	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q52952	MRAY_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q52953	MURD_RHIME	cytoplasm (by similarity).	cytoplasm
Q52955	ILVC_RHIME		cytoplasm
Q52965	PHF2_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52966	Y998_RHIME		inner membrane
Q52967	YA00_RHIME		cytoplasm
Q52968	Y1001_RHIME		inner membrane
Q52969	Y1002_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52975	NOLS_RHIME		cytoplasm
Q52977	NTRB_RHIME		inner membrane
Q52978	PHAAB_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52980	PHAC_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52981	PHAD_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52982	PHAE_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52983	PHAF_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q52984	PHNG_RHIME		cytoplasm
Q52985	PHNH_RHIME		cytoplasm
Q52986	PHNI_RHIME		cytoplasm
Q52987	PHNJ_RHIME		cytoplasm
Q52989	PHOU_RHIME	cytoplasm (by similarity).	cytoplasm
Q52992	YNF7_RHIME		outer membrane

Q52993	NFE1_RHIME		cytoplasm
Q52994	NFE2_RHIME		cytoplasm
Q52995	ECHH_RHIME		cytoplasm
Q52996	Y369_RHIME		cytoplasm
Q52997	Y370_RHIME		cytoplasm
Q52998	TDH_RHIME	cytoplasm (by similarity).	cytoplasm
Q52999	Y2095_RHIME		cytoplasm
Q53009	HADB_BURPI		cytoplasm
Q53016	YREP_BUCRP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q53044	GLNB_RHORT		cytoplasm
Q53046	NIFJ_RHORT		cytoplasm
Q53091	MOAE_RHOS4		cytoplasm
Q53120	PYP_RHOSA		cytoplasm
Q53135	CHEA_RHOSH	cytoplasm (potential).	inner membrane
Q53151	FLIF_RHOSH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q53158	HIS3_RHOS4	cytoplasm (by similarity).	cytoplasm
Q53191	Y4TP_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q53192	Y4TQ_RHISN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q53193	Y4TR_RHISN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q53194	Y4TS_RHISN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q53195	Y4UA_RHISN		cytoplasm
Q53196	Y4UB_RHISN		cytoplasm
Q53197	Y4UC_RHISN		cytoplasm
Q53198	Y4UE_RHISN		cytoplasm
Q53199	DHE3_RHISN		cytoplasm
Q53200	Y4UH_RHISN		cytoplasm
Q53201	Y4UI_RHISN		cytoplasm
Q53202	FIXU_RHISN		cytoplasm
Q53203	NIFZ_RHISN		cytoplasm
Q53204	FDXN_RHISN		periplasm
Q53205	NIFB_RHISN		cytoplasm
Q53206	NIFA_RHISN		cytoplasm
Q53207	FIXX_RHISN		cytoplasm
Q53208	FIXC_RHISN		cytoplasm
Q53209	FIXB_RHISN		cytoplasm
Q53210	FIXA_RHISN		cytoplasm
Q53211	Y4VC_RHISN		inner membrane
Q53212	Y4VD_RHISN		cytoplasm
Q53213	NIFQ_RHISN		cytoplasm
Q53215	Y4VG_RHISN		cytoplasm
Q53216	Y4VH_RHISN		cytoplasm
Q53217	Y4VI_RHISN		cytoplasm
Q53218	Y4VJ_RHISN		cytoplasm
Q53222	BCHF_RHOS4		cytoplasm

Q53226	PYP_RHOS4		cytoplasm
Q53228	REGA_RHOS4		cytoplasm
Q53229	Y2845_RHOS4	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q53238	TREZ_RHIS1		cytoplasm
Q53239	NIR_RHOSH	periplasm (by similarity).	periplasm
Q53245	GLND_RHITR		cytoplasm
Q53464	HAD_PSEUY		cytoplasm
Q53514	NODS_RHITR		cytoplasm
Q53515	NODU_RHITR		inner membrane
Q53547	EST2_PSEFL		cytoplasm
Q53552	NHG2_PSEPU		cytoplasm
Q53562	MMOD_METTR		cytoplasm
Q53563	MMOC_METTR		cytoplasm
Q53636	ENTD_SALAS		periplasm
Q53979	VCOM_SHIDY		cytoplasm
Q54155	DSBB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q54177	TGT_SHIFL		cytoplasm
Q54435	KDTX_SERMA		cytoplasm
Q54462	MERT_SHEPU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q54463	MERP_SHEPU	periplasm (probable).	inner membrane
Q54465	MERA_SHEPU		cytoplasm
Q54478	INH_SERMA	periplasm (by similarity).	periplasm
Q55287	IPGF_SHISO		extracellular
Q55288	MXIJ_SHISO	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q55296	SPAK_SHISO		extracellular
Q56016	KSGA_SALTY		cytoplasm
Q56017	APAG_SALTY		inner membrane
Q56022	SPAO_SALTI		extracellular
Q56023	SPAP_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q56029	PAGD_SALTY		extracellular
Q56030	ENVE_SALTY	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q56032	ENVF_SALTY	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q56035	IACP_SALTY	cytoplasm (by similarity).	cytoplasm
Q56052	INVE_SALTI	probably associated with a bacterial membrane-localized protein (by similarity).	outer membrane
Q56056	RNC_SALTY	cytoplasm (by similarity).	cytoplasm
Q56061	SIFA_SALTY		extracellular
Q56062	PRPB_SALTY		cytoplasm
Q56063	PRPC_SALTY		cytoplasm
Q56064	ENTD_SALTY		periplasm
Q56066	MOEA_SALTY		cytoplasm
Q56067	MOEB_SALTY		cytoplasm

Q56072	EAMA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q56073	DNAK_SALTY		cytoplasm
Q56083	RCSA_SALTI		cytoplasm
Q56093	GALE_SALTI		cytoplasm
Q56106	FLIB_SALTY		cytoplasm
Q56108	NRDH_SALTY		cytoplasm
Q56109	NRDI_SALTY		inner membrane
Q56110	OMPS1_SALTI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q56111	OMPS2_SALTI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q56127	RCSB_SALTI		cytoplasm
Q56128	RCSC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q56131	NLPD_SALTI	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
Q56259	RBL1_THIDA		extracellular
Q56260	RBS_THIDA		cytoplasm
Q56275	GLMS_THIFE	cytoplasm (by similarity).	cytoplasm
Q56349	NAPD_PARP	cytoplasm (potential).	cytoplasm
Q56362	HUPD_THIRO		extracellular
Q56460	MAUE_PARVE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q56461	MAUD_PARVE		inner membrane
Q56463	MAUF_PARVE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q56464	MAUJ_PARVE		cytoplasm
Q56581	DCHS_VIBAN		cytoplasm
Q56582	NQRC_VIBAL	bacterial cell inner membrane (potential).	inner membrane
Q56584	NQRF_VIBAL	bacterial cell inner membrane (potential).	inner membrane
Q56585	BOLA_VIBAL		cytoplasm
Q56586	NQRA_VIBAL		inner membrane
Q56587	NQRB_VIBAL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q56589	NQRE_VIBAL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q56592	VIRA_VIBAN		extracellular
Q56598	GM4D2_VIBCH		cytoplasm
Q56623	GALE_VIBCH		cytoplasm
Q56632	VIBA_VIBCH		cytoplasm
Q56646	VIUB_VIBCH	cytoplasm (potential).	cytoplasm
Q56647	RECX_VIBCH	cytoplasm (potential).	cytoplasm
Q56652	LAMB_VIBCH	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane

Q56691	FRP_VIBHA		cytoplasm
Q56693	PABC_VIBHA		cytoplasm
Q56694	ALDH_VIBHA		cytoplasm
Q56704	FLAG_VIBPA		cytoplasm
Q56732	DHAS_SHESP		cytoplasm
Q56734	DHAS_SHEVI		cytoplasm
Q56738	TPIS_VIBSA		cytoplasm
Q56741	COAE_VIBVU	cytoplasm (by similarity).	cytoplasm
Q56743	VIUB_VIBVU	cytoplasm (potential).	cytoplasm
Q56763	LEP4_XANCP	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q56764	COAE_XANCP	cytoplasm (by similarity).	cytoplasm
Q56797	COPC_XANCP	periplasm (by similarity).	periplasm
Q56815	ALF_XANFL		cytoplasm
Q56826	FLIC_XENNE		flagellar
Q56829	VXS2_XANOR		extracellular
Q56837	XECA_XANP2		extracellular
Q56839	XECC_XANP2		cytoplasm
Q56840	HCDR_XANP2		cytoplasm
Q56841	HCDS_XANP2		cytoplasm
Q56854	FLGN_YEREN		flagellar
Q56855	FES_YEREN	cytoplasm (by similarity).	cytoplasm
Q56872	GM4D_YEREN		cytoplasm
Q56886	FLHB_YEREN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q56888	FLHE_YEREN		inner membrane
Q56892	FLGA_YEREN	periplasm (probable).	flagellar
Q56893	FLGB_YEREN		flagellar
Q56894	FLGC_YEREN		flagellar
Q56897	TRA8_YEREN		cytoplasm
Q56902	RFBD_YEREN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q56903	RFBE_YEREN	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q56927	FBPC_YEREN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q56928	HFQ_YEREN		cytoplasm
Q56953	YFEB_YERPE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q56954	YFEC_YERPE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q56955	YFED_YERPE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q56956	YFEE_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q56957	AIL_YERPS	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q56978	PSAF_YERPE		outer membrane
Q56981	PSAF_YERPS		outer membrane
Q56982	PSAA_YERPS		extracellular
Q56983	PSAC_YERPS	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q56990	HMUS_YERPE		cytoplasm
Q56991	HMUT_YERPE	periplasm (potential).	periplasm
Q56993	HMOV_YERPE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q56998	UVRB_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q57001	ZRP_ZYMMO		cytoplasm
Q57004	HIS82_HAEIN		cytoplasm
Q57006	HSCB_HAEIN		inner membrane
Q57007	Y1107_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57017	Y107_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57020	RNFE_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57022	Y868_HAEIN		cytoplasm
Q57039	6PGL_HAEIN		cytoplasm
Q57045	NUDH_HAEIN		cytoplasm
Q57048	Y020_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q57051	Y588_HAEIN		cytoplasm
Q57060	Y095_HAEIN		cytoplasm
Q57065	Y367_HAEIN		inner membrane
Q57066	Y1720_HAEIN		cytoplasm
Q57070	Y1577_HAEIN		cytoplasm
Q57074	NIFU_HAEIN		cytoplasm
Q57083	PERR_ECOLI		cytoplasm
Q57089	Y1251_HAEIN		cytoplasm
Q57092	RLPA_HAEIN		outer membrane
Q57095	NQRD_VIBAL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q57097	Y118_HAEIN		inner membrane
Q57120	Y948_HAEIN		cytoplasm
Q57122	Y322_HAEIN		cytoplasm
Q57124	Y895_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57125	Y765_HAEIN		cytoplasm
Q57127	Y1453_HAEIN		periplasm
Q57130	Y1471_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane

Q57133	Y973_HAEIN		inner membrane
Q57134	Y1008_HAEIN		cytoplasm
Q57140	NEUA_HAEIN	cytoplasm (potential).	cytoplasm
Q57144	Y461_HAEIN		cytoplasm
Q57147	Y976_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57151	Y1013_HAEIN		cytoplasm
Q57152	Y1436_HAEIN		cytoplasm
Q57154	REPB_ECOLI		cytoplasm
Q57160	HPAB_ECOLI		cytoplasm
Q57163	Y019_HAEIN		cytoplasm
Q57168	T1MH_HAEIN		cytoplasm
Q57180	Y1051_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57184	Y137A_HAEIN		cytoplasm
Q57190	THIL_HAEIN		cytoplasm
Q57194	TUSC_HAEIN	cytoplasm (by similarity).	cytoplasm
Q57195	Y1042_HAEIN		cytoplasm
Q57199	Y1012_HAEIN		cytoplasm
Q57201	Y1457_HAEIN		outer membrane
Q57213	Y1474_HAEIN		inner membrane
Q57223	Y1314_HAEIN	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q57233	THIM_HAEIN		cytoplasm
Q57239	KLAA_ECOLI		cytoplasm
Q57242	UUP1_HAEIN		inner membrane
Q57243	Y1272_HAEIN		inner membrane
Q57251	Y1218_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q57252	Y1163_HAEIN		inner membrane
Q57255	Y1329_HAEIN		cytoplasm
Q57256	Y522_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57259	FLIN_AGRT5	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q57261	TRUD_ECOLI		cytoplasm
Q57270	KLEE2_ECOLI		inner membrane
Q57273	Y1119_HAEIN		cytoplasm
Q57278	MODD_HAEIN		cytoplasm
Q57287	Y1578_HAEIN		cytoplasm
Q57288	RNFD_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q57290	Y740_HAEIN		cytoplasm
Q57293	FBPC_ACTPL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q57300	IMMY_ECOLI		cytoplasm
Q57301	GALE_YEREN		cytoplasm

Q57306	Y355_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q57320	Y1307_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57335	Y036_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q57336	Y261_HAEIN		cytoplasm
Q57337	ISCS_HAEIN		cytoplasm
Q57341	FBPB1_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q57347	Y1049_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57354	Y105_HAEIN		extracellular
Q57362	Y195A_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57368	CORC_HAEIN		inner membrane
Q57374	Y1411_HAEIN		cytoplasm
Q57380	Y1475_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57389	Y680_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57390	LIPA_PELCD	cytoplasm (by similarity).	cytoplasm
Q57392	Y414_HAEIN		outer membrane
Q57394	LIC2B_HAEIN		cytoplasm
Q57399	Y1470_HAEIN		inner membrane
Q57401	FLGM_YEREN		cytoplasm
Q57407	Y1083_HAEIN		cytoplasm
Q57408	HGP4_HAEIN	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
Q57409	Y555_HAEIN		cytoplasm
Q57423	KORA1_ECOLI		cytoplasm
Q57424	Y647_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57425	Y107B_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57427	Y193_HAEIN		cytoplasm
Q57431	Y1278_HAEIN		cytoplasm
Q57434	Y896_HAEIN		cytoplasm
Q57457	Y922_HAEIN	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q57471	TRAM_AGRTU		outer membrane
Q57475	VAPA_BACNO		extracellular
Q57476	CSD_HAEIN		cytoplasm

Q57484	Y867_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57486	Y608_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57491	Y872_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57493	Y092_HAEIN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57498	Y1053_HAEIN		cytoplasm
Q57500	Y894_HAEIN		inner membrane
Q57501	HPAC_ECOLI		cytoplasm
Q57508	CHER_VIBAN		cytoplasm
Q57517	Y053_HAEIN		cytoplasm
Q57519	Y1296_HAEIN		extracellular
Q57523	Y696_HAEIN		outer membrane
Q57525	Y1632_HAEIN		cytoplasm
Q57527	Y1170_HAEIN		cytoplasm
Q57534	Y321_HAEIN		cytoplasm
Q57538	Y664_HAEIN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q57544	Y1663_HAEIN		cytoplasm
Q57549	TPX_HAEIN		cytoplasm
Q576P8	MSRA_BRUAB		cytoplasm
Q576R3	GPMA_BRUAB		cytoplasm
Q576R4	DAPB_BRUAB	cytoplasm (by similarity).	cytoplasm
Q576S3	UPPP_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q576S5	LEPA_BRUAB		cytoplasm
Q576T0	FMT_BRUAB		cytoplasm
Q576T5	DAPD_BRUAB	cytoplasm (by similarity).	cytoplasm
Q577C7	NRDI_BRUAB		inner membrane
Q577H0	GUAA_BRUAB		cytoplasm
Q577Y1	GLMS_BRUAB	cytoplasm (by similarity).	cytoplasm
Q577Y2	GLMU_BRUAB	cytoplasm (by similarity).	cytoplasm
Q578E4	ADEC_BRUAB		cytoplasm
Q578H4	RRMJ_BRUAB	cytoplasm (by similarity).	cytoplasm
Q578I2	FDHD_BRUAB	cytoplasm (potential).	cytoplasm
Q578T1	NIKR_BRUAB		cytoplasm
Q578U0	CRCB4_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q578U1	CRCB3_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q579E8	HUTH_BRUAB	cytoplasm (potential).	cytoplasm
Q579M9	LPXK_BRUAB		cytoplasm
Q579Q6	HIS1_BRUAB	cytoplasm (by similarity).	cytoplasm
Q579T6	FLGH_BRUAB	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q579Z7	Y2386_BRUAB		inner membrane

Q57A81	RS20_BRUAB		cytoplasm
Q57A83	FPG_BRUAB		cytoplasm
Q57A98	TRUB_BRUAB		cytoplasm
Q57AA3	Y2135_BRUAB		inner membrane
Q57AA4	TRMB_BRUAB		cytoplasm
Q57AD6	DNAJ_BRUAB	cytoplasm (probable).	cytoplasm
Q57AD7	DNAK_BRUAB		cytoplasm
Q57AF2	TRPA_BRUAB		cytoplasm
Q57AH2	HIS2_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57AH3	HIS6_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57AH4	HIS4_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57AH5	HIS5_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57AJ7	GIDA_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57AJ8	GIDB_BRUAB		cytoplasm
Q57AM5	ACCA_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57AM8	AROK_BRUAB	cytoplasm (probable).	cytoplasm
Q57AM9	AROB_BRUAB	cytoplasm (probable).	cytoplasm
Q57AR7	HIS8_BRUAB		cytoplasm
Q57AS5	Y1955_BRUAB		inner membrane
Q57AV8	ARGJ_BRUAB	cytoplasm (probable).	cytoplasm
Q57AX1	MDH_BRUAB		cytoplasm
Q57AY4	TRMD_BRUAB	cytoplasm (potential).	cytoplasm
Q57AY9	RL19_BRUAB		cytoplasm
Q57B46	PROB_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57B47	PROA_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57B48	NADD_BRUAB		cytoplasm
Q57B50	Y1819_BRUAB		inner membrane
Q57B54	NUDH_BRUAB		cytoplasm
Q57B86	ATPA_BRUAB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q57B87	ATPG_BRUAB		inner membrane
Q57BA5	ISPG_BRUAB		cytoplasm
Q57BA8	RL32_BRUAB		cytoplasm
Q57BG4	RL31_BRUAB		cytoplasm
Q57BH6	RUVC_BRUAB		cytoplasm
Q57BH7	RUVA_BRUAB		cytoplasm
Q57BH8	RUVB_BRUAB		cytoplasm
Q57BK5	DUT_BRUAB		extracellular
Q57BT1	TRMU_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57BY2	RL25_BRUAB		cytoplasm
Q57C70	MRAW_BRUAB		cytoplasm
Q57C75	MRAY_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57C76	MURD_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57C78	MURG_BRUAB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q57C80	MURB_BRUAB	cytoplasm (probable).	inner membrane
Q57C85	LPXC_BRUAB		cytoplasm
Q57CC2	PDXJ_BRUAB	cytoplasm (probable).	cytoplasm
Q57CC4	KUP_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57CC7	ILVC_BRUAB		cytoplasm

Q57CD6	CRCB2_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57CD7	CRCB1_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57CE8	URE12_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57CE9	URE22_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57CF0	URE32_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57CL3	RADC_BRUAB		cytoplasm
Q57CP5	RL1_BRUAB		cytoplasm
Q57CP8	RPOB_BRUAB		cytoplasm
Q57CP9	RPOC_BRUAB		cytoplasm
Q57CQ3	RS12_BRUAB		cytoplasm
Q57CQ4	RS7_BRUAB		cytoplasm
Q57CQ7	RS10_BRUAB		cytoplasm
Q57CR1	RL2_BRUAB		cytoplasm
Q57CR4	RS3_BRUAB		cytoplasm
Q57CR7	RS17_BRUAB		cytoplasm
Q57CS2	RS8_BRUAB		cytoplasm
Q57CS4	RL18_BRUAB		cytoplasm
Q57CS5	RS5_BRUAB		cytoplasm
Q57CS7	RL15_BRUAB		cytoplasm
Q57CT0	RS13_BRUAB		cytoplasm
Q57CT1	RS11_BRUAB		cytoplasm
Q57CT2	RPOA_BRUAB		cytoplasm
Q57CV7	QUEF_BRUAB	cytoplasm (probable).	cytoplasm
Q57CY0	RRF_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57CY1	UPPS_BRUAB		cytoplasm
Q57CZ5	LEXA_BRUAB		cytoplasm
Q57CZ9	TRPD_BRUAB		cytoplasm
Q57D01	TPIS1_BRUAB		cytoplasm
Q57D18	ISPDF_BRUAB		cytoplasm
Q57D45	QUEA_BRUAB	cytoplasm (potential).	cytoplasm
Q57D60	HIS3_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57DI5	SYU_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57DL3	GID_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57DM1	SURE_BRUAB	cytoplasm (potential).	periplasm
Q57DR5	PURQ_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57DR8	PURL_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57DS5	RS4_BRUAB		cytoplasm
Q57DY1	RISB1_BRUAB		cytoplasm
Q57DY4	NRDR_BRUAB		cytoplasm
Q57DY5	GLYA_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57E41	UVRC_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57E46	NDK_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57E78	RECO_BRUAB		cytoplasm
Q57E81	RNC_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57E83	ACPS_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57E91	RPOZ_BRUAB		cytoplasm
Q57E96	DAPA_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57ED2	CCME_BRUAB	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q57EI0	BETB_BRUAB		cytoplasm

Q57EI4	XYLA_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57EP4	RNH_BRUAB	cytoplasm (potential).	extracellular
Q57EP6	ISPH_BRUAB		cytoplasm
Q57ER2	RS6_BRUAB		cytoplasm
Q57ER5	RL9_BRUAB		cytoplasm
Q57EW9	ISPE_BRUAB		cytoplasm
Q57EX6	RNH2_BRUAB	cytoplasm (potential).	extracellular
Q57F30	PANC_BRUAB		cytoplasm
Q57F40	MSCL_BRUAB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57F73	G6PI_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57F97	MURA_BRUAB	cytoplasm (probable).	inner membrane
Q57F99	HISX_BRUAB		cytoplasm
Q57FG5	THIG_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57FM9	MUTS_BRUAB		cytoplasm
Q57FN0	GLND_BRUAB		cytoplasm
Q57FS2	ILVD_BRUAB		cytoplasm
Q57FT6	PHS_BRUAB		cytoplasm
Q57FU7	DNAE2_BRUAB	cytoplasm (probable).	cytoplasm
Q57FY2	RECR_BRUAB		cytoplasm
Q57G08	RECF_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57G10	DNAA_BRUAB	cytoplasm (by similarity).	cytoplasm
Q57G27	YJX_SALCH		inner membrane
Q57G38	DEOD_SALCH		cytoplasm
Q57G40	TYPH_SALCH		cytoplasm
Q57G60	DNAT_SALCH		nucleoid
Q57GC5	SYV_SALCH	cytoplasm (by similarity).	cytoplasm
Q57GI8	RL9_SALCH		cytoplasm
Q57GJ1	RS6_SALCH		cytoplasm
Q57GJ4	ULAF_SALCH		cytoplasm
Q57GJ5	ULAE_SALCH		cytoplasm
Q57GJ6	ULAD_SALCH		cytoplasm
Q57GJ7	ULAC_SALCH	cytoplasm (probable).	cytoplasm
Q57GJ8	ULAB_SALCH	cytoplasm (potential).	cytoplasm
Q57GJ9	ULAA_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57GK0	ULAG_SALCH	cytoplasm (potential).	extracellular
Q57GK1	ULAR_SALCH	cytoplasm (potential).	cytoplasm
Q57GL4	PURA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57GM7	ORN_SALCH	cytoplasm (potential).	cytoplasm
Q57GM9	PSD_SALCH		cytoplasm
Q57GZ0	LEXA_SALCH		cytoplasm
Q57GZ6	LAMB_SALCH	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q57H01	PSIE_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57H06	G6PI_SALCH	cytoplasm (by similarity).	cytoplasm
Q57H42	ACEK_SALCH	cytoplasm (by similarity).	cytoplasm
Q57H53	ZRAP_SALCH	periplasm (by similarity).	periplasm
Q57H57	NFI_SALCH	cytoplasm (by similarity).	cytoplasm
Q57H59	NUDC_SALCH		cytoplasm
Q57H64	THIG_SALCH	cytoplasm (by similarity).	cytoplasm

Q57H68	RPOC_SALCH		cytoplasm
Q57H69	RPOB_SALCH		cytoplasm
Q57H71	RL10_SALCH		cytoplasm
Q57H72	RL1_SALCH		cytoplasm
Q57H81	MURB_SALCH	cytoplasm (probable).	inner membrane
Q57H87	BTUB_SALCH	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q57H97	CAPP_SALCH		cytoplasm
Q57HC1	RL31_SALCH		cytoplasm
Q57HG6	RHAT_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57HG7	RHAR_SALCH	cytoplasm (by similarity).	cytoplasm
Q57HG8	RHAS_SALCH	cytoplasm (by similarity).	cytoplasm
Q57HG9	RHAB_SALCH		cytoplasm
Q57HH9	FDHD_SALCH	cytoplasm (potential).	cytoplasm
Q57HI9	RBN_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57HL7	RDOA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57HM6	FADB_SALCH		cytoplasm
Q57HM7	FADA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57HQ5	CORA_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57HR5	CYAY_SALCH		inner membrane
Q57HS2	WECG_SALCH		cytoplasm
Q57HS3	WZYE_SALCH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57HS4	WECF_SALCH	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q57HT7	GPPA_SALCH		extracellular
Q57HU2	ILVC_SALCH		cytoplasm
Q57HU7	ILVD_SALCH		cytoplasm
Q57HW3	KUP_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57HW4	RAVA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57HW5	VIAA_SALCH		cytoplasm
Q57HW9	GIDA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57HX0	GIDB_SALCH		cytoplasm
Q57HX7	ATPA_SALCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q57HX8	ATPG_SALCH		inner membrane
Q57HY1	GLMU_SALCH	cytoplasm (by similarity).	cytoplasm
Q57HZ5	MDTL_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57I00	DNAA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57I02	RECF_SALCH	cytoplasm (by similarity).	cytoplasm

Q57I19	CCMAE_SALCH	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q57I24	IBPA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57I25	IBPB_SALCH	cytoplasm (by similarity).	cytoplasm
Q57I28	YIDE_SALCH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57I91	RPOZ_SALCH		cytoplasm
Q57IA1	SLMA_SALCH	nucleoid-associated (by similarity).	cytoplasm
Q57IA2	DUT_SALCH		extracellular
Q57IA4	RADC_SALCH		cytoplasm
Q57IA7	FPG_SALCH		cytoplasm
Q57IC5	TDH_SALCH	cytoplasm (by similarity).	cytoplasm
Q57IC9	GPMI_SALCH		cytoplasm
Q57ID2	SECB_SALCH	cytoplasm (by similarity).	cytoplasm
Q57ID8	LLDD_SALCH		cytoplasm
Q57IG0	XYLA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57IN4	YHIQ_SALCH		inner membrane
Q57IN7	USPB_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57IP3	NIKR_SALCH		cytoplasm
Q57IP9	TUSA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57IU0	GLGC_SALCH		cytoplasm
Q57IU1	GLGA_SALCH		cytoplasm
Q57IW5	BIOH_SALCH	cytoplasm (by similarity).	cytoplasm
Q57IW8	FEOB_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57IX4	PPCK_SALCH	cytoplasm (by similarity).	cytoplasm
Q57IX6	HSLO_SALCH	cytoplasm (by similarity).	cytoplasm
Q57IY7	AROK_SALCH	cytoplasm (probable).	cytoplasm
Q57IY8	AROB_SALCH	cytoplasm (probable).	cytoplasm
Q57IZ2	GPH_SALCH		cytoplasm
Q57J00	TSGA_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57J18	SLYX_SALCH		inner membrane
Q57J21	TUSD_SALCH	cytoplasm (by similarity).	cytoplasm
Q57J22	TUSC_SALCH	cytoplasm (by similarity).	cytoplasm
Q57J23	TUSB_SALCH	cytoplasm (by similarity).	cytoplasm
Q57J24	RS12_SALCH		cytoplasm
Q57J25	RS7_SALCH		cytoplasm
Q57J31	RS10_SALCH		cytoplasm
Q57J35	RL2_SALCH		cytoplasm
Q57J38	RS3_SALCH		cytoplasm
Q57J39	RL16_SALCH		cytoplasm
Q57J40	RL29_SALCH		cytoplasm
Q57J41	RS17_SALCH		cytoplasm
Q57J46	RS8_SALCH		cytoplasm
Q57J48	RL18_SALCH		cytoplasm
Q57J49	RS5_SALCH		cytoplasm
Q57J51	RL15_SALCH		cytoplasm
Q57J54	RS11_SALCH		cytoplasm

Q57J55	RS4_SALCH		cytoplasm
Q57J56	RPOA_SALCH		cytoplasm
Q57J60	MSCL_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57J63	FMT_SALCH		cytoplasm
Q57JA2	AAEX_SALCH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57JA3	AAEA_SALCH	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q57JA4	AAEB_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57JA9	MDH_SALCH		cytoplasm
Q57JG1	MURA_SALCH	cytoplasm (probable).	inner membrane
Q57JH1	RRMJ_SALCH	cytoplasm (by similarity).	cytoplasm
Q57JH3	GLMM_SALCH		cytoplasm
Q57JH7	YHBC_SALCH		inner membrane
Q57JI1	TRUB_SALCH		cytoplasm
Q57JI2	RS15_SALCH		cytoplasm
Q57JP7	MUG_SALCH	cytoplasm (potential).	extracellular
Q57JQ2	YGIH_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57JQ4	UPPP_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57JQ5	CCA_SALCH		cytoplasm
Q57JY0	UXUA_SALCH		cytoplasm
Q57K03	MLTC_SALCH	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q57K04	FETP_SALCH		cytoplasm
Q57K06	TRMB_SALCH		cytoplasm
Q57K36	SPEB_SALCH		cytoplasm
Q57K48	ARGO_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57K69	YGFY_SALCH		cytoplasm
Q57K77	IDI_SALCH	cytoplasm (by similarity).	cytoplasm
Q57KB3	NUDH_SALCH		cytoplasm
Q57KD6	MTFA_SALCH		cytoplasm
Q57KE9	QUEF_SALCH	cytoplasm (probable).	cytoplasm
Q57KF9	RUMA_SALCH		cytoplasm
Q57KH7	CYSJ_SALCH		cytoplasm
Q57KH8	CYSI_SALCH		cytoplasm
Q57KJ4	ISPD_SALCH		cytoplasm
Q57KJ5	ISPF_SALCH		extracellular
Q57KJ6	TRUD_SALCH		cytoplasm
Q57KJ7	SURE_SALCH	cytoplasm (potential).	periplasm
Q57KL5	MUTS_SALCH		cytoplasm
Q57KU4	RECA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57KV3	GSH1_SALCH		cytoplasm
Q57KW6	NRDI_SALCH		inner membrane
Q57L24	PPNK_SALCH	cytoplasm (by similarity).	cytoplasm

Q57L31	TRMD_SALCH	cytoplasm (potential).	cytoplasm
Q57L32	RL19_SALCH		cytoplasm
Q57LC8	LEPA_SALCH		cytoplasm
Q57LD0	RNC_SALCH	cytoplasm (by similarity).	cytoplasm
Q57LD2	RECO_SALCH		cytoplasm
Q57LD3	PDXJ_SALCH	cytoplasm (probable).	cytoplasm
Q57LD4	ACPS_SALCH	cytoplasm (by similarity).	cytoplasm
Q57LF5	HMP_SALCH		cytoplasm
Q57LF7	GLYA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57LH1	ISCA_SALCH		cytoplasm
Q57LH2	HSCB_SALCH		inner membrane
Q57LH5	PEPB_SALCH	cytoplasm (by similarity).	cytoplasm
Q57LI3	NDK_SALCH	cytoplasm (by similarity).	cytoplasm
Q57LI6	ISPG_SALCH		cytoplasm
Q57LJ8	GUAA_SALCH		cytoplasm
Q57LN7	TAL2_SALCH	cytoplasm (probable).	cytoplasm
Q57LT0	ZIPA_SALCH	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q57LW5	FADI_SALCH	cytoplasm (by similarity).	cytoplasm
Q57LW6	FADJ_SALCH	cytoplasm (by similarity).	cytoplasm
Q57LX0	AROC_SALCH		cytoplasm
Q57LX1	MEPA_SALCH	periplasm (by similarity).	extracellular
Q57M43	RNZ_SALCH		extracellular
Q57M54	ARNT_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57M56	ARNC_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57M57	ARNB_SALCH		cytoplasm
Q57M90	ECOT_SALCH	periplasm (by similarity).	periplasm
Q57M92	NAPA_SALCH	periplasm (by similarity).	periplasm
Q57M97	CCMA1_SALCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q57MA1	CCME1_SALCH	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q57MB1	YEJL_SALCH		inner membrane
Q57MB4	RL25_SALCH		cytoplasm
Q57MF5	CDD_SALCH		cytoplasm
Q57MR8	HIS6_SALCH	cytoplasm (by similarity).	cytoplasm
Q57MR9	HIS4_SALCH	cytoplasm (by similarity).	cytoplasm
Q57MS0	HIS5_SALCH	cytoplasm (by similarity).	cytoplasm
Q57MS2	HIS8_SALCH		cytoplasm
Q57MS3	HISX_SALCH		cytoplasm
Q57N17	DSRB_SALCH		inner membrane
Q57N56	UVRC_SALCH	cytoplasm (by similarity).	cytoplasm
Q57N57	PGSA_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57N99	RUVC_SALCH		cytoplasm
Q57NA2	RUVA_SALCH		cytoplasm
Q57NA3	RUVB_SALCH		cytoplasm

Q57NB9	YEBF_SALCH	secreted protein (by similarity).	inner membrane
Q57NH7	YEEN_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57NI8	YOA_H_SALCH		inner membrane
Q57NL1	CVRA_SALCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q57NM8	PTH_SALCH	cytoplasm (by similarity).	cytoplasm
Q57NN2	ISPE_SALCH		cytoplasm
Q57NR0	KITH_SALCH	cytoplasm (potential).	cytoplasm
Q57NT2	TRPA_SALCH		cytoplasm
Q57NU9	GCH2_SALCH		extracellular
Q57P04	ZNTB_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57P88	MAO1_SALCH		cytoplasm
Q57PG8	TUS_SALCH	cytoplasm (by similarity).	cytoplasm
Q57PI3	RNFE_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57PI8	SYI_SALCH	cytoplasm (by similarity).	cytoplasm
Q57PI9	PDXH_SALCH		cytoplasm
Q57PL1	MDTK_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57PT8	YDIA_SALCH		cytoplasm
Q57PU4	BTUD_SALCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q57PU6	BTUC_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57QC0	TRMU_SALCH	cytoplasm (by similarity).	cytoplasm
Q57QC3	PHOP_SALCH	cytoplasm (probable).	cytoplasm
Q57QC4	PHOQ_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57QC7	PEPT_SALCH	cytoplasm (by similarity).	extracellular
Q57QE6	NAGZ_SALCH	cytoplasm (by similarity).	extracellular
Q57QG6	RL32_SALCH		cytoplasm
Q57QH7	FLGI_SALCH	periplasm (by similarity).	periplasm
Q57QH8	FLGH_SALCH	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q57QJ1	MDTH_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57QK1	YCEI_SALCH	periplasm (potential).	periplasm
Q57QK2	YCEA_SALCH		inner membrane
Q57QK4	MDTG_SALCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q57QK9	OPGC_SALCH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q57QM1	YCDX_SALCH		cytoplasm
Q57QR9	TUSE_SALCH	cytoplasm (by similarity).	cytoplasm
Q57QU4	PYRD_SALCH	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q57QZ3	PNCB_SALCH	cytoplasm (by similarity).	cytoplasm
Q57R13	LPXK_SALCH		cytoplasm
Q57R24	SERC_SALCH	cytoplasm (by similarity).	cytoplasm
Q57R56	CLPS_SALCH		cytoplasm
Q57R70	YBJQ_SALCH		inner membrane
Q57R87	RIMK_SALCH		cytoplasm
Q57R91	YBJL_SALCH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q57RD7	DING_SALCH		cytoplasm
Q57RF9	UVRB_SALCH	cytoplasm (by similarity).	cytoplasm
Q57RG6	HUTH_SALCH	cytoplasm (potential).	cytoplasm
Q57RI5	GPMA_SALCH		cytoplasm
Q57RL3	SUCC_SALCH		cytoplasm
Q57RM2	YBGL_SALCH		inner membrane
Q57RR6	RIHA_SALCH		extracellular
Q57RT0	NADD_SALCH		cytoplasm
Q57RT4	YBEA_SALCH		inner membrane
Q57RU6	CRCB_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57RY1	ENTS_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57S51	SELU_SALCH		cytoplasm
Q57S77	HTPG_SALCH	cytoplasm (by similarity).	cytoplasm
Q57S78	RECR_SALCH		cytoplasm
Q57S94	RL31B_SALCH		cytoplasm
Q57SB5	CLPP_SALCH	cytoplasm (by similarity).	cytoplasm
Q57SE9	NRDR_SALCH		cytoplasm
Q57SF9	QUEA_SALCH	cytoplasm (potential).	cytoplasm
Q57SG0	ACPH_SALCH		cytoplasm
Q57SH3	YAIE_SALCH		inner membrane
Q57SH6	AROL_SALCH	cytoplasm (probable).	cytoplasm
Q57SH7	YAIL_SALCH		inner membrane
Q57SN0	LEUD2_SALCH		cytoplasm
Q57SN1	LEU22_SALCH		cytoplasm
Q57ST2	PROA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57ST3	PROB_SALCH	cytoplasm (by similarity).	cytoplasm
Q57ST7	XGPT_SALCH	cytoplasm (by similarity).	cytoplasm
Q57T23	ACCA_SALCH	cytoplasm (by similarity).	cytoplasm
Q57T25	RNH2_SALCH	cytoplasm (potential).	extracellular
Q57T28	FABZ_SALCH	cytoplasm (by similarity).	cytoplasm
Q57T30	SKP_SALCH	periplasm (by similarity).	periplasm
Q57T31	YAET_SALCH	bacterial cell outer membrane (by similarity).	outer membrane
Q57T35	DXR_SALCH		cytoplasm
Q57T36	RRF_SALCH	cytoplasm (by similarity).	cytoplasm
Q57T41	GLND_SALCH		cytoplasm
Q57T42	DAPD_SALCH	cytoplasm (by similarity).	cytoplasm
Q57T44	YAEH_SALCH		inner membrane

Q57T49	BTUF_SALCH	periplasm (by similarity).	periplasm
Q57T70	GLUQ_SALCH		cytoplasm
Q57T74	PANC_SALCH		cytoplasm
Q57T75	PAND_SALCH		cytoplasm
Q57TC9	MURC_SALCH	cytoplasm (probable).	cytoplasm
Q57TD0	MURG_SALCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q57TD2	MURD_SALCH	cytoplasm (by similarity).	cytoplasm
Q57TD3	MRAY_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q57TD8	MRAW_SALCH	cytoplasm (by similarity).	cytoplasm
Q57TD9	MRAZ_SALCH		inner membrane
Q57TE7	LEU3_SALCH	cytoplasm (by similarity).	cytoplasm
Q57TE8	LEU21_SALCH		cytoplasm
Q57TE9	LEUD1_SALCH		cytoplasm
Q57TG7	OSTA_SALCH	bacterial cell outer membrane (by similarity).	outer membrane
Q57TJ0	CAIC_SALCH		cytoplasm
Q57TJ2	CAIE_SALCH		cytoplasm
Q57TJ7	DAPB_SALCH	cytoplasm (by similarity).	cytoplasm
Q57TL0	RIHC_SALCH		extracellular
Q57TL2	ISPH_SALCH		cytoplasm
Q57TL5	SYI_SALCH	cytoplasm (by similarity).	cytoplasm
Q57TL8	RS20_SALCH		cytoplasm
Q57TP2	DNAJ_SALCH	cytoplasm (probable).	cytoplasm
Q57TP3	DNAK_SALCH		cytoplasm
Q57TP8	TAL1_SALCH	cytoplasm (probable).	cytoplasm
Q59083	EXOB_AZOBR		cytoplasm
Q59086	QUIA_ACIAD	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q59087	3DHQ_ACIAD		cytoplasm
Q59088	G6PI_ACIAD	cytoplasm (by similarity).	cytoplasm
Q59091	PCAJ_ACIAD		cytoplasm
Q59092	PCAB_ACIAD		cytoplasm
Q59093	ELH1_ACIAD		cytoplasm
Q59094	SODM_ACIAD	periplasm (probable).	periplasm
Q59097	ODP1_RALEU		cytoplasm
Q59098	ODP2_RALEU		inner membrane
Q59100	ALF1_RALEU		cytoplasm
Q59101	ALF2_RALEU		cytoplasm
Q59102	RBSP_RALEU		cytoplasm
Q59105	NOSZ_RALEU	periplasm (by similarity).	periplasm
Q59106	ODO1_RALEU		cytoplasm
Q59159	OOXB_AGRT4		cytoplasm
Q59160	OOXA_AGRT4		cytoplasm
Q59167	ACSA2_ACEXY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q59168	HDDL_ALCXX		extracellular
Q59169	TRPB_BUCSC		cytoplasm
Q59171	GLK_BRUAB	cytoplasm (by similarity).	cytoplasm
Q59175	OCD_BRUAB		cytoplasm
Q59176	CH10_BUCAP	cytoplasm (by similarity).	cytoplasm

Q59177	CH60_BUCAP	cytoplasm (by similarity).	cytoplasm
Q59178	ARO_A_BUCAP	cytoplasm (probable).	cytoplasm
Q59179	TPIS_BUCAP		cytoplasm
Q59203	CPXP_BRAJA		cytoplasm
Q59204	CPXR_BRAJA		cytoplasm
Q59205	CPXU_BRAJA		cytoplasm
Q59269	DCUP_CAUCR	cytoplasm (probable).	cytoplasm
Q59291	DHAS_CAMJE		cytoplasm
Q59296	CATA_CAMJE		periplasm
Q59342	TNAA_ENTAE		cytoplasm
Q59385	ATCU_ECOLI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q59397	DYR9_ECOLI		cytoplasm
Q59408	DYR13_ECOLI		cytoplasm
Q59418	OGL_ERWCT	periplasm (potential).	periplasm
Q59435	DGTP_ESCFE		cytoplasm
Q59437	BGLA_ENTAG		extracellular
Q59448	SODC_FRATH	periplasm (by similarity).	periplasm
Q59450	RL31_HAEDU		cytoplasm
Q59458	RBL1A_HYDMR		extracellular
Q59459	RBS1_HYDMR		cytoplasm
Q59460	RBL1B_HYDMR		extracellular
Q59461	RBS2_HYDMR		cytoplasm
Q59462	RBL2_HYDMR		extracellular
Q59477	DHAT_KLEPN		cytoplasm
Q59482	DEOD_KLEPN		cytoplasm
Q59514	BLA1_MORCA	bacterial cell membrane; lipid-anchor (probable).	extracellular
Q59536	PTRB_MORLA		cytoplasm
Q59544	APHA_MORMO	periplasm (potential).	periplasm
Q59545	XYLD_MORMO		cytoplasm
Q59599	CARB_NEIGO		cytoplasm
Q59602	CATA_NEIGO		periplasm
Q59603	MTB1_NEIGO		cytoplasm
Q59605	MTB5_NEIGO		cytoplasm
Q59606	MTF7_NEIGO		cytoplasm
Q59613	RBL_NITVU		cytoplasm
Q59614	RBS_NITVU		cytoplasm
Q59622	RPOB_NEIMB		cytoplasm
Q59623	SODC_NEIMB	periplasm (by similarity).	periplasm
Q59633	GALU2_PSEAE		cytoplasm
Q59635	CATB_PSEAE	periplasm (probable).	periplasm
Q59637	ODP1_PSEAE		cytoplasm
Q59638	ODP2_PSEAE		inner membrane
Q59641	PPIA_PSEAE	periplasm (probable).	periplasm
Q59643	HEM2_PSEAE		cytoplasm
Q59647	NORB_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q59649	TRPF_PSEAE		cytoplasm
Q59650	MURE_PSEAE	cytoplasm (probable).	cytoplasm
Q59653	PYRB_PSEAE		cytoplasm
Q59654	PYRF_PSEAE		cytoplasm
Q59662	DHSB_PARDE		cytoplasm
Q59666	HAD_PSEFL		cytoplasm

Q59678	GALE_PASHA		cytoplasm
Q59679	SODM_PASHA		periplasm
Q59680	GPDA_PSELE	cytoplasm (probable).	cytoplasm
Q59683	HEM4_PROMI		cytoplasm
Q59684	HEM3_PROMI		cytoplasm
Q59686	CH10_PASMU	cytoplasm (by similarity).	cytoplasm
Q59687	CH60_PASMU	cytoplasm (by similarity).	cytoplasm
Q59689	SODC_PASMU	periplasm (by similarity).	periplasm
Q59695	ACOC_PSEPU		cytoplasm
Q59699	HYDA_PSEPU		extracellular
Q59701	OCD_AGRT4		cytoplasm
Q59711	PYRB_PSEPU		cytoplasm
Q59712	PYRX_PSEPU		cytoplasm
Q59714	CATA_PSEPU		periplasm
Q59723	CPXO_PSEPU		cytoplasm
Q59727	PHTD_COMTE		cytoplasm
Q59728	HAD9_PSEPU		cytoplasm
Q59735	HEMH_RHOCA	cytoplasm (by similarity).	extracellular
Q59738	BFR_RHOCA		cytoplasm
Q59745	EXOB_RHILT		cytoplasm
Q59746	NOSZ_RHIME	periplasm (by similarity).	periplasm
Q59749	PARC_RHIME	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q59750	BGAL_RHIME		cytoplasm
Q59752	FPG_RHIME		cytoplasm
Q59753	RPOD_RHIME		cytoplasm
Q59754	PPDK_RHIME		cytoplasm
Q59758	DNAA_RICPR	cytoplasm (by similarity).	cytoplasm
Q59787	DHSO_RHOSH		cytoplasm
Q59827	DGTP_SHIBO		cytoplasm
Q5DL43	DAPD_ACIBA	cytoplasm (by similarity).	cytoplasm
Q5DYB5	PDXH_VIBF1		cytoplasm
Q5DYV8	DEOD3_VIBF1		cytoplasm
Q5DZB8	GLGB_VIBF1		extracellular
Q5DZC0	GLGC_VIBF1		cytoplasm
Q5DZC1	GLGA_VIBF1		cytoplasm
Q5DZD2	ACKA2_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5DZF2	Y3774_VIBF1		cytoplasm
Q5DZP0	TKT2_VIBF1		extracellular
Q5DZP1	TAL_VIBF1	cytoplasm (probable).	cytoplasm
Q5E022	Y3554_VIBF1		inner membrane
Q5E0C6	HUTH_VIBF1	cytoplasm (potential).	cytoplasm
Q5E0F1	LPXK_VIBF1		cytoplasm
Q5E0F9	TDH_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E0H4	DEOD2_VIBF1		cytoplasm
Q5E0M1	GAL1_VIBF1	cytoplasm (potential).	cytoplasm
Q5E0T7	COBQ_VIBF1		cytoplasm
Q5E0X3	PEPT_VIBF1	cytoplasm (by similarity).	extracellular
Q5E176	Y3150_VIBF1		inner membrane
Q5E183	GBPA_VIBF1	secreted protein (by similarity).	cytoplasm
Q5E1G0	Y3065_VIBF1		cytoplasm
Q5E1M6	GIDA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E1M7	GIDB_VIBF1		cytoplasm

Q5E1N5	ATPA_VIBF1	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5E1N6	ATPG_VIBF1		inner membrane
Q5E1N9	GLMU_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E1P2	ILVD_VIBF1		cytoplasm
Q5E1Q7	FMT_VIBF1		cytoplasm
Q5E1R1	SMG_VIBF1		cytoplasm
Q5E1R3	PURK_VIBF1		cytoplasm
Q5E1R4	PUR6_VIBF1		cytoplasm
Q5E1S3	ILVC_VIBF1		cytoplasm
Q5E1W5	CYAY_VIBF1		inner membrane
Q5E1X3	PPCK_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E1X4	HSLO_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E209	LEXA_VIBF1		cytoplasm
Q5E225	MURB_VIBF1	cytoplasm (probable).	inner membrane
Q5E238	RPOB_VIBF1		cytoplasm
Q5E257	PUR9_VIBF1		cytoplasm
Q5E260	FIS_VIBF1		cytoplasm
Q5E279	GLMS_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E2A2	SECB_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E2C0	PSD_VIBF1		cytoplasm
Q5E2C2	ORN_VIBF1	cytoplasm (potential).	cytoplasm
Q5E2C6	MUTL_VIBF1		cytoplasm
Q5E2D3	PURA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E2D9	RS6_VIBF1		cytoplasm
Q5E2E1	RL9_VIBF1		cytoplasm
Q5E2E3	CAPP_VIBF1		cytoplasm
Q5E2E5	ARGC_VIBF1	cytoplasm (probable).	cytoplasm
Q5E2E7	ASSY_VIBF1	cytoplasm (probable).	cytoplasm
Q5E2E8	ARLY_VIBF1	cytoplasm (probable).	cytoplasm
Q5E2F9	AROK_VIBF1	cytoplasm (probable).	cytoplasm
Q5E2G0	AROB_VIBF1	cytoplasm (probable).	cytoplasm
Q5E2G4	GPH_VIBF1		cytoplasm
Q5E2H8	RL31_VIBF1		cytoplasm
Q5E2K3	Y2248_VIBF1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5E2K6	UPPP_VIBF1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5E2K7	CCA_VIBF1		cytoplasm
Q5E2P2	MRAW_VIBF1		cytoplasm
Q5E2P7	MRAY_VIBF1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5E2P8	MURD_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E2Q0	MURG_VIBF1	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5E2Q6	LPXC_VIBF1		cytoplasm
Q5E2R0	Y2191_VIBF1		cytoplasm
Q5E2R1	Y2190_VIBF1		inner membrane
Q5E2T1	PANC_VIBF1		cytoplasm
Q5E2W6	GSA_VIBF1	cytoplasm (potential).	cytoplasm
Q5E2Z6	RPIA_VIBF1		cytoplasm

Q5E305	Y2096_VIBF1		cytoplasm
Q5E312	LEPA_VIBF1		cytoplasm
Q5E314	RNC_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E316	RECO_VIBF1		cytoplasm
Q5E317	PDXJ_VIBF1	cytoplasm (probable).	cytoplasm
Q5E318	ACPS_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E320	RUMA_VIBF1		cytoplasm
Q5E326	ENO_VIBF1	cytoplasm (by similarity).	extracellular
Q5E328	ISPD_VIBF1		cytoplasm
Q5E329	ISPF_VIBF1		extracellular
Q5E330	TRUD_VIBF1		cytoplasm
Q5E331	SURE_VIBF1	cytoplasm (potential).	periplasm
Q5E399	SSRP_VIBF1	cytoplasm (potential).	cytoplasm
Q5E3A4	PPNK_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E3A7	DNAK2_VIBF1		cytoplasm
Q5E3A8	DNAJ_VIBF1	cytoplasm (probable).	cytoplasm
Q5E3D1	QUEA_VIBF1	cytoplasm (potential).	cytoplasm
Q5E3D2	TGT_VIBF1		cytoplasm
Q5E3D6	Y1965_VIBF1		inner membrane
Q5E3E2	RRF_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E3E3	UPPS_VIBF1		cytoplasm
Q5E3E5	DXR_VIBF1		cytoplasm
Q5E3F0	FABZ_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E3F3	RNH2_VIBF1	cytoplasm (potential).	extracellular
Q5E3F5	ACCA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E3G5	RNH_VIBF1	cytoplasm (potential).	extracellular
Q5E3I3	DAPA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E3J6	NAPA_VIBF1	periplasm (by similarity).	periplasm
Q5E3L0	ZIPA_VIBF1	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q5E3N0	FLGH_VIBF1	bacterial cell outer membrane; lipid- anchor (probable).	inner membrane
Q5E3N1	FLGI_VIBF1	periplasm (by similarity).	periplasm
Q5E3T1	CCME_VIBF1	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5E3U0	FADI_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E3U1	FADJ_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E3U6	AROC_VIBF1		cytoplasm
Q5E3V8	FTHS_VIBF1		cytoplasm
Q5E3W7	TRMU_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E3Y5	CLPS_VIBF1		cytoplasm
Q5E3Z1	KCY_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E407	RL32_VIBF1		cytoplasm
Q5E408	PLSX_VIBF1		cytoplasm
Q5E416	KTHY_VIBF1		cytoplasm
Q5E424	SYI_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E455	TRUA_VIBF1		cytoplasm
Q5E466	RECR_VIBF1		cytoplasm
Q5E4A2	Y1649_VIBF1		inner membrane
Q5E4C5	UVRC_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E4F2	KITH_VIBF1	cytoplasm (potential).	cytoplasm
Q5E4I0	TORD_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E4J3	MAO1_VIBF1		cytoplasm

Q5E4N9	TPMT_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E4R6	CDD_VIBF1		cytoplasm
Q5E4T4	DNAK1_VIBF1		cytoplasm
Q5E4Y6	NORM_VIBF1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5E5B6	PYRD_VIBF1	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5E624	TRPA_VIBF1		cytoplasm
Q5E633	HIS6_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E634	HIS4_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E635	HIS5_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E637	HIS8_VIBF1		cytoplasm
Q5E638	HISX_VIBF1		cytoplasm
Q5E689	TOLB_VIBF1	periplasm (potential).	periplasm
Q5E699	RUVB_VIBF1		cytoplasm
Q5E6A0	RUVA_VIBF1		cytoplasm
Q5E6A1	RUVC_VIBF1		cytoplasm
Q5E6B5	UVRB_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E6E5	LOLA_VIBF1	periplasm (by similarity).	periplasm
Q5E6F2	SERC_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E6G2	Y889_VIBF1	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q5E6I9	RL25_VIBF1		cytoplasm
Q5E6L4	ACKA1_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E6Q5	CLPP_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E6Q6	TIG_VIBF1		cytoplasm
Q5E6Q9	HTPG_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E6T6	ISPE_VIBF1		cytoplasm
Q5E6U4	Y757_VIBF1		inner membrane
Q5E6V2	Y749_VIBF1		inner membrane
Q5E6V8	LIPB_VIBF1	cytoplasm (potential).	cytoplasm
Q5E6W0	PROA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E6W1	PROB_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E6W3	XGPT_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E6Y5	THII_VIBF1	cytoplasm (potential).	cytoplasm
Q5E6Y8	EX7S_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E702	NRDR_VIBF1		cytoplasm
Q5E706	GLYA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E726	URE3_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E727	URE2_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E728	URE1_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E729	UREE_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E735	NANE_VIBF1		cytoplasm
Q5E743	RIMK_VIBF1		cytoplasm
Q5E763	GUAA_VIBF1		cytoplasm
Q5E772	ISPG_VIBF1		cytoplasm
Q5E776	NDK_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E780	HSCA_VIBF1		cytoplasm
Q5E781	HSCB_VIBF1		inner membrane
Q5E7A0	Y601_VIBF1		inner membrane
Q5E7A3	QUEF_VIBF1	cytoplasm (probable).	cytoplasm
Q5E7A8	MTFA_VIBF1		cytoplasm
Q5E7E1	Y560_VIBF1		inner membrane

Q5E7E2	TRPR_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E7E9	RL19_VIBF1		cytoplasm
Q5E7F0	TRMD_VIBF1	cytoplasm (potential).	cytoplasm
Q5E7F8	GSH1_VIBF1		cytoplasm
Q5E7G6	RECA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E7G7	MUTS_VIBF1		cytoplasm
Q5E7J4	DEOD1_VIBF1		cytoplasm
Q5E7J6	TYPH_VIBF1		cytoplasm
Q5E7L2	RS15_VIBF1		cytoplasm
Q5E7L3	TRUB_VIBF1		cytoplasm
Q5E7L4	RBFA_VIBF1	cytoplasm (potential).	cytoplasm
Q5E7L7	Y484_VIBF1		inner membrane
Q5E7M0	GLMM_VIBF1		cytoplasm
Q5E7M3	RRMJ_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E7N0	DAPB_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E7N1	ISPH_VIBF1		cytoplasm
Q5E7N4	SYI_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E7P1	TYSY_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E7P5	NUDH_VIBF1		cytoplasm
Q5E7Q9	PGK_VIBF1	cytoplasm (potential).	cytoplasm
Q5E7R1	TKT1_VIBF1		extracellular
Q5E7R8	RUVX_VIBF1	cytoplasm (potential).	cytoplasm
Q5E7T0	FETP_VIBF1		cytoplasm
Q5E7U0	SYV_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E7U5	PYRB_VIBF1		cytoplasm
Q5E7U6	PYRI_VIBF1		cytoplasm
Q5E7U8	IDI_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E7V0	MURA_VIBF1	cytoplasm (probable).	inner membrane
Q5E7W7	Y384_VIBF1		cytoplasm
Q5E814	METH_VIBF1		cytoplasm
Q5E840	CYSI_VIBF1		cytoplasm
Q5E841	CYSJ_VIBF1		cytoplasm
Q5E847	G6PI_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E857	LEU3_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E858	LEU2_VIBF1		cytoplasm
Q5E859	LEUD_VIBF1		cytoplasm
Q5E861	DJLA_VIBF1	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q5E862	OSTA_VIBF1	bacterial cell outer membrane (by similarity).	outer membrane
Q5E865	KSGA_VIBF1		cytoplasm
Q5E872	RL27_VIBF1		cytoplasm
Q5E875	MDH_VIBF1		cytoplasm
Q5E889	RPOA_VIBF1		cytoplasm
Q5E890	RS4_VIBF1		cytoplasm
Q5E891	RS11_VIBF1		cytoplasm
Q5E892	RS13_VIBF1		cytoplasm
Q5E897	RS5_VIBF1		cytoplasm
Q5E898	RL18_VIBF1		cytoplasm
Q5E8A0	RS8_VIBF1		cytoplasm
Q5E8A6	RS17_VIBF1		cytoplasm
Q5E8A7	RL29_VIBF1		cytoplasm
Q5E8A8	RL16_VIBF1		cytoplasm
Q5E8A9	RS3_VIBF1		cytoplasm
Q5E8B2	RL2_VIBF1		cytoplasm

Q5E8B7	RS10_VIBF1		cytoplasm
Q5E8C0	RS7_VIBF1		cytoplasm
Q5E8C1	RS12_VIBF1		cytoplasm
Q5E8C2	TUSC_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8C3	TUSD_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8C7	SLYX_VIBF1		inner membrane
Q5E8E5	TPIS_VIBF1		cytoplasm
Q5E8E6	CH60_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8E9	GPMI_VIBF1		cytoplasm
Q5E8L9	COAD_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8M1	FPG_VIBF1		cytoplasm
Q5E8M4	RL28_VIBF1		cytoplasm
Q5E8M5	RADC_VIBF1		cytoplasm
Q5E8M6	COABC_VIBF1		cytoplasm
Q5E8M7	SLMA_VIBF1	nucleoid-associated (by similarity).	cytoplasm
Q5E8N3	BIOH_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8P6	RPOZ_VIBF1		cytoplasm
Q5E8Q0	DTD_VIBF1	cytoplasm (probable).	cytoplasm
Q5E8Q1	RBN_VIBF1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5E8R9	Y082_VIBF1		inner membrane
Q5E8U7	GPPA_VIBF1		extracellular
Q5E8V2	TATA_VIBF1	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q5E8W6	THIG_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8X6	FADB_VIBF1		cytoplasm
Q5E8X7	FADA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8Y2	TUSA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8Z0	RECF_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8Z2	DNAA_VIBF1	cytoplasm (by similarity).	cytoplasm
Q5E8Z7	RNPA_VIBF1		extracellular
Q5F1R7	RS12_HAEPR		cytoplasm
Q5F4W3	RNPA_NEIG1		extracellular
Q5F4X1	RL32_NEIG1		cytoplasm
Q5F4X9	GUAA_NEIG1		cytoplasm
Q5F4Z1	ATPG_NEIG1		inner membrane
Q5F4Z2	ATPA_NEIG1	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5F513	RL31_NEIG1		cytoplasm
Q5F536	CYAY_NEIG1		inner membrane
Q5F541	CH60_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F553	FETP_NEIG1		cytoplasm
Q5F554	Y2082_NEIG1		inner membrane
Q5F556	NADD_NEIG1		cytoplasm
Q5F569	LPXC_NEIG1		cytoplasm
Q5F577	GLMU_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F584	GLMS_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F5A9	CAPP_NEIG1		cytoplasm
Q5F5B2	SLYX_NEIG1		inner membrane
Q5F5C2	THIE_NEIG1		cytoplasm
Q5F5C4	THIG_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F5J4	MUTS_NEIG1		cytoplasm

Q5F5K6	MURA_NEIG1	cytoplasm (probable).	inner membrane
Q5F5M1	DNAJ_NEIG1	cytoplasm (probable).	cytoplasm
Q5F5P7	FMT_NEIG1		cytoplasm
Q5F5Q3	SMG_NEIG1		cytoplasm
Q5F5R2	RL1_NEIG1		cytoplasm
Q5F5R3	RL10_NEIG1		cytoplasm
Q5F5R5	RPOB_NEIG1		cytoplasm
Q5F5R6	RPOC_NEIG1		cytoplasm
Q5F5S1	RS12_NEIG1		cytoplasm
Q5F5S2	RS7_NEIG1		cytoplasm
Q5F5S5	RS10_NEIG1		cytoplasm
Q5F5T0	RL2_NEIG1		cytoplasm
Q5F5T3	RS3_NEIG1		cytoplasm
Q5F5T4	RL16_NEIG1		cytoplasm
Q5F5T5	RL29_NEIG1		cytoplasm
Q5F5T6	RS17_NEIG1		cytoplasm
Q5F5U1	RS8_NEIG1		cytoplasm
Q5F5U3	RL18_NEIG1		cytoplasm
Q5F5U4	RS5_NEIG1		cytoplasm
Q5F5U6	RL15_NEIG1		cytoplasm
Q5F5U9	RS13_NEIG1		cytoplasm
Q5F5V0	RS11_NEIG1		cytoplasm
Q5F5V1	RS4_NEIG1		cytoplasm
Q5F5V2	RPOA_NEIG1		cytoplasm
Q5F5V4	MINC_NEIG1		inner membrane
Q5F5V8	TRUA_NEIG1		cytoplasm
Q5F5W0	SYV_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F5W5	FABZ_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F5X0	DXR_NEIG1		cytoplasm
Q5F5X2	UPPS_NEIG1		cytoplasm
Q5F5X3	RRF_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F5X7	GIDB_NEIG1		cytoplasm
Q5F5X9	RNH2_NEIG1	cytoplasm (potential).	extracellular
Q5F5Y0	GIDA_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F5Y7	DAPB_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F600	Y1768_NEIG1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5F605	PYRD_NEIG1	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5F632	EX7S_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F636	RUVA_NEIG1		cytoplasm
Q5F641	BIOH_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F678	QUEF_NEIG1	cytoplasm (probable).	cytoplasm
Q5F682	RL28_NEIG1		cytoplasm
Q5F685	RL27_NEIG1		cytoplasm
Q5F690	Y1672_NEIG1		cytoplasm
Q5F694	G6PI2_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F695	DAPD_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F6A6	DTD_NEIG1	cytoplasm (probable).	cytoplasm
Q5F6E8	GLUQ_NEIG1		cytoplasm
Q5F6K4	UPPP_NEIG1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5F6K6	MRAZ_NEIG1		inner membrane

Q5F6K7	MRAW_NEIG1		cytoplasm
Q5F6L4	MRAY_NEIG1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5F6L6	MURD_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F6L8	MURG_NEIG1	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5F6P1	PDXJ_NEIG1	cytoplasm (probable).	cytoplasm
Q5F6P2	ACPS_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F6Q6	RS20_NEIG1		cytoplasm
Q5F6R3	SPEB_NEIG1		cytoplasm
Q5F6W5	DNAK_NEIG1		cytoplasm
Q5F746	GLMM_NEIG1		cytoplasm
Q5F753	NUDH_NEIG1		cytoplasm
Q5F756	AROC_NEIG1		cytoplasm
Q5F761	GCSP_NEIG1		cytoplasm
Q5F776	RPOZ_NEIG1		cytoplasm
Q5F792	Y1291_NEIG1		inner membrane
Q5F799	Y1284_NEIG1		inner membrane
Q5F7A0	SERC_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F7B7	Y1261_NEIG1		inner membrane
Q5F7C0	GPMA_NEIG1		cytoplasm
Q5F7D7	HIS8_NEIG1		cytoplasm
Q5F7D8	HISX_NEIG1		cytoplasm
Q5F7E0	HIS1_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F7E5	ILVC_NEIG1		cytoplasm
Q5F7G4	TRMU_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F7H2	PSD_NEIG1		cytoplasm
Q5F7H5	TRPD_NEIG1		cytoplasm
Q5F7I3	ARGJ_NEIG1	cytoplasm (probable).	cytoplasm
Q5F7I8	HSLO_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F7K9	RNH_NEIG1	cytoplasm (potential).	extracellular
Q5F7N6	GCH2_NEIG1		extracellular
Q5F7V9	SURE_NEIG1	cytoplasm (potential).	periplasm
Q5F7W4	GPH_NEIG1		cytoplasm
Q5F816	SSRP_NEIG1	cytoplasm (potential).	cytoplasm
Q5F829	ISPD_NEIG1		cytoplasm
Q5F830	ISPF_NEIG1		extracellular
Q5F831	RPIA_NEIG1		cytoplasm
Q5F836	PNCB_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F849	DAPA_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F858	METX_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F861	RL31B_NEIG1		cytoplasm
Q5F8B4	TRPE_NEIG1		cytoplasm
Q5F8C0	GLYA_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F8C7	Y858_NEIG1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5F8D0	CRCB_NEIG1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5F8D3	PROA_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F8E8	HSCA_NEIG1		cytoplasm
Q5F8F5	ACCA_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F8G6	ILVD_NEIG1		cytoplasm

Q5F8H5	GLND_NEIG1		cytoplasm
Q5F8K5	RECR_NEIG1		cytoplasm
Q5F8K9	CCA_NEIG1		cytoplasm
Q5F8L2	RUVB_NEIG1		cytoplasm
Q5F8M9	RECA_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F8P8	G6PI1_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F8S9	RADC_NEIG1		cytoplasm
Q5F8T1	LEU2_NEIG1		cytoplasm
Q5F8T3	LEUD_NEIG1		cytoplasm
Q5F8T6	LEU3_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F8V2	PDXH_NEIG1		cytoplasm
Q5F8W8	TRUB_NEIG1		cytoplasm
Q5F8Y9	PAND_NEIG1		cytoplasm
Q5F8Z2	ENO_NEIG1	cytoplasm (by similarity).	extracellular
Q5F8Z9	FPG_NEIG1		cytoplasm
Q5F913	ISPG_NEIG1		cytoplasm
Q5F914	CLPP_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F915	TIG_NEIG1		cytoplasm
Q5F922	RL9_NEIG1		cytoplasm
Q5F925	RS6_NEIG1		cytoplasm
Q5F928	UVRC_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F929	TRMB_NEIG1		cytoplasm
Q5F931	UVRB_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F936	RUVX_NEIG1	cytoplasm (potential).	cytoplasm
Q5F9E0	DUT_NEIG1		extracellular
Q5F9F4	RL25_NEIG1		cytoplasm
Q5F9F6	ISPE_NEIG1		cytoplasm
Q5F9F8	PANC_NEIG1		cytoplasm
Q5F9I5	CLPS_NEIG1		cytoplasm
Q5F9J5	PURA_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F9J9	MURB_NEIG1	cytoplasm (probable).	inner membrane
Q5F9K3	PPNK_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F9K6	Y387_NEIG1		inner membrane
Q5F9L0	RRMJ_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F9L4	PTH_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F9P9	LEPA_NEIG1		cytoplasm
Q5F9S5	Y315_NEIG1		cytoplasm
Q5F9U5	TGT_NEIG1		cytoplasm
Q5F9W4	KSGA_NEIG1		cytoplasm
Q5F9X7	RNC_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5F9Y6	TRPA_NEIG1		cytoplasm
Q5F9Z2	LPXK_NEIG1		cytoplasm
Q5F9Z5	KTHY_NEIG1		cytoplasm
Q5FA21	HIS5_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5FA22	HIS4_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5FA23	HIS6_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5FA24	HIS3_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5FA32	Y202_NEIG1		cytoplasm
Q5FA41	RS15_NEIG1		cytoplasm
Q5FA47	HIS2_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5FA58	RIMM_NEIG1	cytoplasm (potential).	cytoplasm
Q5FA59	TRMD_NEIG1	cytoplasm (potential).	cytoplasm
Q5FA60	RL19_NEIG1		cytoplasm
Q5FA76	RUVC_NEIG1		cytoplasm
Q5FA94	NAGZ_NEIG1	cytoplasm (by similarity).	extracellular
Q5FAA9	ARGC_NEIG1	cytoplasm (probable).	cytoplasm

Q5FAB1	SECB_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5FAC7	PBPA_NEIG1	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q5FAD3	AROK_NEIG1	cytoplasm (probable).	cytoplasm
Q5FAD4	AROB_NEIG1	cytoplasm (probable).	cytoplasm
Q5FAF2	ISPH_NEIG1		cytoplasm
Q5FAF5	SYI_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5FAF7	SYI_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5FAG1	FTHS_NEIG1		cytoplasm
Q5FAH3	QUEA_NEIG1	cytoplasm (potential).	cytoplasm
Q5FAH8	ORN_NEIG1	cytoplasm (potential).	cytoplasm
Q5FAH9	GSA_NEIG1	cytoplasm (potential).	cytoplasm
Q5FAJ2	DNAA_NEIG1	cytoplasm (by similarity).	cytoplasm
Q5FB23	URE1_CAMLA	cytoplasm (by similarity).	cytoplasm
Q5FB24	URE23_CAMLA	cytoplasm (by similarity).	cytoplasm
Q5FEH6	RRMJ_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FF10	RUVV_EHRRG		cytoplasm
Q5FF11	Y002_EHRRG	bacterial cell inner membrane; multi- pass membrane protein (by similarity).	inner membrane
Q5FF20	DAPD_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FF40	SYI_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FF55	RS20_EHRRG		cytoplasm
Q5FF66	ATPA_EHRRG	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5FF72	SYV_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FF75	GUAA_EHRRG		cytoplasm
Q5FF92	ISPF_EHRRG		extracellular
Q5FFA2	RL25_EHRRG		cytoplasm
Q5FFA3	PTH_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FFB2	NRDR_EHRRG		cytoplasm
Q5FFD4	RNH2_EHRRG	cytoplasm (potential).	extracellular
Q5FFD8	RPOC_EHRRG		cytoplasm
Q5FFD9	RPOB_EHRRG		cytoplasm
Q5FFE8	RS7_EHRRG		cytoplasm
Q5FFG2	Y202_EHRRG		cytoplasm
Q5FFG3	Y201_EHRRG		inner membrane
Q5FFG4	FMT_EHRRG		cytoplasm
Q5FFG7	CLPP_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FFG8	TIG_EHRRG		cytoplasm
Q5FFL5	ISPH_EHRRG		cytoplasm
Q5FFL8	GPMI_EHRRG		cytoplasm
Q5FFM4	DNAK_EHRRG		cytoplasm
Q5FFQ2	RUVX_EHRRG	cytoplasm (potential).	cytoplasm
Q5FFQ3	PURA_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FFR4	RL18_EHRRG		cytoplasm
Q5FFR9	RS13_EHRRG		cytoplasm
Q5FFS1	RPOA_EHRRG		cytoplasm
Q5FFT0	DAPB_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FFT3	RL32_EHRRG		cytoplasm
Q5FFT9	RS10_EHRRG		cytoplasm
Q5FFU3	RL2_EHRRG		cytoplasm
Q5FFU6	RS3_EHRRG		cytoplasm
Q5FFU7	RL16_EHRRG		cytoplasm

Q5FFU8	RL29_EHRRG		cytoplasm
Q5FFY7	GIDA_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FFZ1	CH60_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FG27	RS6_EHRRG		cytoplasm
Q5FG30	GLYA_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FG35	THYX_EHRRG		cytoplasm
Q5FG39	RUVB_EHRRG		cytoplasm
Q5FG40	RUVA_EHRRG		cytoplasm
Q5FG66	RRF_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FG91	KTHY_EHRRG		cytoplasm
Q5FG94	SECB_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FGD5	ARGC_EHRRG	cytoplasm (probable).	cytoplasm
Q5FGH0	RNC_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FGJ5	RECA_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FGK0	LGT_EHRRG	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5FGN1	NDK_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FGP4	RL19_EHRRG		cytoplasm
Q5FGP5	TRMD_EHRRG	cytoplasm (potential).	cytoplasm
Q5FGP9	Y924_EHRRG		inner membrane
Q5FGQ8	DNAJ_EHRRG	cytoplasm (probable).	cytoplasm
Q5FGS1	TRUA_EHRRG		cytoplasm
Q5FGT9	MDH_EHRRG		cytoplasm
Q5FGW9	PYRB_EHRRG		cytoplasm
Q5FH15	RRMJ_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FH30	KSGA_EHRRG		cytoplasm
Q5FH81	PDXH_EHRRG		cytoplasm
Q5FH91	SYI_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FH93	MRAW_EHRRG		cytoplasm
Q5FHA4	DXR_EHRRG		cytoplasm
Q5FHA6	ISPG_EHRRG		cytoplasm
Q5FHC4	HTPG_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FHE8	MUTS_EHRRG		cytoplasm
Q5FHG0	CCME_EHRRG	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5FHG1	SURE_EHRRG	cytoplasm (potential).	periplasm
Q5FHG7	RPOZ_EHRRG		cytoplasm
Q5FHH4	PDXJ_EHRRG	cytoplasm (probable).	cytoplasm
Q5FHH8	DNAA_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FHK6	RS15_EHRRG		cytoplasm
Q5FHK7	TRUB_EHRRG		cytoplasm
Q5FHL6	ACPS_EHRRG	cytoplasm (by similarity).	cytoplasm
Q5FHM5	ISPE_EHRRG		cytoplasm
Q5FHN8	Y372_EHRRG		inner membrane
Q5FHQ1	LEPA_EHRRG		cytoplasm
Q5FN16	RS20_GLUOX		cytoplasm
Q5FN17	FPG_GLUOX		cytoplasm
Q5FN20	DUT_GLUOX		extracellular
Q5FN26	ILVD_GLUOX		cytoplasm
Q5FN28	SYV_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FNA2	NRDR_GLUOX		cytoplasm

Q5FNB7	MSCL_GLUOX	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5FNB8	RL31_GLUOX		cytoplasm
Q5FND1	RUVX_GLUOX	cytoplasm (potential).	cytoplasm
Q5FNK4	GLYA_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FNL0	PURL_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FNL1	PURQ_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FNM7	TRPD_GLUOX		cytoplasm
Q5FNN5	ENO_GLUOX	cytoplasm (by similarity).	extracellular
Q5FNN8	LEXA_GLUOX		cytoplasm
Q5FNQ1	RRMJ_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FNS2	PDXJ_GLUOX	cytoplasm (probable).	cytoplasm
Q5FNS6	SECB2_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FNT5	THIG_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FNY6	ATPA2_GLUOX	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5FPD9	HIS3_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FPF3	Y2007_GLUOX		inner membrane
Q5FPG9	AROC_GLUOX		cytoplasm
Q5FPH2	PDXH_GLUOX		cytoplasm
Q5FPH6	LPXK_GLUOX		cytoplasm
Q5FPL0	GUA_A_GLUOX		cytoplasm
Q5FPN1	NDK_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FPQ6	CH60_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FPS3	SECB1_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FPT6	GLND_GLUOX		cytoplasm
Q5FPX2	FMT_GLUOX		cytoplasm
Q5FPX3	TRUA_GLUOX		cytoplasm
Q5FPX6	DAPD_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FPY1	RNPA_GLUOX		extracellular
Q5FPY6	FABZ_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FPZ1	DXR_GLUOX		cytoplasm
Q5FPZ4	RRF_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FPZ9	ACPS_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FQ27	RS4_GLUOX		cytoplasm
Q5FQ87	ADEC_GLUOX		cytoplasm
Q5FQ88	PANC_GLUOX		cytoplasm
Q5FQ96	Y1710_GLUOX		inner membrane
Q5FQ98	RPIA_GLUOX		cytoplasm
Q5FQA6	HIS82_GLUOX		cytoplasm
Q5FQB4	GLMM_GLUOX		cytoplasm
Q5FQB6	TILS_GLUOX	cytoplasm (probable).	cytoplasm
Q5FQB9	TOLB_GLUOX	periplasm (potential).	periplasm
Q5FQC4	RUVB_GLUOX		cytoplasm
Q5FQC5	RUVA_GLUOX		cytoplasm
Q5FQC6	RUVC_GLUOX		cytoplasm
Q5FQC7	Y1679_GLUOX		inner membrane
Q5FQD0	ARGJ_GLUOX	cytoplasm (probable).	cytoplasm
Q5FQD6	ISPDF_GLUOX		cytoplasm
Q5FQF4	CCME_GLUOX	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane

Q5FQL0	KUP_GLUOX	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5FQM0	RS15_GLUOX		cytoplasm
Q5FQM1	TRUB_GLUOX		cytoplasm
Q5FQM2	RBFA_GLUOX	cytoplasm (potential).	cytoplasm
Q5FQM6	Y1579_GLUOX		inner membrane
Q5FQP6	ISPE_GLUOX		cytoplasm
Q5FQT4	CLPP2_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FQZ9	PURA_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FR06	QUEA_GLUOX	cytoplasm (potential).	cytoplasm
Q5FR18	THYX_GLUOX		cytoplasm
Q5FR80	SYI_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FRC6	ATPG_GLUOX		inner membrane
Q5FRC7	ATPA1_GLUOX	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5FRE4	FLGH_GLUOX	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q5FRF4	HAM1_GLUOX		extracellular
Q5FRG8	PYRB_GLUOX		cytoplasm
Q5FRH0	Y1265_GLUOX	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5FRI6	SYI_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FRN3	TRPA_GLUOX		cytoplasm
Q5FRR4	HIS81_GLUOX		cytoplasm
Q5FRR8	HUTH_GLUOX	cytoplasm (potential).	cytoplasm
Q5FRS9	Y1150_GLUOX		inner membrane
Q5FRT1	NADD_GLUOX		cytoplasm
Q5FRT2	PROA_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FRT7	PTH_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FRT8	RL25_GLUOX		cytoplasm
Q5FRU4	LGT_GLUOX	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5FRY0	GCSP_GLUOX		cytoplasm
Q5FRY8	ILVC_GLUOX		cytoplasm
Q5FS12	GIDA_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FS13	GIDB_GLUOX		cytoplasm
Q5FS51	HTPG_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FS83	HISX_GLUOX		cytoplasm
Q5FS84	HIS1_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FSH9	Y895_GLUOX		inner membrane
Q5FSK8	DAPB_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FSL4	DNAJ_GLUOX	cytoplasm (probable).	cytoplasm
Q5FSL5	DNAK_GLUOX		cytoplasm
Q5FSM2	MURA_GLUOX	cytoplasm (probable).	inner membrane
Q5FSQ7	Y815_GLUOX		cytoplasm
Q5FT39	TPMT_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FT82	QUEF_GLUOX	cytoplasm (probable).	cytoplasm
Q5FT97	FLGI_GLUOX	periplasm (by similarity).	periplasm
Q5FTB0	CLPS_GLUOX		cytoplasm
Q5FTB6	RECR_GLUOX		cytoplasm
Q5FTN2	HIS2_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FTN3	HIS6_GLUOX	cytoplasm (by similarity).	cytoplasm

Q5FTN5	HIS5_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FTX4	RL1_GLUOX		cytoplasm
Q5FTX5	RL10_GLUOX		cytoplasm
Q5FTX7	RPOB_GLUOX		cytoplasm
Q5FTX8	RPOC_GLUOX		cytoplasm
Q5FTX9	RS12_GLUOX		cytoplasm
Q5FTY0	RS7_GLUOX		cytoplasm
Q5FTY2	RS10_GLUOX		cytoplasm
Q5FTY6	RL2_GLUOX		cytoplasm
Q5FTY9	RS3_GLUOX		cytoplasm
Q5FTZ0	RL16_GLUOX		cytoplasm
Q5FTZ1	RL29_GLUOX		cytoplasm
Q5FTZ2	RS17_GLUOX		cytoplasm
Q5FTZ7	RS8_GLUOX		cytoplasm
Q5FTZ9	RL18_GLUOX		cytoplasm
Q5FU00	RS5_GLUOX		cytoplasm
Q5FU05	RS13_GLUOX		cytoplasm
Q5FU06	RS11_GLUOX		cytoplasm
Q5FU07	RPOA_GLUOX		cytoplasm
Q5FU18	RNH2_GLUOX	cytoplasm (potential).	extracellular
Q5FU29	NUDH_GLUOX		cytoplasm
Q5FU33	GPMI_GLUOX		cytoplasm
Q5FU57	RS6_GLUOX		cytoplasm
Q5FU59	RL9_GLUOX		cytoplasm
Q5FU61	KSGA_GLUOX		cytoplasm
Q5FUB3	EX7S_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FUC2	LEPA_GLUOX		cytoplasm
Q5FUE6	Y211_GLUOX		inner membrane
Q5FUF4	METX_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FUG1	TRMD_GLUOX	cytoplasm (potential).	cytoplasm
Q5FUG2	RL19_GLUOX		cytoplasm
Q5FUG3	LEU2_GLUOX		cytoplasm
Q5FUG4	LEUD_GLUOX		cytoplasm
Q5FUG5	LEU3_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FUH7	ISPH_GLUOX		cytoplasm
Q5FUI1	XGPT_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FUJ3	MURB_GLUOX	cytoplasm (probable).	inner membrane
Q5FUJ5	MURG_GLUOX	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5FUJ7	MURD_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FUJ8	MRAY_GLUOX	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5FUK3	MRAW_GLUOX		cytoplasm
Q5FUK4	MRAZ_GLUOX		inner membrane
Q5FUL1	PROB_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FUL3	RL27_GLUOX		cytoplasm
Q5FUN5	RL32_GLUOX		cytoplasm
Q5FUP5	RL28_GLUOX		cytoplasm
Q5FUR2	TIG_GLUOX		cytoplasm
Q5FUR3	CLPP1_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FUR7	ISPG_GLUOX		cytoplasm
Q5FUS6	UPPP_GLUOX	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q5FUU1	DNAA_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FUX0	SSRP_GLUOX	cytoplasm (potential).	cytoplasm
Q5FUY5	GLMS_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5FUY6	GLMU_GLUOX	cytoplasm (by similarity).	cytoplasm
Q5GRI1	RRF_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GRK0	SYV_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GRK1	DNAJ_WOLTR	cytoplasm (probable).	cytoplasm
Q5GRK4	ISPG_WOLTR		cytoplasm
Q5GRK8	MURB_WOLTR	cytoplasm (probable).	inner membrane
Q5GRL3	UVRB_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GRP0	RECR_WOLTR		cytoplasm
Q5GRP6	MURA_WOLTR	cytoplasm (probable).	inner membrane
Q5GRQ2	RS20_WOLTR		cytoplasm
Q5GRR9	NDK_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GRT1	ATPG_WOLTR		inner membrane
Q5GRT9	RUVC_WOLTR		cytoplasm
Q5GRW6	Y670_WOLTR		inner membrane
Q5GRW9	LEPA_WOLTR		cytoplasm
Q5GRY9	RPOBC_WOLTR		cytoplasm
Q5GRZ3	MRAY_WOLTR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5GRZ4	RNH2_WOLTR	cytoplasm (potential).	extracellular
Q5GS03	Y633_WOLTR		cytoplasm
Q5GS17	CCME_WOLTR	bacterial cell membrane; single-pass type ii membrane protein (potential).	inner membrane
Q5GS20	RL32_WOLTR		cytoplasm
Q5GS25	GIDA_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GS47	FPG_WOLTR		cytoplasm
Q5GS59	RRMJ_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GS82	TIG_WOLTR		cytoplasm
Q5GS83	CLPP_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GS87	SURE_WOLTR	cytoplasm (potential).	periplasm
Q5GS95	SECB_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GSD4	RL9_WOLTR		cytoplasm
Q5GSD6	RS6_WOLTR		cytoplasm
Q5GSE1	DNAK_WOLTR		cytoplasm
Q5GSI7	DAPD_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GSJ3	GUAA_WOLTR		cytoplasm
Q5GSK9	RECA_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GSL3	Y423_WOLTR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5GSM7	ISPDF_WOLTR		cytoplasm
Q5GSN3	GPMI_WOLTR		cytoplasm
Q5GSP9	RPOZ_WOLTR		cytoplasm
Q5GSS3	SYI_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GST1	DUT_WOLTR		extracellular
Q5GST9	RS12_WOLTR		cytoplasm
Q5GSU0	RS7_WOLTR		cytoplasm
Q5GSU7	RL2_WOLTR		cytoplasm
Q5GSV0	RS3_WOLTR		cytoplasm
Q5GSV1	RL16_WOLTR		cytoplasm
Q5GSV3	RS17_WOLTR		cytoplasm
Q5GSW0	RL18_WOLTR		cytoplasm

Q5GSW7	RPOA_WOLTR		cytoplasm
Q5GSX1	ATPA_WOLTR	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5GSY3	RL28_WOLTR		cytoplasm
Q5GT09	DNAA_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GT12	PURA_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GT33	RUVB_WOLTR		cytoplasm
Q5GT37	SYU_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GT39	NUDH_WOLTR		cytoplasm
Q5GT41	MDH_WOLTR		cytoplasm
Q5GT54	RL19_WOLTR		cytoplasm
Q5GT55	TRMD_WOLTR	cytoplasm (potential).	cytoplasm
Q5GTA4	DXR_WOLTR		cytoplasm
Q5GTA6	DAPB_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GTB0	ISPE_WOLTR		cytoplasm
Q5GTE5	HTPG_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GTF3	RIMM_WOLTR	cytoplasm (potential).	cytoplasm
Q5GTH5	MRAW_WOLTR		cytoplasm
Q5GTI3	RNC_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GTI9	PTH_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GTJ0	RL25_WOLTR		cytoplasm
Q5GTJ4	GLMU_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GTK2	SSRP_WOLTR	cytoplasm (potential).	cytoplasm
Q5GTK4	ACPS_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GTN0	FABZ_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GTN6	ISPH_WOLTR		cytoplasm
Q5GTP4	Y038_WOLTR		inner membrane
Q5GTR2	RS15_WOLTR		cytoplasm
Q5GTS7	GLYA_WOLTR	cytoplasm (by similarity).	cytoplasm
Q5GU80	UPPP_XANOR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5GUF2	XYLA2_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GUR8	RHLB_XANOR		cytoplasm
Q5GUT1	CH60_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GUY8	ILVD_XANOR		cytoplasm
Q5GV22	SECB_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GV34	QUEF_XANOR	cytoplasm (probable).	cytoplasm
Q5GV62	SPED_XANOR		cytoplasm
Q5GV68	NUDH_XANOR		cytoplasm
Q5GVE9	HGD_XANOR		cytoplasm
Q5GW07	GLYA_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GW09	NRDR_XANOR		cytoplasm
Q5GW33	MRAZ_XANOR		inner membrane
Q5GW34	MRAW_XANOR		cytoplasm
Q5GW39	MRAY_XANOR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5GW41	MURG_XANOR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5GW46	LPXC_XANOR		cytoplasm
Q5GW71	CAPP_XANOR		cytoplasm

Q5GWB4	LGT_XANOR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5GWB5	TYSY_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GWB9	KSGA_XANOR		cytoplasm
Q5GWC2	OSTA_XANOR	bacterial cell outer membrane (by similarity).	outer membrane
Q5GWE6	MTFA_XANOR		cytoplasm
Q5GWI3	TDH_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GWL8	TAL_XANOR	cytoplasm (probable).	cytoplasm
Q5GWQ2	RBN_XANOR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5GWR3	ISPE_XANOR		cytoplasm
Q5GWR5	RL25_XANOR		cytoplasm
Q5GWR6	PTH_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GWS3	RL1_XANOR		cytoplasm
Q5GWS4	RL10_XANOR		cytoplasm
Q5GWS6	RPOB_XANOR		cytoplasm
Q5GWS8	RS12_XANOR		cytoplasm
Q5GWT2	RS10_XANOR		cytoplasm
Q5GWT7	RL2_XANOR		cytoplasm
Q5GWU0	RS3_XANOR		cytoplasm
Q5GWU1	RL16_XANOR		cytoplasm
Q5GWU2	RL29_XANOR		cytoplasm
Q5GWU3	RS17_XANOR		cytoplasm
Q5GWV0	RL18_XANOR		cytoplasm
Q5GWV1	RS5_XANOR		cytoplasm
Q5GWV6	RS13_XANOR		cytoplasm
Q5GWV8	RS4_XANOR		cytoplasm
Q5GWV9	RPOA_XANOR		cytoplasm
Q5GX75	LEXA1_XANOR		cytoplasm
Q5GX78	DNAE2_XANOR	cytoplasm (probable).	cytoplasm
Q5GXD6	MSCL_XANOR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5GXE4	TRMB_XANOR		cytoplasm
Q5GXE5	THIG_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GXJ5	Y3322_XANOR		inner membrane
Q5GXL9	EX7S_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GXQ3	PSD_XANOR		cytoplasm
Q5GXQ6	AROC_XANOR		cytoplasm
Q5GXR2	TRUA_XANOR		cytoplasm
Q5GXR6	TRPA_XANOR		cytoplasm
Q5GXR8	GLMM_XANOR		cytoplasm
Q5GXU6	Y3221_XANOR		inner membrane
Q5GXV0	RBFA_XANOR	cytoplasm (potential).	cytoplasm
Q5GXV1	TRUB_XANOR		cytoplasm
Q5GXV2	RS15_XANOR		cytoplasm
Q5GXX4	UVRB_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GY22	FETP_XANOR		cytoplasm
Q5GYA0	PHS_XANOR		cytoplasm
Q5GYG0	CCME1_XANOR	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane

Q5GYK4	ENO_XANOR	cytoplasm (by similarity).	extracellular
Q5GYK6	ISPD_XANOR		cytoplasm
Q5GYK7	ISPF_XANOR		extracellular
Q5GYK8	TRUD_XANOR		cytoplasm
Q5GYL1	SURE_XANOR	cytoplasm (potential).	periplasm
Q5GYL7	RRMJ_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GYM5	LEXA2_XANOR		cytoplasm
Q5GYQ7	XYLA1_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GYR2	EFPL_XANOR		cytoplasm
Q5GZ27	FDHD_XANOR	cytoplasm (potential).	cytoplasm
Q5GZ48	Y2769_XANOR		inner membrane
Q5GZ90	GLUQ_XANOR		cytoplasm
Q5GZ95	Y2722_XANOR		inner membrane
Q5GZF1	PROB_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GZF2	PROA_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GZP0	FLGI_XANOR	periplasm (by similarity).	periplasm
Q5GZP1	FLGH_XANOR	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q5GZR3	CLPS_XANOR		cytoplasm
Q5GZS7	CRCB_XANOR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5GZT2	NDK_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GZV7	HTPG_XANOR	cytoplasm (by similarity).	cytoplasm
Q5GZY2	QUEA_XANOR	cytoplasm (potential).	cytoplasm
Q5GZY3	TGT_XANOR		cytoplasm
Q5H051	RS6_XANOR		cytoplasm
Q5H053	RL9_XANOR		cytoplasm
Q5H079	SERC_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H0A0	Y2367_XANOR		inner membrane
Q5H0A4	PANC_XANOR		cytoplasm
Q5H0A6	PAND_XANOR		cytoplasm
Q5H0A7	G6PI_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H0G1	Y2306_XANOR		cytoplasm
Q5H0H1	LPXK_XANOR		cytoplasm
Q5H0H5	UVRC_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H0K6	HIS6_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H0K7	HIS4_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H0K8	HIS5_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H0L0	HIS8_XANOR		cytoplasm
Q5H0L1	HISX_XANOR		cytoplasm
Q5H0N0	PYRD_XANOR	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5H0N1	MURB_XANOR	cytoplasm (probable).	inner membrane
Q5H0N8	ISPG_XANOR		cytoplasm
Q5H0Q5	Y2212_XANOR		cytoplasm
Q5H0Q8	ORN_XANOR	cytoplasm (potential).	cytoplasm
Q5H0S2	GUAA_XANOR		cytoplasm
Q5H0W6	CCME2_XANOR	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5H124	METX_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H185	DNAJ_XANOR	cytoplasm (probable).	cytoplasm
Q5H186	DNAK_XANOR		cytoplasm

Q5H194	SSRP_XANOR	cytoplasm (potential).	cytoplasm
Q5H1D3	GLND_XANOR		cytoplasm
Q5H1E4	RRF_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H1E5	UPPS_XANOR		cytoplasm
Q5H1E7	DXR_XANOR		cytoplasm
Q5H1F1	FABZ_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H1F4	RNH2_XANOR	cytoplasm (potential).	extracellular
Q5H1F6	ACCA_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H1Q0	NAGZ_XANOR	cytoplasm (by similarity).	extracellular
Q5H1Q8	RUMA_XANOR		cytoplasm
Q5H1R0	RECO_XANOR		cytoplasm
Q5H1R2	RNC_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H1R5	LEPA_XANOR		cytoplasm
Q5H233	PQQC_XANOR		cytoplasm
Q5H234	PQQE_XANOR		cytoplasm
Q5H299	TOLB_XANOR	periplasm (potential).	periplasm
Q5H2A4	RUVB_XANOR		cytoplasm
Q5H2A5	KUP_XANOR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5H2A6	RUVA_XANOR		cytoplasm
Q5H2A7	RUVC_XANOR		cytoplasm
Q5H2A8	Y1659_XANOR		inner membrane
Q5H2C5	MUTS_XANOR		cytoplasm
Q5H2D9	ISPH_XANOR		cytoplasm
Q5H2E1	SYI_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H2E4	RS20_XANOR		cytoplasm
Q5H2E6	RL27_XANOR		cytoplasm
Q5H2V7	GPMA_XANOR		cytoplasm
Q5H2Y1	MURD_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H2Z0	PYRB_XANOR		cytoplasm
Q5H2Z1	RUVX_XANOR	cytoplasm (potential).	cytoplasm
Q5H389	RL19_XANOR		cytoplasm
Q5H390	TRMD_XANOR	cytoplasm (potential).	cytoplasm
Q5H391	RIMM_XANOR	cytoplasm (potential).	cytoplasm
Q5H3C7	MURA_XANOR	cytoplasm (probable).	inner membrane
Q5H3D7	Y1280_XANOR		cytoplasm
Q5H3H2	PDXH_XANOR		cytoplasm
Q5H3H3	AROK_XANOR	cytoplasm (probable).	cytoplasm
Q5H3H4	AROB_XANOR	cytoplasm (probable).	cytoplasm
Q5H3K7	RIMK_XANOR		cytoplasm
Q5H3R4	RL31B_XANOR		cytoplasm
Q5H3R9	RPOZ_XANOR		cytoplasm
Q5H3S6	HAM1_XANOR		extracellular
Q5H3U0	RPIA_XANOR		cytoplasm
Q5H401	RECR_XANOR		cytoplasm
Q5H426	RNH_XANOR	cytoplasm (potential).	extracellular
Q5H434	CLPP_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H435	TIG_XANOR		cytoplasm
Q5H496	MDH_XANOR		cytoplasm
Q5H4C1	ILVC_XANOR		cytoplasm
Q5H4C7	LEU3_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H4D0	LEUD_XANOR		cytoplasm
Q5H4D1	LEU2_XANOR		cytoplasm
Q5H4F2	PURA_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H4N5	SYV_XANOR	cytoplasm (by similarity).	cytoplasm

Q5H4Y0	GLMU_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H4Y5	ATPG_XANOR		inner membrane
Q5H4Y6	ATPA_XANOR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5H5B9	DTD_XANOR	cytoplasm (probable).	cytoplasm
Q5H5D0	FMT_XANOR		cytoplasm
Q5H5D4	SMG_XANOR		cytoplasm
Q5H5K1	SYX_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H5L9	DUT_XANOR		extracellular
Q5H5T5	KTHY_XANOR		cytoplasm
Q5H629	Y337_XANOR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5H667	FPG_XANOR		cytoplasm
Q5H6C5	KITH_XANOR	cytoplasm (potential).	cytoplasm
Q5H6D1	BIOH_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H6D9	GIDA_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H6H2	GLGB1_XANOR		extracellular
Q5H6Q3	GLGB2_XANOR		extracellular
Q5H6Q4	GLGA_XANOR		cytoplasm
Q5H6Z4	LLDD_XANOR		cytoplasm
Q5H704	PDXJ_XANOR	cytoplasm (probable).	cytoplasm
Q5H713	RECF_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H715	DNAA_XANOR	cytoplasm (by similarity).	cytoplasm
Q5H9Z6	RL19_EHRRW		cytoplasm
Q5H9Z7	TRMD_EHRRW	cytoplasm (potential).	cytoplasm
Q5HA01	Y882_EHRRW		inner membrane
Q5HA25	NDK_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HA31	RECA_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HA74	RNC_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HA98	ARGC_EHRRW	cytoplasm (probable).	cytoplasm
Q5HAD6	KTHY_EHRRW		cytoplasm
Q5HAD9	SECB_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HAF9	RRF_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HAJ4	RS6_EHRRW		cytoplasm
Q5HAJ7	GLYA_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HAK2	THYX_EHRRW		cytoplasm
Q5HAK4	RUVB_EHRRW		cytoplasm
Q5HAK5	RUVA_EHRRW		cytoplasm
Q5HAN4	GIDA_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HAS1	RS10_EHRRW		cytoplasm
Q5HAS5	RL2_EHRRW		cytoplasm
Q5HAS8	RS3_EHRRW		cytoplasm
Q5HAS9	RL16_EHRRW		cytoplasm
Q5HAT0	RL29_EHRRW		cytoplasm
Q5HAT8	RL18_EHRRW		cytoplasm
Q5HAU3	RS13_EHRRW		cytoplasm
Q5HAU5	RPOA_EHRRW		cytoplasm
Q5HAV4	DAPB_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HAV7	RL32_EHRRW		cytoplasm
Q5HAW7	RUVX_EHRRW	cytoplasm (potential).	cytoplasm
Q5HAW8	PURA_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HAY1	DNAK_EHRRW		cytoplasm
Q5HB13	ISPH_EHRRW		cytoplasm
Q5HB16	GPMI_EHRRW		cytoplasm

Q5HB43	SYI_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HB44	MRAW_EHRRW		cytoplasm
Q5HB55	DXR_EHRRW		cytoplasm
Q5HB57	ISPG_EHRRW		cytoplasm
Q5HBA3	PYRB_EHRRW		cytoplasm
Q5HBA4	TRUA_EHRRW		cytoplasm
Q5HBC6	KSGA_EHRRW		cytoplasm
Q5HBD0	ATPG_EHRRW		inner membrane
Q5HBG4	Y377_EHRRW		inner membrane
Q5HBH4	LEPA_EHRRW		cytoplasm
Q5HBH7	RS15_EHRRW		cytoplasm
Q5HBH8	TRUB_EHRRW		cytoplasm
Q5HBI7	ACPS_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HBJ6	ISPE_EHRRW		cytoplasm
Q5HBM4	CCME_EHRRW	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5HBM5	SURE_EHRRW	cytoplasm (potential).	periplasm
Q5HBN0	RPOZ_EHRRW		cytoplasm
Q5HBN7	PDXJ_EHRRW	cytoplasm (probable).	cytoplasm
Q5HBP0	DNAA_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HBQ7	MUTS_EHRRW		cytoplasm
Q5HBT2	HTPG_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HBX0	Y205_EHRRW		cytoplasm
Q5HBX1	Y204_EHRRW		inner membrane
Q5HBX2	FMT_EHRRW		cytoplasm
Q5HBX5	CLPP_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HBX6	TIG_EHRRW		cytoplasm
Q5HBZ2	PDXH_EHRRW		cytoplasm
Q5HC01	RNH2_EHRRW	cytoplasm (potential).	extracellular
Q5HC04	RPOC_EHRRW		cytoplasm
Q5HC05	RPOB_EHRRW		cytoplasm
Q5HC13	RS7_EHRRW		cytoplasm
Q5HC48	NRDR_EHRRW		cytoplasm
Q5HC74	ISPF_EHRRW		extracellular
Q5HC84	RL25_EHRRW		cytoplasm
Q5HC85	PTH_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HC95	ATPA_EHRRW	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5HC99	SYV_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HCA2	GUAA_EHRRW		cytoplasm
Q5HCB5	SYV_EHRRW	cytoplasm (by similarity).	cytoplasm
Q5HCC9	RS20_EHRRW		cytoplasm
Q5HCG1	RUVC_EHRRW		cytoplasm
Q5HCG2	Y015_EHRRW	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5HCG4	DNAJ_EHRRW	cytoplasm (probable).	cytoplasm
Q5HS68	RUVC_CAMJR		cytoplasm
Q5HS73	QUEF_CAMJR	cytoplasm (probable).	cytoplasm
Q5HS77	LEU3_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HS78	LEU2_CAMJR		cytoplasm
Q5HS79	LEUD_CAMJR		cytoplasm
Q5HS85	KSGA_CAMJR		cytoplasm

Q5HS89	RS10_CAMJR		cytoplasm
Q5HS93	RL2_CAMJR		cytoplasm
Q5HS96	RS3_CAMJR		cytoplasm
Q5HS97	RL16_CAMJR		cytoplasm
Q5HS98	RL29_CAMJR		cytoplasm
Q5HS99	RS17_CAMJR		cytoplasm
Q5HSA4	RS8_CAMJR		cytoplasm
Q5HSA6	RL18_CAMJR		cytoplasm
Q5HSA7	RS5_CAMJR		cytoplasm
Q5HSB7	MURB_CAMJR	cytoplasm (probable).	inner membrane
Q5HSC0	RECA_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HSC1	ENO_CAMJR	cytoplasm (by similarity).	extracellular
Q5HSF7	RNH_CAMJR	cytoplasm (potential).	extracellular
Q5HSF8	RNC_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HSF9	AROC_CAMJR		cytoplasm
Q5HSI0	RS20_CAMJR		cytoplasm
Q5HSI4	ISPDF_CAMJR		cytoplasm
Q5HSI8	HIS6_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HSJ0	HIS4_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HSJ1	HIS52_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HSJ3	HISX_CAMJR		cytoplasm
Q5HSJ4	HIS1_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HSJ6	RPOA_CAMJR		cytoplasm
Q5HSJ7	RS4_CAMJR		cytoplasm
Q5HSJ8	RS11_CAMJR		cytoplasm
Q5HSJ9	RS13_CAMJR		cytoplasm
Q5HSP5	Y1712_CAMJR		inner membrane
Q5HST3	PURA_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HSW8	FLGI_CAMJR	periplasm (by similarity).	periplasm
Q5HSX3	TRUD_CAMJR		cytoplasm
Q5HSX8	TILS_CAMJR	cytoplasm (probable).	cytoplasm
Q5HT08	ACPS_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HT13	NADD_CAMJR		cytoplasm
Q5HT15	PGK_CAMJR	cytoplasm (potential).	cytoplasm
Q5HT16	TPIS_CAMJR		cytoplasm
Q5HT36	HAM1_CAMJR		extracellular
Q5HT48	RUVB_CAMJR		cytoplasm
Q5HT65	DXR_CAMJR		cytoplasm
Q5HT90	HIS51_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HTB5	DCD_CAMJR		extracellular
Q5HTI5	TRMB_CAMJR		cytoplasm
Q5HTJ0	RPOZ_CAMJR		cytoplasm
Q5HTJ2	SYI_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HTK0	RECR_CAMJR		cytoplasm
Q5HTK3	DNAJ_CAMJR	cytoplasm (probable).	cytoplasm
Q5HTL1	PUR2_CAMJR		cytoplasm
Q5HTL3	GUAA_CAMJR		cytoplasm
Q5HTL6	UVRC_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HTM5	PDXJ_CAMJR	cytoplasm (probable).	cytoplasm
Q5HTP2	CH60_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HTP3	CH10_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HTR6	LUXS_CAMJR		extracellular
Q5HTR7	GATB_CAMJR		cytoplasm
Q5HTS6	GIDA_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HTU2	Y1306_CAMJR		inner membrane
Q5HTZ7	CLPS_CAMJR		cytoplasm

Q5HTZ9	SSRP_CAMJR	cytoplasm (potential).	cytoplasm
Q5HU00	ISPE_CAMJR		cytoplasm
Q5HU02	TRUB_CAMJR		cytoplasm
Q5HU06	PYRB_CAMJR		cytoplasm
Q5HU23	THIE_CAMJR		cytoplasm
Q5HU34	RS6_CAMJR		cytoplasm
Q5HU41	SYI_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HU43	GATA_CAMJR		cytoplasm
Q5HU47	MURC_CAMJR	cytoplasm (probable).	cytoplasm
Q5HU51	NPD_CAMJR	cytoplasm (probable).	cytoplasm
Q5HU56	THIG_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HU62	MURG_CAMJR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5HU70	LEPA_CAMJR		cytoplasm
Q5HUF2	TGT_CAMJR		cytoplasm
Q5HUF4	AROB_CAMJR	cytoplasm (probable).	cytoplasm
Q5HUG5	GIDB_CAMJR		cytoplasm
Q5HUU9	RNPA_CAMJR		extracellular
Q5HUK4	PURL_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HUK6	PUR9_CAMJR		cytoplasm
Q5HUM7	PPCK_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HUM8	ARLY_CAMJR	cytoplasm (probable).	cytoplasm
Q5HUR4	ISPH_CAMJR		cytoplasm
Q5HUS3	RS15_CAMJR		cytoplasm
Q5HUT8	MURA_CAMJR	cytoplasm (probable).	inner membrane
Q5HUU3	GSA_CAMJR	cytoplasm (potential).	cytoplasm
Q5HUW9	TRUA_CAMJR		cytoplasm
Q5HUX2	UPPS_CAMJR		cytoplasm
Q5HUX5	GLMU_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HUY1	LPXK_CAMJR		cytoplasm
Q5HUY6	DAPA_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HUZ3	RUVA_CAMJR		cytoplasm
Q5HV12	NAPA_CAMJR	periplasm (by similarity).	periplasm
Q5HV17	SYV_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HV25	COAD_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HV26	KTHY_CAMJR		cytoplasm
Q5HV33	DNAK_CAMJR		cytoplasm
Q5HV55	CORA_CAMJR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5HV67	RL19_CAMJR		cytoplasm
Q5HV68	TRMD_CAMJR	cytoplasm (potential).	cytoplasm
Q5HV69	RIMM_CAMJR	cytoplasm (potential).	cytoplasm
Q5HV88	MRAW_CAMJR		cytoplasm
Q5HV92	ACKA_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HV94	FLGH_CAMJR	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q5HV95	ISPG_CAMJR		cytoplasm
Q5HVA1	UVRB_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HVA9	ASSY_CAMJR	cytoplasm (probable).	cytoplasm
Q5HVB0	RL9_CAMJR		cytoplasm
Q5HVB2	HSLU_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HVD0	PPNK_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HVD4	MSRA_CAMJR		cytoplasm
Q5HVD6	RUVX_CAMJR	cytoplasm (potential).	cytoplasm

Q5HVD9	ILVC_CAMJR		cytoplasm
Q5HVM4	HEM1_CAMJR		cytoplasm
Q5HVP5	HTPG_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HVP6	CRCB_CAMJR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5HVQ1	PUR7_CAMJR		cytoplasm
Q5HVR3	TRPC_CAMJR		cytoplasm
Q5HVX7	RS7_CAMJR		cytoplasm
Q5HVX8	RS12_CAMJR		cytoplasm
Q5HVV8	RPOC_CAMJR		cytoplasm
Q5HVV9	RPOB_CAMJR		cytoplasm
Q5HVZ1	RL10_CAMJR		cytoplasm
Q5HVZ2	RL1_CAMJR		cytoplasm
Q5HW14	THIC_CAMJR		cytoplasm
Q5HW17	RL28_CAMJR		cytoplasm
Q5HW21	ACCA_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HW32	GPMI_CAMJR		cytoplasm
Q5HW33	MRAY_CAMJR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5HW34	MURD_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HW60	LGT_CAMJR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5HW65	GLYA_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HW69	GATC_CAMJR		cytoplasm
Q5HW83	KDSA_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HW85	NUSB_CAMJR		cytoplasm
Q5HW96	Y420_CAMJR	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q5HWA7	GLMM_CAMJR		cytoplasm
Q5HWB0	Y406_CAMJR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5HWB8	TRPA_CAMJR		cytoplasm
Q5HWB9	TRPB_CAMJR		cytoplasm
Q5HWD9	NDK_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HWE2	PLSX_CAMJR		cytoplasm
Q5HWE5	SERC_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HWF4	HIS8_CAMJR		cytoplasm
Q5HWF9	PTH_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HWG0	RL25_CAMJR		cytoplasm
Q5HWH3	PANB_CAMJR		cytoplasm
Q5HWH4	PANC_CAMJR		cytoplasm
Q5HWH5	PAND_CAMJR		cytoplasm
Q5HWH7	SURE_CAMJR	cytoplasm (potential).	periplasm
Q5HWI0	GREA_CAMJR		cytoplasm
Q5HWJ3	FABZ_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HWM8	RRF_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HWN6	ARGB_CAMJR	cytoplasm (probable).	cytoplasm
Q5HWN8	ARGC_CAMJR	cytoplasm (probable).	cytoplasm
Q5HWW4	UPPP_CAMJR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5HWX1	DAPB_CAMJR	cytoplasm (by similarity).	cytoplasm

Q5HWX5	TIG_CAMJR		cytoplasm
Q5HWX6	CLPP_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HWZ9	Y162_CAMJR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5HX10	RL31_CAMJR		cytoplasm
Q5HX17	Y144_CAMJR		inner membrane
Q5HX28	Y133_CAMJR		inner membrane
Q5HX29	RBFA_CAMJR	cytoplasm (potential).	cytoplasm
Q5HX34	LPXC_CAMJR		cytoplasm
Q5HX40	Y121_CAMJR		inner membrane
Q5HX45	Y116_CAMJR		inner membrane
Q5HX54	TOLB_CAMJR	periplasm (potential).	periplasm
Q5HX60	ATPG_CAMJR		inner membrane
Q5HX61	ATPA_CAMJR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5HX68	FMT_CAMJR		cytoplasm
Q5HX71	RL27_CAMJR		cytoplasm
Q5HX80	UNG_CAMJR	cytoplasm (by similarity).	extracellular
Q5HXB2	TRMU_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HXD2	THYX_CAMJR		cytoplasm
Q5HXE4	ILVD_CAMJR		cytoplasm
Q5HXE7	RNH2_CAMJR	cytoplasm (potential).	extracellular
Q5HXF5	DNAA_CAMJR	cytoplasm (by similarity).	cytoplasm
Q5HXN6	CRCB_GLUOX	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5HXU1	DNAE2_GLUOX	cytoplasm (probable).	cytoplasm
Q5LL27	HISX3_SILPO		cytoplasm
Q5LL36	CCME2_SILPO	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5LL90	MURA_SILPO	cytoplasm (probable).	inner membrane
Q5LLE7	Y4450_SILPO		cytoplasm
Q5LLI6	CRCB_SILPO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5LLK8	TRMB_SILPO		cytoplasm
Q5LLL4	Y4008_SILPO		inner membrane
Q5LLN4	SECB_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LLP0	HSLU_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LLT1	RS15_SILPO		cytoplasm
Q5LLT4	TRUB_SILPO		cytoplasm
Q5LLT6	RBFA_SILPO	cytoplasm (potential).	cytoplasm
Q5LLT7	DAPB_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LLU4	Y3827_SILPO		inner membrane
Q5LLU5	RS10_SILPO		cytoplasm
Q5LLW1	GPMI_SILPO		cytoplasm
Q5LLW5	SSRP_SILPO	cytoplasm (potential).	cytoplasm
Q5LLZ3	UPPP_SILPO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5LM03	SFSA_SILPO		cytoplasm
Q5LM10	QUEF_SILPO	cytoplasm (probable).	cytoplasm

Q5LM18	ADEC_SILPO		cytoplasm
Q5LM31	PHS_SILPO		cytoplasm
Q5LMA4	Y3659_SILPO		inner membrane
Q5LMA7	CLPS_SILPO		cytoplasm
Q5LMC7	UVRC_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LME7	ACCA_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LMH8	NUDH_SILPO		cytoplasm
Q5LMJ4	RECR_SILPO		cytoplasm
Q5LMK7	DAPA_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LMQ0	RL1_SILPO		cytoplasm
Q5LMQ3	RL10_SILPO		cytoplasm
Q5LMQ5	RPOB_SILPO		cytoplasm
Q5LMQ6	RPOC_SILPO		cytoplasm
Q5LMR2	RS12_SILPO		cytoplasm
Q5LMR3	RS7_SILPO		cytoplasm
Q5LMR7	MSCL_SILPO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5LMW0	RNH2_SILPO	cytoplasm (potential).	extracellular
Q5LMW7	LPXK_SILPO		cytoplasm
Q5LN53	TDH_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LN75	DAPD_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LN98	ILVD_SILPO		cytoplasm
Q5LNE6	TRMD_SILPO	cytoplasm (potential).	cytoplasm
Q5LNE8	RL19_SILPO		cytoplasm
Q5LNE9	RL31_SILPO		cytoplasm
Q5LNF0	RIMM_SILPO	cytoplasm (potential).	cytoplasm
Q5LNI8	FMT_SILPO		cytoplasm
Q5LNJ2	RNH_SILPO	cytoplasm (potential).	extracellular
Q5LNJ7	ISPH_SILPO		cytoplasm
Q5LNK0	RPOZ_SILPO		cytoplasm
Q5LNK3	ACPS_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LNK5	RNC_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LNK7	RECO_SILPO		cytoplasm
Q5LNM2	RS4_SILPO		cytoplasm
Q5LNM6	HIS8_SILPO		cytoplasm
Q5LNN9	ATPA_SILPO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5LNP0	ATPG_SILPO		inner membrane
Q5LNR0	TRUA_SILPO		cytoplasm
Q5LNR7	SYI_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LNT6	RUVC_SILPO		cytoplasm
Q5LNT7	RUVA_SILPO		cytoplasm
Q5LNT8	RUVB_SILPO		cytoplasm
Q5LNU4	TOLB_SILPO	periplasm (potential).	periplasm
Q5LNU7	TILS_SILPO	cytoplasm (probable).	cytoplasm
Q5LNV0	FTHS_SILPO		cytoplasm
Q5LP30	SYV_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LP58	DTD_SILPO	cytoplasm (probable).	cytoplasm
Q5LPA8	GLYA_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LPG8	ASPD_SILPO		cytoplasm
Q5LPQ1	GLMU_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LQ08	SURE_SILPO	cytoplasm (potential).	periplasm
Q5LQ80	TGT_SILPO		cytoplasm
Q5LQ99	ISPG_SILPO		cytoplasm

Q5LQJ6	PLSX_SILPO		cytoplasm
Q5LQJ7	RL32_SILPO		cytoplasm
Q5LQK5	SYI_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LQL4	ENO_SILPO	cytoplasm (by similarity).	extracellular
Q5LQN0	KSGA_SILPO		cytoplasm
Q5LQP4	NDK_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LQY2	RUVX_SILPO	cytoplasm (potential).	cytoplasm
Q5LR15	CCMA_SILPO	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5LR43	TIG_SILPO		cytoplasm
Q5LR48	RL9_SILPO		cytoplasm
Q5LR50	RS6_SILPO		cytoplasm
Q5LR59	HIS42_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LRA3	QUEA_SILPO	cytoplasm (potential).	cytoplasm
Q5LRC0	CAAL_SILPO		cytoplasm
Q5LRD8	HUTH_SILPO	cytoplasm (potential).	cytoplasm
Q5LRH4	LEXA_SILPO		cytoplasm
Q5LRH8	TRPD_SILPO		cytoplasm
Q5LRI5	XGPT_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LRI7	PDXH_SILPO		cytoplasm
Q5LRL6	GUAA_SILPO		cytoplasm
Q5LRN5	ISPDF_SILPO		cytoplasm
Q5LRS9	G6PI_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LRU0	RECA_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LRV2	Y2021_SILPO		inner membrane
Q5LRY1	RL27_SILPO		cytoplasm
Q5LRY5	PROB_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LRY6	PROA_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LRZ4	RS2_SILPO		cytoplasm
Q5LS78	PURQ_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LS91	CCME1_SILPO	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5LS92	ARGC_SILPO	cytoplasm (probable).	cytoplasm
Q5LS98	PURL_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LSB0	RRMJ_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LSL4	NRDR_SILPO		cytoplasm
Q5LSQ1	UREE_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LSQ2	URE1_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LSQ3	URE2_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LSQ4	URE3_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LST0	GID_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LST2	HIS3_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LSU4	FABZ_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LSU9	DXR_SILPO		cytoplasm
Q5LSV2	RRF_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LSX9	AROB_SILPO	cytoplasm (probable).	cytoplasm
Q5LSY0	AROK_SILPO	cytoplasm (probable).	cytoplasm
Q5LTB1	PQQB_SILPO		cytoplasm
Q5LTB2	PQQC_SILPO		cytoplasm
Q5LTB3	PQQD_SILPO		cytoplasm
Q5LTB4	PQQE_SILPO		cytoplasm
Q5LTP7	ILVC_SILPO		cytoplasm
Q5LTP9	GLMM_SILPO		cytoplasm

Q5LTR1	Y1352_SILPO		inner membrane
Q5LTT5	RPIA_SILPO		cytoplasm
Q5LTU4	PURA_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LTY7	THYX_SILPO		cytoplasm
Q5LU58	MURB_SILPO	cytoplasm (probable).	inner membrane
Q5LU63	MURG_SILPO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5LU71	MURD_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LU74	MRAY_SILPO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5LU79	MRAW_SILPO		cytoplasm
Q5LU80	MRAZ_SILPO		inner membrane
Q5LU93	HIS5_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LU97	HIS41_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LU99	HIS6_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LUA0	HIS2_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LUI2	Y1072_SILPO		inner membrane
Q5LUQ0	CLPP_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LUS0	LEPA_SILPO		cytoplasm
Q5LUS2	HISX2_SILPO		cytoplasm
Q5LUS9	RL28_SILPO		cytoplasm
Q5LUY8	KITH_SILPO	cytoplasm (potential).	cytoplasm
Q5LUZ4	LGT_SILPO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5LV15	CH60_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LV16	CH10_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LV46	XYLA_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LV87	TRPA_SILPO		cytoplasm
Q5LV90	RL25_SILPO		cytoplasm
Q5LV91	PTH_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LVI8	Y713_SILPO		cytoplasm
Q5LVJ2	PPCK_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LVM5	TAUB_SILPO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5LVN0	DNAE2_SILPO	cytoplasm (probable).	cytoplasm
Q5LVV1	HISX1_SILPO		cytoplasm
Q5LW06	RNPA_SILPO		extracellular
Q5LW32	RPOA_SILPO		cytoplasm
Q5LW33	RS11_SILPO		cytoplasm
Q5LW34	RS13_SILPO		cytoplasm
Q5LW38	RL15_SILPO		cytoplasm
Q5LW41	RS5_SILPO		cytoplasm
Q5LW42	RL18_SILPO		cytoplasm
Q5LW44	RS8_SILPO		cytoplasm
Q5LW49	RS17_SILPO		cytoplasm
Q5LW50	RL29_SILPO		cytoplasm
Q5LW55	RL16_SILPO		cytoplasm
Q5LW56	RS3_SILPO		cytoplasm
Q5LW59	RL2_SILPO		cytoplasm
Q5LW93	NADD_SILPO		cytoplasm
Q5LWD3	DUT_SILPO		extracellular
Q5LWE5	GLND_SILPO		cytoplasm

Q5LWF2	GIDB_SILPO		cytoplasm
Q5LWF3	GIDA_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LWF7	HAM1_SILPO		extracellular
Q5LWG3	ASSY_SILPO	cytoplasm (probable).	cytoplasm
Q5LWH0	MUTS_SILPO		cytoplasm
Q5LWJ5	DNAJ_SILPO	cytoplasm (probable).	cytoplasm
Q5LWJ6	DNAK_SILPO		cytoplasm
Q5LWK3	RADC_SILPO		cytoplasm
Q5LWL1	Y064_SILPO		inner membrane
Q5LWL6	ARGJ_SILPO	cytoplasm (probable).	cytoplasm
Q5LWN8	Y091_SILPO		inner membrane
Q5LWT6	IDI_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LWT9	FPG_SILPO		cytoplasm
Q5LWV4	DNAA_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LWV5	RS20_SILPO		cytoplasm
Q5LWX7	FLGH_SILPO	bacterial cell outer membrane; lipid- anchor (probable).	inner membrane
Q5LWZ2	FLGI_SILPO	periplasm (by similarity).	periplasm
Q5LWZ5	LEU3_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LX06	LEU2_SILPO		cytoplasm
Q5LX07	LEUD_SILPO		cytoplasm
Q5LX20	Y220_SILPO		inner membrane
Q5LX40	EX7S_SILPO	cytoplasm (by similarity).	cytoplasm
Q5LX60	AROC_SILPO		cytoplasm
Q5LX62	Y283_SILPO	bacterial cell inner membrane; multi- pass membrane protein (by similarity).	inner membrane
Q5LX75	PYRB_SILPO		cytoplasm
Q5LX98	ISPE_SILPO		cytoplasm
Q5LXE1	MDH_SILPO		cytoplasm
Q5NE80	TRPA_FRATT		cytoplasm
Q5NE98	RECA_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NE99	SECB2_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NEF0	GIDB_FRATT		cytoplasm
Q5NEF7	RS20_FRATT		cytoplasm
Q5NEF8	LEPA_FRATT		cytoplasm
Q5NEF9	TRMU_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NEH3	TPMT_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NEM3	RL28_FRATT		cytoplasm
Q5NEP2	UNG_FRATT	cytoplasm (by similarity).	extracellular
Q5NEP6	DXR_FRATT		cytoplasm
Q5NEQ0	FABZ_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NEQ9	HAM1_FRATT		extracellular
Q5NER3	RNC_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NER4	TRUB_FRATT		cytoplasm
Q5NEV7	SECB1_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NEV8	MUTS_FRATT		cytoplasm
Q5NEV9	ACCA_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NEY6	KGUA_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NF19	PPNK_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NF58	PANB_FRATT		cytoplasm
Q5NF71	PLSX_FRATT		cytoplasm
Q5NF72	RL32_FRATT		cytoplasm
Q5NFB1	GPMI_FRATT		cytoplasm
Q5NFC4	G6PI_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NFD4	MURB_FRATT	cytoplasm (probable).	inner membrane

Q5NFE1	UBIE_FRATT		cytoplasm
Q5NFF8	RNH2_FRATT	cytoplasm (potential).	extracellular
Q5NFG7	DNAK_FRATT		cytoplasm
Q5NFG8	DNAJ_FRATT	cytoplasm (probable).	cytoplasm
Q5NFJ3	GLYA_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NFK3	TILS_FRATT	cytoplasm (probable).	cytoplasm
Q5NFK5	TYSY_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NFK6	LGT_FRATT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5NFL9	QUEA_FRATT	cytoplasm (potential).	cytoplasm
Q5NFM5	RPIA_FRATT		cytoplasm
Q5NFM8	GIDA_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NFN4	UVRB_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NFP4	SSRP_FRATT	cytoplasm (potential).	cytoplasm
Q5NFS0	AROK_FRATT	cytoplasm (probable).	cytoplasm
Q5NFS1	AROB_FRATT	cytoplasm (probable).	cytoplasm
Q5NFS4	Y1151_FRATT		cytoplasm
Q5NFT3	HEMH_FRATT	cytoplasm (by similarity).	extracellular
Q5NFU1	ISPF_FRATT		extracellular
Q5NFU6	Y1123_FRATT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5NFU8	TGT_FRATT		cytoplasm
Q5NFX8	Y1085_FRATT		inner membrane
Q5NFZ9	RS6_FRATT		cytoplasm
Q5NG01	RL9_FRATT		cytoplasm
Q5NG36	TRUA_FRATT		cytoplasm
Q5NG38	GUAA_FRATT		cytoplasm
Q5NG44	RUVB_FRATT		cytoplasm
Q5NG68	RUVX_FRATT	cytoplasm (potential).	cytoplasm
Q5NG98	Y954_FRATT		inner membrane
Q5NGC1	FMT_FRATT		cytoplasm
Q5NGD0	SYI_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NGD3	RRMJ_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NGG6	AROC_FRATT		cytoplasm
Q5NGH1	GPDA_FRATT	cytoplasm (probable).	cytoplasm
Q5NGI2	RNPH_FRATT		cytoplasm
Q5NGJ7	TOLB_FRATT	periplasm (potential).	periplasm
Q5NGK4	ISPH_FRATT		cytoplasm
Q5NGK7	Y829_FRATT	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q5NGM4	MURG_FRATT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5NGM6	RECR_FRATT		cytoplasm
Q5NGQ8	UVRC_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NGR1	RL27_FRATT		cytoplasm
Q5NGW1	UPP_FRATT		cytoplasm
Q5NGW4	TDH_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NGW8	ENO_FRATT	cytoplasm (by similarity).	extracellular
Q5NGX3	RPOZ_FRATT		cytoplasm
Q5NGX8	RS15_FRATT		cytoplasm
Q5NGY1	MRAW_FRATT		cytoplasm
Q5NGY3	FPG_FRATT		cytoplasm

Q5NGY5	SYU_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NGY9	HSLU_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NGZ6	PTH_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NH01	RL25_FRATT		cytoplasm
Q5NH17	RUVA_FRATT		cytoplasm
Q5NH18	RUVC_FRATT		cytoplasm
Q5NH19	Y655_FRATT		inner membrane
Q5NH32	ILVD_FRATT		cytoplasm
Q5NH41	HFQ_FRATT		cytoplasm
Q5NH42	MIAA_FRATT		cytoplasm
Q5NH46	CLPX_FRATT		cytoplasm
Q5NH47	CLPP_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NH48	TIG_FRATT		cytoplasm
Q5NH50	KITH_FRATT	cytoplasm (potential).	cytoplasm
Q5NH55	Y616_FRATT		inner membrane
Q5NH64	ISPG_FRATT		cytoplasm
Q5NH87	COAD_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NHE5	PDXT_FRATT		cytoplasm
Q5NHE6	PDXS_FRATT		extracellular
Q5NHI5	KSGA_FRATT		cytoplasm
Q5NHJ8	FETP_FRATT		cytoplasm
Q5NHK3	MURD_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NHK4	MRAY_FRATT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5NHN2	GLGA_FRATT		cytoplasm
Q5NHN4	GLGB_FRATT		extracellular
Q5NHP0	GCST_FRATT		cytoplasm
Q5NHQ9	GLMS_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NHR0	GLMU_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NHR3	PSD_FRATT		cytoplasm
Q5NHS2	NDK_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NHS9	RL31_FRATT		cytoplasm
Q5NHT8	HTPG_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NHU4	RS4_FRATT		cytoplasm
Q5NHU5	RS11_FRATT		cytoplasm
Q5NHU6	RS13_FRATT		cytoplasm
Q5NHU9	RL15_FRATT		cytoplasm
Q5NHV1	RS5_FRATT		cytoplasm
Q5NHV2	RL18_FRATT		cytoplasm
Q5NHV4	RS8_FRATT		cytoplasm
Q5NHV9	RS17_FRATT		cytoplasm
Q5NHW0	RL29_FRATT		cytoplasm
Q5NHW1	RL16_FRATT		cytoplasm
Q5NHW2	RS3_FRATT		cytoplasm
Q5NHW4	RS19_FRATT		cytoplasm
Q5NHW5	RL2_FRATT		cytoplasm
Q5NHW9	RS10_FRATT		cytoplasm
Q5NHX1	RS7_FRATT		cytoplasm
Q5NHX2	RS12_FRATT		cytoplasm
Q5NHX4	DUT_FRATT		extracellular
Q5NHX6	UPPS_FRATT		cytoplasm
Q5NHX7	RRF_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NHY0	RS2_FRATT		cytoplasm
Q5NHZ4	SYV_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NI19	ISPE_FRATT		cytoplasm

Q5NI30	CRCB_FRATT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5NI31	HEM3_FRATT		cytoplasm
Q5NI50	MURC_FRATT	cytoplasm (probable).	cytoplasm
Q5NI61	ORN_FRATT	cytoplasm (potential).	cytoplasm
Q5NI78	PURA_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NIB6	NUDH_FRATT		cytoplasm
Q5NIC3	RL19_FRATT		cytoplasm
Q5NIC4	TRMD_FRATT	cytoplasm (potential).	cytoplasm
Q5NIC5	RIMM_FRATT	cytoplasm (potential).	cytoplasm
Q5NID1	RPOC_FRATT		cytoplasm
Q5NID2	RPOB_FRATT		cytoplasm
Q5NID4	RL10_FRATT		cytoplasm
Q5NID5	RL1_FRATT		cytoplasm
Q5NID6	RL11_FRATT		cytoplasm
Q5NIF5	KTHY_FRATT		cytoplasm
Q5NIG4	CCA_FRATT		cytoplasm
Q5NII8	GLMM_FRATT		cytoplasm
Q5NIK4	ATPG_FRATT		inner membrane
Q5NIK5	ATPA_FRATT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5NIL6	RBFA_FRATT	cytoplasm (potential).	cytoplasm
Q5NIL9	Y048_FRATT		inner membrane
Q5NIQ8	DNAA_FRATT	cytoplasm (by similarity).	cytoplasm
Q5NL06	GIDB_ZYMMO		cytoplasm
Q5NL75	PTH_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NL76	RL25_ZYMMO		cytoplasm
Q5NL79	MUTS_ZYMMO		cytoplasm
Q5NL81	GPDA_ZYMMO	cytoplasm (probable).	cytoplasm
Q5NL86	PLSX_ZYMMO		cytoplasm
Q5NLA8	SYV_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NLF0	NIFW_ZYMMO		cytoplasm
Q5NLJ4	ILVD_ZYMMO		cytoplasm
Q5NLP5	LEPA_ZYMMO		cytoplasm
Q5NLR1	CLPS_ZYMMO		cytoplasm
Q5NLR2	MURA_ZYMMO	cytoplasm (probable).	inner membrane
Q5NLR6	RPOZ_ZYMMO		cytoplasm
Q5NLS3	RS21_ZYMMO		cytoplasm
Q5NLS7	ACPS_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NLS8	PDXJ_ZYMMO	cytoplasm (probable).	cytoplasm
Q5NLU3	AROC_ZYMMO		cytoplasm
Q5NLU9	PURA_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NLV5	Y1681_ZYMMO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q5NLX2	Y1664_ZYMMO		inner membrane
Q5NLX5	PROA_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NLX8	TILS_ZYMMO	cytoplasm (probable).	cytoplasm
Q5NLZ3	SYV_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NMA4	PURQ_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NMD4	HIS5_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NMD5	HIS4_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NMD6	HIS6_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NMD7	HIS2_ZYMMO	cytoplasm (by similarity).	cytoplasm

Q5NMV7	PPNK_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NMW1	Y1325_ZYMMO		cytoplasm
Q5NMX2	KSGA_ZYMMO		cytoplasm
Q5NN19	GUAA_ZYMMO		cytoplasm
Q5NN30	CCME_ZYMMO	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5NN59	RL9_ZYMMO		cytoplasm
Q5NN61	RS6_ZYMMO		cytoplasm
Q5NN77	KUP_ZYMMO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5NN84	NRDR_ZYMMO		cytoplasm
Q5NN85	GLYA_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NN91	Y1195_ZYMMO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5NNE1	RL31_ZYMMO		cytoplasm
Q5NNK7	RL19_ZYMMO		cytoplasm
Q5NNK8	TRMD_ZYMMO	cytoplasm (potential).	cytoplasm
Q5NNK9	RIMM_ZYMMO	cytoplasm (potential).	cytoplasm
Q5NNP2	MSCL_ZYMMO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5NNQ0	ASSY_ZYMMO	cytoplasm (probable).	cytoplasm
Q5NNS5	Y1011_ZYMMO		cytoplasm
Q5NNT2	RNH2_ZYMMO	cytoplasm (potential).	extracellular
Q5NNT4	GLMM_ZYMMO		cytoplasm
Q5NNV1	SURE_ZYMMO	cytoplasm (potential).	periplasm
Q5NNY7	CLPX_ZYMMO		cytoplasm
Q5NNY8	CLPP_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NNZ0	TIG_ZYMMO		cytoplasm
Q5NP13	ARGJ_ZYMMO	cytoplasm (probable).	cytoplasm
Q5NP61	ISPH_ZYMMO		cytoplasm
Q5NP84	QUEA_ZYMMO	cytoplasm (potential).	cytoplasm
Q5NPA7	MURD_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NPC4	RECR_ZYMMO		cytoplasm
Q5NPC5	FMT_ZYMMO		cytoplasm
Q5NPE5	PYRB_ZYMMO		cytoplasm
Q5NPF1	RUVX_ZYMMO	cytoplasm (potential).	cytoplasm
Q5NPH0	GLND_ZYMMO		cytoplasm
Q5NPJ8	THIG_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NPK4	RPOC_ZYMMO		cytoplasm
Q5NPK5	RPOB_ZYMMO		cytoplasm
Q5NPK9	RL10_ZYMMO		cytoplasm
Q5NPL0	RL1_ZYMMO		cytoplasm
Q5NPL1	RL11_ZYMMO		cytoplasm
Q5NPL5	SSRP_ZYMMO	cytoplasm (potential).	cytoplasm
Q5NPL6	DAPA_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NPM9	DAPB_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NPP5	GID_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NPQ9	LEU3_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NPR4	UVRC_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NPS5	DNAJ_ZYMMO	cytoplasm (probable).	cytoplasm
Q5NPS6	DNAK_ZYMMO		cytoplasm

Q5NPY2	ACCA_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NPY7	AROK_ZYMMO	cytoplasm (probable).	cytoplasm
Q5NPY8	AROB_ZYMMO	cytoplasm (probable).	cytoplasm
Q5NPZ7	TRPA_ZYMMO		cytoplasm
Q5NQ24	Y557_ZYMMO		inner membrane
Q5NQ28	RBFA_ZYMMO	cytoplasm (potential).	cytoplasm
Q5NQ29	KITH_ZYMMO	cytoplasm (potential).	cytoplasm
Q5NQ30	TRUB_ZYMMO		cytoplasm
Q5NQ31	RS15_ZYMMO		cytoplasm
Q5NQ40	RPOA_ZYMMO		cytoplasm
Q5NQ41	RS11_ZYMMO		cytoplasm
Q5NQ42	RS13_ZYMMO		cytoplasm
Q5NQ45	RL15_ZYMMO		cytoplasm
Q5NQ48	RS5_ZYMMO		cytoplasm
Q5NQ49	RL18_ZYMMO		cytoplasm
Q5NQ51	RS8_ZYMMO		cytoplasm
Q5NQ56	RS17_ZYMMO		cytoplasm
Q5NQ57	RL29_ZYMMO		cytoplasm
Q5NQ58	RL16_ZYMMO		cytoplasm
Q5NQ59	RS3_ZYMMO		cytoplasm
Q5NQ61	RL2_ZYMMO		cytoplasm
Q5NQ67	RS7_ZYMMO		cytoplasm
Q5NQ83	GLMU_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NQC7	FTHS_ZYMMO		cytoplasm
Q5NQE8	KGUA_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NQF0	DAPD_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NQH8	RRMJ_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NQQ4	QUEF_ZYMMO	cytoplasm (probable).	cytoplasm
Q5NQQ7	SYI_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NQX0	CCMA_ZYMMO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5NQZ0	ATPG_ZYMMO		inner membrane
Q5NQZ1	ATPA_ZYMMO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5NR05	METX_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NR21	RL27_ZYMMO		cytoplasm
Q5NR24	PROB_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NR31	LEXA_ZYMMO		cytoplasm
Q5NR50	ISPG_ZYMMO		cytoplasm
Q5NR65	TOLB_ZYMMO	periplasm (potential).	periplasm
Q5NR72	RUVB_ZYMMO		cytoplasm
Q5NR75	RUVA_ZYMMO		cytoplasm
Q5NR76	RUVC_ZYMMO		cytoplasm
Q5NR77	Y153_ZYMMO		inner membrane
Q5NR81	TRMB_ZYMMO		cytoplasm
Q5NRC4	LEUD_ZYMMO		cytoplasm
Q5NRC5	LEU2_ZYMMO		cytoplasm
Q5NRF6	Y074_ZYMMO		inner membrane
Q5NRH4	GLMS_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q5NRL5	HRCa_ZYMMO		cytoplasm
Q5NRL6	RNPH_ZYMMO		cytoplasm
Q5NRL7	HAM1_ZYMMO		extracellular
Q5NRM1	CYSJ_ZYMMO		cytoplasm
Q5NRM2	CYSI_ZYMMO		cytoplasm

Q5NSW9	DNAJ_BURCE	cytoplasm (probable).	cytoplasm
Q5NWX5	IDI2_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5NXL5	SYV_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5NXM4	RL19_AZOSE		cytoplasm
Q5NXM5	TRMD_AZOSE	cytoplasm (potential).	cytoplasm
Q5NXM6	RIMM_AZOSE	cytoplasm (potential).	cytoplasm
Q5NXP4	ILVC_AZOSE		cytoplasm
Q5NXV7	GLGB_AZOSE		extracellular
Q5NXZ4	GLGC_AZOSE		cytoplasm
Q5NXZ7	GLGA_AZOSE		cytoplasm
Q5NY71	ILVD3_AZOSE		cytoplasm
Q5NYA9	MDH_AZOSE		cytoplasm
Q5NYE0	SSRP_AZOSE	cytoplasm (potential).	cytoplasm
Q5NYE5	GUAA_AZOSE		cytoplasm
Q5NYJ9	ISPD_AZOSE		cytoplasm
Q5NYK0	ISPF_AZOSE		extracellular
Q5NYP9	MUTS_AZOSE		cytoplasm
Q5NYT0	HSCB_AZOSE		inner membrane
Q5NYT1	HSCA_AZOSE		cytoplasm
Q5NYT9	DAPD_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5NYX9	NORM_AZOSE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5NYY1	Y3608_AZOSE		inner membrane
Q5NZ69	ENO_AZOSE	cytoplasm (by similarity).	extracellular
Q5NZ93	GCSP_AZOSE		cytoplasm
Q5NZF6	BIOH_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5NZG1	RNH2_AZOSE	cytoplasm (potential).	extracellular
Q5NZG4	FABZ_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5NZG9	DXR_AZOSE		cytoplasm
Q5NZH2	RRF_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5NZH5	RS2_AZOSE		cytoplasm
Q5NZH8	GLND_AZOSE		cytoplasm
Q5NZR8	RS15_AZOSE		cytoplasm
Q5NZR9	TRUB_AZOSE		cytoplasm
Q5NZS0	RBFA_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5NZS3	Y3316_AZOSE		inner membrane
Q5NZS9	Y3310_AZOSE		cytoplasm
Q5NZX8	HUTH_AZOSE	cytoplasm (potential).	cytoplasm
Q5P011	IDI1_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P080	UVRC_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P081	NAGZ_AZOSE	cytoplasm (by similarity).	extracellular
Q5P082	ACPS_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P083	PDXJ_AZOSE	cytoplasm (probable).	cytoplasm
Q5P084	RECO_AZOSE		cytoplasm
Q5P086	RNC_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P089	LEPA_AZOSE		cytoplasm
Q5P0D7	PLSX_AZOSE		cytoplasm
Q5P0D8	RL32_AZOSE		cytoplasm
Q5P0E7	UVRB_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P0T2	RL31_AZOSE		cytoplasm
Q5P0T4	G6PI_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P0T5	ORN_AZOSE	cytoplasm (potential).	cytoplasm
Q5P0Z5	Y2894_AZOSE		inner membrane
Q5P0Z7	ASSY_AZOSE	cytoplasm (probable).	cytoplasm
Q5P102	RS20_AZOSE		cytoplasm

Q5P108	LPXK_AZOSE		cytoplasm
Q5P160	CLPX_AZOSE		cytoplasm
Q5P161	CLPP_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P162	TIG_AZOSE		cytoplasm
Q5P1A9	PYRD_AZOSE	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5P1C5	HTPG_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P1F7	GLMM_AZOSE		cytoplasm
Q5P1G0	RRMJ_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P1G6	DAPB_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P1G9	PPNK_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P1H5	DNAK_AZOSE		cytoplasm
Q5P1H7	DNAJ2_AZOSE	cytoplasm (probable).	cytoplasm
Q5P1I9	TRPA_AZOSE		cytoplasm
Q5P1J2	TRUA_AZOSE		cytoplasm
Q5P1J6	LEU3_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P1J7	LEUD_AZOSE		cytoplasm
Q5P1J8	LEU2_AZOSE		cytoplasm
Q5P1K1	AROC_AZOSE		cytoplasm
Q5P1Y4	MTFA_AZOSE		cytoplasm
Q5P1Z7	RL28_AZOSE		cytoplasm
Q5P201	CLPS_AZOSE		cytoplasm
Q5P221	SYI_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P224	ISPH_AZOSE		cytoplasm
Q5P229	Y2510_AZOSE		inner membrane
Q5P233	TYSY_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P255	PROA_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P260	Y2479_AZOSE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P262	RS21_AZOSE		cytoplasm
Q5P2G2	TRPD_AZOSE		cytoplasm
Q5P2I9	TILS_AZOSE	cytoplasm (probable).	cytoplasm
Q5P2J0	ACCA_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P2J9	GLUQ_AZOSE		cytoplasm
Q5P2L5	Y2324_AZOSE		inner membrane
Q5P2M2	RL9_AZOSE		cytoplasm
Q5P2M5	RS6_AZOSE		cytoplasm
Q5P2N0	CRCB_AZOSE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P2P8	PPCK_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P2U2	Y2247_AZOSE		inner membrane
Q5P2U6	RUVA_AZOSE		cytoplasm
Q5P2U7	RUVB_AZOSE		cytoplasm
Q5P307	RPOA_AZOSE		cytoplasm
Q5P308	RS4_AZOSE		cytoplasm
Q5P309	RS11_AZOSE		cytoplasm
Q5P310	RS13_AZOSE		cytoplasm
Q5P313	RL15_AZOSE		cytoplasm
Q5P315	RS5_AZOSE		cytoplasm
Q5P316	RL18_AZOSE		cytoplasm
Q5P318	RS8_AZOSE		cytoplasm
Q5P323	RS17_AZOSE		cytoplasm
Q5P324	RL29_AZOSE		cytoplasm

Q5P325	RL16_AZOSE		cytoplasm
Q5P326	RS3_AZOSE		cytoplasm
Q5P329	RL2_AZOSE		cytoplasm
Q5P333	RS10_AZOSE		cytoplasm
Q5P336	RS7_AZOSE		cytoplasm
Q5P337	RS12_AZOSE		cytoplasm
Q5P338	RPOC_AZOSE		cytoplasm
Q5P339	RPOB_AZOSE		cytoplasm
Q5P341	RL10_AZOSE		cytoplasm
Q5P342	RL1_AZOSE		cytoplasm
Q5P3B7	Y2072_AZOSE		inner membrane
Q5P3C5	RUVC_AZOSE		cytoplasm
Q5P3C8	Y2061_AZOSE		cytoplasm
Q5P3E3	KUP_AZOSE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P3K6	CCME_AZOSE	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5P3L0	CCMA_AZOSE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5P3L9	RPOZ_AZOSE		cytoplasm
Q5P3M1	DNAJ1_AZOSE	cytoplasm (probable).	cytoplasm
Q5P3R2	MURB_AZOSE	cytoplasm (probable).	inner membrane
Q5P3R8	RECA_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P3T0	CCA_AZOSE		cytoplasm
Q5P4A1	LEXA_AZOSE		cytoplasm
Q5P4A4	DNAE2_AZOSE	cytoplasm (probable).	cytoplasm
Q5P4E3	ATPG_AZOSE		inner membrane
Q5P4E4	ATPA_AZOSE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5P4H2	SMG_AZOSE		cytoplasm
Q5P4H6	FMT_AZOSE		cytoplasm
Q5P4J6	GIDA_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P4J7	GIDB_AZOSE		cytoplasm
Q5P4L8	Y1619_AZOSE		inner membrane
Q5P4P0	DNAA_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P4P2	RNPA_AZOSE		extracellular
Q5P4R1	METX_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P4W4	MSCL_AZOSE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P4Z6	TOLB_AZOSE	periplasm (potential).	periplasm
Q5P503	HSLU_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P5J1	THIG_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P5J5	HAM1_AZOSE		extracellular
Q5P5P2	AROK_AZOSE	cytoplasm (probable).	cytoplasm
Q5P5P3	AROB_AZOSE	cytoplasm (probable).	cytoplasm
Q5P5P9	GLMU_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P6E9	LGT_AZOSE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P6F1	ILVD2_AZOSE		cytoplasm

Q5P6G9	QUEF_AZOSE	cytoplasm (probable).	cytoplasm
Q5P6I4	PYRB_AZOSE		cytoplasm
Q5P6I6	RUVX_AZOSE	cytoplasm (potential).	cytoplasm
Q5P6K4	UPPP_AZOSE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P6Y8	MRAZ_AZOSE		inner membrane
Q5P6Y9	MRAW_AZOSE		cytoplasm
Q5P6Z3	MRAY_AZOSE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P6Z4	MURD_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P6Z6	MURG_AZOSE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5P706	ARGJ_AZOSE	cytoplasm (probable).	cytoplasm
Q5P720	TGT_AZOSE		cytoplasm
Q5P721	QUEA_AZOSE	cytoplasm (potential).	cytoplasm
Q5P722	PTH_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P723	RL25_AZOSE		cytoplasm
Q5P725	ISPE_AZOSE		cytoplasm
Q5P728	FPG_AZOSE		cytoplasm
Q5P762	Y26_AZOSE		inner membrane
Q5P787	MURA_AZOSE	cytoplasm (probable).	inner membrane
Q5P789	HIS1_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P790	HISX_AZOSE		cytoplasm
Q5P791	HIS8_AZOSE		cytoplasm
Q5P793	HIS5_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P794	HIS4_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P795	HIS6_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P796	HIS3_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P797	HIS2_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7A9	NDK_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7B3	ISPG_AZOSE		cytoplasm
Q5P7C4	PURA_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7F1	RECR_AZOSE		cytoplasm
Q5P7G2	CH60_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7H5	CAPP_AZOSE		cytoplasm
Q5P7I0	HEM6_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7J1	KSGA_AZOSE		cytoplasm
Q5P7L9	RPIA_AZOSE		cytoplasm
Q5P7N1	SECB_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7N4	GPMA_AZOSE		cytoplasm
Q5P7P0	NRDR_AZOSE		cytoplasm
Q5P7P1	GLYA_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7S1	GLNE_AZOSE		cytoplasm
Q5P7T2	NUDH_AZOSE		cytoplasm
Q5P7T9	ARGC_AZOSE	cytoplasm (probable).	cytoplasm
Q5P7U2	SYU_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7U7	SERC_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7Z3	PROB_AZOSE	cytoplasm (by similarity).	cytoplasm
Q5P7Z5	RL27_AZOSE		cytoplasm
Q5P7Z9	DUT_AZOSE		extracellular
Q5P832	SURE_AZOSE	cytoplasm (potential).	periplasm
Q5P841	RUMA_AZOSE		cytoplasm
Q5P8J4	ILVD1_AZOSE		cytoplasm

Q5P905	PHS_AZOSE		cytoplasm
Q5P951	DTD_AZOSE	cytoplasm (probable).	cytoplasm
Q5P993	ISPF_ANAMM		extracellular
Q5P9A5	RL25_ANAMM		cytoplasm
Q5P9A6	PTH_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5P9D7	RUVC_ANAMM		cytoplasm
Q5P9D8	Y1298_ANAMM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P9E0	DNAJ_ANAMM	cytoplasm (probable).	cytoplasm
Q5P9F9	TRMD_ANAMM	cytoplasm (potential).	cytoplasm
Q5P9G0	RIMM_ANAMM	cytoplasm (potential).	cytoplasm
Q5P9G3	Y1264_ANAMM		inner membrane
Q5P9H3	UVRB_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5P9I2	RS20_ANAMM		cytoplasm
Q5P9I4	LGT_ANAMM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P9K3	MRAY_ANAMM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5P9K5	SYV_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5P9L5	GUAA_ANAMM		cytoplasm
Q5P9M4	ATPA_ANAMM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5P9M7	SSRP_ANAMM	cytoplasm (potential).	cytoplasm
Q5P9M8	SYV_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5P9P7	NDK_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5P9R1	TILS_ANAMM	cytoplasm (probable).	cytoplasm
Q5P9S6	FABZ_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5P9U8	RNC_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PA19	GIDA_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PA30	CH60_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PA34	GLMM_ANAMM		cytoplasm
Q5PA57	RS10_ANAMM		cytoplasm
Q5PA61	RL2_ANAMM		cytoplasm
Q5PA64	RS3_ANAMM		cytoplasm
Q5PA65	RL16_ANAMM		cytoplasm
Q5PA72	RS8_ANAMM		cytoplasm
Q5PA74	RL18_ANAMM		cytoplasm
Q5PA79	RS13_ANAMM		cytoplasm
Q5PA81	RPOA_ANAMM		cytoplasm
Q5PA93	DAPB_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PA96	RL32_ANAMM		cytoplasm
Q5PAA6	RUVX_ANAMM	cytoplasm (potential).	cytoplasm
Q5PAA8	PURA_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PAB8	DNAK_ANAMM		cytoplasm
Q5PAC9	RL28_ANAMM		cytoplasm
Q5PAE6	DUT_ANAMM		extracellular
Q5PAE7	ISPH_ANAMM		cytoplasm
Q5PAE9	GPMI_ANAMM		cytoplasm
Q5PAI9	DXR_ANAMM		cytoplasm
Q5PAJ1	ISPG_ANAMM		cytoplasm
Q5PAJ6	RBFA_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PAK9	MURB_ANAMM	cytoplasm (probable).	inner membrane

Q5PAL3	Y694_ANAMM		inner membrane
Q5PAL9	SYI_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PAN0	RRMJ_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PAN5	PYRB_ANAMM		cytoplasm
Q5PAN6	TRUA_ANAMM		cytoplasm
Q5PAS3	MRAW_ANAMM		cytoplasm
Q5PAS8	RL27_ANAMM		cytoplasm
Q5PAV3	MDH_ANAMM		cytoplasm
Q5PAV9	KSGA_ANAMM		cytoplasm
Q5PAX8	LEPA_ANAMM		cytoplasm
Q5PAY1	RS15_ANAMM		cytoplasm
Q5PAY2	TRUB_ANAMM		cytoplasm
Q5PAZ6	ACPS_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PB05	ISPE_ANAMM		cytoplasm
Q5PB34	SURE_ANAMM	cytoplasm (potential).	periplasm
Q5PB35	CCME_ANAMM	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5PB48	DNAA_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PB75	RECR_ANAMM		cytoplasm
Q5PB81	UVRC_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PB86	HTPG_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PBC5	Y316_ANAMM		cytoplasm
Q5PBC6	Y315_ANAMM		inner membrane
Q5PBC7	FMT_ANAMM		cytoplasm
Q5PBD0	CLPP_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PBD1	TIG_ANAMM		cytoplasm
Q5PBF2	PDXH_ANAMM		cytoplasm
Q5PBG2	RNH2_ANAMM	cytoplasm (potential).	extracellular
Q5PBG3	RPOC_ANAMM		cytoplasm
Q5PBG4	RPOB_ANAMM		cytoplasm
Q5PBH3	RS7_ANAMM		cytoplasm
Q5PBH8	TOLB_ANAMM	periplasm (potential).	periplasm
Q5PBH9	MURD_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PBI8	SUCC_ANAMM		cytoplasm
Q5PBM1	RUVB_ANAMM		cytoplasm
Q5PBM3	THYX_ANAMM		cytoplasm
Q5PBM8	GLYA_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PBN1	RS6_ANAMM		cytoplasm
Q5PBP4	MURA_ANAMM	cytoplasm (probable).	inner membrane
Q5PBP5	CCMA_ANAMM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5PBR8	SECB_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PBT6	RECA_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PBV0	GLMU_ANAMM	cytoplasm (by similarity).	cytoplasm
Q5PBZ2	GPMI_SALPA		cytoplasm
Q5PBZ5	SECB_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PC07	TDH_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PC09	FPG_SALPA		cytoplasm
Q5PC27	SLMA_SALPA	nucleoid-associated (by similarity).	cytoplasm
Q5PC28	DUT_SALPA		extracellular
Q5PC30	RADC_SALPA		cytoplasm
Q5PC31	RL28_SALPA		cytoplasm

Q5PC39	RPOZ_SALPA		cytoplasm
Q5PC79	YGIH_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PC81	UPPP_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PC82	CCA_SALPA		cytoplasm
Q5PC85	GLNE_SALPA		cytoplasm
Q5PCH1	SELU_SALPA		cytoplasm
Q5PCJ8	ATKA_SALPA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q5PCK6	YBGL_SALPA		inner membrane
Q5PCM7	SUCC_SALPA		cytoplasm
Q5PCN2	TOLB_SALPA	periplasm (potential).	periplasm
Q5PCR2	ISPE_SALPA		cytoplasm
Q5PCR7	PTH_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PCT4	KITH_SALPA	cytoplasm (potential).	cytoplasm
Q5PCT5	HNS_SALPA		cytoplasm
Q5PCU4	TAL2_SALPA	cytoplasm (probable).	cytoplasm
Q5PCV6	TRUA_SALPA		cytoplasm
Q5PCW7	YFCK_SALPA		cytoplasm
Q5PCX1	MEPA_SALPA	periplasm (by similarity).	extracellular
Q5PCX2	AROC_SALPA		cytoplasm
Q5PCX6	FADJ_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PCX7	FADI_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PCY8	GCH2_SALPA		extracellular
Q5PD31	SFSA_SALPA		cytoplasm
Q5PD33	GLUQ_SALPA		cytoplasm
Q5PD43	GSA_SALPA	cytoplasm (potential).	cytoplasm
Q5PD47	BTUF_SALPA	periplasm (by similarity).	periplasm
Q5PD50	CLCA_SALPA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q5PD51	YAEH_SALPA		inner membrane
Q5PD56	GLND_SALPA		cytoplasm
Q5PD57	DAPD_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PD59	DXR_SALPA		cytoplasm
Q5PD60	RRF_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PD63	RS2_SALPA		cytoplasm
Q5PD64	SKP_SALPA	periplasm (by similarity).	periplasm
Q5PD65	YAET_SALPA	bacterial cell outer membrane (by similarity).	outer membrane
Q5PD68	UPPS_SALPA		cytoplasm
Q5PD69	ACCA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PD71	RNH2_SALPA	cytoplasm (potential).	extracellular
Q5PD74	FABZ_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PD76	TILS_SALPA	cytoplasm (probable).	cytoplasm
Q5PDA1	SPED_SALPA		cytoplasm
Q5PDB8	PANC_SALPA		cytoplasm
Q5PDB9	PAND_SALPA		cytoplasm
Q5PDC2	MURD_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PDC3	MRAY_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q5PDC7	MURC_SALPA	cytoplasm (probable).	cytoplasm
Q5PDC8	MURG_SALPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5PDD0	SECM_SALPA		periplasm
Q5PDD1	LPXC_SALPA		cytoplasm
Q5PDD7	YACF_SALPA		inner membrane
Q5PDD8	YACG_SALPA		cytoplasm
Q5PDD9	KSGA_SALPA		cytoplasm
Q5PDE4	DJLA_SALPA	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q5PDE5	OSTA_SALPA	bacterial cell outer membrane (by similarity).	outer membrane
Q5PDG2	LEU3_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PDG3	LEU2_SALPA		cytoplasm
Q5PDG4	LEUD_SALPA		cytoplasm
Q5PDH4	MRAW_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PDH5	MRAZ_SALPA		inner membrane
Q5PDJ4	DNAJ_SALPA	cytoplasm (probable).	cytoplasm
Q5PDJ5	DNAK_SALPA		cytoplasm
Q5PDL9	SYI_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PDM2	RS20_SALPA		cytoplasm
Q5PDM6	TAL1_SALPA	cytoplasm (probable).	cytoplasm
Q5PDN6	HIS6_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PDP3	HISX_SALPA		cytoplasm
Q5PDP4	HIS8_SALPA		cytoplasm
Q5PDP5	HIS7_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PDP6	HIS5_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PDP7	HIS4_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PE20	RL25_SALPA		cytoplasm
Q5PE23	YEJL_SALPA		inner membrane
Q5PE39	EFPL_SALPA		cytoplasm
Q5PE68	CDD_SALPA		cytoplasm
Q5PEA8	SICP_SALPA	cytoplasm (probable).	cytoplasm
Q5PEB1	HILA_SALPA		cytoplasm
Q5PEE6	MUTS_SALPA		cytoplasm
Q5PEG0	FTSB_SALPA	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q5PEG1	ISPD_SALPA		cytoplasm
Q5PEG2	ISPF_SALPA		extracellular
Q5PEG3	TRUD_SALPA		cytoplasm
Q5PEG4	SURE_SALPA	cytoplasm (potential).	periplasm
Q5PEH4	ENO_SALPA	cytoplasm (by similarity).	extracellular
Q5PEH7	CYSJ_SALPA		cytoplasm
Q5PEH8	CYSI_SALPA		cytoplasm
Q5PEJ4	RUMA_SALPA		cytoplasm
Q5PEK0	QUEF_SALPA	cytoplasm (probable).	cytoplasm
Q5PEK5	MTFA_SALPA		cytoplasm
Q5PEN3	NUDH_SALPA		cytoplasm
Q5PEN5	LGT_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q5PEN6	TYSY_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PEN7	AAS_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PEQ4	RCNA_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PF04	NRDI_SALPA		inner membrane
Q5PF10	GSH1_SALPA		cytoplasm
Q5PF11	LUXS_SALPA		extracellular
Q5PF15	RECA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PF36	NORW_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PF68	PROB_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PF69	PROA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PF80	XGPT_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PFD8	RNH_SALPA	cytoplasm (potential).	extracellular
Q5PFF7	RIMM_SALPA	cytoplasm (potential).	cytoplasm
Q5PFF8	TRMD_SALPA	cytoplasm (potential).	cytoplasm
Q5PFF9	RL19_SALPA		cytoplasm
Q5PFG2	SSRP_SALPA	cytoplasm (potential).	cytoplasm
Q5PFG7	PPNK_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PFK6	RECR_SALPA		cytoplasm
Q5PFK7	HTPG_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PFL9	RL31B_SALPA		cytoplasm
Q5PFN3	TIG_SALPA		cytoplasm
Q5PFN4	CLPP_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PFR8	EX7S_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PFS6	NRDR_SALPA		cytoplasm
Q5PFT4	ACPH_SALPA		cytoplasm
Q5PFT5	QUEA_SALPA	cytoplasm (potential).	cytoplasm
Q5PFT6	TGT_SALPA		cytoplasm
Q5PFU0	YAIE_SALPA		inner membrane
Q5PFV4	YAII_SALPA		inner membrane
Q5PFV5	AROL_SALPA	cytoplasm (probable).	cytoplasm
Q5PG19	DING_SALPA		cytoplasm
Q5PG52	UVRB_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PG56	6PGL_SALPA		cytoplasm
Q5PG61	HUTH_SALPA	cytoplasm (potential).	cytoplasm
Q5PG75	GPMA_SALPA		cytoplasm
Q5PG85	NANE1_SALPA		cytoplasm
Q5PG91	WRBA_SALPA		cytoplasm
Q5PGA2	CBPA_SALPA	nucleoid (by similarity).	inner membrane
Q5PGB1	SIGE_SALPA	cytoplasm (probable).	cytoplasm
Q5PGB9	TUSE_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PGD2	FABA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PGD8	PNCB_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PGE0	PYRD_SALPA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5PGH1	LPXK_SALPA		cytoplasm
Q5PGK0	CLPS_SALPA		cytoplasm
Q5PGL4	YBJQ_SALPA		inner membrane
Q5PGM5	RIMK_SALPA		cytoplasm
Q5PGN8	YBJL_SALPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q5PGS9	THIK_SALPA		cytoplasm
Q5PGT0	NAGZ_SALPA	cytoplasm (by similarity).	extracellular
Q5PGT4	KTHY_SALPA		cytoplasm
Q5PGT9	RL32_SALPA		cytoplasm
Q5PGU6	FLGH_SALPA	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q5PGU7	FLGI_SALPA	periplasm (by similarity).	periplasm
Q5PGW7	MDTH_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PGX0	YCEA_SALPA		inner membrane
Q5PGX1	YCEI_SALPA	periplasm (potential).	periplasm
Q5PGX5	OPGC_SALPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q5PGY0	MDTG_SALPA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q5PGZ4	YCDX_SALPA		cytoplasm
Q5PH16	MDTK_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PH77	YDIA_SALPA		cytoplasm
Q5PH81	BTUD_SALPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5PH87	BTUC_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PHH8	DMSD_SALPA		cytoplasm
Q5PHR6	ZNTB_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PHY7	MAO1_SALPA		cytoplasm
Q5PI10	UVRC_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PI18	PGSA_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PI28	HEM6_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PI36	ECOT_SALPA	periplasm (by similarity).	periplasm
Q5PI52	GUAA_SALPA		cytoplasm
Q5PI55	CCMA1_SALPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5PI61	NAPA_SALPA	periplasm (by similarity).	periplasm
Q5PI77	YEBN_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PI99	YEBF_SALPA	secreted protein (by similarity).	inner membrane
Q5PIA7	RUVB_SALPA		cytoplasm
Q5PIB8	TUS_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PIC3	HNSB2_SALPA		cytoplasm
Q5PIC6	RNFE_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q5PIE3	YEEN_SALPA		inner membrane
Q5PIG3	RECO_SALPA		cytoplasm
Q5PIG5	RNC_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PIH6	HMP_SALPA		cytoplasm
Q5PII3	GLYA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PIJ4	ACPS_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PIK6	PDXH_SALPA		cytoplasm
Q5PIK7	SYU_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PIL0	CAIC_SALPA		cytoplasm
Q5PIL2	CAIE_SALPA		cytoplasm
Q5PIL3	KEFC_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PIL9	DAPB_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PIN4	FIXA_SALPA		cytoplasm
Q5PIP0	FIXB_SALPA		cytoplasm
Q5PIR7	FIEF_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PIR9	HSLU_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PIT4	MSCL_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PIT7	FMT_SALPA		cytoplasm
Q5PIU0	SMG_SALPA		cytoplasm
Q5PIU1	RL29_SALPA		cytoplasm
Q5PIU2	RS17_SALPA		cytoplasm
Q5PIU7	RS8_SALPA		cytoplasm
Q5PIU9	RL18_SALPA		cytoplasm
Q5PIV0	RL16_SALPA		cytoplasm
Q5PIV1	RS10_SALPA		cytoplasm
Q5PIV5	RL2_SALPA		cytoplasm
Q5PIV8	RS3_SALPA		cytoplasm
Q5PIW0	TUSB_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PIW1	RS12_SALPA		cytoplasm
Q5PIW2	RS7_SALPA		cytoplasm
Q5PJ05	ACTP_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PJ47	ULAB_SALPA	cytoplasm (potential).	cytoplasm
Q5PJ48	ULAA_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PJ49	ULAG_SALPA	cytoplasm (potential).	extracellular
Q5PJ50	ULAR_SALPA	cytoplasm (potential).	cytoplasm
Q5PJ55	RL9_SALPA		cytoplasm
Q5PJ58	RS6_SALPA		cytoplasm
Q5PJ61	ULAF_SALPA		cytoplasm
Q5PJ62	ULAE_SALPA		cytoplasm
Q5PJ63	ULAD_SALPA		cytoplasm
Q5PJ64	ULAC_SALPA	cytoplasm (probable).	cytoplasm
Q5PJ71	RCMNS_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PJ81	MSRA_SALPA		cytoplasm
Q5PJB2	PYRB_SALPA		cytoplasm
Q5PJB3	PYRI_SALPA		cytoplasm
Q5PJB8	SYV_SALPA	cytoplasm (by similarity).	cytoplasm

Q5PJF5	YGFY_SALPA		cytoplasm
Q5PJG4	GCST_SALPA		cytoplasm
Q5PJH1	RPIA_SALPA		cytoplasm
Q5PJH7	SPEB_SALPA		cytoplasm
Q5PJI9	ARGO_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PJJ7	RUVX_SALPA	cytoplasm (potential).	cytoplasm
Q5PJN1	NIKR_SALPA		cytoplasm
Q5PJN7	TUSA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PJN8	YHIQ_SALPA		inner membrane
Q5PJP1	USPB_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PJT2	OADG2_SALPA	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q5PJT7	AAEX_SALPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q5PJT8	AAEA_SALPA	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q5PJT9	AAEB_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PJW3	FIS_SALPA		cytoplasm
Q5PJX1	GIDA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PJX2	GIDB_SALPA		cytoplasm
Q5PJX7	KUP_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PJX8	RAVA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PJX9	VIAA_SALPA		cytoplasm
Q5PJZ5	ILVC_SALPA		cytoplasm
Q5PK00	ILVD_SALPA		cytoplasm
Q5PK01	RS5_SALPA		cytoplasm
Q5PK03	RL15_SALPA		cytoplasm
Q5PK07	RS13_SALPA		cytoplasm
Q5PK08	RS11_SALPA		cytoplasm
Q5PK09	RS4_SALPA		cytoplasm
Q5PK10	RPOA_SALPA		cytoplasm
Q5PK20	DEOD_SALPA		cytoplasm
Q5PK22	TYPH_SALPA		cytoplasm
Q5PK30	YJX_SALPA		inner membrane
Q5PK31	TRPR_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PK51	RL31_SALPA		cytoplasm
Q5PK55	CAPP_SALPA		cytoplasm
Q5PK67	BTUB_SALPA	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q5PK75	ARGC_SALPA	cytoplasm (probable).	cytoplasm
Q5PK78	MURB_SALPA	cytoplasm (probable).	inner membrane
Q5PK87	THIG_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PK92	RPOC_SALPA		cytoplasm
Q5PK93	RPOB_SALPA		cytoplasm

Q5PK95	RL10_SALPA		cytoplasm
Q5PK96	RL1_SALPA		cytoplasm
Q5PK98	ZRAP_SALPA	periplasm (by similarity).	periplasm
Q5PKA2	NFI_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKA4	NUDC_SALPA		cytoplasm
Q5PKB8	RDOA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKD5	OMPL_SALPA	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q5PKE2	DTD_SALPA	cytoplasm (probable).	cytoplasm
Q5PKG0	FDHD_SALPA	cytoplasm (potential).	cytoplasm
Q5PKG1	RHAR_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKG2	RHAS_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKG3	RHAB_SALPA		cytoplasm
Q5PKG4	RHAA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKG5	RHAD_SALPA	cytoplasm (by similarity).	extracellular
Q5PKI0	RHAT_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PKI2	RIHC_SALPA		extracellular
Q5PKI4	ISPH_SALPA		cytoplasm
Q5PKI8	OADG1_SALPA	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q5PKJ6	CITD2_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKK4	GPPA_SALPA		extracellular
Q5PKK7	WECG_SALPA		cytoplasm
Q5PKK8	WZYE_SALPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q5PKK9	WECF_SALPA	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q5PKM1	CORA_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PKM9	CYAY_SALPA		inner membrane
Q5PKQ2	FADB_SALPA		cytoplasm
Q5PKQ3	FADA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKS5	IBPA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKS6	IBPB_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKS7	YIDE_SALPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q5PKT4	TORD_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKT6	CCMA2_SALPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5PKU4	RNPA_SALPA		extracellular
Q5PKU6	DNAA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKU8	RECF_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKV5	YIDZ_SALPA		cytoplasm
Q5PKV6	MDTL_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PKV8	GLMU_SALPA	cytoplasm (by similarity).	cytoplasm

Q5PKV9	GLMS_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PKX0	ATPA_SALPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5PKX1	ATPG_SALPA		inner membrane
Q5PKX5	MUG_SALPA	cytoplasm (potential).	extracellular
Q5PKZ7	LAMB_SALPA	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q5PL02	PSIE_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PL07	G6PI_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PL14	LEXA_SALPA		cytoplasm
Q5PL20	KEFG_SALPA		cytoplasm
Q5PL21	KEFB_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PL24	SLYX_SALPA		inner membrane
Q5PL31	IDI_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PL58	PURA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PL62	CH60_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PL63	CH10_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PL69	CUTA_SALPA	cytoplasm (by similarity).	periplasm
Q5PLA8	YHBC_SALPA		inner membrane
Q5PLB1	RBFA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PLB2	TRUB_SALPA		cytoplasm
Q5PLB3	RS15_SALPA		cytoplasm
Q5PLB5	RL27_SALPA		cytoplasm
Q5PLC1	RRMJ_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PLC4	GLMM_SALPA		cytoplasm
Q5PLD1	MURA_SALPA	cytoplasm (probable).	inner membrane
Q5PLD3	YHBJ_SALPA		cytoplasm
Q5PLF1	NANE2_SALPA		cytoplasm
Q5PLG7	ORN_SALPA	cytoplasm (potential).	cytoplasm
Q5PLG9	PSD_SALPA		cytoplasm
Q5PLH7	GRCA_SALPA		cytoplasm
Q5PLI6	DSRB_SALPA		inner membrane
Q5PLM6	XYLA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PLN6	DLGD_SALPA	cytoplasm (potential).	cytoplasm
Q5PLQ7	LLDD_SALPA		cytoplasm
Q5PLS8	YBEY_SALPA		inner membrane
Q5PLW2	TSGA_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PLX1	AROK_SALPA	cytoplasm (probable).	cytoplasm
Q5PLX2	AROB_SALPA	cytoplasm (probable).	cytoplasm
Q5PLX6	GPH_SALPA		cytoplasm
Q5PLX8	PPCK_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PLY0	HSLO_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PLY8	BIOH_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PLZ1	FEOB_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PLZ7	GLPE_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PM06	GLGB_SALPA		extracellular

Q5PM08	GLGC_SALPA		cytoplasm
Q5PM09	GLGA_SALPA		cytoplasm
Q5PM73	DNAT_SALPA		nucleoid
Q5PM78	RIHA_SALPA		extracellular
Q5PM81	YBEA_SALPA		inner membrane
Q5PM85	NADD_SALPA		cytoplasm
Q5PM94	CRCB_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PMB1	CITD1_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PME0	ENTS_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PMF1	CAAL_SALPA		cytoplasm
Q5PMJ0	PHOQ_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PMJ1	PHOP_SALPA	cytoplasm (probable).	cytoplasm
Q5PMJ4	TRMU_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PMK2	PEPT_SALPA	cytoplasm (by similarity).	extracellular
Q5PMK3	TRMB_SALPA		cytoplasm
Q5PMK9	HAM1_SALPA		extracellular
Q5PMM0	MLTC_SALPA	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q5PMM1	FETP_SALPA		cytoplasm
Q5PMP2	UXUA_SALPA		cytoplasm
Q5PMP8	UXAC_SALPA		cytoplasm
Q5PMV3	CCME_SALPA	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5PMZ4	YEBC_SALPA		inner membrane
Q5PN00	RUVC_SALPA		cytoplasm
Q5PN01	RUVA_SALPA		cytoplasm
Q5PN10	CVRA_SALPA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q5PN20	TUSD_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PN21	TUSC_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PN46	YFBV_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PN69	RNZ_SALPA		extracellular
Q5PNA4	ARNB_SALPA		cytoplasm
Q5PNA5	ARNC_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PNA6	ARNA_SALPA		cytoplasm
Q5PNA8	ARNT_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5PNC0	LEPA_SALPA		cytoplasm
Q5PND3	CYSZ_SALPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q5PND8	ZIPA_SALPA	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q5PNG5	ISCA_SALPA		cytoplasm
Q5PNH1	HSCA_SALPA		cytoplasm
Q5PNH2	HSCB_SALPA		inner membrane
Q5PNH7	NDK_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PNI2	ISPG_SALPA		cytoplasm
Q5PNK8	KDSA_SALPA	cytoplasm (by similarity).	cytoplasm
Q5PNL7	YOAH_SALPA		inner membrane
Q5PNM0	TRPA_SALPA		cytoplasm
Q5PWZ8	1A1D_PSEPU		cytoplasm
Q5QTW7	PDXH_IDILO		cytoplasm
Q5QTY6	Y970_IDILO		inner membrane
Q5QTY9	RBFA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QTZ0	TRUB_IDILO		cytoplasm
Q5QTZ1	RS15_IDILO		cytoplasm
Q5QU03	DAPA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QU19	PHS_IDILO		cytoplasm
Q5QU37	LPXK_IDILO		cytoplasm
Q5QUB6	MUTS_IDILO		cytoplasm
Q5QUB8	RECA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QUC2	FTSB_IDILO	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q5QUC3	ISPD_IDILO		cytoplasm
Q5QUC4	ISPF_IDILO		extracellular
Q5QUC5	TRUD_IDILO		cytoplasm
Q5QUC6	SURE_IDILO	cytoplasm (potential).	periplasm
Q5QUE5	TRUA_IDILO		cytoplasm
Q5QUF4	DXR_IDILO		cytoplasm
Q5QUN4	PPCK_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QUN6	COAD_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QUN8	TDH_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QUP5	LEXA_IDILO		cytoplasm
Q5QUS3	HEM3_IDILO		cytoplasm
Q5QUT9	DNAE2_IDILO	cytoplasm (probable).	cytoplasm
Q5QUU1	RL19_IDILO		cytoplasm
Q5QUU9	RIMM_IDILO	cytoplasm (potential).	cytoplasm
Q5QUV0	TRMD_IDILO	cytoplasm (potential).	cytoplasm
Q5QUY5	UPPP_IDILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5QUZ5	NAGZ_IDILO	cytoplasm (by similarity).	extracellular
Q5QV03	PTH_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QV04	RL25_IDILO		cytoplasm
Q5QV06	ISPE_IDILO		cytoplasm
Q5QV30	HUTH_IDILO	cytoplasm (potential).	cytoplasm
Q5QV32	DTD_IDILO	cytoplasm (probable).	cytoplasm
Q5QV33	RBN_IDILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5QV36	HSLU_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QV38	RRAA_IDILO		cytoplasm

Q5QV44	RL31_IDILO		cytoplasm
Q5QV48	AROK_IDILO	cytoplasm (probable).	cytoplasm
Q5QV55	Y2477_IDILO		inner membrane
Q5QV59	AROB_IDILO	cytoplasm (probable).	cytoplasm
Q5QVA8	GCST_IDILO		cytoplasm
Q5QVB3	RPIA_IDILO		cytoplasm
Q5QVD9	EX7S_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QVL1	QUEA_IDILO	cytoplasm (potential).	cytoplasm
Q5QVL2	TGT_IDILO		cytoplasm
Q5QVL5	RS21_IDILO		cytoplasm
Q5QVM3	RUVX_IDILO	cytoplasm (potential).	cytoplasm
Q5QVN7	KSGA_IDILO		cytoplasm
Q5QVP5	SYI_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QVQ6	GLUQ_IDILO		cytoplasm
Q5QVQ8	SFSA_IDILO		cytoplasm
Q5QVR4	PANC_IDILO		cytoplasm
Q5QVR5	PANB_IDILO		cytoplasm
Q5QVT4	CH60_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QVW0	PSD_IDILO		cytoplasm
Q5QVW2	ORN_IDILO	cytoplasm (potential).	cytoplasm
Q5QVW5	SLYX_IDILO		inner membrane
Q5QVZ8	HEMH_IDILO	cytoplasm (by similarity).	extracellular
Q5QW04	Y2341_IDILO		inner membrane
Q5QW08	QUEF_IDILO	cytoplasm (probable).	cytoplasm
Q5QW15	MTFA_IDILO		cytoplasm
Q5QW40	AROC_IDILO		cytoplasm
Q5QW45	XGPT_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QW47	SSRP_IDILO	cytoplasm (potential).	cytoplasm
Q5QW95	PURA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QW99	HFQ_IDILO		cytoplasm
Q5QWA0	MIAA_IDILO		cytoplasm
Q5QWA4	RPOC_IDILO		cytoplasm
Q5QWA5	RPOB_IDILO		cytoplasm
Q5QWA7	RL10_IDILO		cytoplasm
Q5QWA8	RL1_IDILO		cytoplasm
Q5QWA9	RL11_IDILO		cytoplasm
Q5QWB5	RS7_IDILO		cytoplasm
Q5QWB6	RS12_IDILO		cytoplasm
Q5QWC7	KITH_IDILO	cytoplasm (potential).	cytoplasm
Q5QWI9	Y164_IDILO		inner membrane
Q5QWP8	HISX_IDILO		cytoplasm
Q5QWQ5	HIS6_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QWQ6	HIS4_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QWQ7	HIS5_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QWQ9	HIS82_IDILO		cytoplasm
Q5QWR1	RECR_IDILO		cytoplasm
Q5QWR2	HTPG_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QWR6	NORM_IDILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5QWW0	G6PI_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QWW5	GUAA_IDILO		cytoplasm
Q5QWY8	MAO1_IDILO		cytoplasm
Q5QWZ9	ASSY_IDILO	cytoplasm (probable).	cytoplasm
Q5QX02	ARGC_IDILO	cytoplasm (probable).	cytoplasm

Q5QX65	PYRD_IDILO	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5QX79	TRPA_IDILO		cytoplasm
Q5QX82	TRPD_IDILO		cytoplasm
Q5QXB8	UVRB_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXE0	CRYD_IDILO		cytoplasm
Q5QXH7	FADB_IDILO		cytoplasm
Q5QXH8	FADA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXH9	TUSA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXI3	SMG_IDILO		cytoplasm
Q5QXI6	FMT_IDILO		cytoplasm
Q5QXJ4	HEM6_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXK8	RRMJ_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXL1	DNAK_IDILO		cytoplasm
Q5QXL2	DNAJ_IDILO	cytoplasm (probable).	cytoplasm
Q5QXL5	DAPB_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXM1	FADJ_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXM6	PPNK_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXN5	FADI_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXN7	TIG_IDILO		cytoplasm
Q5QXN8	CLPP_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXN9	CLPX_IDILO		cytoplasm
Q5QXS0	RS2_IDILO		cytoplasm
Q5QXS3	RRF_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXS9	DAPD_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXT0	GLND_IDILO		cytoplasm
Q5QXT4	GLYA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QXT5	NRDR_IDILO		cytoplasm
Q5QXT8	TYPH_IDILO		cytoplasm
Q5QXV5	RS13_IDILO		cytoplasm
Q5QXV6	RS11_IDILO		cytoplasm
Q5QXV7	RS4_IDILO		cytoplasm
Q5QXV8	RPOA_IDILO		cytoplasm
Q5QXV9	RS8_IDILO		cytoplasm
Q5QXW1	RL18_IDILO		cytoplasm
Q5QXW2	RS5_IDILO		cytoplasm
Q5QXW4	RL15_IDILO		cytoplasm
Q5QXX2	RL29_IDILO		cytoplasm
Q5QXX3	RS17_IDILO		cytoplasm
Q5QXY0	RL2_IDILO		cytoplasm
Q5QXY1	RS19_IDILO		cytoplasm
Q5QXY3	RS3_IDILO		cytoplasm
Q5QXY4	RL16_IDILO		cytoplasm
Q5QXY9	RS10_IDILO		cytoplasm
Q5QY01	RS6_IDILO		cytoplasm
Q5QY04	RL9_IDILO		cytoplasm
Q5QY15	SYV_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QY18	GLNE_IDILO		cytoplasm
Q5QY23	CCA_IDILO		cytoplasm
Q5QY37	RECF_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QY39	DNAA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QY47	Y1969_IDILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5QY51	HAM1_IDILO		extracellular

Q5QY57	PROB_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QY58	FETP_IDILO		cytoplasm
Q5QY61	TRMB_IDILO		cytoplasm
Q5QY68	PROA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QY76	MURB_IDILO	cytoplasm (probable).	inner membrane
Q5QYA9	ISPG_IDILO		cytoplasm
Q5QYB9	NDK_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QYC3	Y943_IDILO		inner membrane
Q5QYD9	Y952_IDILO		inner membrane
Q5QYF8	RHLB_IDILO		cytoplasm
Q5QYG2	UBIE_IDILO		cytoplasm
Q5QYH9	KGUA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QYI0	RPOZ_IDILO		cytoplasm
Q5QYL5	RNPH_IDILO		cytoplasm
Q5QYN9	Y2351_IDILO		inner membrane
Q5QYR6	Y1051_IDILO		cytoplasm
Q5QYU0	TOLB_IDILO	periplasm (potential).	periplasm
Q5QYU4	RUVA_IDILO		cytoplasm
Q5QYU5	RUVB_IDILO		cytoplasm
Q5QYV0	RUVC_IDILO		cytoplasm
Q5QYV3	Y1088_IDILO		inner membrane
Q5QYY0	FABA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QYZ7	TRMU_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QZ06	Y1317_IDILO		cytoplasm
Q5QZ32	KTHY_IDILO		cytoplasm
Q5QZ37	RL32_IDILO		cytoplasm
Q5QZ38	PLSX_IDILO		cytoplasm
Q5QZ48	SERC_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QZ49	HIS81_IDILO		cytoplasm
Q5QZA7	SECB_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QZA9	GPMI_IDILO		cytoplasm
Q5QZB3	RL28_IDILO		cytoplasm
Q5QZB4	RADC_IDILO		cytoplasm
Q5QZB6	DUT_IDILO		extracellular
Q5QZB7	SLMA_IDILO	nucleoid-associated (by similarity).	cytoplasm
Q5QZB8	GPDA_IDILO	cytoplasm (probable).	cytoplasm
Q5QZC0	BIOH_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QZC1	FPG_IDILO		cytoplasm
Q5QZH4	GLMU_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QZH5	GLMS_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QZI4	ATPA_IDILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5QZI5	ATPG_IDILO		inner membrane
Q5QZI8	GIDA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QZI9	GIDB_IDILO		cytoplasm
Q5QZK1	RNPA_IDILO		extracellular
Q5QZK7	RNH2_IDILO	cytoplasm (potential).	extracellular
Q5QZL3	RNH_IDILO	cytoplasm (potential).	extracellular
Q5QZP1	ACCA_IDILO	cytoplasm (by similarity).	cytoplasm
Q5QZP7	CCMA_IDILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q5QZQ1	CCME_IDILO	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5QZR7	ISPH_IDILO		cytoplasm
Q5QZS3	RS20_IDILO		cytoplasm
Q5QZS6	SYI_IDILO	cytoplasm (by similarity).	cytoplasm
Q5R030	MDH_IDILO		cytoplasm
Q5R033	RL27_IDILO		cytoplasm
Q5R041	Y489_IDILO		inner membrane
Q5R053	NUDH_IDILO		cytoplasm
Q5R059	RIMK_IDILO		cytoplasm
Q5R064	TYSY_IDILO	cytoplasm (by similarity).	cytoplasm
Q5R065	LGT_IDILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5R080	UNG_IDILO	cytoplasm (by similarity).	extracellular
Q5R0A6	UVRC_IDILO	cytoplasm (by similarity).	cytoplasm
Q5R0A9	CRCB_IDILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5R0B5	LFTR_IDILO	cytoplasm (by similarity).	cytoplasm
Q5R0C3	CLPS_IDILO		cytoplasm
Q5R0I2	Y393_IDILO		cytoplasm
Q5R0J5	MURA_IDILO	cytoplasm (probable).	inner membrane
Q5R0L7	MRAW_IDILO		cytoplasm
Q5R0L8	MRAZ_IDILO		inner membrane
Q5R0M1	MURG_IDILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5R0M3	MURD_IDILO	cytoplasm (by similarity).	cytoplasm
Q5R0M4	MRAY_IDILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5R0N4	Y448_IDILO		inner membrane
Q5R0N5	Y447_IDILO		cytoplasm
Q5R0P9	FLGH_IDILO	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q5R0Q0	FLGI_IDILO	periplasm (by similarity).	periplasm
Q5R0Q7	TILS_IDILO	cytoplasm (probable).	cytoplasm
Q5R0R2	GLMM_IDILO		cytoplasm
Q5R0Z2	FABZ_IDILO	cytoplasm (by similarity).	cytoplasm
Q5R0Z7	Y819_IDILO		cytoplasm
Q5R104	LEPA_IDILO		cytoplasm
Q5R106	RNC_IDILO	cytoplasm (by similarity).	cytoplasm
Q5R108	RECO_IDILO		cytoplasm
Q5R109	PDXJ_IDILO	cytoplasm (probable).	cytoplasm
Q5R110	ACPS_IDILO	cytoplasm (by similarity).	cytoplasm
Q5R143	ENO_IDILO	cytoplasm (by similarity).	extracellular
Q5R146	THIG_IDILO	cytoplasm (by similarity).	cytoplasm
Q5R192	GCSP_IDILO		cytoplasm
Q5U907	MDH_ACTSC		cytoplasm
Q5WSE7	RNPA_LEGPL		extracellular
Q5WSG6	ATPA_LEGPL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q5WSG7	ATPG_LEGPL		inner membrane
Q5WSH5	PSD_LEGPL		cytoplasm
Q5WSM2	PAND_LEGPL		cytoplasm
Q5WSM3	KSGA_LEGPL		cytoplasm
Q5WSM5	APAH_LEGPL		extracellular
Q5WSQ8	UBIE_LEGPL		cytoplasm
Q5WSS3	GIDB_LEGPL		cytoplasm
Q5WSS4	GIDA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WST8	GLMU_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WSU1	NUDH_LEGPL		cytoplasm
Q5WSU4	LGT_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5WSU5	TYSY_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WSX5	SSRP_LEGPL	cytoplasm (potential).	cytoplasm
Q5WSX6	Y2750_LEGPL		inner membrane
Q5WSX8	GLMS_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WSX9	TPMT_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WSY8	PPNK_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WT13	RRMJ_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WT16	GLMM_LEGPL		cytoplasm
Q5WT34	Y2691_LEGPL		inner membrane
Q5WT37	RBFA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WT38	TRUB_LEGPL		cytoplasm
Q5WT39	RS15_LEGPL		cytoplasm
Q5WT52	RECR_LEGPL		cytoplasm
Q5WT58	CCA_LEGPL		cytoplasm
Q5WT59	ORN_LEGPL	cytoplasm (potential).	cytoplasm
Q5WT65	HEM3_LEGPL		cytoplasm
Q5WT78	CAAL2_LEGPL		cytoplasm
Q5WTA1	MIAA_LEGPL		cytoplasm
Q5WTE1	FEOB_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5WTE7	PTH_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WTE8	RL25_LEGPL		cytoplasm
Q5WTF0	RL27_LEGPL		cytoplasm
Q5WTF5	UVRC_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WTG3	RS20_LEGPL		cytoplasm
Q5WTI4	MRAY_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5WTI5	MURD_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WTI7	MURC_LEGPL	cytoplasm (probable).	cytoplasm
Q5WTI8	MURB_LEGPL	cytoplasm (probable).	inner membrane
Q5WTK7	FMT_LEGPL		cytoplasm
Q5WTR2	MSCL_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5WTW6	DUT_LEGPL		extracellular
Q5WTW9	HAM1_LEGPL		extracellular
Q5WU69	KUP3_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q5WU90	Y2278_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5WU94	MDH_LEGPL		cytoplasm
Q5WUB9	EX7S_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WUE0	GPDA_LEGPL	cytoplasm (probable).	cytoplasm
Q5WUE1	SECB_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WUE6	AROC_LEGPL		cytoplasm
Q5WUK0	UPP_LEGPL		cytoplasm
Q5WUW3	KUP2_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5WV02	ENO_LEGPL	cytoplasm (by similarity).	extracellular
Q5WV15	DNAK_LEGPL		cytoplasm
Q5WV16	DNAJ_LEGPL	cytoplasm (probable).	cytoplasm
Q5WV28	RNPH_LEGPL		cytoplasm
Q5WV30	KGUA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WV31	RPOZ_LEGPL		cytoplasm
Q5WV37	QUEA_LEGPL	cytoplasm (potential).	cytoplasm
Q5WV42	PHS_LEGPL		cytoplasm
Q5WV43	HIS82_LEGPL		cytoplasm
Q5WVC4	FETP_LEGPL		cytoplasm
Q5WVI1	LEPA_LEGPL		cytoplasm
Q5WVI4	RNC_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WVI9	TIG_LEGPL		cytoplasm
Q5WVJ0	CLPP_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WVJ1	CLPX_LEGPL		cytoplasm
Q5WVK6	DTD_LEGPL	cytoplasm (probable).	cytoplasm
Q5WVL3	Y1802_LEGPL		inner membrane
Q5WVN0	PYRD_LEGPL	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5WVN2	MSBA_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5WVN3	LPXK_LEGPL		cytoplasm
Q5WVP6	MUTS_LEGPL		cytoplasm
Q5WVQ0	RECA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WVQ1	RECX_LEGPL	cytoplasm (potential).	cytoplasm
Q5WVS9	LFTR_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WVX4	GUAA_LEGPL		cytoplasm
Q5WVX6	GLND_LEGPL		cytoplasm
Q5WVY7	RS2_LEGPL		cytoplasm
Q5WVZ0	RRF_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WW59	Y1601_LEGPL		inner membrane
Q5WW81	RUMA_LEGPL		cytoplasm
Q5WW99	BIOH_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WWF5	Y1498_LEGPL		cytoplasm
Q5WWH5	NDK_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WWJ9	TOLB_LEGPL	periplasm (potential).	periplasm
Q5WWK4	RUVB_LEGPL		cytoplasm
Q5WWK6	Y1447_LEGPL		cytoplasm
Q5WWL7	RL9_LEGPL		cytoplasm
Q5WWM0	RS6_LEGPL		cytoplasm
Q5WWN7	PROA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WWQ2	TRUD_LEGPL		cytoplasm

Q5WWT0	SERC_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WVV6	PLSX_LEGPL		cytoplasm
Q5WVV7	RL32_LEGPL		cytoplasm
Q5WWW8	HUTH_LEGPL	cytoplasm (potential).	cytoplasm
Q5WWX2	Y1327_LEGPL		inner membrane
Q5WWX5	RNH2_LEGPL	cytoplasm (potential).	extracellular
Q5WWX9	HTPG_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WX31	TRPA_LEGPL		cytoplasm
Q5WX34	TRUA_LEGPL		cytoplasm
Q5WX47	RUVA_LEGPL		cytoplasm
Q5WX48	RUVC_LEGPL		cytoplasm
Q5WX49	Y1249_LEGPL		inner membrane
Q5WX50	HGD_LEGPL		cytoplasm
Q5WX53	SURE_LEGPL	cytoplasm (potential).	periplasm
Q5WX67	FLGI_LEGPL	periplasm (by similarity).	periplasm
Q5WX68	FLGH_LEGPL	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q5WX75	HEM6_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WX91	HISX_LEGPL		cytoplasm
Q5WX92	HIS81_LEGPL		cytoplasm
Q5WX93	HIS7_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WX94	HIS52_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WX95	HIS4_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WXA2	KUP1_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5WXA7	KDSA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WXX1	SYI_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WXX5	AROB_LEGPL	cytoplasm (probable).	cytoplasm
Q5WXX6	AROK_LEGPL	cytoplasm (probable).	cytoplasm
Q5WXZ4	MRAW_LEGPL		cytoplasm
Q5WXZ5	MRAZ_LEGPL		inner membrane
Q5WY20	DAPD_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WY48	CCME_LEGPL	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5WY52	CCMA_LEGPL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5WY61	MURA2_LEGPL	cytoplasm (probable).	inner membrane
Q5WY74	TRPD_LEGPL		cytoplasm
Q5WY89	CLPS_LEGPL		cytoplasm
Q5WY98	MURG_LEGPL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5WYB4	TILS_LEGPL	cytoplasm (probable).	cytoplasm
Q5WYB5	ACCA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WYE0	G6PI_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WYE9	HIS51_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WYH3	NRDR_LEGPL		cytoplasm
Q5WYH4	GLYA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WYI0	SYV_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WYJ0	HUTG_LEGPL		cytoplasm
Q5WYJ7	TDH_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WYL2	CH60_LEGPL	cytoplasm (by similarity).	cytoplasm

Q5WYQ0	RL31_LEGPL		cytoplasm
Q5WYQ8	HSLU_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WYR3	KITH_LEGPL	cytoplasm (potential).	cytoplasm
Q5WYT2	QUEF_LEGPL	cytoplasm (probable).	cytoplasm
Q5WYT4	CAAL1_LEGPL		cytoplasm
Q5WYW3	PYRB_LEGPL		cytoplasm
Q5WYW4	RUVX_LEGPL	cytoplasm (potential).	cytoplasm
Q5WYX3	MURA1_LEGPL	cytoplasm (probable).	inner membrane
Q5WYX4	SYU_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WYY6	FPG_LEGPL		cytoplasm
Q5WYZ4	COAD_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WZ02	PDXH_LEGPL		cytoplasm
Q5WZ32	CRCB_LEGPL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5WZ34	FABZ_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WZ40	UPPS_LEGPL		cytoplasm
Q5WZ43	GPMI_LEGPL		cytoplasm
Q5WZ50	ASSY_LEGPL	cytoplasm (probable).	cytoplasm
Q5WZ56	PURA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WZ61	TRMB_LEGPL		cytoplasm
Q5WZ62	RL28_LEGPL		cytoplasm
Q5WZ79	PRMA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5WZB4	HEMH_LEGPL	cytoplasm (by similarity).	extracellular
Q5WZC5	RIMK_LEGPL		cytoplasm
Q5WZE1	RIMM_LEGPL	cytoplasm (potential).	cytoplasm
Q5WZE2	TRMD_LEGPL	cytoplasm (potential).	cytoplasm
Q5WZE3	RL19_LEGPL		cytoplasm
Q5WZI7	RPOA_LEGPL		cytoplasm
Q5WZI8	RS4_LEGPL		cytoplasm
Q5WZI9	RS11_LEGPL		cytoplasm
Q5WZJ0	RS13_LEGPL		cytoplasm
Q5WZJ3	RL15_LEGPL		cytoplasm
Q5WZJ5	RS5_LEGPL		cytoplasm
Q5WZJ6	RL18_LEGPL		cytoplasm
Q5WZK3	RS17_LEGPL		cytoplasm
Q5WZK4	RL29_LEGPL		cytoplasm
Q5WZK5	RL16_LEGPL		cytoplasm
Q5WZK6	RS3_LEGPL		cytoplasm
Q5WZK8	RS19_LEGPL		cytoplasm
Q5WZK9	RL2_LEGPL		cytoplasm
Q5WZL3	RS10_LEGPL		cytoplasm
Q5WZL6	RS7_LEGPL		cytoplasm
Q5WZL7	RS12_LEGPL		cytoplasm
Q5WZL8	RPOC_LEGPL		cytoplasm
Q5WZL9	RPOB_LEGPL		cytoplasm
Q5WZM1	RL10_LEGPL		cytoplasm
Q5WZM2	RL1_LEGPL		cytoplasm
Q5WZM3	RL11_LEGPL		cytoplasm
Q5WZW1	Y270_LEGPL		cytoplasm
Q5X090	DAPB_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5X0A4	GCST_LEGPL		cytoplasm
Q5X0C7	RPIA_LEGPL		cytoplasm
Q5X0D2	Y088_LEGPL		inner membrane
Q5X0E6	UVRB_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5X0L0	HFQ_LEGPL		cytoplasm

Q5X0L4	AGUA_LEGPL		extracellular
Q5X0L8	DNAA_LEGPL	cytoplasm (by similarity).	cytoplasm
Q5X0M0	RNPA_LEGPA		extracellular
Q5X0P1	ATPA2_LEGPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5X0P2	ATPG_LEGPA		inner membrane
Q5X0Q0	PSD_LEGPA		cytoplasm
Q5X0V0	PAND_LEGPA		cytoplasm
Q5X0V1	KSGA_LEGPA		cytoplasm
Q5X0V3	APAH_LEGPA		extracellular
Q5X0X6	UBIE_LEGPA		cytoplasm
Q5X0Z7	GIDB_LEGPA		cytoplasm
Q5X0Z8	GIDA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X112	GLMU_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X115	NUDH_LEGPA		cytoplasm
Q5X118	LGT_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X119	TYSY_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X148	SSRP_LEGPA	cytoplasm (potential).	cytoplasm
Q5X149	Y2897_LEGPA		inner membrane
Q5X153	GLMS_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X154	TPMT_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X168	PPNK_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X1A0	RRMJ_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X1A3	GLMM_LEGPA		cytoplasm
Q5X1C1	Y2822_LEGPA		inner membrane
Q5X1C4	RBFA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X1C5	TRUB_LEGPA		cytoplasm
Q5X1C6	RS15_LEGPA		cytoplasm
Q5X1D9	RECR_LEGPA		cytoplasm
Q5X1E5	CCA_LEGPA		cytoplasm
Q5X1E6	ORN_LEGPA	cytoplasm (potential).	cytoplasm
Q5X1F2	HEM3_LEGPA		cytoplasm
Q5X1G7	CAAL2_LEGPA		cytoplasm
Q5X1J2	MIAA_LEGPA		cytoplasm
Q5X1N2	FEOB_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X1N7	PTH_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X1N8	RL25_LEGPA		cytoplasm
Q5X1P0	RL27_LEGPA		cytoplasm
Q5X1P5	UVRC_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X1Q4	RS20_LEGPA		cytoplasm
Q5X1S3	MRAY_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X1S4	MURD_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X1S6	MURC_LEGPA	cytoplasm (probable).	cytoplasm
Q5X1S7	MURB_LEGPA	cytoplasm (probable).	inner membrane
Q5X1U6	FMT_LEGPA		cytoplasm
Q5X1Y7	MSCL_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X242	DUT_LEGPA		extracellular

Q5X245	HAM1_LEGPA		extracellular
Q5X2E7	KUP3_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X2M8	TYPH_LEGPA		cytoplasm
Q5X2Q3	ATPA1_LEGPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5X2T2	Y2305_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X2T6	MDH_LEGPA		cytoplasm
Q5X2W0	EX7S_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X2Y0	GPDA_LEGPA	cytoplasm (probable).	cytoplasm
Q5X2Y1	SECB_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X2Y6	AROC_LEGPA		cytoplasm
Q5X340	UPP_LEGPA		cytoplasm
Q5X3H2	KUP2_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X3L4	ENO_LEGPA	cytoplasm (by similarity).	extracellular
Q5X3M7	DNAK_LEGPA		cytoplasm
Q5X3M8	DNAJ_LEGPA	cytoplasm (probable).	cytoplasm
Q5X3P0	RNPH_LEGPA		cytoplasm
Q5X3P2	KGUA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X3P3	RPOZ_LEGPA		cytoplasm
Q5X3P9	QUEA_LEGPA	cytoplasm (potential).	cytoplasm
Q5X3Q4	PHS_LEGPA		cytoplasm
Q5X3Q5	HIS82_LEGPA		cytoplasm
Q5X3X9	FETP_LEGPA		cytoplasm
Q5X443	LEPA_LEGPA		cytoplasm
Q5X446	RNC_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X450	TIG_LEGPA		cytoplasm
Q5X451	CLPP_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X452	CLPX_LEGPA		cytoplasm
Q5X472	DTD_LEGPA	cytoplasm (probable).	cytoplasm
Q5X479	Y1801_LEGPA		inner membrane
Q5X496	PYRD_LEGPA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5X498	MSBA_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X499	LPXK_LEGPA		cytoplasm
Q5X4B2	MUTS_LEGPA		cytoplasm
Q5X4B5	RECA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X4B6	RECX_LEGPA	cytoplasm (potential).	cytoplasm
Q5X4E4	LFTR_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X4I9	GUAA_LEGPA		cytoplasm
Q5X4J1	GLND_LEGPA		cytoplasm
Q5X4J7	RS2_LEGPA		cytoplasm
Q5X4K0	RRF_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X4T2	TRUD_LEGPA		cytoplasm
Q5X4U9	PROA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X4X0	RS6_LEGPA		cytoplasm
Q5X4X3	RL9_LEGPA		cytoplasm

Q5X4Y4	Y1536_LEGPA		cytoplasm
Q5X4Y6	RUVB_LEGPA		cytoplasm
Q5X4Z1	TOLB_LEGPA	periplasm (potential).	periplasm
Q5X515	NDK_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X535	Y1485_LEGPA		cytoplasm
Q5X590	BIOH_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X5A8	RUMA_LEGPA		cytoplasm
Q5X5C5	Y1395_LEGPA		inner membrane
Q5X5E7	SERC_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X5G6	KTHY_LEGPA		cytoplasm
Q5X5H3	PLSX_LEGPA		cytoplasm
Q5X5H4	RL32_LEGPA		cytoplasm
Q5X5I5	HUTH_LEGPA	cytoplasm (potential).	cytoplasm
Q5X5I9	Y1331_LEGPA		inner membrane
Q5X5J7	HTPG_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X5Q1	TRPA_LEGPA		cytoplasm
Q5X5Q4	TRUA_LEGPA		cytoplasm
Q5X5R9	RUVA_LEGPA		cytoplasm
Q5X5S0	RUVC_LEGPA		cytoplasm
Q5X5S1	Y1249_LEGPA		inner membrane
Q5X5S2	HGD_LEGPA		cytoplasm
Q5X5S5	SURE_LEGPA	cytoplasm (potential).	periplasm
Q5X5T9	FLGI_LEGPA	periplasm (by similarity).	periplasm
Q5X5U0	FLGH_LEGPA	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q5X5U7	HEM6_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X5W9	HISX_LEGPA		cytoplasm
Q5X5X0	HIS81_LEGPA		cytoplasm
Q5X5X1	HIS7_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X5X2	HIS52_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X5X3	HIS4_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X5Y0	KUP1_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X5Y5	KDSA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X6G6	SYI_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X6H0	AROB_LEGPA	cytoplasm (probable).	cytoplasm
Q5X6H1	AROK_LEGPA	cytoplasm (probable).	cytoplasm
Q5X6J0	MRAW_LEGPA		cytoplasm
Q5X6J1	MRAZ_LEGPA		inner membrane
Q5X6L6	DAPD_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X6P3	CCME_LEGPA	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5X6P7	CCMA_LEGPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5X6Q6	MURA_LEGPA	cytoplasm (probable).	inner membrane
Q5X6R9	TRPD_LEGPA		cytoplasm
Q5X6T6	CLPS_LEGPA		cytoplasm
Q5X6U5	MURG_LEGPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5X6W4	TILS_LEGPA	cytoplasm (probable).	cytoplasm
Q5X6W5	ACCA_LEGPA	cytoplasm (by similarity).	cytoplasm

Q5X6Y8	G6PI_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X6Z7	HIS51_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X721	NRDR_LEGPA		cytoplasm
Q5X722	GLYA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X728	SYV_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X741	HUTG_LEGPA		cytoplasm
Q5X748	TDH_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X762	CH60_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X7A1	RL31_LEGPA		cytoplasm
Q5X7B0	HSLU_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X7B5	KITH_LEGPA	cytoplasm (potential).	cytoplasm
Q5X7D5	QUEF_LEGPA	cytoplasm (probable).	cytoplasm
Q5X7D7	CAAL1_LEGPA		cytoplasm
Q5X7G7	PYRB_LEGPA		cytoplasm
Q5X7G8	RUVX_LEGPA	cytoplasm (potential).	cytoplasm
Q5X7H7	SYV_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X7I9	FPG_LEGPA		cytoplasm
Q5X7J6	COAD_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X7K4	PDXH_LEGPA		cytoplasm
Q5X7N1	CRCB_LEGPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5X7N3	FABZ_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X7N9	UPPS_LEGPA		cytoplasm
Q5X7P2	GPMI_LEGPA		cytoplasm
Q5X7P9	ASSY_LEGPA	cytoplasm (probable).	cytoplasm
Q5X7Q5	PURA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X7R0	TRMB_LEGPA		cytoplasm
Q5X7R1	RL28_LEGPA		cytoplasm
Q5X7S8	PRMA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X7W3	HEMH_LEGPA	cytoplasm (by similarity).	extracellular
Q5X7X4	RIMK_LEGPA		cytoplasm
Q5X7Z0	RIMM_LEGPA	cytoplasm (potential).	cytoplasm
Q5X7Z1	TRMD_LEGPA	cytoplasm (potential).	cytoplasm
Q5X7Z2	RL19_LEGPA		cytoplasm
Q5X834	RPOA_LEGPA		cytoplasm
Q5X835	RS4_LEGPA		cytoplasm
Q5X836	RS11_LEGPA		cytoplasm
Q5X837	RS13_LEGPA		cytoplasm
Q5X840	RL15_LEGPA		cytoplasm
Q5X842	RS5_LEGPA		cytoplasm
Q5X843	RL18_LEGPA		cytoplasm
Q5X850	RS17_LEGPA		cytoplasm
Q5X851	RL29_LEGPA		cytoplasm
Q5X852	RL16_LEGPA		cytoplasm
Q5X853	RS3_LEGPA		cytoplasm
Q5X855	RS19_LEGPA		cytoplasm
Q5X856	RL2_LEGPA		cytoplasm
Q5X860	RS10_LEGPA		cytoplasm
Q5X863	RS7_LEGPA		cytoplasm
Q5X864	RS12_LEGPA		cytoplasm
Q5X865	RPOC_LEGPA		cytoplasm
Q5X866	RPOB_LEGPA		cytoplasm
Q5X868	RL10_LEGPA		cytoplasm
Q5X869	RL1_LEGPA		cytoplasm
Q5X870	RL11_LEGPA		cytoplasm

Q5X8H6	Y275_LEGPA		cytoplasm
Q5X8U7	DAPB_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X8W1	GCST_LEGPA		cytoplasm
Q5X8Y4	RPIA_LEGPA		cytoplasm
Q5X8Y9	Y103_LEGPA		inner membrane
Q5X906	UVRB_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5X982	HFQ_LEGPA		cytoplasm
Q5X986	AGUA_LEGPA		extracellular
Q5X990	DNAA_LEGPA	cytoplasm (by similarity).	cytoplasm
Q5XWD9	TUSB_PECCC	cytoplasm (by similarity).	cytoplasm
Q5ZR99	ATPA2_LEGPH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5ZRA0	ATPG_LEGPH		inner membrane
Q5ZRA9	PSD_LEGPH		cytoplasm
Q5ZRF3	PAND_LEGPH		cytoplasm
Q5ZRF4	KSGA_LEGPH		cytoplasm
Q5ZRF6	APAH_LEGPH		extracellular
Q5ZRH9	UBIE_LEGPH		cytoplasm
Q5ZRJ1	GIDB_LEGPH		cytoplasm
Q5ZRJ2	GIDA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZRK6	GLMU_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZRK9	NUDH_LEGPH		cytoplasm
Q5ZRL2	LGT_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5ZRL3	TYSY_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZRP1	SSRP_LEGPH	cytoplasm (potential).	cytoplasm
Q5ZRP2	Y2838_LEGPH		inner membrane
Q5ZRP4	GLMS_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZRP5	TPMT_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZRQ7	PPNK_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZRT1	RRMJ_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZRT4	GLMM_LEGPH		cytoplasm
Q5ZRV2	Y2774_LEGPH		inner membrane
Q5ZRV5	RBFA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZRV6	TRUB_LEGPH		cytoplasm
Q5ZRV7	RS15_LEGPH		cytoplasm
Q5ZRX0	RECR_LEGPH		cytoplasm
Q5ZRX9	CCA_LEGPH		cytoplasm
Q5ZRY0	ORN_LEGPH	cytoplasm (potential).	cytoplasm
Q5ZRY6	HEM3_LEGPH		cytoplasm
Q5ZS01	CAAL_LEGPH		cytoplasm
Q5ZS23	MIAA_LEGPH		cytoplasm
Q5ZS58	PANB_LEGPH		cytoplasm
Q5ZS62	FEOB_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5ZS66	PTH_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZS67	RL25_LEGPH		cytoplasm
Q5ZS69	RL27_LEGPH		cytoplasm
Q5ZS74	UVRC_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZS83	RS20_LEGPH		cytoplasm
Q5ZSA2	MRAY_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q5ZSA3	MURD_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZSA5	MURC_LEGPH	cytoplasm (probable).	cytoplasm
Q5ZSA6	MURB_LEGPH	cytoplasm (probable).	inner membrane
Q5ZSC5	FMT_LEGPH		cytoplasm
Q5ZSH7	MSCL_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5ZSN0	DUT_LEGPH		extracellular
Q5ZSN3	HAM1_LEGPH		extracellular
Q5ZSY2	KUP3_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5ZT09	Y2356_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5ZT13	MDH_LEGPH		cytoplasm
Q5ZT17	SODC_LEGPH	periplasm (by similarity).	periplasm
Q5ZT36	EX7S_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZT56	GPDA_LEGPH	cytoplasm (probable).	cytoplasm
Q5ZT57	SECB_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZT62	AROC_LEGPH		cytoplasm
Q5ZT84	LLY_LEGPH		extracellular
Q5ZTB9	UPP_LEGPH		cytoplasm
Q5ZTN5	KUP2_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5ZTX1	ENO_LEGPH	cytoplasm (by similarity).	extracellular
Q5ZTY3	DNAK_LEGPH		cytoplasm
Q5ZTY4	DNAJ_LEGPH	cytoplasm (probable).	cytoplasm
Q5ZTZ6	RNPH_LEGPH		cytoplasm
Q5ZTZ8	KGUA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZU04	QUEA_LEGPH	cytoplasm (potential).	cytoplasm
Q5ZU09	PHS_LEGPH		cytoplasm
Q5ZU10	HIS82_LEGPH		cytoplasm
Q5ZU80	FETP_LEGPH		cytoplasm
Q5ZUD2	LEPA_LEGPH		cytoplasm
Q5ZUD5	RNC_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZUD8	TIG_LEGPH		cytoplasm
Q5ZUD9	CLPP_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZUE0	CLPX_LEGPH		cytoplasm
Q5ZUF4	DTD_LEGPH	cytoplasm (probable).	cytoplasm
Q5ZUG1	Y1837_LEGPH		inner membrane
Q5ZUH7	PYRD_LEGPH	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q5ZUH9	MSBA_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5ZUI0	LPXK_LEGPH		cytoplasm
Q5ZUJ3	MUTS_LEGPH		cytoplasm
Q5ZUJ7	RECA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZUM5	LFTR_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZUS0	GUAA_LEGPH		cytoplasm
Q5ZUS2	GLND_LEGPH		cytoplasm
Q5ZUS8	RS2_LEGPH		cytoplasm
Q5ZUT1	RRF_LEGPH	cytoplasm (by similarity).	cytoplasm

Q5ZV19	TRUD_LEGPH		cytoplasm
Q5ZV33	PROA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZV48	RS6_LEGPH		cytoplasm
Q5ZV51	RL9_LEGPH		cytoplasm
Q5ZV64	RUVB_LEGPH		cytoplasm
Q5ZV69	TOLB_LEGPH	periplasm (potential).	periplasm
Q5ZV92	NDK_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZVB2	Y1528_LEGPH		cytoplasm
Q5ZVG6	BIOH_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZVI4	RUMA_LEGPH		cytoplasm
Q5ZVK0	Y1440_LEGPH		inner membrane
Q5ZVM2	SERC_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZVP1	KTHY_LEGPH		cytoplasm
Q5ZVP8	PLSX_LEGPH		cytoplasm
Q5ZVP9	RL32_LEGPH		cytoplasm
Q5ZVR0	HUTH_LEGPH	cytoplasm (potential).	cytoplasm
Q5ZVR4	Y1376_LEGPH		inner membrane
Q5ZVS1	HTPG_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZVY3	TRPA_LEGPH		cytoplasm
Q5ZVY6	TRUA_LEGPH		cytoplasm
Q5ZW00	RUVA_LEGPH		cytoplasm
Q5ZW01	RUVC_LEGPH		cytoplasm
Q5ZW02	Y1286_LEGPH		inner membrane
Q5ZW64	FLGI_LEGPH	periplasm (by similarity).	periplasm
Q5ZW65	FLGH_LEGPH	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q5ZW72	HEM6_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZW87	HISX_LEGPH		cytoplasm
Q5ZW88	HIS81_LEGPH		cytoplasm
Q5ZW89	HIS7_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZW90	HIS52_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZW91	HIS4_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZW98	KUP1_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5ZWA3	KDSA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZWN7	ATPA1_LEGPH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5ZWR3	TYPH_LEGPH		cytoplasm
Q5ZWZ6	SYI_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZX00	AROB_LEGPH	cytoplasm (probable).	cytoplasm
Q5ZX01	AROK_LEGPH	cytoplasm (probable).	cytoplasm
Q5ZX19	MRAW_LEGPH		cytoplasm
Q5ZX20	MRAZ_LEGPH		inner membrane
Q5ZX45	DAPD_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZX72	CCME_LEGPH	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q5ZX76	CCMA_LEGPH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5ZX85	MURA_LEGPH	cytoplasm (probable).	inner membrane
Q5ZX98	TRPD_LEGPH		cytoplasm
Q5ZXB5	CLPS_LEGPH		cytoplasm

Q5ZXC4	MURG_LEGPH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q5ZXE5	TILS_LEGPH	cytoplasm (probable).	cytoplasm
Q5ZXE6	ACCA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZXH2	G6PI_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZXI1	HIS51_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZXK5	NRDR_LEGPH		cytoplasm
Q5ZXK6	GLYA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZXL2	SYV_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZXM2	HUTG_LEGPH		cytoplasm
Q5ZXM9	TDH_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZXP3	CH60_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZXT1	RL31_LEGPH		cytoplasm
Q5ZXU0	HSLU_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZXU5	KITH_LEGPH	cytoplasm (potential).	cytoplasm
Q5ZXZ2	PYRB_LEGPH		cytoplasm
Q5ZXZ3	RUVX_LEGPH	cytoplasm (potential).	cytoplasm
Q5ZY07	SYV_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZY18	FPG_LEGPH		cytoplasm
Q5ZY26	COAD_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZY37	PDXH_LEGPH		cytoplasm
Q5ZY60	CRCB_LEGPH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q5ZY62	FABZ_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZY69	UPPS_LEGPH		cytoplasm
Q5ZY71	GPMI_LEGPH		cytoplasm
Q5ZY78	ASSY_LEGPH	cytoplasm (probable).	cytoplasm
Q5ZY86	PURA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZY91	TRMB_LEGPH		cytoplasm
Q5ZY93	RL28_LEGPH		cytoplasm
Q5ZYP1	PRMA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZYE7	HEMH_LEGPH	cytoplasm (by similarity).	extracellular
Q5ZYF8	RIMK_LEGPH		cytoplasm
Q5ZYH5	RIMM_LEGPH	cytoplasm (potential).	cytoplasm
Q5ZYH6	TRMD_LEGPH	cytoplasm (potential).	cytoplasm
Q5ZYH7	RL19_LEGPH		cytoplasm
Q5ZYL8	RPOA_LEGPH		cytoplasm
Q5ZYL9	RS4_LEGPH		cytoplasm
Q5ZYM0	RS11_LEGPH		cytoplasm
Q5ZYM1	RS13_LEGPH		cytoplasm
Q5ZYM4	RL15_LEGPH		cytoplasm
Q5ZYM6	RS5_LEGPH		cytoplasm
Q5ZYM7	RL18_LEGPH		cytoplasm
Q5ZYN4	RS17_LEGPH		cytoplasm
Q5ZYN5	RL29_LEGPH		cytoplasm
Q5ZYN6	RL16_LEGPH		cytoplasm
Q5ZYN7	RS3_LEGPH		cytoplasm
Q5ZYP0	RL2_LEGPH		cytoplasm
Q5ZYP4	RS10_LEGPH		cytoplasm
Q5ZYP7	RS7_LEGPH		cytoplasm
Q5ZYP8	RS12_LEGPH		cytoplasm
Q5ZYP9	RPOC_LEGPH		cytoplasm
Q5ZYQ0	RPOB_LEGPH		cytoplasm
Q5ZYQ2	RL10_LEGPH		cytoplasm

Q5ZYQ3	RL1_LEGPH		cytoplasm
Q5ZYQ4	RL11_LEGPH		cytoplasm
Q5ZYZ5	Y216_LEGPH		cytoplasm
Q5ZZ14	Y197_LEGPH		inner membrane
Q5ZZ80	DAPB_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZZ93	GCST_LEGPH		cytoplasm
Q5ZZB7	RPIA_LEGPH		cytoplasm
Q5ZZC2	Y089_LEGPH		inner membrane
Q5ZZD9	UVRB_LEGPH	cytoplasm (by similarity).	cytoplasm
Q5ZZK1	HFQ_LEGPH		cytoplasm
Q5ZZK4	AGUA_LEGPH		extracellular
Q5ZZK8	DNAA_LEGPH	cytoplasm (by similarity).	cytoplasm
Q60006	FOLD_SALTI		cytoplasm
Q60010	NUOH_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q60028	RBL2_THIDA		extracellular
Q60099	HAD_XANAU		cytoplasm
Q60101	RECA_XANCP	cytoplasm (by similarity).	cytoplasm
Q60107	CLPP_YEREN	cytoplasm (by similarity).	cytoplasm
Q60112	AROA_YERPE	cytoplasm (probable).	cytoplasm
Q60115	INVB_ZYMMO		cytoplasm
Q60132	T2C9_CITFR		extracellular
Q60153	TCPA1_VIBCH	secreted protein (by similarity).	extracellular
Q60169	HEM3_PSEAE		cytoplasm
Q60214	NIR_RHIHE	periplasm (by similarity).	periplasm
Q60216	TRAN4_ECOLI		cytoplasm
Q60228	RELX_ECOLI		cytoplasm
Q60235	HRCQB_PSESY	cytoplasm (by similarity). or loosely associated with a membrane component, probably hrcqa (by similarity).	inner membrane
Q60236	HRPW_PSESY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q60251	CHEW_RHOSH	cytoplasm (by similarity).	cytoplasm
Q602J1	FPG_METCA		cytoplasm
Q602K3	HEM3_METCA		cytoplasm
Q602L8	TAL_METCA	cytoplasm (probable).	cytoplasm
Q602M8	RNPA_METCA		extracellular
Q602N0	DNAA_METCA	cytoplasm (by similarity).	cytoplasm
Q602N2	RECF_METCA	cytoplasm (by similarity).	cytoplasm
Q602N9	RNPH_METCA		cytoplasm
Q602P0	HAM1_METCA		extracellular
Q602S2	Y2989_METCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q602T6	KGUA_METCA	cytoplasm (by similarity).	cytoplasm
Q603C1	NDK_METCA	cytoplasm (by similarity).	cytoplasm
Q603C9	FABA_METCA	cytoplasm (by similarity).	cytoplasm
Q603G2	FMT_METCA		cytoplasm
Q603K0	HIS7_METCA	cytoplasm (by similarity).	cytoplasm
Q603K1	HIS5_METCA	cytoplasm (by similarity).	cytoplasm
Q603K2	HIS4_METCA	cytoplasm (by similarity).	cytoplasm
Q603K3	HIS6_METCA	cytoplasm (by similarity).	cytoplasm

Q603K4	HIS3_METCA	cytoplasm (by similarity).	cytoplasm
Q603K5	HIS2_METCA	cytoplasm (by similarity).	cytoplasm
Q603L4	HEM6_METCA	cytoplasm (by similarity).	cytoplasm
Q603M1	DUT_METCA		extracellular
Q603M3	ARGB_METCA	cytoplasm (probable).	cytoplasm
Q603Q7	RBL_METCA		extracellular
Q603S2	TYSY_METCA	cytoplasm (by similarity).	cytoplasm
Q603S3	LGT_METCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q603U2	ATPA2_METCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q604C6	SSRP_METCA	cytoplasm (potential).	cytoplasm
Q604D9	GLGA2_METCA		cytoplasm
Q604F8	TRPD_METCA		cytoplasm
Q604K1	GPDA_METCA	cytoplasm (probable).	cytoplasm
Q604K2	SECB_METCA	cytoplasm (by similarity).	cytoplasm
Q604M0	TRUD_METCA		cytoplasm
Q604M1	ISPF_METCA		extracellular
Q604M2	ISPD_METCA		cytoplasm
Q604M4	ENO2_METCA	cytoplasm (by similarity).	extracellular
Q604M5	KDSA_METCA	cytoplasm (by similarity).	cytoplasm
Q604P1	TRUA_METCA		cytoplasm
Q604P4	TRPA_METCA		cytoplasm
Q604Q5	ISPG_METCA		cytoplasm
Q604U1	FABZ_METCA	cytoplasm (by similarity).	cytoplasm
Q604U3	RNH2_METCA	cytoplasm (potential).	extracellular
Q604U5	QUEF_METCA	cytoplasm (probable).	cytoplasm
Q604U9	MRAZ_METCA		inner membrane
Q604V0	MRAW_METCA		cytoplasm
Q604V4	MRAY_METCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q604V5	MURD_METCA	cytoplasm (by similarity).	cytoplasm
Q604V7	MURG_METCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q604V8	MURC_METCA	cytoplasm (probable).	cytoplasm
Q604V9	MURB_METCA	cytoplasm (probable).	inner membrane
Q604W8	SURE_METCA	cytoplasm (potential).	periplasm
Q605A7	RS12_METCA		cytoplasm
Q605A8	RS7_METCA		cytoplasm
Q605B1	RS10_METCA		cytoplasm
Q605B5	RL2_METCA		cytoplasm
Q605B6	RS19_METCA		cytoplasm
Q605B8	RS3_METCA		cytoplasm
Q605B9	RL16_METCA		cytoplasm
Q605C0	RL29_METCA		cytoplasm
Q605C1	RS17_METCA		cytoplasm
Q605C6	RS8_METCA		cytoplasm
Q605C8	RL18_METCA		cytoplasm
Q605C9	RS5_METCA		cytoplasm
Q605D1	RL15_METCA		cytoplasm
Q605D4	RS13_METCA		cytoplasm
Q605D5	RS11_METCA		cytoplasm

Q605D6	RS4_METCA		cytoplasm
Q605D7	RPOA_METCA		cytoplasm
Q605D9	GLNE_METCA		cytoplasm
Q605E5	RL28_METCA		cytoplasm
Q605E9	RUVX_METCA	cytoplasm (potential).	cytoplasm
Q605F1	PYRB_METCA		cytoplasm
Q605G6	Y2318_METCA		inner membrane
Q605G8	PAND_METCA		cytoplasm
Q605G9	PANC_METCA		cytoplasm
Q605H0	PANB_METCA		cytoplasm
Q605I0	COBQ_METCA		cytoplasm
Q605K9	ILVC_METCA		cytoplasm
Q605M7	SYI_METCA	cytoplasm (by similarity).	cytoplasm
Q605N0	RS20_METCA		cytoplasm
Q605N1	PROB_METCA	cytoplasm (by similarity).	cytoplasm
Q605N3	RL27_METCA		cytoplasm
Q605Q9	FTHS_METCA		cytoplasm
Q605R1	ATKA_METCA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q605S7	UVRB_METCA	cytoplasm (by similarity).	cytoplasm
Q605V8	LEXA_METCA		cytoplasm
Q605W1	DNAE2_METCA	cytoplasm (probable).	cytoplasm
Q606C1	SYV_METCA	cytoplasm (by similarity).	cytoplasm
Q606C7	Y2091_METCA		inner membrane
Q606C8	Y2090_METCA		cytoplasm
Q606D2	NUDH_METCA		cytoplasm
Q606D6	ILVD_METCA		cytoplasm
Q606F1	KAD_METCA	cytoplasm (by similarity).	cytoplasm
Q606F2	LEU2_METCA		cytoplasm
Q606F3	LEUD_METCA		cytoplasm
Q606F4	LEU3_METCA	cytoplasm (by similarity).	cytoplasm
Q606H8	RS6_METCA		cytoplasm
Q606H9	RS18_METCA		cytoplasm
Q606I1	RL9_METCA		cytoplasm
Q606J4	RPOZ_METCA		cytoplasm
Q606J9	UBIE_METCA		cytoplasm
Q606K1	HSLU_METCA	cytoplasm (by similarity).	cytoplasm
Q606K2	HSLV_METCA	cytoplasm (by similarity).	cytoplasm
Q606L1	RL32_METCA		cytoplasm
Q606L2	PLSX_METCA		cytoplasm
Q606L6	ACP_METCA	cytoplasm (by similarity).	cytoplasm
Q606M0	KTHY_METCA		cytoplasm
Q606N2	NAGZ_METCA	cytoplasm (by similarity).	extracellular
Q606N3	HFQ_METCA		cytoplasm
Q606N8	PURA_METCA	cytoplasm (by similarity).	cytoplasm
Q606P3	CRCB_METCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q606Q0	MURA_METCA	cytoplasm (probable).	inner membrane
Q606Q1	HIS1_METCA	cytoplasm (by similarity).	cytoplasm
Q606Q2	HISX_METCA		cytoplasm
Q606T2	ENO1_METCA	cytoplasm (by similarity).	extracellular
Q606W5	RUMA_METCA		cytoplasm
Q606Y1	PROA_METCA	cytoplasm (by similarity).	cytoplasm
Q607A2	PPNK_METCA	cytoplasm (by similarity).	cytoplasm

Q607A3	HRC_A_METCA		cytoplasm
Q607A5	DNAK_METCA		cytoplasm
Q607A6	DNAJ_METCA	cytoplasm (probable).	cytoplasm
Q607A7	DAPB_METCA	cytoplasm (by similarity).	cytoplasm
Q607B2	RRMJ_METCA	cytoplasm (by similarity).	cytoplasm
Q607B4	GLMM_METCA		cytoplasm
Q607D1	CLPX3_METCA		cytoplasm
Q607E5	ISPH_METCA		cytoplasm
Q607E7	RRAA_METCA		cytoplasm
Q607F5	TILS_METCA	cytoplasm (probable).	cytoplasm
Q607F6	ACCA_METCA	cytoplasm (by similarity).	cytoplasm
Q607G8	LFTR_METCA	cytoplasm (by similarity).	cytoplasm
Q607H2	CLPS_METCA		cytoplasm
Q607L6	DTD_METCA	cytoplasm (probable).	cytoplasm
Q607Q3	CH602_METCA	cytoplasm (by similarity).	cytoplasm
Q607R4	MIAA_METCA		cytoplasm
Q607S4	UVRC_METCA	cytoplasm (by similarity).	cytoplasm
Q607S6	ARGJ_METCA	cytoplasm (probable).	cytoplasm
Q607T4	HEMH_METCA	cytoplasm (by similarity).	extracellular
Q607U4	GLYA_METCA	cytoplasm (by similarity).	cytoplasm
Q607U5	NRDR_METCA		cytoplasm
Q607X6	NPD_METCA	cytoplasm (probable).	cytoplasm
Q608B0	PHK_METCA		cytoplasm
Q608F1	Y1541_METCA		cytoplasm
Q608G0	TRMB_METCA		cytoplasm
Q608K0	DAPD_METCA	cytoplasm (by similarity).	cytoplasm
Q608L4	GLGA1_METCA		cytoplasm
Q608L5	GLGB_METCA		extracellular
Q608L6	GLGC_METCA		cytoplasm
Q608M4	LEPA_METCA		cytoplasm
Q608M7	RNC_METCA	cytoplasm (by similarity).	cytoplasm
Q608M9	RECO_METCA		cytoplasm
Q608N3	Y1457_METCA		inner membrane
Q608P0	PQQE_METCA		cytoplasm
Q608P1	PQQD_METCA		cytoplasm
Q608P3	PQQB_METCA		cytoplasm
Q608S3	HIS82_METCA		cytoplasm
Q608S6	KCY_METCA	cytoplasm (by similarity).	cytoplasm
Q608T0	PSD_METCA		cytoplasm
Q608V9	MODC_METCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q609A9	RECR_METCA		cytoplasm
Q609B8	Y1317_METCA		inner membrane
Q609C1	RBFA_METCA	cytoplasm (by similarity).	cytoplasm
Q609C2	TRUB_METCA		cytoplasm
Q609C5	RS15_METCA		cytoplasm
Q609G2	UBIG_METCA		cytoplasm
Q609I6	ARLY_METCA	cytoplasm (probable).	cytoplasm
Q609I7	G6PI_METCA	cytoplasm (by similarity).	cytoplasm
Q609J2	PYRD_METCA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q609K4	TOLB_METCA	periplasm (potential).	periplasm
Q609L0	RUVB_METCA		cytoplasm
Q609L1	RUVA_METCA		cytoplasm

Q609L2	RUVC_METCA		cytoplasm
Q609L3	Y1220_METCA		inner membrane
Q609M6	Y1207_METCA		inner membrane
Q609Q1	CYSA_METCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q609V0	BIOH_METCA	cytoplasm (by similarity).	cytoplasm
Q609W4	HIS81_METCA		cytoplasm
Q609X7	ASSY_METCA	cytoplasm (probable).	cytoplasm
Q609Z5	THIG_METCA	cytoplasm (by similarity).	cytoplasm
Q609Z8	PSTB_METCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q60A05	RPOC_METCA		cytoplasm
Q60A06	RPOB_METCA		cytoplasm
Q60A08	RL10_METCA		cytoplasm
Q60A09	RL1_METCA		cytoplasm
Q60A10	RL11_METCA		cytoplasm
Q60A14	PTH_METCA	cytoplasm (by similarity).	cytoplasm
Q60A15	RL25_METCA		cytoplasm
Q60A17	ISPE_METCA		cytoplasm
Q60A20	HEM1_METCA		cytoplasm
Q60A92	KUP_METCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q60AA3	MSBA_METCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q60AB3	CCMA_METCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q60AJ7	FETP_METCA		cytoplasm
Q60AK3	HTPG_METCA	cytoplasm (by similarity).	cytoplasm
Q60AK7	Y844_METCA		cytoplasm
Q60AM9	EX7S_METCA	cytoplasm (by similarity).	cytoplasm
Q60AQ2	TPMT_METCA	cytoplasm (by similarity).	cytoplasm
Q60AU5	SYT_METCA	cytoplasm (by similarity).	cytoplasm
Q60AV5	Y739_METCA		cytoplasm
Q60AY0	CH601_METCA	cytoplasm (by similarity).	cytoplasm
Q60AY5	AROC_METCA		cytoplasm
Q60AZ8	QUEA_METCA	cytoplasm (potential).	cytoplasm
Q60B18	UPPP_METCA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q60B49	LPXK_METCA		cytoplasm
Q60B71	MDH_METCA		cytoplasm
Q60B77	KSGA_METCA		cytoplasm
Q60BA1	MUTS_METCA		cytoplasm
Q60BA4	DXR_METCA		cytoplasm
Q60BA7	RRF_METCA	cytoplasm (by similarity).	cytoplasm
Q60BB0	RS2_METCA		cytoplasm
Q60BB2	GLND_METCA		cytoplasm
Q60BD8	ORN_METCA	cytoplasm (potential).	cytoplasm
Q60BE7	CLPX2_METCA		cytoplasm
Q60BE8	CLPP2_METCA	cytoplasm (by similarity).	cytoplasm
Q60BF9	GLUQ_METCA		cytoplasm

Q60BH8	PHS_METCA		cytoplasm
Q60BJ2	Y476_METCA		inner membrane
Q60BP6	CCME_METCA	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q60BS0	RL19_METCA		cytoplasm
Q60BS1	TRMD_METCA	cytoplasm (potential).	cytoplasm
Q60BS2	RIMM_METCA	cytoplasm (potential).	cytoplasm
Q60BS8	RECA_METCA	cytoplasm (by similarity).	cytoplasm
Q60BT8	Y377_METCA		inner membrane
Q60BV9	RPIA_METCA		cytoplasm
Q60BW3	GCST_METCA		cytoplasm
Q60BY2	AROB_METCA	cytoplasm (probable).	cytoplasm
Q60BY3	AROK_METCA	cytoplasm (probable).	cytoplasm
Q60BZ1	RL31_METCA		cytoplasm
Q60C21	GUAA_METCA		cytoplasm
Q60C56	NIFW_METCA		cytoplasm
Q60C59	HSCA_METCA		cytoplasm
Q60C60	HSCB_METCA		inner membrane
Q60C67	CLPX1_METCA		cytoplasm
Q60C68	CLPP1_METCA	cytoplasm (by similarity).	cytoplasm
Q60CG8	SAHH_METCA	cytoplasm (by similarity).	cytoplasm
Q60CJ7	LIPB_METCA	cytoplasm (potential).	cytoplasm
Q60CL4	MTFA_METCA		cytoplasm
Q60CN9	COAD_METCA	cytoplasm (by similarity).	cytoplasm
Q60CQ4	CCA_METCA		cytoplasm
Q60CR2	GLMU_METCA	cytoplasm (by similarity).	cytoplasm
Q60CR5	ATPG_METCA		inner membrane
Q60CR6	ATPA1_METCA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q60CS4	GIDB_METCA		cytoplasm
Q60CS5	GIDA_METCA	cytoplasm (by similarity).	cytoplasm
Q60CT7	NFI_METCA	cytoplasm (by similarity).	cytoplasm
Q60CV4	TRUA_BRUSU		cytoplasm
Q629Z7	ISPH1_BURMA		cytoplasm
Q62AG8	MDH_BURMA		cytoplasm
Q62AI6	LEU2_BURMA		cytoplasm
Q62AI8	LEUD_BURMA		cytoplasm
Q62AI9	LEU3_BURMA	cytoplasm (by similarity).	cytoplasm
Q62AJ2	TRUA_BURMA		cytoplasm
Q62AJ6	TRPA_BURMA		cytoplasm
Q62AW4	TAUB_BURMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q62BI3	ASPD_BURMA		cytoplasm
Q62CD5	SYI2_BURMA	cytoplasm (by similarity).	cytoplasm
Q62CE3	1A1D_BURMA		cytoplasm
Q62CH7	BETB_BURMA		cytoplasm
Q62CJ3	RS21C_BURMA		cytoplasm
Q62CK6	UVRB_BURMA	cytoplasm (by similarity).	cytoplasm
Q62DC9	TRPD_BURMA		cytoplasm
Q62DF2	HEM1_BURMA		cytoplasm
Q62DI5	GLYA2_BURMA	cytoplasm (by similarity).	cytoplasm
Q62DT5	RS21B_BURMA		cytoplasm

Q62DU0	Y4331_BURMA		inner membrane
Q62DU3	EX7S_BURMA	cytoplasm (by similarity).	cytoplasm
Q62EB0	ATPA2_BURMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q62EC5	TYPH_BURMA		cytoplasm
Q62EL5	TDH_BURMA	cytoplasm (by similarity).	cytoplasm
Q62EM2	RNPA_BURMA		extracellular
Q62EP0	GLMU_BURMA	cytoplasm (by similarity).	cytoplasm
Q62ES6	FLGI_BURMA	periplasm (by similarity).	periplasm
Q62ES7	FLGH_BURMA	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q62EU1	CCA_BURMA		cytoplasm
Q62EU9	Y3309_BURMA		inner membrane
Q62EZ9	HSLV_BURMA	cytoplasm (by similarity).	cytoplasm
Q62F00	HSLU_BURMA	cytoplasm (by similarity).	cytoplasm
Q62F04	ARGB_BURMA	cytoplasm (probable).	cytoplasm
Q62F07	METX_BURMA	cytoplasm (by similarity).	cytoplasm
Q62F43	GPMA_BURMA		cytoplasm
Q62F46	SECB_BURMA	cytoplasm (by similarity).	cytoplasm
Q62F47	GPDA_BURMA	cytoplasm (probable).	cytoplasm
Q62F99	DNAE2_BURMA	cytoplasm (probable).	cytoplasm
Q62FB8	COAD_BURMA	cytoplasm (by similarity).	cytoplasm
Q62FC0	HIS82_BURMA		cytoplasm
Q62FC1	PTH_BURMA	cytoplasm (by similarity).	cytoplasm
Q62FC2	RL25_BURMA		cytoplasm
Q62FC4	ISPE_BURMA		cytoplasm
Q62FC7	FPG_BURMA		cytoplasm
Q62FD0	Y3112_BURMA		cytoplasm
Q62FI7	PPCK_BURMA	cytoplasm (by similarity).	cytoplasm
Q62FM9	GCST_BURMA		cytoplasm
Q62FN1	GCSP_BURMA		cytoplasm
Q62FR4	ATPE_BURMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q62FR6	ATPG_BURMA		inner membrane
Q62FR7	ATPA1_BURMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q62FS8	GIDA_BURMA	cytoplasm (by similarity).	cytoplasm
Q62FU4	PHS_BURMA		cytoplasm
Q62FV3	CAAL_BURMA		cytoplasm
Q62G22	SAHH_BURMA	cytoplasm (by similarity).	cytoplasm
Q62G98	Y2758_BURMA		cytoplasm
Q62GA2	CYAY_BURMA		inner membrane
Q62GA9	AROK_BURMA	cytoplasm (probable).	cytoplasm
Q62GB0	AROB_BURMA	cytoplasm (probable).	cytoplasm
Q62GC6	THIG_BURMA	cytoplasm (by similarity).	cytoplasm
Q62GD7	MURA_BURMA	cytoplasm (probable).	inner membrane
Q62GD8	HIS1_BURMA	cytoplasm (by similarity).	cytoplasm
Q62GD9	HISX_BURMA		cytoplasm
Q62GE0	HIS81_BURMA		cytoplasm
Q62GE1	HIS7_BURMA	cytoplasm (by similarity).	cytoplasm
Q62GE3	HIS5_BURMA	cytoplasm (by similarity).	cytoplasm
Q62GE4	HIS4_BURMA	cytoplasm (by similarity).	cytoplasm
Q62GE5	HIS6_BURMA	cytoplasm (by similarity).	cytoplasm

Q62GE6	HIS3_BURMA	cytoplasm (by similarity).	cytoplasm
Q62GE7	HIS2_BURMA	cytoplasm (by similarity).	cytoplasm
Q62GJ3	RL11_BURMA		cytoplasm
Q62GJ4	RL1_BURMA		cytoplasm
Q62GJ5	RL10_BURMA		cytoplasm
Q62GJ7	RPOC_BURMA		cytoplasm
Q62GK0	RS12_BURMA		cytoplasm
Q62GK1	RS7_BURMA		cytoplasm
Q62GK4	RS10_BURMA		cytoplasm
Q62GK8	RL2_BURMA		cytoplasm
Q62GK9	RS19_BURMA		cytoplasm
Q62GL1	RS3_BURMA		cytoplasm
Q62GL2	RL16_BURMA		cytoplasm
Q62GL4	RS17_BURMA		cytoplasm
Q62GL9	RS8_BURMA		cytoplasm
Q62GM1	RL18_BURMA		cytoplasm
Q62GM2	RS5_BURMA		cytoplasm
Q62GM4	RL15_BURMA		cytoplasm
Q62GM8	RS13_BURMA		cytoplasm
Q62GM9	RS11_BURMA		cytoplasm
Q62GN0	RS4_BURMA		cytoplasm
Q62GN1	RPOA_BURMA		cytoplasm
Q62GR8	MRAZ_BURMA		inner membrane
Q62GR9	MRAW_BURMA		cytoplasm
Q62GS4	MRAY_BURMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q62GS5	MURD_BURMA	cytoplasm (by similarity).	cytoplasm
Q62GS7	MURG_BURMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q62GS8	MURC_BURMA	cytoplasm (probable).	cytoplasm
Q62GT9	ARGJ_BURMA	cytoplasm (probable).	cytoplasm
Q62GU2	Y2535_BURMA		inner membrane
Q62GV3	RL27_BURMA		cytoplasm
Q62GV5	PROB_BURMA	cytoplasm (by similarity).	cytoplasm
Q62GV7	NUDH_BURMA		cytoplasm
Q62GX2	PRMA_BURMA	cytoplasm (by similarity).	cytoplasm
Q62H14	ALLC2_BURMA		extracellular
Q62H17	DAPB_BURMA	cytoplasm (by similarity).	cytoplasm
Q62H23	PROA_BURMA	cytoplasm (by similarity).	cytoplasm
Q62H59	PHNC_BURMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q62H77	QUEA_BURMA	cytoplasm (potential).	cytoplasm
Q62HA7	RUVC_BURMA		cytoplasm
Q62HA8	RUVA_BURMA		cytoplasm
Q62HA9	RUVB_BURMA		cytoplasm
Q62HB3	DTD_BURMA	cytoplasm (probable).	cytoplasm
Q62HB4	SYI_BURMA	cytoplasm (by similarity).	cytoplasm
Q62HC7	GLNE_BURMA		cytoplasm
Q62HC9	PPNK_BURMA	cytoplasm (by similarity).	cytoplasm
Q62HD0	HRCB_BURMA		cytoplasm
Q62HD1	HEMH_BURMA	cytoplasm (by similarity).	extracellular
Q62HD5	DNAK_BURMA		cytoplasm
Q62HD6	DNAJ_BURMA	cytoplasm (probable).	cytoplasm

Q62HD8	PANB_BURMA		cytoplasm
Q62HE6	MIAA_BURMA		cytoplasm
Q62HH7	CLPS_BURMA		cytoplasm
Q62HI0	KAD_BURMA	cytoplasm (by similarity).	cytoplasm
Q62HL1	DUT_BURMA		extracellular
Q62HL4	SY11_BURMA	cytoplasm (by similarity).	cytoplasm
Q62HM5	RL28_BURMA		cytoplasm
Q62HM8	ISPH2_BURMA		cytoplasm
Q62HR9	UREE_BURMA	cytoplasm (by similarity).	cytoplasm
Q62HS0	URE1_BURMA	cytoplasm (by similarity).	cytoplasm
Q62HS1	URE2_BURMA	cytoplasm (by similarity).	cytoplasm
Q62HS2	URE3_BURMA	cytoplasm (by similarity).	cytoplasm
Q62HT6	UPPP_BURMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q62HT7	TRMB_BURMA		cytoplasm
Q62HU2	CRCB_BURMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q62HZ7	HAM1_BURMA		extracellular
Q62HZ8	RNPH_BURMA		cytoplasm
Q62I00	KGUA_BURMA	cytoplasm (by similarity).	cytoplasm
Q62I01	RPOZ_BURMA		cytoplasm
Q62I10	TOLB_BURMA	periplasm (potential).	periplasm
Q62I16	GLYA1_BURMA	cytoplasm (by similarity).	cytoplasm
Q62I17	NRDR_BURMA		cytoplasm
Q62I34	HGD_BURMA		cytoplasm
Q62I82	CH60_BURMA	cytoplasm (by similarity).	cytoplasm
Q62I87	RUVX_BURMA	cytoplasm (potential).	cytoplasm
Q62I89	PYRB_BURMA		cytoplasm
Q62ID1	HTPG_BURMA	cytoplasm (by similarity).	cytoplasm
Q62ID7	TAL_BURMA	cytoplasm (probable).	cytoplasm
Q62IG0	GLUQ_BURMA		cytoplasm
Q62IG1	Y1914_BURMA		inner membrane
Q62IG3	MSBA_BURMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q62II5	Y1889_BURMA		inner membrane
Q62II8	HEM6_BURMA	cytoplasm (by similarity).	cytoplasm
Q62IJ0	Y1884_BURMA		inner membrane
Q62IJ1	UPP_BURMA		cytoplasm
Q62IJ3	MGSA_BURMA		cytoplasm
Q62IJ6	ATKA_BURMA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q62IM0	ILVC_BURMA		cytoplasm
Q62IM1	PSD_BURMA		cytoplasm
Q62IN0	RS15_BURMA		cytoplasm
Q62IU9	FETP_BURMA		cytoplasm
Q62IZ4	HSCB_BURMA		inner membrane
Q62IZ5	HSCA_BURMA		cytoplasm
Q62J09	KDSA_BURMA	cytoplasm (by similarity).	cytoplasm
Q62J10	ENO_BURMA	cytoplasm (by similarity).	extracellular
Q62J26	MUTS_BURMA		cytoplasm
Q62J39	ACCA_BURMA	cytoplasm (by similarity).	cytoplasm
Q62J40	TILS_BURMA	cytoplasm (probable).	cytoplasm

Q62J86	RRAA_BURMA		cytoplasm
Q62JB3	DAPD_BURMA	cytoplasm (by similarity).	cytoplasm
Q62JC2	GLND_BURMA		cytoplasm
Q62JC4	RS2_BURMA		cytoplasm
Q62JC7	RRF_BURMA	cytoplasm (by similarity).	cytoplasm
Q62JD0	DXR_BURMA		cytoplasm
Q62JD4	LPXD_BURMA		cytoplasm
Q62JD5	FABZ_BURMA	cytoplasm (by similarity).	cytoplasm
Q62JD8	RNH2_BURMA	cytoplasm (potential).	extracellular
Q62JE0	Y1539_BURMA		cytoplasm
Q62JE7	SSRP_BURMA	cytoplasm (by similarity).	cytoplasm
Q62JF6	GUAA_BURMA		cytoplasm
Q62JH0	ALLC1_BURMA		extracellular
Q62JH4	MSCL_BURMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q62JI5	ISPD_BURMA		cytoplasm
Q62JI6	ISPF_BURMA		extracellular
Q62JK6	TIG_BURMA		cytoplasm
Q62JK7	CLPP_BURMA	cytoplasm (by similarity).	cytoplasm
Q62JK8	CLPX_BURMA		cytoplasm
Q62JL8	G6PI_BURMA	cytoplasm (by similarity).	cytoplasm
Q62JN9	KTHY_BURMA		cytoplasm
Q62JQ9	RS6_BURMA		cytoplasm
Q62JR1	RS18_BURMA		cytoplasm
Q62JR2	RL9_BURMA		cytoplasm
Q62JU1	RL31B_BURMA		cytoplasm
Q62JV0	RECR_BURMA		cytoplasm
Q62JV2	SURE_BURMA	cytoplasm (potential).	periplasm
Q62JV9	RUMA_BURMA		cytoplasm
Q62JW1	NDK_BURMA	cytoplasm (by similarity).	cytoplasm
Q62JW4	ISPG_BURMA		cytoplasm
Q62JW9	HFQ_BURMA		cytoplasm
Q62JX6	PURA_BURMA	cytoplasm (by similarity).	cytoplasm
Q62JX8	KUP_BURMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q62K38	RPIA_BURMA		cytoplasm
Q62K45	PYRD_BURMA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q62K79	LEXA_BURMA		cytoplasm
Q62K82	CYSA_BURMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q62KD8	OTCC_BURMA	cytoplasm (probable).	cytoplasm
Q62KD9	ARCA_BURMA	cytoplasm (by similarity).	extracellular
Q62KI5	RL20_BURMA		cytoplasm
Q62KK7	Y1063_BURMA		inner membrane
Q62KL0	RBFA_BURMA	cytoplasm (by similarity).	cytoplasm
Q62KL1	TRUB_BURMA		cytoplasm
Q62KU7	AROC_BURMA		cytoplasm
Q62KW5	SYV_BURMA	cytoplasm (by similarity).	cytoplasm
Q62KY7	Y904_BURMA		inner membrane
Q62L02	Y885_BURMA		cytoplasm

Q62L74	PSTB_BURMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q62L77	GLMM_BURMA		cytoplasm
Q62L80	RRMJ_BURMA	cytoplasm (by similarity).	cytoplasm
Q62LC0	HEM3_BURMA		cytoplasm
Q62LC1	CAPP_BURMA		cytoplasm
Q62LD1	ARLY_BURMA	cytoplasm (probable).	cytoplasm
Q62LE7	PANC_BURMA		cytoplasm
Q62LE8	PAND_BURMA		cytoplasm
Q62LF1	COBQ_BURMA		cytoplasm
Q62LG5	LGT_BURMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q62LG7	ILVD_BURMA		cytoplasm
Q62LJ6	HUTH_BURMA	cytoplasm (potential).	cytoplasm
Q62LS6	ACPS_BURMA	cytoplasm (by similarity).	cytoplasm
Q62LS7	RECO_BURMA		cytoplasm
Q62LT1	LEPA_BURMA		cytoplasm
Q62LT9	ACP_BURMA	cytoplasm (by similarity).	cytoplasm
Q62LU3	PLSX_BURMA		cytoplasm
Q62LU4	RL32_BURMA		cytoplasm
Q62LW1	PNCB_BURMA	cytoplasm (by similarity).	cytoplasm
Q62LW6	NORM_BURMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q62M18	UBIG_BURMA		cytoplasm
Q62M26	KCY_BURMA	cytoplasm (by similarity).	cytoplasm
Q62M51	RIMM_BURMA	cytoplasm (potential).	cytoplasm
Q62M52	TRMD_BURMA	cytoplasm (potential).	cytoplasm
Q62M53	RL19_BURMA		cytoplasm
Q62M61	ORN_BURMA	cytoplasm (potential).	cytoplasm
Q62M74	RS20_BURMA		cytoplasm
Q62M76	OTC_BURMA	cytoplasm (probable).	cytoplasm
Q62M77	MURB_BURMA	cytoplasm (probable).	inner membrane
Q62M79	Y372_BURMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q62M91	PDXH_BURMA		cytoplasm
Q62MH0	RECA_BURMA	cytoplasm (by similarity).	cytoplasm
Q62ML1	Y223_BURMA		inner membrane
Q62MM2	KSGA_BURMA		cytoplasm
Q62MN2	FDHD_BURMA	cytoplasm (potential).	cytoplasm
Q62MP2	UBIB_BURMA	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q62MP4	UBIE_BURMA		cytoplasm
Q62MP8	QUEF_BURMA	cytoplasm (probable).	cytoplasm
Q62MT4	FMT_BURMA		cytoplasm
Q62MX7	ACEK_BURMA	cytoplasm (by similarity).	cytoplasm
Q62N15	LIPB_BURMA	cytoplasm (potential).	cytoplasm
Q63IF8	SDHD_BURPS		cytoplasm
Q63IT3	DNS2_BURPS		extracellular
Q63IV6	TYPH_BURPS		cytoplasm

Q63IX0	ATPA2_BURPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q63JD9	KHSE_BURPS		cytoplasm
Q63JF2	EX7S_BURPS	cytoplasm (by similarity).	cytoplasm
Q63JF5	Y5761_BURPS		inner membrane
Q63JK9	LEU2_BURPS		cytoplasm
Q63JL1	LEUD_BURPS		cytoplasm
Q63JL2	LEU3_BURPS	cytoplasm (by similarity).	cytoplasm
Q63JL6	TRUA_BURPS		cytoplasm
Q63JM0	TRPA_BURPS		cytoplasm
Q63JZ3	TAUB_BURPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q63KK8	BETB_BURPS		cytoplasm
Q63KN7	1A1D_BURPS		cytoplasm
Q63KP6	SYI2_BURPS	cytoplasm (by similarity).	cytoplasm
Q63LV9	ASPD_BURPS		cytoplasm
Q63MV1	GLYA2_BURPS	cytoplasm (by similarity).	cytoplasm
Q63N20	CH602_BURPS	cytoplasm (by similarity).	cytoplasm
Q63NE3	UVRB_BURPS	cytoplasm (by similarity).	cytoplasm
Q63NL2	SELD_BURPS		cytoplasm
Q63NV1	Y4198_BURPS		inner membrane
Q63PD9	TDH_BURPS	cytoplasm (by similarity).	cytoplasm
Q63PF1	PHS_BURPS		cytoplasm
Q63PG8	GIDA_BURPS	cytoplasm (by similarity).	cytoplasm
Q63PH8	ATPA1_BURPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q63PH9	ATPG_BURPS		inner membrane
Q63PI1	ATPE_BURPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q63PL2	GCSP_BURPS		cytoplasm
Q63PL4	GCST_BURPS		cytoplasm
Q63PT2	SAHH_BURPS	cytoplasm (by similarity).	cytoplasm
Q63PZ9	RL11_BURPS		cytoplasm
Q63Q00	RL1_BURPS		cytoplasm
Q63Q01	RL10_BURPS		cytoplasm
Q63Q03	RPOB_BURPS		cytoplasm
Q63Q04	RPOC_BURPS		cytoplasm
Q63Q06	RS12_BURPS		cytoplasm
Q63Q07	RS7_BURPS		cytoplasm
Q63Q10	RS10_BURPS		cytoplasm
Q63Q14	RL2_BURPS		cytoplasm
Q63Q15	RS19_BURPS		cytoplasm
Q63Q17	RS3_BURPS		cytoplasm
Q63Q18	RL16_BURPS		cytoplasm
Q63Q20	RS17_BURPS		cytoplasm
Q63Q25	RS8_BURPS		cytoplasm
Q63Q27	RL18_BURPS		cytoplasm
Q63Q28	RS5_BURPS		cytoplasm
Q63Q30	RL15_BURPS		cytoplasm
Q63Q34	RS13_BURPS		cytoplasm
Q63Q35	RS11_BURPS		cytoplasm
Q63Q36	RS4_BURPS		cytoplasm

Q63Q37	RPOA_BURPS		cytoplasm
Q63Q46	Y3178_BURPS		cytoplasm
Q63Q50	CYAY_BURPS		inner membrane
Q63Q56	AROK_BURPS	cytoplasm (probable).	cytoplasm
Q63Q57	AROB_BURPS	cytoplasm (probable).	cytoplasm
Q63Q73	THIG_BURPS	cytoplasm (by similarity).	cytoplasm
Q63Q84	MURA_BURPS	cytoplasm (probable).	inner membrane
Q63Q85	HIS1_BURPS	cytoplasm (by similarity).	cytoplasm
Q63Q86	HISX_BURPS		cytoplasm
Q63Q87	HIS82_BURPS		cytoplasm
Q63Q88	HIS7_BURPS	cytoplasm (by similarity).	cytoplasm
Q63Q90	HIS5_BURPS	cytoplasm (by similarity).	cytoplasm
Q63Q91	HIS4_BURPS	cytoplasm (by similarity).	cytoplasm
Q63Q92	HIS6_BURPS	cytoplasm (by similarity).	cytoplasm
Q63Q93	HIS3_BURPS	cytoplasm (by similarity).	cytoplasm
Q63Q94	HIS2_BURPS	cytoplasm (by similarity).	cytoplasm
Q63QF1	HEM1_BURPS		cytoplasm
Q63QH1	TRPD_BURPS		cytoplasm
Q63QI8	MRAZ_BURPS		inner membrane
Q63QI9	MRAW_BURPS		cytoplasm
Q63QJ4	MRAY_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63QJ5	MURD_BURPS	cytoplasm (by similarity).	cytoplasm
Q63QJ7	MURG_BURPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q63QJ8	MURC_BURPS	cytoplasm (probable).	cytoplasm
Q63QK7	ARGJ_BURPS	cytoplasm (probable).	cytoplasm
Q63QL1	Y3012_BURPS		inner membrane
Q63QM1	PROB_BURPS	cytoplasm (by similarity).	cytoplasm
Q63QM4	NUDH_BURPS		cytoplasm
Q63QN9	PRMA_BURPS	cytoplasm (by similarity).	cytoplasm
Q63QS9	ALLC2_BURPS		extracellular
Q63QT3	DAPB_BURPS	cytoplasm (by similarity).	cytoplasm
Q63QT9	PROA_BURPS	cytoplasm (by similarity).	cytoplasm
Q63QX0	SYX_BURPS	cytoplasm (by similarity).	cytoplasm
Q63QX1	DTD_BURPS	cytoplasm (probable).	cytoplasm
Q63QX5	RUVB_BURPS		cytoplasm
Q63QX6	RUVA_BURPS		cytoplasm
Q63QX7	RUVC_BURPS		cytoplasm
Q63R06	QUEA_BURPS	cytoplasm (potential).	cytoplasm
Q63R24	PHNC_BURPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q63R39	GLNE_BURPS		cytoplasm
Q63R41	PPNK_BURPS	cytoplasm (by similarity).	cytoplasm
Q63R42	HRCA_BURPS		cytoplasm
Q63R43	HEMH_BURPS	cytoplasm (by similarity).	extracellular
Q63R47	DNAJ_BURPS	cytoplasm (probable).	cytoplasm
Q63R49	PANB_BURPS		cytoplasm
Q63R58	MIAA_BURPS		cytoplasm
Q63RA1	KDSA2_BURPS	cytoplasm (by similarity).	cytoplasm
Q63RA8	TOLB_BURPS	periplasm (potential).	periplasm
Q63RB4	GLYA1_BURPS	cytoplasm (by similarity).	cytoplasm
Q63RB5	NRDR_BURPS		cytoplasm

Q63RD3	HGD_BURPS		cytoplasm
Q63RI0	RUVX_BURPS	cytoplasm (potential).	cytoplasm
Q63RI2	PYRB_BURPS		cytoplasm
Q63RL2	UREE_BURPS	cytoplasm (by similarity).	cytoplasm
Q63RL3	URE1_BURPS	cytoplasm (by similarity).	cytoplasm
Q63RL4	URE2_BURPS	cytoplasm (by similarity).	cytoplasm
Q63RL5	URE3_BURPS	cytoplasm (by similarity).	cytoplasm
Q63RM9	UPPP2_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63RN0	TRMB_BURPS		cytoplasm
Q63RN4	CRCB_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63RV4	HAM1_BURPS		extracellular
Q63RV5	RNPH_BURPS		cytoplasm
Q63RV7	KGUA_BURPS	cytoplasm (by similarity).	cytoplasm
Q63RV8	RPOZ_BURPS		cytoplasm
Q63RZ8	UBIG_BURPS		cytoplasm
Q63S05	KCY_BURPS	cytoplasm (by similarity).	cytoplasm
Q63S31	RIMM_BURPS	cytoplasm (potential).	cytoplasm
Q63S32	TRMD_BURPS	cytoplasm (potential).	cytoplasm
Q63S33	RL19_BURPS		cytoplasm
Q63S41	ORN_BURPS	cytoplasm (potential).	cytoplasm
Q63S57	NORM_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63S63	PNCB_BURPS	cytoplasm (by similarity).	cytoplasm
Q63S81	RL32_BURPS		cytoplasm
Q63S82	PLSX_BURPS		cytoplasm
Q63S86	ACP_BURPS	cytoplasm (by similarity).	cytoplasm
Q63S94	LEPA_BURPS		cytoplasm
Q63S98	RECO_BURPS		cytoplasm
Q63S99	ACPS_BURPS	cytoplasm (by similarity).	cytoplasm
Q63SH6	HUTH_BURPS	cytoplasm (potential).	cytoplasm
Q63SJ4	FETP_BURPS		cytoplasm
Q63SN4	HSCB_BURPS		inner membrane
Q63SN5	HSCA_BURPS		cytoplasm
Q63SP9	KDSA1_BURPS	cytoplasm (by similarity).	cytoplasm
Q63SQ0	ENO_BURPS	cytoplasm (by similarity).	extracellular
Q63SR9	MUTS_BURPS		cytoplasm
Q63ST0	ACCA_BURPS	cytoplasm (by similarity).	cytoplasm
Q63ST1	TILS_BURPS	cytoplasm (probable).	cytoplasm
Q63SX7	RRAA_BURPS		cytoplasm
Q63T02	DAPD_BURPS	cytoplasm (by similarity).	cytoplasm
Q63T10	GLND_BURPS		cytoplasm
Q63T12	RS2_BURPS		cytoplasm
Q63T15	RRF_BURPS	cytoplasm (by similarity).	cytoplasm
Q63T18	DXR_BURPS		cytoplasm
Q63T22	LPXD_BURPS		cytoplasm
Q63T23	FABZ_BURPS	cytoplasm (by similarity).	cytoplasm
Q63T26	RNH2_BURPS	cytoplasm (potential).	extracellular
Q63T28	Y2143_BURPS		cytoplasm
Q63T34	SSRP_BURPS	cytoplasm (potential).	cytoplasm
Q63T42	GUAA_BURPS		cytoplasm
Q63T53	ALLC1_BURPS		extracellular

Q63T62	MSCL_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63T70	ISPD_BURPS		cytoplasm
Q63T71	ISPF_BURPS		extracellular
Q63T91	GLGB_BURPS		extracellular
Q63TE9	Y2019_BURPS		cytoplasm
Q63TG5	Y2003_BURPS		inner membrane
Q63TI8	SYV_BURPS	cytoplasm (by similarity).	cytoplasm
Q63TK6	AROC_BURPS		cytoplasm
Q63TM5	RL20_BURPS		cytoplasm
Q63TP6	Y1920_BURPS		inner membrane
Q63TP9	RBFA_BURPS	cytoplasm (by similarity).	cytoplasm
Q63TQ0	TRUB_BURPS		cytoplasm
Q63TU6	RPIA_BURPS		cytoplasm
Q63TV1	PYRD_BURPS	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q63TX7	LEXA_BURPS		cytoplasm
Q63TY1	CYSA_BURPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q63U72	OTCC_BURPS	cytoplasm (probable).	cytoplasm
Q63U73	ARCA_BURPS	cytoplasm (potential).	extracellular
Q63U95	ASSY_BURPS	cytoplasm (probable).	cytoplasm
Q63US0	KUP_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63US2	PURA_BURPS	cytoplasm (by similarity).	cytoplasm
Q63US8	HFQ_BURPS		cytoplasm
Q63UT3	ISPG_BURPS		cytoplasm
Q63UT6	NDK_BURPS	cytoplasm (by similarity).	cytoplasm
Q63UT8	RUMA_BURPS		cytoplasm
Q63UU4	SURE_BURPS	cytoplasm (potential).	periplasm
Q63UU6	RECR_BURPS		cytoplasm
Q63UV5	RL31B_BURPS		cytoplasm
Q63UY3	RL9_BURPS		cytoplasm
Q63UY4	RS18_BURPS		cytoplasm
Q63UY6	RS6_BURPS		cytoplasm
Q63V08	KTHY_BURPS		cytoplasm
Q63V31	G6PI_BURPS	cytoplasm (by similarity).	cytoplasm
Q63V40	CLPX_BURPS		cytoplasm
Q63V41	CLPP_BURPS	cytoplasm (by similarity).	cytoplasm
Q63V42	TIG_BURPS		cytoplasm
Q63V58	UPPP1_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63V79	PSTB_BURPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q63V83	GLMM_BURPS		cytoplasm
Q63V86	RRMJ_BURPS	cytoplasm (by similarity).	cytoplasm
Q63VB7	PPCK_BURPS	cytoplasm (by similarity).	cytoplasm
Q63VN8	RS15_BURPS		cytoplasm
Q63VP5	PSD_BURPS		cytoplasm
Q63VP6	ILVC_BURPS		cytoplasm

Q63VS3	ATKA_BURPS	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q63VS6	MGSA_BURPS		cytoplasm
Q63VS8	UPP_BURPS		cytoplasm
Q63VS9	Y1165_BURPS		inner membrane
Q63VT1	HEM6_BURPS	cytoplasm (by similarity).	cytoplasm
Q63VT4	Y1160_BURPS		inner membrane
Q63VX7	MSBA_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63VX9	Y1116_BURPS		inner membrane
Q63VY0	GLUQ_BURPS		cytoplasm
Q63W00	TAL_BURPS	cytoplasm (probable).	cytoplasm
Q63W07	HTPG_BURPS	cytoplasm (by similarity).	cytoplasm
Q63W73	HEM3_BURPS		cytoplasm
Q63W75	CAPP_BURPS		cytoplasm
Q63W97	PANC_BURPS		cytoplasm
Q63W98	PAND_BURPS		cytoplasm
Q63WA1	COBQ_BURPS		cytoplasm
Q63WB7	LGT_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63WB9	ILVD_BURPS		cytoplasm
Q63WC3	AMPA_BURPS	cytoplasm (by similarity).	cytoplasm
Q63WH0	ISPH1_BURPS		cytoplasm
Q63WH3	RL28_BURPS		cytoplasm
Q63WI3	SY11_BURPS	cytoplasm (by similarity).	cytoplasm
Q63WI6	DUT_BURPS		extracellular
Q63WJ1	CLPS_BURPS		cytoplasm
Q63WL3	LPXK_BURPS		cytoplasm
Q63WL6	KAD_BURPS	cytoplasm (by similarity).	cytoplasm
Q63WM0	RS20_BURPS		cytoplasm
Q63WM2	OTC_BURPS	cytoplasm (probable).	cytoplasm
Q63WM3	MURB_BURPS	cytoplasm (probable).	inner membrane
Q63WM5	Y866_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q63WN7	PDXH_BURPS		cytoplasm
Q63WX9	KPTA_BURPS		cytoplasm
Q63X54	ARO1_BURPS	cytoplasm (probable).	cytoplasm
Q63X65	Y672_BURPS		inner membrane
Q63X76	KSGA_BURPS		cytoplasm
Q63X85	FDHD_BURPS	cytoplasm (potential).	cytoplasm
Q63X97	UBIB_BURPS	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q63XA0	UBIE_BURPS		cytoplasm
Q63XA4	QUEF_BURPS	cytoplasm (probable).	cytoplasm
Q63XL1	Y529_BURPS		cytoplasm
Q63XL4	FPG_BURPS		cytoplasm
Q63XL7	ISPE_BURPS		cytoplasm
Q63XL9	RL25_BURPS		cytoplasm
Q63XM0	PTH_BURPS	cytoplasm (by similarity).	cytoplasm
Q63XM1	HIS81_BURPS		cytoplasm
Q63XM3	COAD_BURPS	cytoplasm (by similarity).	cytoplasm

Q63XR8	DNAE2_BURPS	cytoplasm (probable).	cytoplasm
Q63XU3	GPDA_BURPS	cytoplasm (probable).	cytoplasm
Q63XU4	SECB_BURPS	cytoplasm (by similarity).	cytoplasm
Q63XU7	GPMA_BURPS		cytoplasm
Q63XX6	LIPB_BURPS	cytoplasm (potential).	cytoplasm
Q63Y16	ACEK_BURPS	cytoplasm (by similarity).	cytoplasm
Q63Y75	GLMU_BURPS	cytoplasm (by similarity).	cytoplasm
Q63YB1	FLGI_BURPS	periplasm (by similarity).	periplasm
Q63YB2	FLGH_BURPS	bacterial cell outer membrane (probable).	inner membrane
Q63YC3	CCA_BURPS		cytoplasm
Q63YD1	Y257_BURPS		inner membrane
Q63YI3	HSLV_BURPS	cytoplasm (by similarity).	cytoplasm
Q63YI4	HSLU_BURPS	cytoplasm (by similarity).	cytoplasm
Q63YI7	ARGB_BURPS	cytoplasm (probable).	cytoplasm
Q63YJ0	METX_BURPS	cytoplasm (by similarity).	cytoplasm
Q63YR6	FMT_BURPS		cytoplasm
Q63YW3	RNPA_BURPS		extracellular
Q63YW5	DNAA_BURPS	cytoplasm (by similarity).	cytoplasm
Q63Z36	CAAL_BURPS		cytoplasm
Q659V3	RL25_PASPI		cytoplasm
Q65PY0	XYLA_MANSM	cytoplasm (by similarity).	cytoplasm
Q65PY5	Y2368_MANSM		inner membrane
Q65PY6	TRMA_MANSM		cytoplasm
Q65PZ6	OSTA_MANSM	bacterial cell outer membrane (by similarity).	outer membrane
Q65PZ8	TAL_MANSM	cytoplasm (probable).	cytoplasm
Q65PZ9	GIDA_MANSM	cytoplasm (by similarity).	cytoplasm
Q65Q00	GIDB_MANSM		cytoplasm
Q65Q05	ATPA_MANSM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q65Q06	ATPG_MANSM		inner membrane
Q65Q08	ATPE_MANSM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q65Q22	GPH_MANSM		cytoplasm
Q65Q24	RHAB_MANSM		cytoplasm
Q65Q25	RHAA_MANSM	cytoplasm (by similarity).	cytoplasm
Q65Q26	RHAD_MANSM	cytoplasm (by similarity).	extracellular
Q65Q27	RHAT_MANSM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65Q30	RHAR_MANSM	cytoplasm (by similarity).	cytoplasm
Q65Q31	RHAS_MANSM	cytoplasm (by similarity).	cytoplasm
Q65Q32	GPMA_MANSM		cytoplasm
Q65Q41	CCA_MANSM		cytoplasm
Q65Q60	PPCK_MANSM	cytoplasm (by similarity).	cytoplasm
Q65Q72	NAPA_MANSM	periplasm (by similarity).	periplasm
Q65Q77	FTSB_MANSM	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsI to the division site (by similarity).	inner membrane
Q65Q78	ISPD_MANSM		cytoplasm
Q65Q79	ISPF_MANSM		extracellular

Q65Q80	TRUD_MANS		cytoplasm
Q65Q81	SURE_MANS	cytoplasm (potential).	periplasm
Q65Q91	HSLO_MANS	cytoplasm (by similarity).	cytoplasm
Q65QA9	MUTS_MANS		cytoplasm
Q65QB0	RECA_MANS	cytoplasm (by similarity).	cytoplasm
Q65QC6	Y2227_MANS		cytoplasm
Q65QC7	CRCB_MANS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65QD4	ILVD_MANS		cytoplasm
Q65QD9	SECB_MANS	cytoplasm (by similarity).	cytoplasm
Q65QE0	GPDA_MANS	cytoplasm (probable).	cytoplasm
Q65QF1	FMT_MANS		cytoplasm
Q65QF7	MSCL_MANS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65QH0	GLPE_MANS	cytoplasm (by similarity).	cytoplasm
Q65QN6	XGPT_MANS	cytoplasm (by similarity).	cytoplasm
Q65QS1	CYAY_MANS		inner membrane
Q65QT9	PDXH_MANS		cytoplasm
Q65QV4	RS10_MANS		cytoplasm
Q65QV8	RL2_MANS		cytoplasm
Q65QV9	RS19_MANS		cytoplasm
Q65QW1	RS3_MANS		cytoplasm
Q65QW2	RL16_MANS		cytoplasm
Q65QW3	RL29_MANS		cytoplasm
Q65QW4	RS17_MANS		cytoplasm
Q65QW9	RS8_MANS		cytoplasm
Q65QX1	RL18_MANS		cytoplasm
Q65QX2	RS5_MANS		cytoplasm
Q65QX4	RL15_MANS		cytoplasm
Q65QX7	RS13_MANS		cytoplasm
Q65QX8	RS11_MANS		cytoplasm
Q65QX9	RS4_MANS		cytoplasm
Q65QY0	RPOA_MANS		cytoplasm
Q65R34	AROK_MANS	cytoplasm (probable).	cytoplasm
Q65R35	AROB_MANS	cytoplasm (probable).	cytoplasm
Q65R52	COAD_MANS	cytoplasm (by similarity).	cytoplasm
Q65R54	GLMU_MANS	cytoplasm (by similarity).	cytoplasm
Q65R59	FPG_MANS		cytoplasm
Q65R61	RL28_MANS		cytoplasm
Q65R66	DUT_MANS		extracellular
Q65R67	SLMA_MANS	nucleoid-associated (by similarity).	cytoplasm
Q65R70	RS2_MANS		cytoplasm
Q65R74	RRF_MANS	cytoplasm (by similarity).	cytoplasm
Q65R75	DXR_MANS		cytoplasm
Q65R81	LPXD_MANS		cytoplasm
Q65R84	ISPG_MANS		cytoplasm
Q65R92	NDPA_MANS	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q65R93	Y1910_MANS		inner membrane
Q65RA1	DJLA_MANS	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane

Q65RA4	DEOD_MANSM		cytoplasm
Q65RB1	HISX_MANSM		cytoplasm
Q65RB2	HIS82_MANSM		cytoplasm
Q65RB3	HIS7_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RB8	HIS5_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RC0	HIS4_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RC1	HIS6_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RC3	UPP_MANSM		cytoplasm
Q65RC8	ACP_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RD3	PLSX_MANSM		cytoplasm
Q65RD4	RL32_MANSM		cytoplasm
Q65RD9	PSD_MANSM		cytoplasm
Q65RE1	PROB_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RF4	TIG_MANSM		cytoplasm
Q65RF6	CLPP_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RF7	CLPX_MANSM		cytoplasm
Q65RG4	RRAA_MANSM		cytoplasm
Q65RG7	RHLB_MANSM		cytoplasm
Q65RG8	CDD_MANSM		cytoplasm
Q65RH0	Y1833_MANSM		inner membrane
Q65RH6	LIPB_MANSM	cytoplasm (potential).	cytoplasm
Q65RI0	Y1823_MANSM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65RN5	ZIPA_MANSM	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q65RN9	KITH_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RP7	RS20_MANSM		cytoplasm
Q65RQ2	SYI_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RQ4	ISPH_MANSM		cytoplasm
Q65RQ6	HAM1_MANSM		extracellular
Q65RQ9	RPIA_MANSM		cytoplasm
Q65RR5	KGUA_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RR6	RPOZ_MANSM		cytoplasm
Q65RT1	HSCB_MANSM		inner membrane
Q65RT2	HSCA_MANSM		cytoplasm
Q65RT5	Y1718_MANSM		cytoplasm
Q65RU6	MURA_MANSM	cytoplasm (probable).	inner membrane
Q65RV1	KHSE_MANSM	cytoplasm (potential).	cytoplasm
Q65RV7	HTPG_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RX0	GSHAB_MANSM		cytoplasm
Q65RX3	Y1680_MANSM		inner membrane
Q65RX7	MRAZ_MANSM		inner membrane
Q65RX8	MRAW_MANSM		cytoplasm
Q65RY3	MRAY_MANSM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65RY4	MURD_MANSM	cytoplasm (by similarity).	cytoplasm
Q65RY6	MURG_MANSM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q65RY7	MURC_MANSM	cytoplasm (probable).	cytoplasm
Q65RY9	DDL_MANSM	cytoplasm (by similarity).	cytoplasm
Q65S32	PURA_MANSM	cytoplasm (by similarity).	cytoplasm
Q65S49	PROA_MANSM	cytoplasm (by similarity).	cytoplasm

Q65S55	RL27_MANS		cytoplasm
Q65S61	Y1592_MANS		cytoplasm
Q65S66	FBPC_MANS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q65S79	HIS81_MANS		cytoplasm
Q65S80	SERC_MANS	cytoplasm (by similarity).	cytoplasm
Q65S87	TRPR_MANS	cytoplasm (by similarity).	cytoplasm
Q65S94	TGT_MANS		cytoplasm
Q65S95	QUEA_MANS	cytoplasm (potential).	cytoplasm
Q65S97	SYV_MANS	cytoplasm (by similarity).	cytoplasm
Q65SB8	ISPE_MANS		cytoplasm
Q65SD5	HFQ_MANS		cytoplasm
Q65SD6	MIAA_MANS		cytoplasm
Q65SE1	ORN_MANS	cytoplasm (potential).	cytoplasm
Q65SE7	RECR_MANS		cytoplasm
Q65SE8	SSRP_MANS	cytoplasm (by similarity).	cytoplasm
Q65SH6	KCY_MANS	cytoplasm (by similarity).	cytoplasm
Q65SK7	Y1446_MANS		inner membrane
Q65SL1	RBFA_MANS	cytoplasm (by similarity).	cytoplasm
Q65SL2	TRUB_MANS		cytoplasm
Q65SS8	NRDR_MANS		cytoplasm
Q65ST2	UVRB_MANS	cytoplasm (by similarity).	cytoplasm
Q65SV7	HEMH_MANS	cytoplasm (by similarity).	extracellular
Q65SW3	MODC_MANS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q65SX9	HEM1_MANS		cytoplasm
Q65SY9	NORM_MANS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65SZ8	GLND_MANS		cytoplasm
Q65T08	GLYA_MANS	cytoplasm (by similarity).	cytoplasm
Q65T14	Y1289_MANS		inner membrane
Q65T25	GLNE_MANS		cytoplasm
Q65T36	ARGR_MANS	cytoplasm (probable).	cytoplasm
Q65T37	MDH_MANS		cytoplasm
Q65T42	CYSA_MANS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q65T53	CYSJ_MANS		cytoplasm
Q65T54	CYSI_MANS		cytoplasm
Q65T71	SYI_MANS	cytoplasm (by similarity).	cytoplasm
Q65T73	SFSA_MANS		cytoplasm
Q65TA9	FABA_MANS	cytoplasm (by similarity).	cytoplasm
Q65TB4	KDSA_MANS	cytoplasm (by similarity).	cytoplasm
Q65TC2	G6PI_MANS	cytoplasm (by similarity).	cytoplasm
Q65TC6	DAPD_MANS	cytoplasm (by similarity).	cytoplasm
Q65TC8	TRUA_MANS		cytoplasm
Q65TE9	TRPA_MANS		cytoplasm
Q65TF2	TRPD_MANS		cytoplasm
Q65TI0	GLGB_MANS		extracellular
Q65TI2	GLGC_MANS		cytoplasm
Q65TI3	GLGA_MANS		cytoplasm
Q65TI7	RL25_MANS		cytoplasm
Q65TL1	Y1092_MANS		cytoplasm

Q65TM9	Y1074_MANS		inner membrane
Q65TN0	OTC_MANS	cytoplasm (probable).	cytoplasm
Q65TN4	QUEF_MANS	cytoplasm (probable).	cytoplasm
Q65TP1	THII_MANS	cytoplasm (potential).	cytoplasm
Q65TP2	EX7S_MANS	cytoplasm (by similarity).	cytoplasm
Q65TP7	RL20_MANS		cytoplasm
Q65TR8	PYRD_MANS	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q65TV4	ACKA_MANS	cytoplasm (by similarity).	cytoplasm
Q65TY2	DAPB_MANS	cytoplasm (by similarity).	cytoplasm
Q65TY6	GLMM_MANS		cytoplasm
Q65TZ0	RRMJ_MANS	cytoplasm (by similarity).	cytoplasm
Q65U20	LPXK_MANS		cytoplasm
Q65U21	MSBA_MANS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65U29	METX_MANS	cytoplasm (by similarity).	cytoplasm
Q65U54	DNAJ_MANS	cytoplasm (probable).	cytoplasm
Q65U55	DNAK_MANS		cytoplasm
Q65U59	MTFA_MANS		cytoplasm
Q65U60	FDHD_MANS	cytoplasm (potential).	cytoplasm
Q65U87	AROC_MANS		cytoplasm
Q65UA9	Y844_MANS		inner membrane
Q65UB4	MGSA_MANS		cytoplasm
Q65UE9	TILS_MANS	cytoplasm (probable).	cytoplasm
Q65UF7	KAD_MANS	cytoplasm (by similarity).	cytoplasm
Q65UI1	GUAA_MANS		cytoplasm
Q65UI5	ACCA_MANS	cytoplasm (by similarity).	cytoplasm
Q65UJ6	RIMK_MANS		cytoplasm
Q65UK9	LEXA_MANS		cytoplasm
Q65UL1	PPNK_MANS	cytoplasm (by similarity).	cytoplasm
Q65UN0	TOLB_MANS	periplasm (potential).	periplasm
Q65UP0	RUVB_MANS		cytoplasm
Q65UP1	RUVA_MANS		cytoplasm
Q65UP2	RUVC_MANS		cytoplasm
Q65UP3	Y710_MANS		inner membrane
Q65UQ4	RS15_MANS		cytoplasm
Q65UT5	NDK_MANS	cytoplasm (by similarity).	cytoplasm
Q65UX2	APAH_MANS		extracellular
Q65UX3	KSGA_MANS		cytoplasm
Q65UZ8	CCME_MANS	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q65V02	CCMA_MANS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q65V05	LEU3_MANS	cytoplasm (by similarity).	cytoplasm
Q65V07	LEU22_MANS		cytoplasm
Q65V08	LEUD2_MANS		cytoplasm
Q65V34	KTHY_MANS		cytoplasm
Q65V47	PTH_MANS	cytoplasm (by similarity).	cytoplasm
Q65V59	UXAC_MANS		cytoplasm
Q65V66	UXUA_MANS		cytoplasm
Q65V70	PRMA_MANS	cytoplasm (by similarity).	cytoplasm

Q65V72	FIS_MANSM		cytoplasm
Q65VB6	RECF_MANSM	cytoplasm (by similarity).	cytoplasm
Q65VB8	DNAA_MANSM	cytoplasm (by similarity).	cytoplasm
Q65VC0	RNPA_MANSM		extracellular
Q65VD1	RL9_MANSM		cytoplasm
Q65VD2	RS18_MANSM		cytoplasm
Q65VD4	RS6_MANSM		cytoplasm
Q65VE3	FABZ_MANSM	cytoplasm (by similarity).	cytoplasm
Q65VE4	CH60_MANSM	cytoplasm (by similarity).	cytoplasm
Q65VF5	RL31_MANSM		cytoplasm
Q65VG0	RL19_MANSM		cytoplasm
Q65VG1	TRMD_MANSM	cytoplasm (potential).	cytoplasm
Q65VG2	RIMM_MANSM	cytoplasm (potential).	cytoplasm
Q65VI0	RNH2_MANSM	cytoplasm (potential).	extracellular
Q65VI4	LUXS_MANSM		extracellular
Q65VJ5	NUDH_MANSM		cytoplasm
Q65VJ7	LGT_MANSM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65VJ8	TYSY_MANSM	cytoplasm (by similarity).	cytoplasm
Q65VK7	NAGZ_MANSM	cytoplasm (by similarity).	extracellular
Q65VK8	RUMB_MANSM		cytoplasm
Q65VN1	GRCA_MANSM		cytoplasm
Q65VN2	LEPA_MANSM		cytoplasm
Q65VN5	RNC_MANSM	cytoplasm (by similarity).	cytoplasm
Q65VP5	Y358_MANSM		cytoplasm
Q65VR9	LEUD1_MANSM		cytoplasm
Q65VS0	LEU21_MANSM		cytoplasm
Q65VT3	CORA_MANSM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65VT5	TRMB_MANSM		cytoplasm
Q65VT7	FETP_MANSM		cytoplasm
Q65VT8	MLTC_MANSM	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q65VX7	HEM3_MANSM		cytoplasm
Q65VY1	HSLU_MANSM	cytoplasm (by similarity).	cytoplasm
Q65VY2	HSLV_MANSM	cytoplasm (by similarity).	cytoplasm
Q65VZ2	RUVX_MANSM	cytoplasm (potential).	cytoplasm
Q65VZ7	ENO_MANSM	cytoplasm (by similarity).	extracellular
Q65W01	RNPH_MANSM		cytoplasm
Q65W13	RUMA_MANSM		cytoplasm
Q65W14	RECO_MANSM		cytoplasm
Q65W16	ARLY_MANSM	cytoplasm (probable).	cytoplasm
Q65W17	ARGB_MANSM	cytoplasm (probable).	cytoplasm
Q65W40	RPOC_MANSM		cytoplasm
Q65W41	RPOB_MANSM		cytoplasm
Q65W44	RL10_MANSM		cytoplasm
Q65W45	RL1_MANSM		cytoplasm
Q65W46	RL11_MANSM		cytoplasm
Q65W61	Y192_MANSM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65W71	NUDC_MANSM		cytoplasm
Q65W80	TUSA_MANSM	cytoplasm (by similarity).	cytoplasm
Q65W90	RS7_MANSM		cytoplasm

Q65W91	RS12_MANSMS		cytoplasm
Q65W97	SLYX_MANSMS		inner membrane
Q65WA1	DEOC_MANSMS	cytoplasm (by similarity).	cytoplasm
Q65WJ9	DLGD_MANSMS	cytoplasm (potential).	cytoplasm
Q65WK8	ILVC_MANSMS		cytoplasm
Q65WL3	OADG_MANSMS	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q65WM1	RBN_MANSMS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q65WM2	DTD_MANSMS	cytoplasm (probable).	cytoplasm
Q65WM5	MURB_MANSMS	cytoplasm (probable).	inner membrane
Q663I2	YSCH_YERPS		extracellular
Q663J7	YSCP_YERPS	cytoplasm (potential).	extracellular
Q663P9	GIDA_YERPS	cytoplasm (by similarity).	cytoplasm
Q663Q0	GIDB_YERPS		cytoplasm
Q663Q6	ATPA_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q663Q7	ATPG_YERPS		inner membrane
Q663Q9	ATPE_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q663R0	GLMU_YERPS	cytoplasm (by similarity).	cytoplasm
Q663R1	GLMS_YERPS	cytoplasm (by similarity).	cytoplasm
Q663R5	PSTB2_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q663S5	Y3950_YERPS		cytoplasm
Q663S9	RNPA_YERPS		extracellular
Q663T2	DNAA_YERPS	cytoplasm (by similarity).	cytoplasm
Q663T4	RECF_YERPS	cytoplasm (by similarity).	cytoplasm
Q663U8	FDHD_YERPS	cytoplasm (potential).	cytoplasm
Q663W8	Y3906_YERPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q663W9	IBPB_YERPS	cytoplasm (by similarity).	cytoplasm
Q663X0	IBPA_YERPS	cytoplasm (by similarity).	cytoplasm
Q663Y3	XYLA_YERPS	cytoplasm (by similarity).	cytoplasm
Q664B8	HUTH_YERPS	cytoplasm (potential).	cytoplasm
Q664F4	Y3815_YERPS		inner membrane
Q664F8	USPB_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q664I2	GLGB_YERPS		extracellular
Q664I4	GLGC_YERPS		cytoplasm
Q664I5	GLGA_YERPS		cytoplasm
Q664J2	GLPE_YERPS	cytoplasm (by similarity).	cytoplasm
Q664J8	BIOH_YERPS	cytoplasm (by similarity).	cytoplasm
Q664K7	PPCK_YERPS	cytoplasm (by similarity).	cytoplasm
Q664K8	HSLO_YERPS	cytoplasm (by similarity).	cytoplasm
Q664L9	AROK_YERPS	cytoplasm (probable).	cytoplasm
Q664M0	AROB_YERPS	cytoplasm (probable).	cytoplasm
Q664M4	GPH_YERPS		cytoplasm

Q664N1	TSGA_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q664P8	TAUB_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q664Q4	KEFG_YERPS		cytoplasm
Q664Q5	KEFB_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q664Q8	SLYX_YERPS		inner membrane
Q664R1	TUSD_YERPS	cytoplasm (by similarity).	cytoplasm
Q664R2	TUSC_YERPS	cytoplasm (by similarity).	cytoplasm
Q664R3	TUSB_YERPS	cytoplasm (by similarity).	cytoplasm
Q664R4	RS12_YERPS		cytoplasm
Q664R5	RS7_YERPS		cytoplasm
Q664S0	RS10_YERPS		cytoplasm
Q664S7	RS3_YERPS		cytoplasm
Q664S8	RL16_YERPS		cytoplasm
Q664S9	RL29_YERPS		cytoplasm
Q664T0	RS17_YERPS		cytoplasm
Q664T5	RS8_YERPS		cytoplasm
Q664T7	RL18_YERPS		cytoplasm
Q664T8	RS5_YERPS		cytoplasm
Q664U0	RL15_YERPS		cytoplasm
Q664U3	RS13_YERPS		cytoplasm
Q664U4	RS11_YERPS		cytoplasm
Q664U5	RS4_YERPS		cytoplasm
Q664U6	RPOA_YERPS		cytoplasm
Q664V0	MSCL_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q664V3	FMT_YERPS		cytoplasm
Q664V6	SMG_YERPS		cytoplasm
Q664W1	META_YERPS	cytoplasm (by similarity).	cytoplasm
Q664W4	ACEK_YERPS	cytoplasm (by similarity).	cytoplasm
Q664W9	G6PI_YERPS	cytoplasm (by similarity).	cytoplasm
Q664X0	PSIE_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q664X6	LAMB1_YERPS	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q665B4	SSUD_YERPS		cytoplasm
Q665E1	FIS_YERPS		cytoplasm
Q665E3	PRMA_YERPS	cytoplasm (by similarity).	cytoplasm
Q665F0	Y3568_YERPS		cytoplasm
Q665H0	AAEX_YERPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q665H1	AAEA_YERPS	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q665H2	AAEB_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q665I5	PYRI_YERPS		cytoplasm
Q665I6	PYRB_YERPS		cytoplasm
Q665I8	Y3529_YERPS		cytoplasm
Q665K4	MURA_YERPS	cytoplasm (probable).	inner membrane
Q665M1	DIAA_YERPS		cytoplasm
Q665N8	UXAC_YERPS		cytoplasm
Q665U4	RS21_YERPS		cytoplasm
Q665U6	Y3414_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q665U8	UPPP_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q665U9	CCA_YERPS		cytoplasm
Q665V2	GLNE_YERPS		cytoplasm
Q666C4	FLGH2_YERPS	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q666C5	FLGI2_YERPS	periplasm (by similarity).	periplasm
Q666M2	MLTC_YERPS	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q666M3	FETP_YERPS		cytoplasm
Q666M5	TRMB_YERPS		cytoplasm
Q666N1	HAM1_YERPS		extracellular
Q666N7	AGUA_YERPS		extracellular
Q666N9	RUVX_YERPS	cytoplasm (potential).	cytoplasm
Q666Q4	ARGO_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q666Q7	RPIA1_YERPS		cytoplasm
Q666R5	GCST_YERPS		cytoplasm
Q666R7	GCSP_YERPS		cytoplasm
Q666S3	Y3173_YERPS		cytoplasm
Q666Z8	LAMB2_YERPS	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q667F1	AAS_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q667F6	NUDH_YERPS		cytoplasm
Q667F8	LGT_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q667F9	TYSY_YERPS	cytoplasm (by similarity).	cytoplasm
Q667H7	MTFA_YERPS		cytoplasm
Q667I0	QUEF_YERPS	cytoplasm (probable).	cytoplasm
Q667I5	Y3007_YERPS		cytoplasm
Q667I6	DAPD_YERPS	cytoplasm (by similarity).	cytoplasm
Q667I7	GLND_YERPS		cytoplasm
Q667I9	RS2_YERPS		cytoplasm
Q667J2	RRF_YERPS	cytoplasm (by similarity).	cytoplasm
Q667J3	DXR_YERPS		cytoplasm
Q667J4	UPPS_YERPS		cytoplasm
Q667J7	YAET_YERPS	bacterial cell outer membrane (by similarity).	outer membrane
Q667J9	LPXD_YERPS		cytoplasm
Q667K0	FABZ_YERPS	cytoplasm (by similarity).	cytoplasm

Q667K3	RNH2_YERPS	cytoplasm (potential).	extracellular
Q667K5	ACCA_YERPS	cytoplasm (by similarity).	cytoplasm
Q667K7	TILS_YERPS	cytoplasm (probable).	cytoplasm
Q667M7	RNH_YERPS	cytoplasm (potential).	extracellular
Q667N6	NRDI_YERPS		inner membrane
Q667Q9	CRCB3_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q667R0	CRCB2_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q667S5	ATKA_YERPS	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q667T3	Y2908_YERPS		inner membrane
Q667T6	PCP_YERPS	cytoplasm (by similarity).	cytoplasm
Q667T8	GRCA_YERPS		cytoplasm
Q667U9	LEPA_YERPS		cytoplasm
Q667V1	RNC_YERPS	cytoplasm (by similarity).	cytoplasm
Q667V3	RECO_YERPS		cytoplasm
Q667V4	ACPS_YERPS	cytoplasm (by similarity).	cytoplasm
Q667X1	GLYA_YERPS	cytoplasm (by similarity).	cytoplasm
Q667Y3	ISCA_YERPS		cytoplasm
Q667Y4	HSCB_YERPS		inner membrane
Q667Y5	HSCA_YERPS		cytoplasm
Q667Y8	PEPB_YERPS	cytoplasm (by similarity).	cytoplasm
Q667Z5	NDK_YERPS	cytoplasm (by similarity).	cytoplasm
Q667Z9	ISPG_YERPS		cytoplasm
Q668A8	GUAA_YERPS		cytoplasm
Q668C5	MDTC_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q668C6	MDTB_YERPS	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q668C7	MDTA_YERPS	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q668E1	PSTB1_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q668E6	UPP_YERPS		cytoplasm
Q668E8	HDA_YERPS		nucleoid
Q668F7	DAPA_YERPS	cytoplasm (by similarity).	cytoplasm
Q668I0	NAPA_YERPS	periplasm (by similarity).	periplasm
Q668I5	Y2755_YERPS		inner membrane
Q668I6	HEM6_YERPS	cytoplasm (by similarity).	cytoplasm
Q668J7	NANE_YERPS		cytoplasm
Q668J9	NANK_YERPS		cytoplasm
Q668K2	NANT_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q668K6	CYSA_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q668M5	ZIPA_YERPS	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q668N7	GLK_YERPS	cytoplasm (by similarity).	cytoplasm
Q668Q3	FBPC_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q668T8	CCMA_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q668U2	CCME_YERPS	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q668V0	FADI_YERPS	cytoplasm (by similarity).	cytoplasm
Q668V1	FADJ_YERPS	cytoplasm (by similarity).	cytoplasm
Q668V5	AROC_YERPS		cytoplasm
Q668V6	MEPA_YERPS	periplasm (by similarity).	extracellular
Q668W0	Y2627_YERPS		cytoplasm
Q668W7	PDXB_YERPS		cytoplasm
Q668W9	TRUA_YERPS		cytoplasm
Q668Z0	ACKA_YERPS	cytoplasm (by similarity).	cytoplasm
Q668Z2	Y2595_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q668Z3	Y2594_YERPS		inner membrane
Q668Z7	Y2590_YERPS		cytoplasm
Q669I8	CVRA_YERPS	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q669J6	Y2488_YERPS		inner membrane
Q669J9	MTOX_YERPS		cytoplasm
Q669K9	RL32_YERPS		cytoplasm
Q669L0	PLSX_YERPS		cytoplasm
Q669L4	ACP_YERPS	cytoplasm (by similarity).	cytoplasm
Q669L8	KTHY_YERPS		cytoplasm
Q669N5	NAGZ_YERPS	cytoplasm (by similarity).	extracellular
Q669P6	NPD_YERPS	cytoplasm (probable).	cytoplasm
Q669P7	PEPT_YERPS	cytoplasm (by similarity).	extracellular
Q669Q4	TRMU_YERPS	cytoplasm (by similarity).	cytoplasm
Q669S1	DSRB_YERPS		inner membrane
Q669S5	FLHD_YERPS	cytoplasm (by similarity).	cytoplasm
Q669Z3	RL20_YERPS		cytoplasm
Q669Z9	BTUC_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66A01	BTUD_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q66A06	ARNT_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66A13	Y2319_YERPS		cytoplasm
Q66A23	SUFE_YERPS	cytoplasm (by similarity).	cytoplasm

Q66A27	MDTK_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66A48	PDXH_YERPS		cytoplasm
Q66A49	SYU_YERPS	cytoplasm (by similarity).	cytoplasm
Q66A75	ZNTB_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66A90	ACPD_YERPS		extracellular
Q66A98	Y2234_YERPS	periplasm (potential).	periplasm
Q66AE3	TUS_YERPS	cytoplasm (by similarity).	cytoplasm
Q66AF7	ARAB_YERPS		cytoplasm
Q66AF8	ARAA_YERPS		cytoplasm
Q66AI6	GCH2_YERPS		extracellular
Q66AK2	TRPD_YERPS		cytoplasm
Q66AK5	TRPA_YERPS		cytoplasm
Q66AM8	KITH_YERPS	cytoplasm (by similarity).	cytoplasm
Q66AP0	SELD_YERPS		cytoplasm
Q66AP6	MSRB_YERPS		cytoplasm
Q66AT9	RUVB_YERPS		cytoplasm
Q66AU0	RUVA_YERPS		cytoplasm
Q66AU1	RUVC_YERPS		cytoplasm
Q66AU2	Y2038_YERPS		inner membrane
Q66AU9	CUTC_YERPS	cytoplasm (probable).	periplasm
Q66AV8	MDTH_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66AW1	Y2019_YERPS		cytoplasm
Q66AX1	KDSA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66AX6	HEM1_YERPS		cytoplasm
Q66AX8	ISPE_YERPS		cytoplasm
Q66AY1	PTH_YERPS	cytoplasm (by similarity).	cytoplasm
Q66BN4	PGSA_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66BN5	UVRC_YERPS	cytoplasm (by similarity).	cytoplasm
Q66BP3	WRBA_YERPS		cytoplasm
Q66BT8	FLG11_YERPS	periplasm (by similarity).	periplasm
Q66BT9	FLGH1_YERPS	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q66BW1	YEBF_YERPS	secreted protein (by similarity).	inner membrane
Q66BW6	Y1650_YERPS		inner membrane
Q66BY6	Y1630_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66C32	LLDD_YERPS		cytoplasm
Q66C49	HISX_YERPS		cytoplasm
Q66C50	HIS8_YERPS		cytoplasm
Q66C51	HIS7_YERPS	cytoplasm (by similarity).	cytoplasm
Q66C52	HIS5_YERPS	cytoplasm (by similarity).	cytoplasm
Q66C53	HIS4_YERPS	cytoplasm (by similarity).	cytoplasm
Q66C54	HIS6_YERPS	cytoplasm (by similarity).	cytoplasm
Q66C79	CDD_YERPS		cytoplasm
Q66C80	MAO1_YERPS		cytoplasm
Q66CD8	TUSE_YERPS	cytoplasm (by similarity).	cytoplasm

Q66CE3	Y1460_YERPS		inner membrane
Q66CE4	MGSA_YERPS		cytoplasm
Q66CF3	FABA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66CG4	PYRD_YERPS	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q66CG6	PNCB_YERPS	cytoplasm (by similarity).	cytoplasm
Q66CH5	MUKF_YERPS	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q66CI2	LPXK_YERPS		cytoplasm
Q66CI3	MSBA_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66CI7	KCY_YERPS	cytoplasm (by similarity).	cytoplasm
Q66CI9	SERC_YERPS	cytoplasm (by similarity).	cytoplasm
Q66CK7	LFTR_YERPS	cytoplasm (by similarity).	cytoplasm
Q66CL0	CLPS_YERPS		cytoplasm
Q66CN7	RUMB_YERPS		cytoplasm
Q66CP5	Y1357_YERPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q66CP8	DEOC1_YERPS	cytoplasm (by similarity).	cytoplasm
Q66CR5	END4_YERPS		extracellular
Q66CS8	RPIA2_YERPS		cytoplasm
Q66CT6	EFPL_YERPS		cytoplasm
Q66CT7	UXUA_YERPS		cytoplasm
Q66CV3	RL25_YERPS		cytoplasm
Q66CV4	NDPA_YERPS	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q66CV5	Y1297_YERPS		inner membrane
Q66CZ4	UBIG_YERPS		cytoplasm
Q66CZ8	ECOT_YERPS	periplasm (by similarity).	periplasm
Q66D26	PHNC_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q66D46	ZRAP_YERPS	periplasm (by similarity).	periplasm
Q66D53	BETB_YERPS		cytoplasm
Q66D62	UVRB_YERPS	cytoplasm (by similarity).	cytoplasm
Q66D69	6PGL_YERPS		cytoplasm
Q66D71	MODC_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q66D83	GPMA_YERPS		cytoplasm
Q66D85	ZITB_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66D90	TOLB_YERPS	periplasm (potential).	periplasm
Q66DA0	SUCC_YERPS		cytoplasm
Q66DA9	PPNK_YERPS	cytoplasm (by similarity).	cytoplasm
Q66DB4	SSRP_YERPS	cytoplasm (by similarity).	cytoplasm
Q66DD5	Y1111_YERPS		inner membrane
Q66DE8	Y1098_YERPS		inner membrane
Q66DF4	LIPB_YERPS	cytoplasm (potential).	cytoplasm

Q66DF7	CRCB1_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66DP5	ASCD_YERPS		cytoplasm
Q66DP7	KAD_YERPS	cytoplasm (by similarity).	cytoplasm
Q66DP8	HTPG_YERPS	cytoplasm (by similarity).	cytoplasm
Q66DP9	RECR_YERPS		cytoplasm
Q66DR1	RL31B_YERPS		cytoplasm
Q66DT3	CLPX_YERPS		cytoplasm
Q66DT4	CLPP_YERPS	cytoplasm (by similarity).	cytoplasm
Q66DT5	TIG_YERPS		cytoplasm
Q66DV2	EX7S_YERPS	cytoplasm (by similarity).	cytoplasm
Q66DW0	NRDR_YERPS		cytoplasm
Q66DW5	TGT_YERPS		cytoplasm
Q66DW6	QUEA_YERPS	cytoplasm (potential).	cytoplasm
Q66DW7	ACPH_YERPS		cytoplasm
Q66DY1	Y912_YERPS		inner membrane
Q66DY2	AROL_YERPS	cytoplasm (probable).	cytoplasm
Q66DY8	PROA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66DY9	PROB_YERPS	cytoplasm (by similarity).	cytoplasm
Q66DZ2	XGPT_YERPS	cytoplasm (by similarity).	cytoplasm
Q66E56	RL19_YERPS		cytoplasm
Q66E57	TRMD_YERPS	cytoplasm (potential).	cytoplasm
Q66E58	RIMM_YERPS	cytoplasm (potential).	cytoplasm
Q66E63	LUXS_YERPS		extracellular
Q66E64	GSH1_YERPS		cytoplasm
Q66E67	CSRA_YERPS		cytoplasm
Q66E69	RECX_YERPS	cytoplasm (by similarity).	cytoplasm
Q66E70	RECA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66EB5	MUTS_YERPS		cytoplasm
Q66EC0	SURE_YERPS	cytoplasm (potential).	periplasm
Q66EC1	TRUD_YERPS		cytoplasm
Q66EC2	ISPF_YERPS		extracellular
Q66EC3	ISPD_YERPS		cytoplasm
Q66ED3	CYSI_YERPS		cytoplasm
Q66ED4	CYSJ_YERPS		cytoplasm
Q66ED8	ENO_YERPS	cytoplasm (by similarity).	extracellular
Q66EE5	DGTP_YERPS		cytoplasm
Q66EE7	BTUF_YERPS	periplasm (by similarity).	periplasm
Q66EF1	GSA_YERPS	cytoplasm (potential).	cytoplasm
Q66EF8	SFSA_YERPS		cytoplasm
Q66EG0	GLUQ_YERPS		cytoplasm
Q66EG3	PANB_YERPS		cytoplasm
Q66EG4	PANC_YERPS		cytoplasm
Q66EG5	PAND_YERPS		cytoplasm
Q66EH4	SPED_YERPS		cytoplasm
Q66EJ2	Y701_YERPS		inner membrane
Q66EJ3	Y700_YERPS		cytoplasm
Q66EJ9	LPXC_YERPS		cytoplasm
Q66EK4	MURC_YERPS	cytoplasm (probable).	cytoplasm
Q66EK5	MURG_YERPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q66EK7	MURD_YERPS	cytoplasm (by similarity).	cytoplasm

Q66EK8	MRAY_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66EL3	MRAW_YERPS	cytoplasm (by similarity).	cytoplasm
Q66EL4	MRAZ_YERPS		inner membrane
Q66EM1	LEU1_YERPS		cytoplasm
Q66EM2	LEU3_YERPS	cytoplasm (by similarity).	cytoplasm
Q66EM3	LEU2_YERPS		cytoplasm
Q66EM4	LEUD_YERPS		cytoplasm
Q66EN4	RAPA_YERPS		cytoplasm
Q66EQ5	DJLA_YERPS	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q66EQ6	OSTA_YERPS	bacterial cell outer membrane (by similarity).	outer membrane
Q66EQ8	KSGA_YERPS		cytoplasm
Q66ER0	APAH_YERPS		extracellular
Q66ER8	CARA_YERPS		cytoplasm
Q66ER9	DAPB_YERPS	cytoplasm (by similarity).	cytoplasm
Q66ES1	ISPH_YERPS		cytoplasm
Q66ES4	SYI_YERPS	cytoplasm (by similarity).	cytoplasm
Q66ES6	RS20_YERPS		cytoplasm
Q66ES9	DNAJ_YERPS	cytoplasm (probable).	cytoplasm
Q66ET0	DNAK_YERPS		cytoplasm
Q66ET5	TAL_YERPS	cytoplasm (probable).	cytoplasm
Q66ET8	KHSE_YERPS	cytoplasm (potential).	cytoplasm
Q66EU4	Y597_YERPS		inner membrane
Q66EU5	TRPR_YERPS	cytoplasm (by similarity).	cytoplasm
Q66EV7	DEOD_YERPS		cytoplasm
Q66EV9	TYPH_YERPS		cytoplasm
Q66EW0	DEOC2_YERPS	cytoplasm (by similarity).	cytoplasm
Q66F09	AMPA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66F11	SYV_YERPS	cytoplasm (by similarity).	cytoplasm
Q66F14	OTC_YERPS	cytoplasm (probable).	cytoplasm
Q66F57	RS15_YERPS		cytoplasm
Q66F58	TRUB_YERPS		cytoplasm
Q66F59	RBFA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66F62	Y478_YERPS		inner membrane
Q66F64	GLMM_YERPS		cytoplasm
Q66F67	RRMJ_YERPS	cytoplasm (by similarity).	cytoplasm
Q66F75	RL27_YERPS		cytoplasm
Q66F81	ARGR_YERPS	cytoplasm (probable).	cytoplasm
Q66F89	MSRA_YERPS		cytoplasm
Q66F95	RCMNS_YERPS	cytoplasm (by similarity).	cytoplasm
Q66F99	RL9_YERPS		cytoplasm
Q66FA0	RS18_YERPS		cytoplasm
Q66FA2	RS6_YERPS		cytoplasm
Q66FB0	PURA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66FB5	HFQ_YERPS		cytoplasm
Q66FB6	MIAA_YERPS		cytoplasm
Q66FC2	ORN_YERPS	cytoplasm (potential).	cytoplasm
Q66FC4	PSD_YERPS		cytoplasm
Q66FD1	SUGE_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66FD5	CH60_YERPS	cytoplasm (by similarity).	cytoplasm

Q66FD6	CH10_YERPS	cytoplasm (by similarity).	cytoplasm
Q66FE0	CUTA_YERPS	cytoplasm (by similarity).	periplasm
Q66FF0	RHAT_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66FF1	RHAR_YERPS	cytoplasm (by similarity).	cytoplasm
Q66FF2	RHAS_YERPS	cytoplasm (by similarity).	cytoplasm
Q66FF3	RHAB_YERPS		cytoplasm
Q66FF4	RHAA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66FF5	RHAD_YERPS	cytoplasm (by similarity).	extracellular
Q66FG7	LEXA_YERPS		cytoplasm
Q66FN0	ACTP_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66FP0	NFI_YERPS	cytoplasm (by similarity).	cytoplasm
Q66FP3	NUDC_YERPS		cytoplasm
Q66FP9	THIG_YERPS	cytoplasm (by similarity).	cytoplasm
Q66FQ1	RPOC_YERPS		cytoplasm
Q66FQ2	RPOB_YERPS		cytoplasm
Q66FQ4	RL10_YERPS		cytoplasm
Q66FQ5	RL1_YERPS		cytoplasm
Q66FQ6	RL11_YERPS		cytoplasm
Q66FR3	MURB_YERPS	cytoplasm (probable).	inner membrane
Q66FR8	FADB_YERPS		cytoplasm
Q66FR9	FADA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66FS8	UBIB_YERPS	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q66FT0	UBIE_YERPS		cytoplasm
Q66FX1	TUSA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66FY7	CORA_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66FZ7	CYAY_YERPS		inner membrane
Q66G00	HEM3_YERPS		cytoplasm
Q66G05	WECG_YERPS		cytoplasm
Q66G06	WZYE_YERPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q66G07	WECF_YERPS	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q66G19	RHLB_YERPS		cytoplasm
Q66G20	GPPA_YERPS		extracellular
Q66G37	ILVC_YERPS		cytoplasm
Q66G45	ILVD_YERPS		cytoplasm
Q66G57	BTUB_YERPS	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q66G58	TRMA_YERPS		cytoplasm
Q66G70	ARLY_YERPS	cytoplasm (probable).	cytoplasm
Q66G71	ARGB_YERPS	cytoplasm (probable).	cytoplasm
Q66G72	ARGC_YERPS	cytoplasm (probable).	cytoplasm
Q66G74	CAPP_YERPS		cytoplasm
Q66G80	RL31_YERPS		cytoplasm
Q66G84	HSLV_YERPS	cytoplasm (by similarity).	cytoplasm

Q66G85	HSLU_YERPS	cytoplasm (by similarity).	cytoplasm
Q66G87	RRAA_YERPS		cytoplasm
Q66GA9	FIEF_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66GB8	GPDA_YERPS	cytoplasm (probable).	cytoplasm
Q66GB9	SECB_YERPS	cytoplasm (by similarity).	cytoplasm
Q66GC2	GPMI_YERPS		cytoplasm
Q66GC5	TDH_YERPS	cytoplasm (by similarity).	cytoplasm
Q66GD2	COAD_YERPS	cytoplasm (by similarity).	cytoplasm
Q66GD3	FPG_YERPS		cytoplasm
Q66GD5	RL28_YERPS		cytoplasm
Q66GD6	RADC_YERPS		cytoplasm
Q66GD8	DUT_YERPS		extracellular
Q66GD9	SLMA_YERPS	nucleoid-associated (by similarity).	cytoplasm
Q66GE1	RNPH_YERPS		cytoplasm
Q66GE5	KGUA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66GE6	RPOZ_YERPS		cytoplasm
Q66GF4	DTD_YERPS	cytoplasm (probable).	cytoplasm
Q66GF5	RBN_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66GG2	Y020_YERPS		inner membrane
Q66GH5	KUP_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q66GH6	RAVA_YERPS	cytoplasm (by similarity).	cytoplasm
Q66GH7	VIAA_YERPS		inner membrane
Q66X92	RECA_BURPS	cytoplasm (by similarity).	cytoplasm
Q673V5	RBL_SOVGS		extracellular
Q684Q1	LUXS_SERMA		extracellular
Q68VM9	HEMH_RICTY	cytoplasm (by similarity).	extracellular
Q68VN7	LIPB_RICTY	cytoplasm (potential).	cytoplasm
Q68VQ6	TRUA_RICTY		cytoplasm
Q68VR6	HEMK_RICTY		cytoplasm
Q68VS2	TIG_RICTY		cytoplasm
Q68VU6	ATPA_RICTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q68VU7	ATPG_RICTY		inner membrane
Q68VU9	ATPE_RICTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q68VV5	CCMA_RICTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q68VX6	RL32_RICTY		cytoplasm
Q68VX7	FABH_RICTY	cytoplasm (probable).	cytoplasm
Q68VY3	KGUA_RICTY	cytoplasm (by similarity).	cytoplasm
Q68VY8	RECA_RICTY	cytoplasm (by similarity).	cytoplasm
Q68VZ8	RL27_RICTY		cytoplasm
Q68W04	END3_RICTY		cytoplasm
Q68W07	GLYA_RICTY	cytoplasm (by similarity).	cytoplasm
Q68W09	Y726_RICTY		inner membrane
Q68W20	RNH_RICTY	cytoplasm (potential).	extracellular

Q68W26	TGT_RICTY		cytoplasm
Q68W45	CLPX_RICTY		cytoplasm
Q68W50	SYV_RICTY	cytoplasm (by similarity).	cytoplasm
Q68W53	KTHY_RICTY		cytoplasm
Q68W57	UBIE_RICTY		cytoplasm
Q68W66	KSGA_RICTY		cytoplasm
Q68W77	RS10_RICTY		cytoplasm
Q68W81	RL2_RICTY		cytoplasm
Q68W82	RS19_RICTY		cytoplasm
Q68W84	RS3_RICTY		cytoplasm
Q68W85	RL16_RICTY		cytoplasm
Q68W86	RL29_RICTY		cytoplasm
Q68W87	RS17_RICTY		cytoplasm
Q68W92	RS8_RICTY		cytoplasm
Q68W94	RL18_RICTY		cytoplasm
Q68W95	RS5_RICTY		cytoplasm
Q68W97	RL15_RICTY		cytoplasm
Q68W99	KAD_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WA0	RS13_RICTY		cytoplasm
Q68WA2	RPOA_RICTY		cytoplasm
Q68WA4	RS20_RICTY		cytoplasm
Q68WB5	UBIG_RICTY		cytoplasm
Q68WC2	SYI_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WC4	RS21_RICTY		cytoplasm
Q68WD0	RL20_RICTY		cytoplasm
Q68WD3	RL25_RICTY		cytoplasm
Q68WD4	PTH_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WD8	DNAA_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WF0	CCME_RICTY	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q68WF7	MURA_RICTY	cytoplasm (probable).	inner membrane
Q68WF8	RPOZ_RICTY		cytoplasm
Q68WF9	ACPS_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WG4	UVRC_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WG6	MRAZ_RICTY		inner membrane
Q68WG7	MRAW_RICTY		cytoplasm
Q68WH9	SYI_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WI2	Y541_RICTY		inner membrane
Q68WJ4	PRIA_RICTY		cytoplasm
Q68WL3	KCY_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WL5	CLPP_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WM5	MIAA_RICTY		cytoplasm
Q68WN2	RS15_RICTY		cytoplasm
Q68WN3	TRUB_RICTY		cytoplasm
Q68WR3	HEM3_RICTY		cytoplasm
Q68WS4	Y442_RICTY		inner membrane
Q68WS6	IDI2_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WT8	PPNK_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WU0	RECR_RICTY		cytoplasm
Q68WU3	RBFA_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WU5	SUCC_RICTY		cytoplasm
Q68WV4	UPPS_RICTY		cytoplasm

Q68WW7	MURG_RICTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q68WW9	MURD_RICTY	cytoplasm (by similarity).	cytoplasm
Q68WX8	DUT_RICTY		extracellular
Q68WZ0	RUVB_RICTY		cytoplasm
Q68WZ1	RUVA_RICTY		cytoplasm
Q68WZ8	MDH_RICTY		cytoplasm
Q68X23	EX7S_RICTY	cytoplasm (by similarity).	cytoplasm
Q68X25	RIMM_RICTY	cytoplasm (potential).	cytoplasm
Q68X27	RS4_RICTY		cytoplasm
Q68X41	RUVX_RICTY	cytoplasm (potential).	cytoplasm
Q68X52	HSLU_RICTY	cytoplasm (by similarity).	cytoplasm
Q68X53	HSLV_RICTY	cytoplasm (by similarity).	cytoplasm
Q68X66	TRMU_RICTY	cytoplasm (by similarity).	cytoplasm
Q68X69	TOLB_RICTY	periplasm (potential).	periplasm
Q68X70	THYX_RICTY		cytoplasm
Q68X73	MUTS_RICTY		cytoplasm
Q68X95	LEPA_RICTY		cytoplasm
Q68XA5	IDH_RICTY		cytoplasm
Q68XB5	LPXC_RICTY		cytoplasm
Q68XC1	MURB_RICTY	cytoplasm (probable).	inner membrane
Q68XC2	MURC_RICTY	cytoplasm (probable).	cytoplasm
Q68XC8	PSD_RICTY		cytoplasm
Q68XD3	NUDH_RICTY		cytoplasm
Q68XF6	QUEA_RICTY	cytoplasm (potential).	cytoplasm
Q68XG5	UVRB_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XG6	RNH2_RICTY	cytoplasm (potential).	extracellular
Q68XG8	HSCA_RICTY		cytoplasm
Q68XH5	DAPD_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XI2	DNAK_RICTY		cytoplasm
Q68XI3	DNAJ_RICTY	cytoplasm (probable).	cytoplasm
Q68XK5	RRMJ_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XL4	RRF_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XM0	DAPB_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XM6	AMPA_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XM7	RPOC_RICTY		cytoplasm
Q68XN0	RL10_RICTY		cytoplasm
Q68XN1	RL1_RICTY		cytoplasm
Q68XN2	RL11_RICTY		cytoplasm
Q68XN6	RS7_RICTY		cytoplasm
Q68XN7	RS12_RICTY		cytoplasm
Q68XP2	Y114_RICTY		inner membrane
Q68XQ6	RECF_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XR5	RS6_RICTY		cytoplasm
Q68XR6	RS18_RICTY		cytoplasm
Q68XR7	RL9_RICTY		cytoplasm
Q68XR8	TILS_RICTY	cytoplasm (probable).	cytoplasm
Q68XS2	LGT_RICTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q68XS9	NDK_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XT0	GIDA_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XT1	GIDB_RICTY		cytoplasm
Q68XT6	KDSA_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XU4	SECB_RICTY	cytoplasm (by similarity).	cytoplasm

Q68XU6	QUEF_RICTY	cytoplasm (probable).	cytoplasm
Q68XV5	RS2_RICTY		cytoplasm
Q68XW7	RL28_RICTY		cytoplasm
Q68XX8	ACKA_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XX9	TRMD_RICTY	cytoplasm (potential).	cytoplasm
Q68XY0	RL19_RICTY		cytoplasm
Q68XY5	RNC_RICTY	cytoplasm (by similarity).	cytoplasm
Q68XY7	RUVC_RICTY		cytoplasm
Q68XZ4	LPXD_RICTY		cytoplasm
Q68XZ5	FABZ_RICTY	cytoplasm (by similarity).	cytoplasm
Q68Y01	Y001_RICTY		cytoplasm
Q68Y08	HTPG_RICTY	cytoplasm (by similarity).	cytoplasm
Q69AA9	CPS12_ACTPL		cytoplasm
Q69AY8	CCME_ACEDI	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q6AIM2	HEM1_DESPS		cytoplasm
Q6AIP3	AROC_DESPS		cytoplasm
Q6AIZ3	RECA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJ07	FABZ_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJ19	TILS_DESPS	cytoplasm (probable).	cytoplasm
Q6AJ39	Y2912_DESPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6AJ42	RRMJ_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJ43	Y2908_DESPS		inner membrane
Q6AJ44	PANB_DESPS		cytoplasm
Q6AJ46	MRAZ_DESPS		inner membrane
Q6AJ47	MRAW_DESPS		cytoplasm
Q6AJ51	MRAY_DESPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6AJ52	MURD_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJ53	MURG_DESPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6AJ54	MURC_DESPS	cytoplasm (probable).	cytoplasm
Q6AJ55	MURB_DESPS	cytoplasm (probable).	inner membrane
Q6AJ82	SYI_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJ90	Y2861_DESPS		inner membrane
Q6AJ91	KGUA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJA8	QUEA_DESPS	cytoplasm (potential).	cytoplasm
Q6AJE2	KTHY_DESPS		cytoplasm
Q6AJE5	RNH2_DESPS	cytoplasm (potential).	extracellular
Q6AJE6	RL19_DESPS		cytoplasm
Q6AJE8	TRMD_DESPS	cytoplasm (potential).	cytoplasm
Q6AJG5	NRDR_DESPS		cytoplasm
Q6AJH7	COAD_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJI0	PSD_DESPS		cytoplasm
Q6AJL0	ARGJ_DESPS	cytoplasm (probable).	cytoplasm
Q6AJL6	ISPE_DESPS		cytoplasm
Q6AJL8	RL25_DESPS		cytoplasm
Q6AJL9	PTH_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJM4	RL31_DESPS		cytoplasm
Q6AJM7	MURA_DESPS	cytoplasm (probable).	inner membrane

Q6AJN1	Y2720_DESPS		cytoplasm
Q6AJQ3	EX7S_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJQ6	HSLU_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJR0	CSRA_DESPS		cytoplasm
Q6AJR7	FLGI_DESPS	periplasm (by similarity).	periplasm
Q6AJR8	FLGH_DESPS	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q6AJV8	HIS3_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJW3	MSBA_DESPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6AJY2	Y2615_DESPS		inner membrane
Q6AJY5	RBFA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJY7	TRUB_DESPS		cytoplasm
Q6AJY8	RS15_DESPS		cytoplasm
Q6AJZ0	DUT_DESPS		extracellular
Q6AJZ3	AMPA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AJZ9	RL9_DESPS		cytoplasm
Q6AK01	RS18_DESPS		cytoplasm
Q6AK02	RS6_DESPS		cytoplasm
Q6AK06	RL27_DESPS		cytoplasm
Q6AK08	PROB_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AK09	PROA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AK45	SYI_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AK58	TIG_DESPS		cytoplasm
Q6AK59	CLPP_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AK60	CLPX_DESPS		cytoplasm
Q6AK68	Y2529_DESPS		cytoplasm
Q6AKB8	UVRC_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AKI4	TOLB_DESPS	periplasm (potential).	periplasm
Q6AKK4	RIMK_DESPS		cytoplasm
Q6AKL2	UPPP_DESPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6AKP0	TRUA2_DESPS		cytoplasm
Q6AKP3	HUTH_DESPS	cytoplasm (potential).	cytoplasm
Q6AL12	PPNK_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AL19	FTHS_DESPS		cytoplasm
Q6AL30	Y2216_DESPS		inner membrane
Q6AL31	Y2215_DESPS		inner membrane
Q6AL53	LEPA_DESPS		cytoplasm
Q6AL71	KSGA_DESPS		cytoplasm
Q6AL80	ISPH_DESPS		cytoplasm
Q6ALS3	DEOC_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ALU8	NDK_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ALV4	PAND_DESPS		cytoplasm
Q6ALV8	LPXK_DESPS		cytoplasm
Q6ALW3	SERC_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ALW6	MIAA_DESPS		cytoplasm
Q6ALZ3	LFTR_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AM16	PSTB_DESPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6AM21	GLYA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AM97	ENO_DESPS	cytoplasm (by similarity).	extracellular
Q6AMA0	SYV_DESPS	cytoplasm (by similarity).	cytoplasm

Q6AMF7	THIG_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AMF9	GLMU_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AMG3	KAD_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AML3	RL28_DESPS		cytoplasm
Q6AMQ3	DNAK_DESPS		cytoplasm
Q6AMQ5	GLMM_DESPS		cytoplasm
Q6AMR6	TRPA_DESPS		cytoplasm
Q6AMS5	TRPD_DESPS		cytoplasm
Q6AMS8	RS20_DESPS		cytoplasm
Q6AMT5	Y1611_DESPS		inner membrane
Q6AN63	DNAJ_DESPS	cytoplasm (probable).	cytoplasm
Q6AN65	PURA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ANC0	RL20_DESPS		cytoplasm
Q6ANH1	ADEC2_DESPS		cytoplasm
Q6ANH2	ADEC1_DESPS		cytoplasm
Q6ANH6	Y1369_DESPS		cytoplasm
Q6ANL3	UBIE_DESPS		cytoplasm
Q6ANL7	HIS7_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ANM1	TRUA1_DESPS		cytoplasm
Q6ANP1	Y1304_DESPS		inner membrane
Q6ANR1	LEU3_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ANR2	GIDB_DESPS		cytoplasm
Q6ANR3	HISX_DESPS		cytoplasm
Q6ANS2	RNPH_DESPS		cytoplasm
Q6ANV0	RNC_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ANV2	METX_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ANW2	GSHAB_DESPS		cytoplasm
Q6AP26	CRCB_DESPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6AP32	ISPG_DESPS		cytoplasm
Q6AP35	DXR_DESPS		cytoplasm
Q6AP38	RRF_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AP43	RPOA_DESPS		cytoplasm
Q6AP44	RS4_DESPS		cytoplasm
Q6AP45	RS11_DESPS		cytoplasm
Q6AP46	RS13_DESPS		cytoplasm
Q6AP51	RL15_DESPS		cytoplasm
Q6AP53	RS5_DESPS		cytoplasm
Q6AP54	RL18_DESPS		cytoplasm
Q6AP62	RS17_DESPS		cytoplasm
Q6AP64	RL16_DESPS		cytoplasm
Q6AP65	RS3_DESPS		cytoplasm
Q6AP67	RS19_DESPS		cytoplasm
Q6AP68	RL2_DESPS		cytoplasm
Q6AP72	RS10_DESPS		cytoplasm
Q6AP75	RS7_DESPS		cytoplasm
Q6AP76	RS12_DESPS		cytoplasm
Q6AP77	RPOC_DESPS		cytoplasm
Q6AP78	RPOB_DESPS		cytoplasm
Q6AP80	RL10_DESPS		cytoplasm
Q6AP81	RL1_DESPS		cytoplasm
Q6AP82	RL11_DESPS		cytoplasm
Q6APB0	HEMH_DESPS	cytoplasm (by similarity).	extracellular

Q6APL9	Y976_DESPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6APT2	FPG_DESPS		cytoplasm
Q6APU2	GUAA_DESPS		cytoplasm
Q6APV5	Y890_DESPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6APV7	CLPS_DESPS		cytoplasm
Q6APZ0	RNPA_DESPS		extracellular
Q6APZ2	GIDA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6APZ3	LGT_DESPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6AQ04	MUTS_DESPS		cytoplasm
Q6AQ09	ATPE_DESPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6AQ11	ATPG_DESPS		inner membrane
Q6AQ12	ATPA_DESPS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6AQ48	G6PI_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AQ79	KDSA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AQ97	FMT_DESPS		cytoplasm
Q6AQD7	HAM1_DESPS		extracellular
Q6AQF1	PRMA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AQG4	XGPT_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AQH2	UPP_DESPS		cytoplasm
Q6AQI3	MDH_DESPS		cytoplasm
Q6AQJ3	GPDA_DESPS	cytoplasm (probable).	cytoplasm
Q6AQK2	HIS8_DESPS		cytoplasm
Q6AQT4	HSLO_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AQT5	ACKA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AQV6	Y538_DESPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6AQW6	LUXS_DESPS		extracellular
Q6AQX3	SSRP_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AR56	ARGB_DESPS	cytoplasm (probable).	cytoplasm
Q6AR58	OTC_DESPS	cytoplasm (probable).	cytoplasm
Q6AR59	ASSY_DESPS	cytoplasm (probable).	cytoplasm
Q6AR60	ARLY_DESPS	cytoplasm (probable).	cytoplasm
Q6AR64	DAPB_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ARA9	Y386_DESPS	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6ARJ9	LIPB_DESPS	cytoplasm (potential).	cytoplasm
Q6ARL8	DNAA_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ARM0	HTPG_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ARN9	ISPDF_DESPS		cytoplasm
Q6ARV6	CH60_DESPS	cytoplasm (by similarity).	cytoplasm
Q6ARX8	QUEF_DESPS	cytoplasm (probable).	cytoplasm
Q6AS26	DTD_DESPS	cytoplasm (probable).	cytoplasm
Q6AS31	SECB_DESPS	cytoplasm (by similarity).	cytoplasm
Q6AS40	PYRB_DESPS		cytoplasm

Q6AS42	SFSA_DESPS		cytoplasm
Q6AS54	NIKR_DESPS		cytoplasm
Q6ASE4	END4_DESPS		extracellular
Q6AW21	MDH_MORJA		cytoplasm
Q6AW23	MDH_VIBMA		cytoplasm
Q6BEX0	YTFR_ECOLI		inner membrane
Q6BEX5	YJDP_ECOLI		nucleoid
Q6BF16	DGOA_ECOLI		cytoplasm
Q6BF17	DGOD_ECOLI		cytoplasm
Q6CYI6	GIDA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CYI7	GIDB_ERWCT		cytoplasm
Q6CYJ3	ATPA_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6CYJ4	ATPG_ERWCT		inner membrane
Q6CYJ6	ATPE_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6CYJ8	GLMU_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CYJ9	GLMS_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CYM1	Y4486_ERWCT		inner membrane
Q6CYN3	PSTB2_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6CYR1	RNPA_ERWCT		extracellular
Q6CYR4	DNAA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CYR6	RECF_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CYU4	SSUD_ERWCT		cytoplasm
Q6CYV2	IBPA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CYV3	IBPB_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CYV4	Y4401_ERWCT	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6CZ00	TUSA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZ45	FIEF_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6CZ80	AGUA_ERWCT		extracellular
Q6CZ89	RRAA_ERWCT		cytoplasm
Q6CZ91	HSLU_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZ92	HSLV_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZ96	RL31_ERWCT		cytoplasm
Q6CZB4	TRMA_ERWCT		cytoplasm
Q6CZB5	BTUB_ERWCT	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q6CZC7	ILVD_ERWCT		cytoplasm
Q6CZD1	ILVC_ERWCT		cytoplasm
Q6CZD8	GPPA_ERWCT		extracellular
Q6CZD9	RHLB_ERWCT		cytoplasm
Q6CZF0	WECF_ERWCT	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q6CZF1	WZYE_ERWCT	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q6CZF2	WECG_ERWCT		cytoplasm
Q6CZG3	HEM3_ERWCT		cytoplasm
Q6CZG6	CYAY_ERWCT		inner membrane
Q6CZH4	CORA_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6CZK0	GLGB_ERWCT		extracellular
Q6CZK2	GLGC_ERWCT		cytoplasm
Q6CZK3	GLGA_ERWCT		cytoplasm
Q6CZL2	GLPE_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZL9	BIOH_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZP5	PPCK_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZP6	HSLO_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZQ8	AROK_ERWCT	cytoplasm (probable).	cytoplasm
Q6CZQ9	AROB_ERWCT	cytoplasm (probable).	cytoplasm
Q6CZR3	GPH_ERWCT		cytoplasm
Q6CZU4	KEFG_ERWCT		cytoplasm
Q6CZU5	KEFB_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6CZV1	SLYX_ERWCT		inner membrane
Q6CZV5	GLYA2_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZW0	TUSD_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZW1	TUSC_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZW2	TUSB_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6CZW3	RS12_ERWCT		cytoplasm
Q6CZW4	RS7_ERWCT		cytoplasm
Q6CZW9	RS10_ERWCT		cytoplasm
Q6CZX3	RL2_ERWCT		cytoplasm
Q6CZX4	RS19_ERWCT		cytoplasm
Q6CZX6	RS3_ERWCT		cytoplasm
Q6CZX7	RL16_ERWCT		cytoplasm
Q6CZX8	RL29_ERWCT		cytoplasm
Q6CZX9	RS17_ERWCT		cytoplasm
Q6CZY4	RS8_ERWCT		cytoplasm
Q6CZY6	RL18_ERWCT		cytoplasm
Q6CZY7	RS5_ERWCT		cytoplasm
Q6CZY9	RL15_ERWCT		cytoplasm
Q6CZZ2	RS13_ERWCT		cytoplasm
Q6CZZ3	RS11_ERWCT		cytoplasm
Q6CZZ4	RS4_ERWCT		cytoplasm
Q6CZZ5	RPOA_ERWCT		cytoplasm
Q6CZZ8	MSCL_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D001	FMT_ERWCT		cytoplasm
Q6D004	SMG_ERWCT		cytoplasm
Q6D009	META_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D012	ACEK_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D022	G6PI_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D035	PSD_ERWCT		cytoplasm
Q6D036	ENGC_ERWCT		cytoplasm
Q6D037	ORN_ERWCT	cytoplasm (potential).	cytoplasm
Q6D066	MIAA_ERWCT		cytoplasm
Q6D067	HFQ_ERWCT		cytoplasm
Q6D072	PURA_ERWCT	cytoplasm (by similarity).	cytoplasm

Q6D075	RUVX_ERWCT	cytoplasm (potential).	cytoplasm
Q6D090	ARGO_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D093	RPIA_ERWCT		cytoplasm
Q6D0A0	TRPR_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D0A1	Y3898_ERWCT		inner membrane
Q6D0A9	KHSE_ERWCT	cytoplasm (potential).	cytoplasm
Q6D0B2	TAL2_ERWCT	cytoplasm (probable).	cytoplasm
Q6D0B7	DNAK_ERWCT		cytoplasm
Q6D0B8	DNAJ_ERWCT	cytoplasm (probable).	cytoplasm
Q6D0C1	RS20_ERWCT		cytoplasm
Q6D0C3	SYI_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D0C6	ISPH_ERWCT		cytoplasm
Q6D0C7	DAPB_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D0C8	CARA_ERWCT		cytoplasm
Q6D0D8	APAH_ERWCT		extracellular
Q6D0E0	KSGA_ERWCT		cytoplasm
Q6D0E3	OSTA_ERWCT	bacterial cell outer membrane (by similarity).	outer membrane
Q6D0E4	DJLA_ERWCT	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q6D0E6	RAPA_ERWCT		cytoplasm
Q6D0G5	LEUD_ERWCT		cytoplasm
Q6D0G6	LEU2_ERWCT		cytoplasm
Q6D0G7	LEU3_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D0H4	MRAZ_ERWCT		inner membrane
Q6D0H5	MRAW_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D0I0	MRAY_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D0I1	MURD_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D0I3	MURG_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6D0I4	MURC_ERWCT	cytoplasm (probable).	cytoplasm
Q6D0I9	LPXC_ERWCT		cytoplasm
Q6D0J4	Y3804_ERWCT		cytoplasm
Q6D0J5	Y3803_ERWCT		inner membrane
Q6D107	AAS_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D117	HAM1_ERWCT		extracellular
Q6D127	RLMB_ERWCT		cytoplasm
Q6D135	RS6_ERWCT		cytoplasm
Q6D137	RS18_ERWCT		cytoplasm
Q6D138	RL9_ERWCT		cytoplasm
Q6D142	RCMNS_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D151	MSRA_ERWCT		cytoplasm
Q6D157	Y3591_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D159	UPPP_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q6D160	CCA_ERWCT		cytoplasm
Q6D163	GLNE_ERWCT		cytoplasm
Q6D178	RUMA_ERWCT		cytoplasm
Q6D182	ENO_ERWCT	cytoplasm (by similarity).	extracellular
Q6D1A1	CYSJ_ERWCT		cytoplasm
Q6D1A2	CYSI_ERWCT		cytoplasm
Q6D1B3	ISPD_ERWCT		cytoplasm
Q6D1B4	ISPF_ERWCT		extracellular
Q6D1B5	TRUD_ERWCT		cytoplasm
Q6D1B6	PIMT_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D1I0	XGPT_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D1I3	PROB_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D1I4	PROA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D1S8	RECA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D1S9	RECX_ERWCT	cytoplasm (potential).	cytoplasm
Q6D1T1	CSRA_ERWCT		cytoplasm
Q6D1T4	GSH1_ERWCT		cytoplasm
Q6D1T5	LUXS_ERWCT		extracellular
Q6D1T9	RIMM_ERWCT	cytoplasm (potential).	cytoplasm
Q6D1U0	TRMD_ERWCT	cytoplasm (potential).	cytoplasm
Q6D1U1	RL19_ERWCT		cytoplasm
Q6D1V7	RNH_ERWCT	cytoplasm (potential).	extracellular
Q6D1W1	NRDI_ERWCT		inner membrane
Q6D1W3	SPED_ERWCT		cytoplasm
Q6D1X4	PAND_ERWCT		cytoplasm
Q6D1X5	PANC_ERWCT		cytoplasm
Q6D1X6	PANB_ERWCT		cytoplasm
Q6D1X9	GLUQ_ERWCT		cytoplasm
Q6D1Y1	SFSA_ERWCT		cytoplasm
Q6D1Z0	GSA_ERWCT	cytoplasm (potential).	cytoplasm
Q6D1Z3	BTUF_ERWCT	periplasm (by similarity).	periplasm
Q6D201	CYSA_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6D208	UNG_ERWCT	cytoplasm (by similarity).	extracellular
Q6D209	GRCA_ERWCT		cytoplasm
Q6D217	LEPA_ERWCT		cytoplasm
Q6D219	RNC_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D221	RECO_ERWCT		cytoplasm
Q6D222	ACPS_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D245	HMP_ERWCT		cytoplasm
Q6D246	GLYA1_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D261	ISCA_ERWCT		cytoplasm
Q6D262	HSCB_ERWCT		inner membrane
Q6D263	HSCA_ERWCT		cytoplasm
Q6D266	PEPB_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D272	NDK_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D276	ISPG_ERWCT		cytoplasm
Q6D288	GUAA_ERWCT		cytoplasm
Q6D2A1	PSIE_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D2B0	MDTC_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q6D2B1	MDTB_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q6D2B2	MDTA_ERWCT	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q6D2D5	PSTB1_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6D2E9	ARNB_ERWCT		cytoplasm
Q6D2F0	ARNC_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D2F1	ARNA_ERWCT		cytoplasm
Q6D2F3	ARNT_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D2F6	FBPC2_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6D2L6	FADI_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D2L7	FADJ_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D2M6	AROC_ERWCT		cytoplasm
Q6D2M7	MEPA_ERWCT	periplasm (by similarity).	extracellular
Q6D2N1	Y3064_ERWCT		cytoplasm
Q6D2N5	PDXB_ERWCT		cytoplasm
Q6D2N7	TRUA_ERWCT		cytoplasm
Q6D2Q7	Y3038_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D2Q8	Y3037_ERWCT		inner membrane
Q6D2R1	Y3034_ERWCT		cytoplasm
Q6D364	PGSA_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D3B3	MAO1_ERWCT		cytoplasm
Q6D3B4	CDD_ERWCT		cytoplasm
Q6D3C4	UVRB_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D3C7	MOAA_ERWCT		cytoplasm
Q6D3F5	IDI_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D3J6	Y2748_ERWCT		inner membrane
Q6D3J7	NDPA_ERWCT	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q6D3J9	RL25_ERWCT		cytoplasm
Q6D3L2	EFPL_ERWCT		cytoplasm
Q6D3L8	END4_ERWCT		extracellular
Q6D3R0	DEOC_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D3R1	Y2683_ERWCT	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6D3R5	RIMK_ERWCT		cytoplasm
Q6D3S2	RUMB_ERWCT		cytoplasm
Q6D3S8	Y2666_ERWCT		inner membrane
Q6D3T7	CLPS_ERWCT		cytoplasm
Q6D3U2	LFTR_ERWCT	cytoplasm (by similarity).	cytoplasm

Q6D400	SERC_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D402	KCY_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D406	HIS6_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D407	HIS4_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D408	HIS5_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D409	HIS7_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D410	HIS8_ERWCT		cytoplasm
Q6D411	HISX_ERWCT		cytoplasm
Q6D421	CITD_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D425	CITG_ERWCT		cytoplasm
Q6D437	MSBA_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D438	LPXK_ERWCT		cytoplasm
Q6D445	MUKF_ERWCT	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q6D454	PNCB_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D456	PYRD_ERWCT	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q6D465	Y2529_ERWCT		cytoplasm
Q6D469	MDTH_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D476	CUTC_ERWCT	cytoplasm (probable).	periplasm
Q6D499	Y2494_ERWCT		inner membrane
Q6D4A0	RUVC_ERWCT		cytoplasm
Q6D4A1	RUVA_ERWCT		cytoplasm
Q6D4A2	RUVB_ERWCT		cytoplasm
Q6D4E3	PEPT_ERWCT	cytoplasm (by similarity).	extracellular
Q6D4E9	TRMU_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D4H1	RL20_ERWCT		cytoplasm
Q6D4K1	Y2389_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D4L3	Y2377_ERWCT		inner membrane
Q6D4P7	MSRB_ERWCT		cytoplasm
Q6D4R3	KITH_ERWCT	cytoplasm (potential).	cytoplasm
Q6D4T9	TRPA_ERWCT		cytoplasm
Q6D4U2	TRPD_ERWCT		cytoplasm
Q6D4W5	ARAA_ERWCT		cytoplasm
Q6D4W6	ARAB_ERWCT		cytoplasm
Q6D547	KDSA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D552	HEM1_ERWCT		cytoplasm
Q6D554	ISPE_ERWCT		cytoplasm
Q6D557	PTH_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D5N6	ACPD_ERWCT		extracellular
Q6D5P8	ZNTB_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D5U5	GCH2_ERWCT		extracellular
Q6D5V2	SYX_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D5V3	PDXH_ERWCT		cytoplasm
Q6D5V7	SLYA_ERWCT		cytoplasm

Q6D5W6	MDTK_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D5Z2	NAPA_ERWCT	periplasm (by similarity).	periplasm
Q6D602	CCME_ERWCT	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q6D606	CCMA_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6D624	SUFE_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D625	SUFS_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D636	Y1852_ERWCT		cytoplasm
Q6D654	BTUD_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6D656	BTUC_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D674	NAGZ_ERWCT	cytoplasm (by similarity).	extracellular
Q6D683	KTHY_ERWCT		cytoplasm
Q6D687	ACP_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D690	FABH_ERWCT	cytoplasm (probable).	cytoplasm
Q6D691	PLSX_ERWCT		cytoplasm
Q6D692	RL32_ERWCT		cytoplasm
Q6D6A3	Y1782_ERWCT	periplasm (potential).	periplasm
Q6D6A4	Y1781_ERWCT		inner membrane
Q6D6C3	TUSE_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D6C8	MGSA_ERWCT		cytoplasm
Q6D6D7	FABA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D6E0	BETB_ERWCT		cytoplasm
Q6D6H2	FLGI_ERWCT	periplasm (by similarity).	periplasm
Q6D6H3	FLGH_ERWCT	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q6D6J5	FLHD_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D6J8	DSRB_ERWCT		inner membrane
Q6D734	FBPC1_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6D7C2	6PGL_ERWCT		cytoplasm
Q6D7C4	GLK_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D7D0	MODC_ERWCT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6D7E3	GPMA_ERWCT		cytoplasm
Q6D7E5	ZITB_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D7F2	TOLB_ERWCT	periplasm (potential).	periplasm
Q6D7H0	END8_ERWCT		cytoplasm
Q6D7H1	PCP_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D7I3	ATKA_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q6D7K8	Y1317_ERWCT		inner membrane

Q6D7L9	NADD_ERWCT		cytoplasm
Q6D7M1	Y1304_ERWCT		inner membrane
Q6D7M7	LIPB_ERWCT	cytoplasm (potential).	cytoplasm
Q6D7N0	CRCB_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D7R8	HDA_ERWCT		nucleoid
Q6D7S0	UPP_ERWCT		cytoplasm
Q6D7U0	HYPB_ERWCT		cytoplasm
Q6D7X5	UBIG_ERWCT		cytoplasm
Q6D7Z4	HEMH_ERWCT	cytoplasm (by similarity).	extracellular
Q6D7Z5	KAD_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D7Z6	HTPG_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D7Z7	RECR_ERWCT		cytoplasm
Q6D807	RL31B_ERWCT		cytoplasm
Q6D826	CLPX_ERWCT		cytoplasm
Q6D827	CLPP_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D828	TIG_ERWCT		cytoplasm
Q6D842	EX7S_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D850	NRDR_ERWCT		cytoplasm
Q6D855	TGT_ERWCT		cytoplasm
Q6D856	QUEA_ERWCT	cytoplasm (potential).	cytoplasm
Q6D857	ACPH_ERWCT		cytoplasm
Q6D871	Y1104_ERWCT		inner membrane
Q6D872	AROL_ERWCT	cytoplasm (probable).	cytoplasm
Q6D882	UXUA_ERWCT		cytoplasm
Q6D8C1	MUTS_ERWCT		cytoplasm
Q6D8C5	TILS_ERWCT	cytoplasm (probable).	cytoplasm
Q6D8C7	ACCA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D8C9	RNH2_ERWCT	cytoplasm (potential).	extracellular
Q6D8D2	FABZ_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D8D3	LPXD_ERWCT		cytoplasm
Q6D8D4	SKP_ERWCT	periplasm (by similarity).	periplasm
Q6D8D5	YAET_ERWCT	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
Q6D8D8	UPPS_ERWCT		cytoplasm
Q6D8D9	DXR_ERWCT		cytoplasm
Q6D8E0	RRF_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D8E3	RS2_ERWCT		cytoplasm
Q6D8E5	GLND_ERWCT		cytoplasm
Q6D8E6	DAPD_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D8E7	Y1027_ERWCT		inner membrane
Q6D8F4	QUEF_ERWCT	cytoplasm (probable).	cytoplasm
Q6D8F7	MTFA_ERWCT		cytoplasm
Q6D8I6	TYSY_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D8I7	LGT_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D8I9	NUDH_ERWCT		cytoplasm
Q6D8J7	TRMB_ERWCT		cytoplasm
Q6D8J9	FETP_ERWCT		cytoplasm
Q6D8K0	MLTC_ERWCT	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular

Q6D8S6	ZIPA_ERWCT	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q6D8S7	CYSZ_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D8U8	HEM6_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D8U9	Y873_ERWCT		inner membrane
Q6D8W0	TAL1_ERWCT	cytoplasm (probable).	cytoplasm
Q6D8X9	GRPE_ERWCT	cytoplasm (probable).	cytoplasm
Q6D8Y0	PPNK_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D8Y5	SSRP_ERWCT	cytoplasm (potential).	cytoplasm
Q6D913	ACTP_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6D960	Y759_ERWCT		cytoplasm
Q6D974	GCSP_ERWCT		cytoplasm
Q6D976	GCST_ERWCT		cytoplasm
Q6D989	DEOD_ERWCT		cytoplasm
Q6D991	TYPH_ERWCT		cytoplasm
Q6D994	Y725_ERWCT		cytoplasm
Q6D9A2	RS15_ERWCT		cytoplasm
Q6D9A3	TRUB_ERWCT		cytoplasm
Q6D9A4	RBFA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D9A7	Y710_ERWCT		inner membrane
Q6D9B6	GLMM_ERWCT		cytoplasm
Q6D9B9	RRMJ_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D9C5	RL27_ERWCT		cytoplasm
Q6D9D1	MDH_ERWCT		cytoplasm
Q6D9D2	ARGR_ERWCT	cytoplasm (probable).	cytoplasm
Q6D9D4	RS21_ERWCT		cytoplasm
Q6D9D7	MUG_ERWCT	cytoplasm (potential).	extracellular
Q6D9G9	UXAB_ERWCT		cytoplasm
Q6D9H0	UXAC_ERWCT		cytoplasm
Q6D9I4	Y631_ERWCT		cytoplasm
Q6D9I5	LEXA_ERWCT		cytoplasm
Q6D9J0	CH60_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D9J1	CH10_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6D9J5	CUTA_ERWCT	cytoplasm (by similarity).	periplasm
Q6DA13	RHAT_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6DA21	RHAR_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DA22	RHAS_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DA23	RHAB_ERWCT		cytoplasm
Q6DA24	RHAA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DA25	RHAD_ERWCT	cytoplasm (by similarity).	extracellular
Q6DA52	AMPA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DA54	SYV_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DA62	KPTA_ERWCT		cytoplasm
Q6DA71	OTC_ERWCT	cytoplasm (probable).	cytoplasm
Q6DA73	PYRB_ERWCT		cytoplasm
Q6DA74	PYRI_ERWCT		cytoplasm
Q6DAD4	DIAA_ERWCT		cytoplasm
Q6DAF2	MURA_ERWCT	cytoplasm (probable).	inner membrane
Q6DAG8	Y285_ERWCT		cytoplasm

Q6DAH4	AAEB_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6DAH5	AAEA_ERWCT	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q6DAH6	AAEX_ERWCT	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6DAI9	Y264_ERWCT		cytoplasm
Q6DAJ5	PRMA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DAJ6	CAH_ERWCT	periplasm (potential).	periplasm
Q6DAJ8	FIS_ERWCT		cytoplasm
Q6DAL8	NUDC_ERWCT		cytoplasm
Q6DAM4	THIG_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DAM9	RPOC_ERWCT		cytoplasm
Q6DAN0	RPOB_ERWCT		cytoplasm
Q6DAN2	RL10_ERWCT		cytoplasm
Q6DAN3	RL1_ERWCT		cytoplasm
Q6DAN4	RL11_ERWCT		cytoplasm
Q6DAP0	MURB_ERWCT	cytoplasm (probable).	inner membrane
Q6DAP5	FADB_ERWCT		cytoplasm
Q6DAP6	FADA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DAQ5	UBIB_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6DAQ7	UBIE_ERWCT		cytoplasm
Q6DAQ9	ARLY_ERWCT	cytoplasm (probable).	cytoplasm
Q6DAR0	ARGB_ERWCT	cytoplasm (probable).	cytoplasm
Q6DAR1	ARGC_ERWCT	cytoplasm (probable).	cytoplasm
Q6DAR6	CAPP_ERWCT		cytoplasm
Q6DAT0	GPDA_ERWCT	cytoplasm (probable).	cytoplasm
Q6DAT1	SECB_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DAT5	TDH_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DAV2	COAD_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DAV4	FPG_ERWCT		cytoplasm
Q6DAV6	RL28_ERWCT		cytoplasm
Q6DAV7	RADC_ERWCT		cytoplasm
Q6DAV9	DUT_ERWCT		extracellular
Q6DAW0	SLMA_ERWCT	nucleoid-associated (by similarity).	cytoplasm
Q6DAW2	RNPH_ERWCT		cytoplasm
Q6DAY3	LLDD_ERWCT		cytoplasm
Q6DB05	XYLA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DB09	FDHD_ERWCT	cytoplasm (potential).	cytoplasm
Q6DB47	Y054_ERWCT		inner membrane
Q6DB52	USPB_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6DB60	KGUA_ERWCT	cytoplasm (by similarity).	cytoplasm
Q6DB61	RPOZ_ERWCT		cytoplasm
Q6DB68	DTD_ERWCT	cytoplasm (probable).	cytoplasm
Q6DB92	KUP_ERWCT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6DB93	RAVA_ERWCT	cytoplasm (by similarity).	cytoplasm

Q6DB94	VIAA_ERWCT		inner membrane
Q6F6K8	RNPA_ACIAD		extracellular
Q6F6N3	DNAK_ACIAD		cytoplasm
Q6F6N4	GRPE_ACIAD	cytoplasm (probable).	cytoplasm
Q6F6P9	FMT_ACIAD		cytoplasm
Q6F6Q0	ILVD2_ACIAD		cytoplasm
Q6F6Q6	CAPP_ACIAD		cytoplasm
Q6F6R1	DNAJ_ACIAD	cytoplasm (probable).	cytoplasm
Q6F6R2	DAPB_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F6U8	GLMS_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F6U9	GLMU_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F6W0	DTD_ACIAD	cytoplasm (probable).	cytoplasm
Q6F6W1	PNCB_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F6W3	PSD_ACIAD		cytoplasm
Q6F6Y1	ALLC_ACIAD		extracellular
Q6F6Z6	PYRE_ACIAD		cytoplasm
Q6F703	MURG_ACIAD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6F704	MURC_ACIAD	cytoplasm (probable).	cytoplasm
Q6F717	GLMM_ACIAD		cytoplasm
Q6F718	PDXH_ACIAD		cytoplasm
Q6F793	KUP_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6F799	HIS6_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F7A5	HIS4_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F7A7	HIS5_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F7A8	HIS7_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F7D1	MRAW_ACIAD		cytoplasm
Q6F7D6	MRAY_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6F7E4	AROK_ACIAD	cytoplasm (probable).	cytoplasm
Q6F7E5	AROB_ACIAD	cytoplasm (probable).	cytoplasm
Q6F7G9	RPOZ_ACIAD		cytoplasm
Q6F7H0	KGUA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F7I0	RIMM_ACIAD	cytoplasm (potential).	cytoplasm
Q6F7I2	RL19_ACIAD		cytoplasm
Q6F7I5	TRUB_ACIAD		cytoplasm
Q6F7N0	HEMH_ACIAD	cytoplasm (by similarity).	extracellular
Q6F7N4	GCH2_ACIAD		extracellular
Q6F7R1	RS10_ACIAD		cytoplasm
Q6F7R5	RL2_ACIAD		cytoplasm
Q6F7R6	RS19_ACIAD		cytoplasm
Q6F7R8	RS3_ACIAD		cytoplasm
Q6F7R9	RL16_ACIAD		cytoplasm
Q6F7S0	RL29_ACIAD		cytoplasm
Q6F7S1	RS17_ACIAD		cytoplasm
Q6F7S6	RS8_ACIAD		cytoplasm
Q6F7S8	RL18_ACIAD		cytoplasm
Q6F7S9	RS5_ACIAD		cytoplasm
Q6F7T1	RL15_ACIAD		cytoplasm
Q6F7T4	RS13_ACIAD		cytoplasm
Q6F7T5	RS11_ACIAD		cytoplasm
Q6F7T6	RS4_ACIAD		cytoplasm

Q6F7T7	RPOA_ACIAD		cytoplasm
Q6F7U1	Y3189_ACIAD		cytoplasm
Q6F7W6	Y3160_ACIAD		inner membrane
Q6F7X1	MDH_ACIAD		cytoplasm
Q6F7Z8	RADC_ACIAD		cytoplasm
Q6F801	SECB_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F805	ORN_ACIAD	cytoplasm (potential).	cytoplasm
Q6F821	ILVC_ACIAD		cytoplasm
Q6F847	RUMA_ACIAD		cytoplasm
Q6F855	PANC_ACIAD		cytoplasm
Q6F856	Y3059_ACIAD		cytoplasm
Q6F867	RL35_ACIAD		cytoplasm
Q6F868	RL20_ACIAD		cytoplasm
Q6F880	TILS_ACIAD	cytoplasm (probable).	cytoplasm
Q6F8A0	KSGA_ACIAD		cytoplasm
Q6F8F0	SYV_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F8F4	Y2946_ACIAD		inner membrane
Q6F8G2	RL27_ACIAD		cytoplasm
Q6F8H3	LIPB_ACIAD	cytoplasm (potential).	cytoplasm
Q6F8H8	TPMT_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F8I3	Y2917_ACIAD		inner membrane
Q6F8I6	PAND_ACIAD		cytoplasm
Q6F8I7	PTH_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F8I8	RL25_ACIAD		cytoplasm
Q6F8J0	ISPE_ACIAD		cytoplasm
Q6F8J3	HEM1_ACIAD		cytoplasm
Q6F8J9	SSRP_ACIAD	cytoplasm (potential).	cytoplasm
Q6F8K0	COAD_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F8L4	SUCC_ACIAD		cytoplasm
Q6F8M7	CARA_ACIAD		cytoplasm
Q6F8N1	RRMJ_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F8P2	PPCK_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F8P5	CH10_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F8P6	CH60_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F961	SERC_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F963	Y2644_ACIAD		inner membrane
Q6F985	TOLB_ACIAD	periplasm (potential).	periplasm
Q6F991	RUVB_ACIAD		cytoplasm
Q6F992	RUVA_ACIAD		cytoplasm
Q6F9A5	DAPD_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F9A8	CYSA_ACIAD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6F9B5	KTHY_ACIAD		cytoplasm
Q6F9B9	LEPA_ACIAD		cytoplasm
Q6F9C2	RNC_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F9D2	UVRB_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F9D9	PROB_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F9H7	Y2520_ACIAD		inner membrane
Q6F9I9	PQQE_ACIAD		cytoplasm
Q6F9J0	PQQD_ACIAD		cytoplasm
Q6F9J1	PQQC_ACIAD		cytoplasm
Q6F9J2	PQQB_ACIAD		cytoplasm
Q6F9P9	PRMA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F9Q1	GIDA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6F9Q8	RL9_ACIAD		cytoplasm

Q6F9Q9	RS18_ACIAD		cytoplasm
Q6F9R0	RS6_ACIAD		cytoplasm
Q6F9R8	Y2422_ACIAD		cytoplasm
Q6F9W1	MIAA_ACIAD		cytoplasm
Q6F9W7	GIDB_ACIAD		cytoplasm
Q6F9X0	MSBA_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6F9Y2	UNG_ACIAD	cytoplasm (by similarity).	extracellular
Q6F9Y6	KCY_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FA18	RECR_ACIAD		cytoplasm
Q6FA38	CCA_ACIAD		cytoplasm
Q6FA43	SAHH_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FA53	RS2_ACIAD		cytoplasm
Q6FA61	QUEF_ACIAD	cytoplasm (probable).	cytoplasm
Q6FA66	GLYA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FA87	PPNK_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FAA5	RL31_ACIAD		cytoplasm
Q6FAM5	GLND_ACIAD		cytoplasm
Q6FAP5	Y2052_ACIAD		inner membrane
Q6FAQ2	RUVC_ACIAD		cytoplasm
Q6FAR2	AROC_ACIAD		cytoplasm
Q6FAT8	KDSA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FAT9	ENO_ACIAD	cytoplasm (by similarity).	extracellular
Q6FAU1	ISPD_ACIAD		cytoplasm
Q6FAU4	ISPF_ACIAD		extracellular
Q6FAZ1	MURB_ACIAD	cytoplasm (probable).	inner membrane
Q6FB19	FDHD_ACIAD	cytoplasm (potential).	cytoplasm
Q6FBI2	Y1740_ACIAD		inner membrane
Q6FBI3	AROQ_ACIAD		cytoplasm
Q6FBS0	TRMA_ACIAD		cytoplasm
Q6FBS3	UPPP3_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6FC54	MUTS_ACIAD		cytoplasm
Q6FCE6	HSCA_ACIAD		cytoplasm
Q6FCF4	RRAA_ACIAD		cytoplasm
Q6FCF6	RS20_ACIAD		cytoplasm
Q6FCG1	RECX_ACIAD	cytoplasm (potential).	cytoplasm
Q6FCG4	FABZ_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FCG5	LPXD_ACIAD		cytoplasm
Q6FCG9	DXR_ACIAD		cytoplasm
Q6FCH1	UPPS_ACIAD		cytoplasm
Q6FCH2	RRF_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FCH8	OTC_ACIAD	cytoplasm (probable).	cytoplasm
Q6FCI2	CLPS_ACIAD		cytoplasm
Q6FCI4	ARGC_ACIAD	cytoplasm (probable).	cytoplasm
Q6FCI6	RPIA_ACIAD		cytoplasm
Q6FCL0	RS21_ACIAD		cytoplasm
Q6FCL9	PYRD_ACIAD	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q6FCN1	THIG_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FCN6	TRMB_ACIAD		cytoplasm
Q6FCR5	PYRB_ACIAD		cytoplasm
Q6FCR9	ILVD1_ACIAD		cytoplasm

Q6FCS7	PURA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FCT4	CSRA_ACIAD		cytoplasm
Q6FCT7	RNH2_ACIAD	cytoplasm (potential).	extracellular
Q6FCV6	SURE_ACIAD	cytoplasm (potential).	periplasm
Q6FCW2	TRMU_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FCW7	PSTB_ACIAD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6FCY1	GSA_ACIAD	cytoplasm (potential).	cytoplasm
Q6FD71	KAD_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FD81	UREE_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FD83	URE1_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FD84	URE2_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FD85	URE3_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FDB8	Y1052_ACIAD		inner membrane
Q6FDC1	SDHD_ACIAD		cytoplasm
Q6FDF8	BETB_ACIAD		cytoplasm
Q6FDH0	ASPD_ACIAD		cytoplasm
Q6FDP8	UPPP2_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6FDQ8	ARGB_ACIAD	cytoplasm (probable).	cytoplasm
Q6FDR0	DUT_ACIAD		extracellular
Q6FDS1	LFTR_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FDS2	ATE_ACIAD	cytoplasm (potential).	cytoplasm
Q6FDS7	RS7_ACIAD		cytoplasm
Q6FDS8	RS12_ACIAD		cytoplasm
Q6FDT7	ACP_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FDU0	RL32_ACIAD		cytoplasm
Q6FE58	UPP_ACIAD		cytoplasm
Q6FE77	Y721_ACIAD		extracellular
Q6FE90	FPG_ACIAD		cytoplasm
Q6FE93	MSCL_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6FEB3	Y679_ACIAD		inner membrane
Q6FEB4	UPPP1_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6FEC7	HIS8_ACIAD		cytoplasm
Q6FEC8	HISX_ACIAD		cytoplasm
Q6FEC9	HIS1_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FED0	MURA_ACIAD	cytoplasm (probable).	inner membrane
Q6FED8	ARGJ_ACIAD	cytoplasm (probable).	cytoplasm
Q6FEE6	TRPA_ACIAD		cytoplasm
Q6FEI9	GLNE_ACIAD		cytoplasm
Q6FEJ3	QUEA_ACIAD	cytoplasm (potential).	cytoplasm
Q6FEJ4	TGT_ACIAD		cytoplasm
Q6FEM3	ISPG_ACIAD		cytoplasm
Q6FEM7	NDK_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FEN5	PROA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FEP7	CLPX_ACIAD		cytoplasm
Q6FEP8	CLPP_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FEP9	TIG_ACIAD		cytoplasm
Q6FEQ3	METX_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FEQ6	HAM1_ACIAD		extracellular

Q6FER4	LGT_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6FER7	TYSY_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FES9	RL28_ACIAD		cytoplasm
Q6FEV2	TRUA_ACIAD		cytoplasm
Q6FEV6	LEU3_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FEV8	LEUD_ACIAD		cytoplasm
Q6FEW0	LEU2_ACIAD		cytoplasm
Q6FEW7	NUDH_ACIAD		cytoplasm
Q6FEY7	NORM_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6FEZ1	RL31B_ACIAD		cytoplasm
Q6FEZ6	Y419_ACIAD		cytoplasm
Q6FF10	CRCB_ACIAD	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6FF13	RS15_ACIAD		cytoplasm
Q6FF39	RBFA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FF42	Y367_ACIAD		inner membrane
Q6FF48	COAE_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FF50	RLMB_ACIAD		cytoplasm
Q6FF51	Y356_ACIAD		inner membrane
Q6FF55	RUVX_ACIAD	cytoplasm (potential).	cytoplasm
Q6FF64	UVRC_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FF68	FADB_ACIAD		cytoplasm
Q6FF69	FADA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FF82	HTPG_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FF89	RPOC_ACIAD		cytoplasm
Q6FF90	RPOB_ACIAD		cytoplasm
Q6FF92	RL10_ACIAD		cytoplasm
Q6FF93	RL1_ACIAD		cytoplasm
Q6FF94	RL11_ACIAD		cytoplasm
Q6FFA9	HEM3_ACIAD		cytoplasm
Q6FFB2	ARLY_ACIAD	cytoplasm (probable).	cytoplasm
Q6FFB3	FETP_ACIAD		cytoplasm
Q6FFC0	GLUQ_ACIAD		cytoplasm
Q6FFC2	MURD_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FFD5	GPMI_ACIAD		cytoplasm
Q6FFD8	AMPA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FFE5	NRDR_ACIAD		cytoplasm
Q6FFJ9	ATPE_ACIAD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6FFK1	ATPG_ACIAD		inner membrane
Q6FFK2	ATPA_ACIAD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6FFL8	MAO1_ACIAD		cytoplasm
Q6FFN2	GUAA_ACIAD		cytoplasm
Q6FFS1	LLDD_ACIAD		cytoplasm
Q6FFX5	RNPH_ACIAD		cytoplasm
Q6FFY1	UBIG_ACIAD		cytoplasm
Q6FG02	SYI_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FG10	SYI_ACIAD	cytoplasm (by similarity).	cytoplasm

Q6FG21	DNAA_ACIAD	cytoplasm (by similarity).	cytoplasm
Q6FYB9	GIDA_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FYC0	GIDB_BARQU		cytoplasm
Q6FYD0	MDH_BARQU		cytoplasm
Q6FYD4	ODO2_BARQU		inner membrane
Q6FYG6	RL28_BARQU		cytoplasm
Q6FYH3	RIMM_BARQU	cytoplasm (potential).	cytoplasm
Q6FYH4	TRMD_BARQU	cytoplasm (potential).	cytoplasm
Q6FYH5	RL19_BARQU		cytoplasm
Q6FYM1	ATPA_BARQU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6FYM2	ATPG_BARQU		inner membrane
Q6FYM4	ATPE_BARQU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6FYM8	Y1207_BARQU		inner membrane
Q6FYN6	RL31_BARQU		cytoplasm
Q6FYP4	RUVC_BARQU		cytoplasm
Q6FYP5	RUVA_BARQU		cytoplasm
Q6FYP6	RUVB_BARQU		cytoplasm
Q6FYQ1	TOLB_BARQU	periplasm (potential).	periplasm
Q6FYQ2	Y1179_BARQU		inner membrane
Q6FYQ5	TILS_BARQU	cytoplasm (probable).	cytoplasm
Q6FYQ7	GLMM_BARQU		cytoplasm
Q6FYR2	DUT_BARQU		extracellular
Q6FYT1	MURA_BARQU	cytoplasm (probable).	inner membrane
Q6FYZ7	GCSP_BARQU		cytoplasm
Q6FZ00	Y1007_BARQU		inner membrane
Q6FZ12	GPMA_BARQU		cytoplasm
Q6FZ13	DAPB_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZ17	RNPA_BARQU		extracellular
Q6FZ91	TYSY_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZ96	MIAA_BARQU		cytoplasm
Q6FZ98	ILVC_BARQU		cytoplasm
Q6FZB1	SYV_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZB7	RS12_BARQU		cytoplasm
Q6FZB8	RS7_BARQU		cytoplasm
Q6FZC1	RS10_BARQU		cytoplasm
Q6FZC5	RL2_BARQU		cytoplasm
Q6FZC6	RS19_BARQU		cytoplasm
Q6FZC8	RS3_BARQU		cytoplasm
Q6FZC9	RL16_BARQU		cytoplasm
Q6FZD0	RL29_BARQU		cytoplasm
Q6FZD1	RS17_BARQU		cytoplasm
Q6FZD6	RS8_BARQU		cytoplasm
Q6FZD8	RL18_BARQU		cytoplasm
Q6FZD9	RS5_BARQU		cytoplasm
Q6FZE1	RL15_BARQU		cytoplasm
Q6FZE3	KAD_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZE4	RS13_BARQU		cytoplasm
Q6FZE5	RS11_BARQU		cytoplasm
Q6FZE6	RPOA_BARQU		cytoplasm
Q6FZF0	RECA_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZG1	COAD_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZH5	GLMU_BARQU	cytoplasm (by similarity).	cytoplasm

Q6FZH6	GLMS_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZI9	PURL_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZJ0	PURQ_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZK6	TIG_BARQU		cytoplasm
Q6FZL5	RL11_BARQU		cytoplasm
Q6FZL6	RL1_BARQU		cytoplasm
Q6FZL7	RL10_BARQU		cytoplasm
Q6FZM0	RPOC_BARQU		cytoplasm
Q6FZM5	CLPS_BARQU		cytoplasm
Q6FZN0	RS2_BARQU		cytoplasm
Q6FZN3	RRF_BARQU	cytoplasm (by similarity).	cytoplasm
Q6FZP1	RPIA_BARQU		cytoplasm
Q6FZP9	HUTH_BARQU	cytoplasm (potential).	cytoplasm
Q6FZQ9	LIPB_BARQU	cytoplasm (potential).	cytoplasm
Q6FZS3	Y640_BARQU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6FZS5	PYRB_BARQU		cytoplasm
Q6FZS7	RUVX_BARQU	cytoplasm (potential).	cytoplasm
Q6FZU1	SFSA_BARQU		cytoplasm
Q6FZU4	RS4_BARQU		cytoplasm
Q6FZU5	LEXA_BARQU		cytoplasm
Q6FZU8	KTHY_BARQU		cytoplasm
Q6FZW4	SYX_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G001	PLSX_BARQU		cytoplasm
Q6G008	NRDR_BARQU		cytoplasm
Q6G009	GLYA_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G014	HFQ_BARQU		cytoplasm
Q6G026	KDSA_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G043	NDK_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G052	KSGA_BARQU		cytoplasm
Q6G053	KGUA_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G056	ACP_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G060	RS6_BARQU		cytoplasm
Q6G061	RS18_BARQU		cytoplasm
Q6G063	RL9_BARQU		cytoplasm
Q6G070	PSD_BARQU		cytoplasm
Q6G078	PANB_BARQU		cytoplasm
Q6G079	PANC_BARQU		cytoplasm
Q6G080	RECO_BARQU		cytoplasm
Q6G082	RNC_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G084	ACPS_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G088	RPOZ_BARQU		cytoplasm
Q6G090	SSRP_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0A2	THIG_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0B0	GLNE_BARQU		cytoplasm
Q6G0C9	ISPH_BARQU		cytoplasm
Q6G0D5	EX7S_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0E0	PDXH_BARQU		cytoplasm
Q6G0E6	RNH2_BARQU	cytoplasm (potential).	extracellular
Q6G0F4	LGT_BARQU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6G0F8	RL25_BARQU		cytoplasm
Q6G0F9	PTH_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0G3	ISPE_BARQU		cytoplasm

Q6G0I1	UBIG_BARQU		cytoplasm
Q6G0J2	LLDD_BARQU		cytoplasm
Q6G0J6	LPXK_BARQU		cytoplasm
Q6G0K6	SYI_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0L3	FPG_BARQU		cytoplasm
Q6G0L7	PSTB_BARQU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6G0M8	RECR_BARQU		cytoplasm
Q6G0N3	Y212_BARQU		inner membrane
Q6G0N8	TRMB_BARQU		cytoplasm
Q6G0N9	Y206_BARQU		inner membrane
Q6G0P3	RBFA_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0P4	TRUB_BARQU		cytoplasm
Q6G0P6	RS15_BARQU		cytoplasm
Q6G0P9	HSLU_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0Q0	HSLV_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0Q6	ARGJ_BARQU	cytoplasm (probable).	cytoplasm
Q6G0S2	NUDH_BARQU		cytoplasm
Q6G0S6	PROA_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0S7	PROB_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0S9	RRMJ_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0T7	RL27_BARQU		cytoplasm
Q6G0T9	G6PI_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0U2	RS21_BARQU		cytoplasm
Q6G0U8	GPDA_BARQU	cytoplasm (probable).	cytoplasm
Q6G0V6	DNAA_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0V7	RS20_BARQU		cytoplasm
Q6G0V9	CCMA_BARQU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6G0W9	SECB_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G0X1	MUTS_BARQU		cytoplasm
Q6G0X4	KCY_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G104	ISPG_BARQU		cytoplasm
Q6G107	RL32_BARQU		cytoplasm
Q6G116	MRAW_BARQU		cytoplasm
Q6G121	MRAY_BARQU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6G122	MURD_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G125	MURC_BARQU	cytoplasm (probable).	cytoplasm
Q6G126	MURB_BARQU	cytoplasm (probable).	inner membrane
Q6G147	CCME_BARQU	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q6G160	MSCL_BARQU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6G164	ISPDF_BARQU		cytoplasm
Q6G173	ENO_BARQU	cytoplasm (by similarity).	extracellular
Q6G177	CLPX_BARQU		cytoplasm
Q6G178	CLPP_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G197	GUAA_BARQU		cytoplasm
Q6G1A4	PURA_BARQU	cytoplasm (by similarity).	cytoplasm

Q6G1B3	COAE_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G1B4	Y001_BARQU		cytoplasm
Q6G1D6	SAHH_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G1D9	CBIO_BARQU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6G1E4	GRPE_BARQU	cytoplasm (probable).	cytoplasm
Q6G1E5	HRC_A_BARQU		cytoplasm
Q6G1E6	HAM1_BARQU		extracellular
Q6G1F4	DAPD_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G1F5	LEPA_BARQU		cytoplasm
Q6G1F7	ARGB_BARQU	cytoplasm (probable).	cytoplasm
Q6G1F8	DNAJ_BARQU	cytoplasm (probable).	cytoplasm
Q6G1F9	DNAK_BARQU		cytoplasm
Q6G1G3	RL35_BARQU		cytoplasm
Q6G1G8	FMT_BARQU		cytoplasm
Q6G1G9	TRUA_BARQU		cytoplasm
Q6G1H4	AROB_BARQU	cytoplasm (probable).	cytoplasm
Q6G1H5	AROK_BARQU	cytoplasm (probable).	cytoplasm
Q6G1I2	UBIE_BARQU		cytoplasm
Q6G1J4	LPXD_BARQU		cytoplasm
Q6G1J5	FABZ_BARQU	cytoplasm (by similarity).	cytoplasm
Q6G1K9	GIDA_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G1L0	GIDB_BARHE		cytoplasm
Q6G1M0	MDH_BARHE		cytoplasm
Q6G1N8	AROB_BARHE	cytoplasm (probable).	cytoplasm
Q6G1N9	AROK_BARHE	cytoplasm (probable).	cytoplasm
Q6G1Q5	Y1619_BARHE		inner membrane
Q6G1R8	RIMM_BARHE	cytoplasm (potential).	cytoplasm
Q6G1R9	TRMD_BARHE	cytoplasm (potential).	cytoplasm
Q6G1S0	RL19_BARHE		cytoplasm
Q6G1W7	ATPA_BARHE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6G1W8	ATPG_BARHE		inner membrane
Q6G1X0	ATPE_BARHE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6G1X4	ISPG_BARHE		cytoplasm
Q6G1X6	RL32_BARHE		cytoplasm
Q6G1Y5	Y1514_BARHE		inner membrane
Q6G1Z4	RL31_BARHE		cytoplasm
Q6G202	DUT_BARHE		extracellular
Q6G226	MURA_BARHE	cytoplasm (probable).	inner membrane
Q6G2F1	GCSP_BARHE		cytoplasm
Q6G2G3	DAPB_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G2G7	RNPA_BARHE		extracellular
Q6G2L1	PTH_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G2L2	RL25_BARHE		cytoplasm
Q6G2L7	LGT_BARHE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6G2P7	MRAW_BARHE		cytoplasm
Q6G2Q2	MRAY_BARHE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q6G2Q3	MURD_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G2Q5	MURG_BARHE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6G2Q6	MURC_BARHE	cytoplasm (probable).	cytoplasm
Q6G2Q7	MURB_BARHE	cytoplasm (probable).	inner membrane
Q6G2S8	TYSY_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G2T3	MIAA_BARHE		cytoplasm
Q6G2T6	ILVC_BARHE		cytoplasm
Q6G2V3	SYV_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G2W1	RS7_BARHE		cytoplasm
Q6G2W4	RS10_BARHE		cytoplasm
Q6G2W8	RL2_BARHE		cytoplasm
Q6G2W9	RS19_BARHE		cytoplasm
Q6G2X1	RS3_BARHE		cytoplasm
Q6G2X2	RL16_BARHE		cytoplasm
Q6G2X3	RL29_BARHE		cytoplasm
Q6G2X4	RS17_BARHE		cytoplasm
Q6G2X9	RS8_BARHE		cytoplasm
Q6G2Y1	RL18_BARHE		cytoplasm
Q6G2Y2	RS5_BARHE		cytoplasm
Q6G2Y4	RL15_BARHE		cytoplasm
Q6G2Y6	KAD_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G2Y7	RS13_BARHE		cytoplasm
Q6G2Y8	RS11_BARHE		cytoplasm
Q6G2Z3	RECA_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G304	COAD_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G321	GLMU_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G322	GLMS_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G335	PURL_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G336	PURQ_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G3B6	SYV_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G3D7	SFSA_BARHE		cytoplasm
Q6G3F1	RUVX_BARHE	cytoplasm (potential).	cytoplasm
Q6G3F3	PYRB_BARHE		cytoplasm
Q6G3F5	Y818_BARHE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6G3I3	LIPB_BARHE	cytoplasm (potential).	cytoplasm
Q6G3K5	PLSX_BARHE		cytoplasm
Q6G3L2	NRDR_BARHE		cytoplasm
Q6G3L3	GLYA_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G3L8	HFQ_BARHE		cytoplasm
Q6G3U8	HUTH_BARHE	cytoplasm (potential).	cytoplasm
Q6G3V6	RPIA_BARHE		cytoplasm
Q6G3X4	RPOC_BARHE		cytoplasm
Q6G3X7	RL10_BARHE		cytoplasm
Q6G3X8	RL1_BARHE		cytoplasm
Q6G3X9	RL11_BARHE		cytoplasm
Q6G3Y8	TIG_BARHE		cytoplasm
Q6G3Y9	GID_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G3Z2	CLPX_BARHE		cytoplasm
Q6G3Z3	CLPP_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G3Z8	ISPDF_BARHE		cytoplasm
Q6G409	KDSA_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G426	NDK_BARHE	cytoplasm (by similarity).	cytoplasm

Q6G438	KSGA_BARHE		cytoplasm
Q6G439	KGUA_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G442	ACP_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G446	RS6_BARHE		cytoplasm
Q6G447	RS18_BARHE		cytoplasm
Q6G449	RL9_BARHE		cytoplasm
Q6G456	PANC_BARHE		cytoplasm
Q6G457	RECO_BARHE		cytoplasm
Q6G459	RNC_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G461	ACPS_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G465	RPOZ_BARHE		cytoplasm
Q6G467	SSRP_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G488	GLNE_BARHE		cytoplasm
Q6G494	CCME_BARHE	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q6G4C5	ISPH_BARHE		cytoplasm
Q6G4D2	EX7S_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G4D4	AROC_BARHE		cytoplasm
Q6G4D7	PDXH_BARHE		cytoplasm
Q6G4E3	RNH2_BARHE	cytoplasm (potential).	extracellular
Q6G4E4	ISPE_BARHE		cytoplasm
Q6G4F8	MSCL_BARHE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6G4G5	PURA_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G4Q8	CBIO_BARHE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6G4R2	LLDD_BARHE		cytoplasm
Q6G4R6	LPXK_BARHE		cytoplasm
Q6G4S5	SYI_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G4T2	FPG_BARHE		cytoplasm
Q6G4T6	PSTB_BARHE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6G4V2	RECR_BARHE		cytoplasm
Q6G4V3	Y229_BARHE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6G4V8	Y224_BARHE		inner membrane
Q6G4W3	TRMB_BARHE		cytoplasm
Q6G4W4	Y218_BARHE		inner membrane
Q6G4W8	RBFA_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G4Y4	NUDH_BARHE		cytoplasm
Q6G4Y7	Y161_BARHE		inner membrane
Q6G4Z0	PROA_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G4Z1	PROB_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G4Z3	RRMJ_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G507	RL27_BARHE		cytoplasm
Q6G512	RS21_BARHE		cytoplasm
Q6G518	GPDA_BARHE	cytoplasm (probable).	cytoplasm
Q6G523	AROQ_BARHE		cytoplasm
Q6G526	DNAA_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G527	RS20_BARHE		cytoplasm

Q6G529	CCMA_BARHE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6G540	SECB_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G542	MUTS_BARHE		cytoplasm
Q6G543	ILVD_BARHE		cytoplasm
Q6G546	KCY_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G547	TRUA_BARHE		cytoplasm
Q6G549	DAPD_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G550	LEPA_BARHE		cytoplasm
Q6G552	ARGB_BARHE	cytoplasm (probable).	cytoplasm
Q6G553	DNAJ_BARHE	cytoplasm (probable).	cytoplasm
Q6G554	DNAK_BARHE		cytoplasm
Q6G563	GRPE_BARHE	cytoplasm (probable).	cytoplasm
Q6G564	HRCB_BARHE		cytoplasm
Q6G565	HAM1_BARHE		extracellular
Q6G577	UBIE_BARHE		cytoplasm
Q6G584	SAHH_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G5A8	COAE_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G5B1	Y001_BARHE		cytoplasm
Q6G5C6	RRF_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G5C9	RS2_BARHE		cytoplasm
Q6G5D4	CLPS_BARHE		cytoplasm
Q6G5D6	KTHY_BARHE		cytoplasm
Q6G5D9	LEXA_BARHE		cytoplasm
Q6G5E2	RS4_BARHE		cytoplasm
Q6G5E5	RL35_BARHE		cytoplasm
Q6G5F1	FMT_BARHE		cytoplasm
Q6G5F5	TRUB_BARHE		cytoplasm
Q6G5F7	RS15_BARHE		cytoplasm
Q6G5G0	HSLU_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G5G1	HSLV_BARHE	cytoplasm (by similarity).	cytoplasm
Q6G5G6	PSD_BARHE		cytoplasm
Q6G5H3	PANB_BARHE		cytoplasm
Q6G5J3	GUAA_BARHE		cytoplasm
Q6G5K3	UBIG_BARHE		cytoplasm
Q6G5N5	Y1276_BARHE		inner membrane
Q6G5P2	Y1481_BARHE		inner membrane
Q6G5P5	TILS_BARHE	cytoplasm (probable).	cytoplasm
Q6G5P7	GLMM_BARHE		cytoplasm
Q6G5Q9	RUVC_BARHE		cytoplasm
Q6G5R0	RUVA_BARHE		cytoplasm
Q6G5R1	RUVB_BARHE		cytoplasm
Q6G5R6	TOLB_BARHE	periplasm (potential).	periplasm
Q6G5T1	RL28_BARHE		cytoplasm
Q6G5T6	ARGJ_BARHE	cytoplasm (probable).	cytoplasm
Q6IT00	GRPE_VIBHA	cytoplasm (probable).	cytoplasm
Q6J171	UREE_EDWIC	cytoplasm (by similarity).	cytoplasm
Q6J256	1A1D_VARPD		cytoplasm
Q6LBA6	RBL_OLICA		extracellular
Q6LFR3	Y6022_PHOPR		cytoplasm
Q6LFS0	PDXH_PHOPR		cytoplasm
Q6LG09	OTC_PHOPR	cytoplasm (probable).	cytoplasm
Q6LG16	AGUA_PHOPR		extracellular
Q6LG56	UXUA_PHOPR		cytoplasm
Q6LGY8	Y5582_PHOPR		inner membrane

Q6LHF5	SDHD_PHOPR		cytoplasm
Q6LHK5	MAO12_PHOPR		cytoplasm
Q6LHL2	MODC_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LHN1	GLGB_PHOPR		extracellular
Q6LHN5	GCSP_PHOPR		cytoplasm
Q6LHN7	GLYA2_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LI45	PHS_PHOPR		cytoplasm
Q6LIL7	COBQ_PHOPR		cytoplasm
Q6LIN8	Y4970_PHOPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6LJ30	NAGZ_PHOPR	cytoplasm (by similarity).	extracellular
Q6LK12	Y4495_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LKA1	GLGA_PHOPR		cytoplasm
Q6LKA2	GLGC_PHOPR		cytoplasm
Q6LKD4	FBPC2_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LKG5	GBPA_PHOPR	secreted protein (by similarity).	cytoplasm
Q6LKQ1	Y4184_PHOPR		inner membrane
Q6LKZ7	ATPG2_PHOPR		inner membrane
Q6LKZ8	ATPA2_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LLA7	DEOD2_PHOPR		cytoplasm
Q6LLD4	ACKA2_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LLF0	TAL_PHOPR	cytoplasm (probable).	cytoplasm
Q6LLF7	GIDA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LLF8	GIDB_PHOPR		cytoplasm
Q6LLG6	ATPA1_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LLG7	ATPG1_PHOPR		inner membrane
Q6LLH1	GLMU_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LLH3	GLMS_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LLH7	ILVD_PHOPR		cytoplasm
Q6LLJ2	FMT_PHOPR		cytoplasm
Q6LLJ6	SMG_PHOPR		cytoplasm
Q6LLK0	HEM6_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LLL4	GPPA_PHOPR		extracellular
Q6LLL5	RHLB_PHOPR		cytoplasm
Q6LLM4	Y3533_PHOPR		inner membrane
Q6LLN3	CYAY_PHOPR		inner membrane
Q6LLQ8	Y3499_PHOPR		inner membrane
Q6LLR8	RBN_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LLR9	DTD_PHOPR	cytoplasm (probable).	cytoplasm
Q6LLS2	PPCK_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LLS3	HSLO_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LLU2	TRMA_PHOPR		cytoplasm

Q6LLU3	BTUB_PHOPR	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q6LLV2	MURB_PHOPR	cytoplasm (probable).	inner membrane
Q6LLV9	RL1_PHOPR		cytoplasm
Q6LLW0	RL10_PHOPR		cytoplasm
Q6LLW2	RPOB_PHOPR		cytoplasm
Q6LLW3	RPOC_PHOPR		cytoplasm
Q6LLX8	Y3416_PHOPR		inner membrane
Q6LLY3	FIS_PHOPR		cytoplasm
Q6LLY5	PRMA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LLZ0	AROQ_PHOPR		cytoplasm
Q6LM05	CH10_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LM06	CH60_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LM07	RIMK_PHOPR		cytoplasm
Q6LM21	PSD_PHOPR		cytoplasm
Q6LM23	ORN_PHOPR	cytoplasm (potential).	cytoplasm
Q6LM29	MIAA_PHOPR		cytoplasm
Q6LM35	PURA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LM37	HMP_PHOPR		cytoplasm
Q6LM40	RLMB_PHOPR		cytoplasm
Q6LM41	RS6_PHOPR		cytoplasm
Q6LM42	PRIB_PHOPR		cytoplasm
Q6LM43	RS18_PHOPR		cytoplasm
Q6LM44	RL9_PHOPR		cytoplasm
Q6LM51	G6PI_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LM58	CYSJ_PHOPR		cytoplasm
Q6LM59	CYSI_PHOPR		cytoplasm
Q6LMB3	Y3258_PHOPR		cytoplasm
Q6LMC9	MURA_PHOPR	cytoplasm (probable).	inner membrane
Q6LMF3	MRAY_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LMF4	MURD_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LMF6	MURG_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LMF7	MURC_PHOPR	cytoplasm (probable).	cytoplasm
Q6LMG1	LPXC_PHOPR		cytoplasm
Q6LMG5	Y3206_PHOPR		cytoplasm
Q6LMG6	Y3205_PHOPR		inner membrane
Q6LMG7	COAE_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LMI4	Y3187_PHOPR		inner membrane
Q6LMJ5	PANC_PHOPR		cytoplasm
Q6LMJ6	PANB_PHOPR		cytoplasm
Q6LMJ9	GLUQ_PHOPR		cytoplasm
Q6LMK1	SFSA_PHOPR		cytoplasm
Q6LMK7	FETP_PHOPR		cytoplasm
Q6LML0	TRMB_PHOPR		cytoplasm
Q6LML4	HAM1_PHOPR		extracellular
Q6LMM2	RUVX_PHOPR	cytoplasm (potential).	cytoplasm
Q6LMP0	RPIA_PHOPR		cytoplasm
Q6LMR2	Y3099_PHOPR		cytoplasm
Q6LMS0	LEPA_PHOPR		cytoplasm
Q6LMS2	RNC_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LMS4	RECO_PHOPR		cytoplasm

Q6LMS5	ACPS_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LMS7	RUMA_PHOPR		cytoplasm
Q6LMT1	ENO_PHOPR	cytoplasm (by similarity).	extracellular
Q6LMT3	ISPD_PHOPR		cytoplasm
Q6LMT4	ISPF_PHOPR		extracellular
Q6LMT5	TRUD_PHOPR		cytoplasm
Q6LMT6	SURE_PHOPR	cytoplasm (potential).	periplasm
Q6LMT7	PIMT_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LMU0	MUTS_PHOPR		cytoplasm
Q6LMU2	RECA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LMU5	CSRA_PHOPR		cytoplasm
Q6LMV1	GSH1_PHOPR		cytoplasm
Q6LMV3	LUXS_PHOPR		extracellular
Q6LMV8	RIMM_PHOPR	cytoplasm (potential).	cytoplasm
Q6LMV9	TRMD_PHOPR	cytoplasm (potential).	cytoplasm
Q6LMW0	RL19_PHOPR		cytoplasm
Q6LMY0	CLPB_PHOPR	cytoplasm (probable).	cytoplasm
Q6LN05	MTFA_PHOPR		cytoplasm
Q6LN10	QUEF_PHOPR	cytoplasm (probable).	cytoplasm
Q6LN21	Y2971_PHOPR		inner membrane
Q6LN22	GLND_PHOPR		cytoplasm
Q6LN24	RS2_PHOPR		cytoplasm
Q6LN27	RRF_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LN28	UPPS_PHOPR		cytoplasm
Q6LN30	DXR_PHOPR		cytoplasm
Q6LN34	LPXD_PHOPR		cytoplasm
Q6LN35	FABZ_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LN38	RNH2_PHOPR	cytoplasm (potential).	extracellular
Q6LN40	ACCA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LN41	TILS_PHOPR	cytoplasm (probable).	cytoplasm
Q6LN45	Y2947_PHOPR		inner membrane
Q6LN74	UPP_PHOPR		cytoplasm
Q6LN85	DAPA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LN88	LIPB_PHOPR	cytoplasm (potential).	cytoplasm
Q6LN94	Y2889_PHOPR		inner membrane
Q6LNA3	Y2880_PHOPR		inner membrane
Q6LNA9	PTH_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LNB1	ISPE_PHOPR		cytoplasm
Q6LNB3	HEM1_PHOPR		cytoplasm
Q6LNB8	KDSA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LNF6	ACKA1_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LNF7	Y2797_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LNJ0	FTHS_PHOPR		cytoplasm
Q6LNN4	AROC_PHOPR		cytoplasm
Q6LNU2	PDXB_PHOPR		cytoplasm
Q6LNU5	TRUA_PHOPR		cytoplasm
Q6LNV9	TIG_PHOPR		cytoplasm
Q6LNW0	CLPP_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LNW1	CLPX_PHOPR		cytoplasm
Q6LNX0	Y2627_PHOPR		cytoplasm
Q6LP05	SELD_PHOPR		cytoplasm
Q6LP06	SELU_PHOPR		cytoplasm
Q6LP11	Y2586_PHOPR		inner membrane

Q6LP12	NDPA_PHOPR	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q6LP25	RNT_PHOPR		cytoplasm
Q6LP47	TOLB_PHOPR	periplasm (potential).	periplasm
Q6LP67	Y2527_PHOPR		cytoplasm
Q6LPA3	TRPA_PHOPR		cytoplasm
Q6LPA6	TRPD_PHOPR		cytoplasm
Q6LPD7	UBIG_PHOPR		cytoplasm
Q6LPD9	SERC_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LPE2	KCY_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LPF8	Y2435_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LPG9	Y2423_PHOPR		cytoplasm
Q6LPH2	Y2420_PHOPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6LPK7	LPXK_PHOPR		cytoplasm
Q6LPV5	NANE_PHOPR		cytoplasm
Q6LPX9	ALLC_PHOPR		extracellular
Q6LQ00	CBIO_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LQ03	UVRC_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LQ49	NORM_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LQ56	HUTH_PHOPR	cytoplasm (potential).	cytoplasm
Q6LQ71	RL20_PHOPR		cytoplasm
Q6LQ76	BTUC_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LQ77	BTUD_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LQ84	Y2144_PHOPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6LQG2	RIHA_PHOPR		extracellular
Q6LQJ3	NAPA2_PHOPR	periplasm (by similarity).	periplasm
Q6LQM6	MAO11_PHOPR		cytoplasm
Q6LQV7	EFPL_PHOPR		cytoplasm
Q6LR20	FBPC1_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LR24	PEPT_PHOPR	cytoplasm (by similarity).	extracellular
Q6LR94	Y1775_PHOPR		inner membrane
Q6LR96	FABA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LRA2	PYRD_PHOPR	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LRA6	PNCB_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LRB9	Y1750_PHOPR		cytoplasm
Q6LRD1	Y1738_PHOPR		inner membrane
Q6LRD9	TDH_PHOPR	cytoplasm (by similarity).	cytoplasm

Q6LRI0	CDD_PHOPR		cytoplasm
Q6LRI5	TPMT_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LRY1	RL25_PHOPR		cytoplasm
Q6LS21	TORD_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LS31	DNAK2_PHOPR		cytoplasm
Q6LSD6	METE_PHOPR		cytoplasm
Q6LSS4	Y1241_PHOPR		inner membrane
Q6LST5	MGSA_PHOPR		cytoplasm
Q6LSW5	KTHY_PHOPR		cytoplasm
Q6LSX3	PLSX_PHOPR		cytoplasm
Q6LSX4	RL32_PHOPR		cytoplasm
Q6LT10	LFTR_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LT14	CLPS_PHOPR		cytoplasm
Q6LT47	UVRB_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LT48	RUVB_PHOPR		cytoplasm
Q6LT49	RUVA_PHOPR		cytoplasm
Q6LT51	RUVC_PHOPR		cytoplasm
Q6LT52	Y1113_PHOPR		inner membrane
Q6LT75	HIS8_PHOPR		cytoplasm
Q6LT81	KITH_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LTE0	HEMH_PHOPR	cytoplasm (by similarity).	extracellular
Q6LTE1	KAD_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LTE2	HTPG_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LTE6	RECR_PHOPR		cytoplasm
Q6LTK3	FADJ_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LTK4	FADI_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LTL4	CCME_PHOPR	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q6LTL7	CCMA_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LTQ6	FLGI2_PHOPR	periplasm (by similarity).	periplasm
Q6LTQ7	FLGH_PHOPR	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q6LTU6	ZIPA_PHOPR	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q6LTV9	NAPA1_PHOPR	periplasm (by similarity).	periplasm
Q6LTX2	PROA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LTX3	PROB_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LTX5	Y837_PHOPR		cytoplasm
Q6LTX6	XGPT_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LU05	EX7S_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LU16	NRDR_PHOPR		cytoplasm
Q6LU17	GLYA1_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LU31	GUAA_PHOPR		cytoplasm
Q6LU49	ISPG_PHOPR		cytoplasm
Q6LU53	NDK_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LU58	HSCA_PHOPR		cytoplasm
Q6LU59	HSCB_PHOPR		inner membrane
Q6LU69	TGT_PHOPR		cytoplasm
Q6LU70	QUEA_PHOPR	cytoplasm (potential).	cytoplasm
Q6LU78	CUTC_PHOPR	cytoplasm (probable).	periplasm

Q6LU82	PSTB_PHOPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6LUA6	DNAJ_PHOPR	cytoplasm (probable).	cytoplasm
Q6LUA7	DNAK1_PHOPR		cytoplasm
Q6LUA8	GRPE_PHOPR	cytoplasm (probable).	cytoplasm
Q6LUA9	PPNK_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUB4	SSRP_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUG1	Y643_PHOPR		inner membrane
Q6LUG2	TRPR_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUH1	DEOD1_PHOPR		cytoplasm
Q6LUH3	TYPH_PHOPR		cytoplasm
Q6LUI9	RS15_PHOPR		cytoplasm
Q6LUJ0	TRUB_PHOPR		cytoplasm
Q6LUJ4	Y610_PHOPR		inner membrane
Q6LUJ6	GLMM_PHOPR		cytoplasm
Q6LUJ9	RRMJ_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUK6	CARA_PHOPR		cytoplasm
Q6LUK7	DAPB_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUK8	ISPH_PHOPR		cytoplasm
Q6LUL1	SYI_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUL5	RS20_PHOPR		cytoplasm
Q6LUM1	TYSY_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUM2	LGT_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LUM5	NUDH_PHOPR		cytoplasm
Q6LUP5	END4_PHOPR		extracellular
Q6LUP6	GRCA_PHOPR		cytoplasm
Q6LUP9	KHSE_PHOPR	cytoplasm (potential).	cytoplasm
Q6LUQ3	RRAA2_PHOPR		cytoplasm
Q6LUR6	BTUF_PHOPR	periplasm (by similarity).	periplasm
Q6LUR9	SYI_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUS3	GSA_PHOPR	cytoplasm (potential).	cytoplasm
Q6LUW0	AMPA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUW1	SYV_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUW9	OTCC_PHOPR	cytoplasm (probable).	cytoplasm
Q6LUX0	PYRB_PHOPR		cytoplasm
Q6LUX1	PYRI_PHOPR		cytoplasm
Q6LUX5	IDI_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUY7	XYLA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LUZ5	HLDE_PHOPR		cytoplasm
Q6LUZ6	GLNE_PHOPR		cytoplasm
Q6LV05	CCA_PHOPR		cytoplasm
Q6LV09	Y435_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LV25	LEU3_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LV26	LEU2_PHOPR		cytoplasm
Q6LV27	LEUD_PHOPR		cytoplasm
Q6LV34	RAPA_PHOPR		cytoplasm
Q6LV37	DJLA_PHOPR	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q6LV41	KSGA_PHOPR		cytoplasm
Q6LV42	APAG_PHOPR		inner membrane

Q6LV43	APAH_PHOPR		extracellular
Q6LV48	RL27_PHOPR		cytoplasm
Q6LV53	ARGR_PHOPR	cytoplasm (probable).	cytoplasm
Q6LV65	MSRA_PHOPR		cytoplasm
Q6LV74	META_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LV91	RPOA_PHOPR		cytoplasm
Q6LV92	RS4_PHOPR		cytoplasm
Q6LV93	RS11_PHOPR		cytoplasm
Q6LV94	RS13_PHOPR		cytoplasm
Q6LV97	RL15_PHOPR		cytoplasm
Q6LV99	RS5_PHOPR		cytoplasm
Q6LVA0	RL18_PHOPR		cytoplasm
Q6LVA2	RS8_PHOPR		cytoplasm
Q6LVA4	RL5_PHOPR		cytoplasm
Q6LVA7	RS17_PHOPR		cytoplasm
Q6LVA8	RL29_PHOPR		cytoplasm
Q6LVA9	RL16_PHOPR		cytoplasm
Q6LVB0	RS3_PHOPR		cytoplasm
Q6LVB1	RL22_PHOPR		cytoplasm
Q6LVB2	RS19_PHOPR		cytoplasm
Q6LVB3	RL2_PHOPR		cytoplasm
Q6LVB7	RS10_PHOPR		cytoplasm
Q6LVC2	RS7_PHOPR		cytoplasm
Q6LVC3	RS12_PHOPR		cytoplasm
Q6LVC6	TUSD_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LVD0	SLYX_PHOPR		inner membrane
Q6LVF1	GPH_PHOPR		cytoplasm
Q6LVF5	AROB_PHOPR	cytoplasm (probable).	cytoplasm
Q6LVF6	AROK_PHOPR	cytoplasm (probable).	cytoplasm
Q6LVG8	ASSY_PHOPR	cytoplasm (probable).	cytoplasm
Q6LVG9	ARGB_PHOPR	cytoplasm (probable).	cytoplasm
Q6LVH1	CAPP_PHOPR		cytoplasm
Q6LVH8	RL31_PHOPR		cytoplasm
Q6LVI2	HSLV_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LVI3	HSLU_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LVI5	RRAA1_PHOPR		cytoplasm
Q6LVK9	GPDA_PHOPR	cytoplasm (probable).	cytoplasm
Q6LVL0	SECB_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LVL2	GPMI_PHOPR		cytoplasm
Q6LVM8	COAD_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LVN0	FPG_PHOPR		cytoplasm
Q6LVN3	RL28_PHOPR		cytoplasm
Q6LVN4	RADC_PHOPR		cytoplasm
Q6LVN6	SLMA_PHOPR	nucleoid-associated (by similarity).	cytoplasm
Q6LVN7	PYRE_PHOPR		cytoplasm
Q6LVN8	RNPH_PHOPR		cytoplasm
Q6LVP5	KGUA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LVP6	RPOZ_PHOPR		cytoplasm
Q6LVQ7	BIOH_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LVQ9	Y177_PHOPR		inner membrane
Q6LVS1	LEXA_PHOPR		cytoplasm
Q6LVT0	GLPE_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LVW8	UBIB_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q6LVX7	THIG_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LVY3	CRCB_PHOPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6LVZ5	ILVC_PHOPR		cytoplasm
Q6LW06	FADB_PHOPR		cytoplasm
Q6LW07	FADA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LW10	TUSA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LW50	DNAA_PHOPR	cytoplasm (by similarity).	cytoplasm
Q6LW54	RNPA_PHOPR		extracellular
Q6LW60	FLGI1_PHOPR	periplasm (by similarity).	periplasm
Q6MGL6	GIDA_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MGM5	ATPA_BDEBA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6MGM6	ATPG_BDEBA		inner membrane
Q6MGM8	ATPE_BDEBA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6MGQ3	GRPE_BDEBA	cytoplasm (probable).	cytoplasm
Q6MGR8	DEOC_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MGS4	RL27_BDEBA		cytoplasm
Q6MGS7	Y3845_BDEBA		inner membrane
Q6MGU3	Y3828_BDEBA		cytoplasm
Q6MGU4	NDK_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MH01	TIG_BDEBA		cytoplasm
Q6MH10	AMPA_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MH11	CLPP_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MH12	CLPX_BDEBA		cytoplasm
Q6MH30	RECR_BDEBA		cytoplasm
Q6MHH9	PAND_BDEBA		cytoplasm
Q6MHI3	PANB_BDEBA		cytoplasm
Q6MHQ3	UBIE_BDEBA		cytoplasm
Q6MHT3	HEMH_BDEBA	cytoplasm (by similarity).	extracellular
Q6MHV9	GLMU_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MHW4	KITH_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MI44	Y3330_BDEBA		cytoplasm
Q6MID2	TYSY_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MIF7	MRAY_BDEBA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6MIF8	MURD_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MIG1	MURG_BDEBA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6MII5	PPNK_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MIK8	COAE_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MIP7	PHNC_BDEBA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6MIV0	CLPB_BDEBA	cytoplasm (probable).	cytoplasm
Q6MJ04	RL1_BDEBA		cytoplasm
Q6MJ05	RL10_BDEBA		cytoplasm
Q6MJ09	RPOB_BDEBA		cytoplasm
Q6MJ10	RPOC_BDEBA		cytoplasm
Q6MJ11	RS12_BDEBA		cytoplasm

Q6MJ12	RS7_BDEBA		cytoplasm
Q6MJ14	RS10_BDEBA		cytoplasm
Q6MJ17	RL2_BDEBA		cytoplasm
Q6MJ18	RS19_BDEBA		cytoplasm
Q6MJ19	RL22_BDEBA		cytoplasm
Q6MJ20	RS3_BDEBA		cytoplasm
Q6MJ21	RL16_BDEBA		cytoplasm
Q6MJ22	RL29_BDEBA		cytoplasm
Q6MJ23	RS17_BDEBA		cytoplasm
Q6MJ25	RL5_BDEBA		cytoplasm
Q6MJ28	RL18_BDEBA		cytoplasm
Q6MJ29	RS5_BDEBA		cytoplasm
Q6MJ31	RL15_BDEBA		cytoplasm
Q6MJ33	RS13_BDEBA		cytoplasm
Q6MJ34	RS11_BDEBA		cytoplasm
Q6MJ35	RS4_BDEBA		cytoplasm
Q6MJ36	RPOA_BDEBA		cytoplasm
Q6MJE4	MAO1_BDEBA		cytoplasm
Q6MJH6	SP5G_BDEBA		inner membrane
Q6MJJ2	NPD_BDEBA	cytoplasm (probable).	cytoplasm
Q6MJL7	FMT_BDEBA		cytoplasm
Q6MJP3	GPMA_BDEBA		cytoplasm
Q6MJR3	PTH_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MJR4	RL25_BDEBA		cytoplasm
Q6MJR8	HAM1_BDEBA		extracellular
Q6MJR9	RNPH_BDEBA		cytoplasm
Q6MJY0	ADEC_BDEBA		cytoplasm
Q6MK42	SELD_BDEBA		cytoplasm
Q6MK43	SELU_BDEBA		cytoplasm
Q6MK45	PDXH_BDEBA		cytoplasm
Q6MK56	APAG_BDEBA		inner membrane
Q6MKB6	RUVC_BDEBA		cytoplasm
Q6MKB7	RUVA_BDEBA		cytoplasm
Q6MKE9	Y2446_BDEBA		inner membrane
Q6MKS2	UVRC_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MKW4	CAAL_BDEBA		cytoplasm
Q6MKX0	SYI_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6ML45	FPG_BDEBA		cytoplasm
Q6ML86	Y2131_BDEBA		inner membrane
Q6ML96	RIMM_BDEBA	cytoplasm (potential).	cytoplasm
Q6ML97	TRMD_BDEBA	cytoplasm (potential).	cytoplasm
Q6ML99	RL19_BDEBA		cytoplasm
Q6MLA0	RNH2_BDEBA	cytoplasm (potential).	extracellular
Q6MLD2	GUAA_BDEBA		cytoplasm
Q6MLF5	Y2056_BDEBA		cytoplasm
Q6MLJ4	RL32_BDEBA		cytoplasm
Q6MLJ8	ACP_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MLK1	GLYA_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MLL0	KUP_BDEBA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6MLR5	RNC_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MLS4	GLMM_BDEBA		cytoplasm
Q6MLS8	TILS_BDEBA	cytoplasm (probable).	cytoplasm

Q6MMH0	PSTB_BDEBA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6MMJ3	GIDB_BDEBA		cytoplasm
Q6MMK2	IDI2_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MMK3	RL20_BDEBA		cytoplasm
Q6MMK4	RL35_BDEBA		cytoplasm
Q6MMK8	DTD_BDEBA	cytoplasm (probable).	cytoplasm
Q6MMQ7	RPOZ_BDEBA		cytoplasm
Q6MMQ9	MIAA_BDEBA		cytoplasm
Q6MMS3	RS15_BDEBA		cytoplasm
Q6MMS4	TRUB_BDEBA		cytoplasm
Q6MMS5	RBFA_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MMS8	Y1545_BDEBA		inner membrane
Q6MMY3	Y1487_BDEBA		inner membrane
Q6MMZ9	RS20_BDEBA		cytoplasm
Q6MN06	PURA_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MN26	Y1437_BDEBA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6MN40	RRMJ_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MNC0	SAHH_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MNF8	DNAK_BDEBA		cytoplasm
Q6MNG0	DNAJ_BDEBA	cytoplasm (probable).	cytoplasm
Q6MNG7	CLPS_BDEBA		cytoplasm
Q6MNN2	Y1212_BDEBA		inner membrane
Q6MNP6	NRDJ_BDEBA		cytoplasm
Q6MNQ2	SYI_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MNX8	NRDR_BDEBA		cytoplasm
Q6MNY2	Y1093_BDEBA		cytoplasm
Q6MP09	KCY_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MP28	KTHY_BDEBA		cytoplasm
Q6MPQ2	ENO_BDEBA	cytoplasm (by similarity).	extracellular
Q6MPS6	LIPB_BDEBA	cytoplasm (potential).	cytoplasm
Q6MPZ6	GCSP_BDEBA		cytoplasm
Q6MQ03	GCST_BDEBA		cytoplasm
Q6MQ46	SSRP_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MQ47	KSGA_BDEBA		cytoplasm
Q6MQ60	COAD_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MQB7	Y559_BDEBA		inner membrane
Q6MQD7	FLGI_BDEBA	periplasm (by similarity).	periplasm
Q6MQH6	TRUA_BDEBA		cytoplasm
Q6MQK8	SYV_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MQS6	DNAE2_BDEBA	cytoplasm (probable).	cytoplasm
Q6MQU0	TRMB_BDEBA		cytoplasm
Q6MR60	RS21_BDEBA		cytoplasm
Q6MR96	EX7S_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MRB2	TOLB_BDEBA	periplasm (potential).	periplasm
Q6MRC6	MSCL_BDEBA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6MRI1	CH60_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6MRJ2	QUEF_BDEBA	cytoplasm (probable).	cytoplasm
Q6MRK7	MURA_BDEBA	cytoplasm (probable).	inner membrane
Q6MRL0	RL31B_BDEBA		cytoplasm
Q6MRM8	DAPB_BDEBA	cytoplasm (by similarity).	cytoplasm

Q6MRP1	RL9_BDEBA		cytoplasm
Q6MRP3	RS6_BDEBA		cytoplasm
Q6MRS1	DNAA_BDEBA	cytoplasm (by similarity).	cytoplasm
Q6N0C7	RS20_RHOPA		cytoplasm
Q6N0I5	PSTB_RHOPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6N0J0	OTC_RHOPA	cytoplasm (probable).	cytoplasm
Q6N0J2	APAG_RHOPA		inner membrane
Q6N0K6	NIKR_RHOPA		cytoplasm
Q6N0L9	ARLY_RHOPA	cytoplasm (probable).	cytoplasm
Q6N0P7	MODC_RHOPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6N0T9	HGD_RHOPA		cytoplasm
Q6N0W9	RBL2_RHOPA		cytoplasm
Q6N103	NIFW_RHOPA		cytoplasm
Q6N176	MURA_RHOPA	cytoplasm (probable).	inner membrane
Q6N1G8	METX_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N1H2	CLPB_RHOPA	cytoplasm (probable).	cytoplasm
Q6N1I5	SLYX_RHOPA		inner membrane
Q6N1I7	Y4418_RHOPA		periplasm
Q6N1M8	SYI_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N1P8	RL25_RHOPA		cytoplasm
Q6N1P9	PTH_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N1T6	GLMM_RHOPA		cytoplasm
Q6N1V9	PURA_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N1Y1	ISPH2_RHOPA		cytoplasm
Q6N1Y2	KHSE_RHOPA		cytoplasm
Q6N233	Y4217_RHOPA		cytoplasm
Q6N260	Y4190_RHOPA		inner membrane
Q6N274	RS21_RHOPA		cytoplasm
Q6N2N5	SAHH_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N2R5	HLDE_RHOPA		cytoplasm
Q6N2Y8	FLGI_RHOPA	periplasm (by similarity).	periplasm
Q6N2Z5	FLGH_RHOPA	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q6N344	GCSP_RHOPA		cytoplasm
Q6N356	Y3838_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6N376	PURQ_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N3G0	ISPH1_RHOPA		cytoplasm
Q6N3N0	URE3_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N3N1	URE2_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N3N3	URE1_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N3P9	GLGB_RHOPA		extracellular
Q6N3W7	THIG_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N408	MRAY_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6N409	MURD_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N411	MURG_RHOPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6N447	TYSY_RHOPA	cytoplasm (by similarity).	cytoplasm

Q6N453	Y3487_RHOPA		inner membrane
Q6N4R5	RL1_RHOPA		cytoplasm
Q6N4R7	RL10_RHOPA		cytoplasm
Q6N4R8	RL7_RHOPA		cytoplasm
Q6N4R9	RPOB_RHOPA		cytoplasm
Q6N4S0	RPOC_RHOPA		cytoplasm
Q6N4T2	RS12_RHOPA		cytoplasm
Q6N4T3	RS7_RHOPA		cytoplasm
Q6N4T6	RS10_RHOPA		cytoplasm
Q6N4T7	RL23_RHOPA		cytoplasm
Q6N4T8	RS19_RHOPA		cytoplasm
Q6N4T9	RL22_RHOPA		cytoplasm
Q6N4U0	RS3_RHOPA		cytoplasm
Q6N4U1	RL16_RHOPA		cytoplasm
Q6N4U2	RL29_RHOPA		cytoplasm
Q6N4U3	RS17_RHOPA		cytoplasm
Q6N4U4	RL14_RHOPA		cytoplasm
Q6N4U5	RL5_RHOPA		cytoplasm
Q6N4U6	RS14_RHOPA		cytoplasm
Q6N4U7	RS8_RHOPA		cytoplasm
Q6N4U8	RL6_RHOPA		cytoplasm
Q6N4U9	RL18_RHOPA		cytoplasm
Q6N4V0	RS5_RHOPA		cytoplasm
Q6N4V1	RL30_RHOPA		cytoplasm
Q6N4V2	RL15_RHOPA		cytoplasm
Q6N4V5	RS13_RHOPA		cytoplasm
Q6N4V6	RS11_RHOPA		cytoplasm
Q6N4V7	RPOA_RHOPA		cytoplasm
Q6N4V8	RL17_RHOPA		cytoplasm
Q6N514	LIPB_RHOPA	cytoplasm (potential).	cytoplasm
Q6N535	CLPS2_RHOPA		cytoplasm
Q6N554	RL33_RHOPA		cytoplasm
Q6N565	PYRB_RHOPA		cytoplasm
Q6N569	RUVX_RHOPA	cytoplasm (potential).	cytoplasm
Q6N583	PANB_RHOPA		cytoplasm
Q6N5A1	RL9_RHOPA		cytoplasm
Q6N5A3	RS18_RHOPA		cytoplasm
Q6N5A4	RS6_RHOPA		cytoplasm
Q6N5B4	KSGA_RHOPA		cytoplasm
Q6N5B9	AMPA_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N5C3	NDK_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N5F2	KUP1_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6N5G6	KUP2_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6N5H0	ATKA_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q6N5L2	TIG_RHOPA		cytoplasm
Q6N5L3	CLPP_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N5L4	CLPX_RHOPA		cytoplasm
Q6N5Q2	RS2_RHOPA		cytoplasm
Q6N5Q6	DXR_RHOPA		cytoplasm
Q6N5Q9	LPXD_RHOPA		cytoplasm

Q6N5T0	TRPD_RHOPA		cytoplasm
Q6N5U3	OPGH_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6N5U4	OPGG_RHOPA	periplasm (by similarity).	periplasm
Q6N5U5	QUEF_RHOPA	cytoplasm (probable).	cytoplasm
Q6N5U6	ENO_RHOPA	cytoplasm (by similarity).	extracellular
Q6N5X7	SURE_RHOPA	cytoplasm (potential).	periplasm
Q6N5Z1	GID_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N622	GLYA2_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N650	RS9_RHOPA		cytoplasm
Q6N651	RL13_RHOPA		cytoplasm
Q6N678	PLSX_RHOPA		cytoplasm
Q6N692	NRDR_RHOPA		cytoplasm
Q6N693	GLYA1_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N6A7	ATE_RHOPA	cytoplasm (potential).	cytoplasm
Q6N6B8	RECO_RHOPA		cytoplasm
Q6N6C1	RNC_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N6C3	ACPS_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N6C5	RPOZ_RHOPA		cytoplasm
Q6N6C9	SSRP_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N6D0	MSCL_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6N6D6	Y2681_RHOPA		inner membrane
Q6N6F8	GLMU_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N6K3	SSUD_RHOPA		cytoplasm
Q6N6L0	QUEA_RHOPA	cytoplasm (potential).	cytoplasm
Q6N6L4	COAD_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N6M5	ISPDF_RHOPA		cytoplasm
Q6N6N1	SYV_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N6U0	NPD_RHOPA	cytoplasm (probable).	cytoplasm
Q6N703	SYV_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N726	AROQ_RHOPA		cytoplasm
Q6N734	LFTR_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N765	METE_RHOPA		cytoplasm
Q6N7Q9	RRMJ_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N7R5	Y2191_RHOPA		inner membrane
Q6N7S7	XGPT_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N7T7	RBLL1_RHOPA		cytoplasm
Q6N869	ILVC_RHOPA		cytoplasm
Q6N874	MIAA_RHOPA		cytoplasm
Q6N893	KUP3_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6N898	PSD_RHOPA		cytoplasm
Q6N8C3	RPIA_RHOPA		cytoplasm
Q6N8F3	PQQE_RHOPA		cytoplasm
Q6N8F4	PQQD_RHOPA		cytoplasm
Q6N8F5	PQQC_RHOPA		cytoplasm
Q6N8F6	PQQB_RHOPA		cytoplasm
Q6N8F8	FCTA_RHOPA		cytoplasm
Q6N8H0	GLNE_RHOPA		cytoplasm

Q6N8I3	CCME_RHOPA	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q6N8V0	DNAE2_RHOPA	cytoplasm (probable).	cytoplasm
Q6N8Y1	Y1770_RHOPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6N976	PHK_RHOPA		cytoplasm
Q6N9F2	PURL_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6N9G0	RS4_RHOPA		cytoplasm
Q6N9G6	Y1583_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6N9J0	RBL1_RHOPA		cytoplasm
Q6N9J7	ACSF_RHOPA		inner membrane
Q6N9S5	ILVD_RHOPA		cytoplasm
Q6NA00	PHNC2_RHOPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6NAI2	CLPS1_RHOPA		cytoplasm
Q6NAI4	AROC_RHOPA		cytoplasm
Q6NAI9	PDXH_RHOPA		cytoplasm
Q6NAM5	LPXK_RHOPA		cytoplasm
Q6NAQ6	TILS_RHOPA	cytoplasm (probable).	cytoplasm
Q6NAR1	TOLB_RHOPA	periplasm (potential).	periplasm
Q6NAT0	RUVA_RHOPA		cytoplasm
Q6NAT1	RUVC_RHOPA		cytoplasm
Q6NAZ1	ISPE_RHOPA		cytoplasm
Q6NAZ6	RNH2_RHOPA	cytoplasm (potential).	extracellular
Q6NB57	HYP_A_RHOPA		cytoplasm
Q6NB75	EX7S_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NB79	NORM_RHOPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6NB80	Y948_RHOPA		cytoplasm
Q6NBB0	RL31_RHOPA		cytoplasm
Q6NBB9	WRBA_RHOPA		cytoplasm
Q6NBF3	HEMH_RHOPA	cytoplasm (by similarity).	extracellular
Q6NBT1	CYSA_RHOPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6NBX6	PHNC1_RHOPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6NC44	ARGB_RHOPA	cytoplasm (probable).	cytoplasm
Q6NC47	DAPD_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NC50	FMT_RHOPA		cytoplasm
Q6NC55	RECR_RHOPA		cytoplasm
Q6NC69	UBIG_RHOPA		cytoplasm
Q6NCE6	RL32_RHOPA		cytoplasm
Q6NCF3	ISPG_RHOPA		cytoplasm
Q6NCG4	ACCA_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NCG8	AROK_RHOPA	cytoplasm (probable).	cytoplasm
Q6NCG9	AROB_RHOPA	cytoplasm (probable).	cytoplasm
Q6NCH9	RL28_RHOPA		cytoplasm

Q6NCM6	Y446_RHOPA		inner membrane
Q6NCM9	TRMB_RHOPA		cytoplasm
Q6NCN2	Y439_RHOPA		inner membrane
Q6NCN6	RBFA_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NCN7	RS15_RHOPA		cytoplasm
Q6NCT7	GLGA_RHOPA		cytoplasm
Q6NCT8	GLGC_RHOPA		cytoplasm
Q6NCU3	PIMT_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NCX7	GPMI_RHOPA		cytoplasm
Q6NCX8	DAPB_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NCY3	DNAJ_RHOPA	cytoplasm (probable).	cytoplasm
Q6NCY4	DNAK_RHOPA		cytoplasm
Q6NCY6	GRPE_RHOPA	cytoplasm (probable).	cytoplasm
Q6NCY7	HRCR_RHOPA		cytoplasm
Q6NCY8	RNPH_RHOPA		cytoplasm
Q6NCY9	HAM1_RHOPA		extracellular
Q6ND03	HSLU_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6ND07	SECB_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6ND09	COAE_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6ND11	Y298_RHOPA		cytoplasm
Q6ND15	GIDA_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6ND16	GIDB_RHOPA		cytoplasm
Q6ND47	RBLL2_RHOPA		cytoplasm
Q6ND66	RIMM_RHOPA	cytoplasm (potential).	cytoplasm
Q6ND67	TRMD_RHOPA	cytoplasm (potential).	cytoplasm
Q6ND68	RL19_RHOPA		cytoplasm
Q6ND69	LEU2_RHOPA		cytoplasm
Q6ND74	LEUD_RHOPA		cytoplasm
Q6ND82	LEU3_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NDA6	CCMA_RHOPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6NDD0	ATPA_RHOPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6NDD1	ATPG_RHOPA		inner membrane
Q6NDD3	ATPE_RHOPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6NDE1	Y167_RHOPA		inner membrane
Q6NDE4	PROA_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NDE5	PROB_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NDE9	RL27_RHOPA		cytoplasm
Q6NDF0	RL21_RHOPA		cytoplasm
Q6NDM1	FPG_RHOPA		cytoplasm
Q6NDM2	UBIE_RHOPA		cytoplasm
Q6NDN5	TRPA_RHOPA		cytoplasm
Q6NDP1	RS1_RHOPA		cytoplasm
Q6NDP2	KCY_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6NDR5	RL35_RHOPA		cytoplasm
Q6NDR6	RL20_RHOPA		cytoplasm
Q6NDV3	DNAA_RHOPA	cytoplasm (by similarity).	cytoplasm
Q6PT86	SUGE_SALTH	bacterial cell inner membrane; multi- pass membrane protein (by similarity).	inner membrane

Q6PT90	SUGE2_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q6Q3H0	DHLA_XANFL		cytoplasm
Q6QNZ6	MTFA_PHOAZ		cytoplasm
Q6QW47	YP85_AZOBR		inner membrane
Q6RCE0	PHNC_PSEST	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6REU8	TYSY_XENNE	cytoplasm (by similarity).	cytoplasm
Q6RH47	TAUB_PARPEN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6RH51	TAUZ_PARPEN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6RH59	TAUZ_PARDE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q6RK33	HRPA_ERWCT	extracellular and secreted via type iii secretion system (by similarity).	extracellular
Q6RSN5	DNAJ_AGRTU	cytoplasm (probable).	cytoplasm
Q6T893	GLPE_ACTPL	cytoplasm (by similarity).	cytoplasm
Q6TG07	GMHB_CAMJE	cytoplasm (by similarity).	cytoplasm
Q6TG09	HLDE_CAMJE		cytoplasm
Q6TP45	LUXS_VIBAL		extracellular
Q6U5S4	Y189_KLEPN		inner membrane
Q6UR32	UREE_YERRO	cytoplasm (by similarity).	cytoplasm
Q6UR35	URE3_YERRO	cytoplasm (by similarity).	cytoplasm
Q6UR41	UREE_YERMO	cytoplasm (by similarity).	cytoplasm
Q6UR44	URE3_YERMO	cytoplasm (by similarity).	cytoplasm
Q6UR50	UREE_YERKR	cytoplasm (by similarity).	cytoplasm
Q6UR53	URE3_YERKR	cytoplasm (by similarity).	cytoplasm
Q6UR59	UREE_YERIN	cytoplasm (by similarity).	cytoplasm
Q6UR62	URE3_YERIN	cytoplasm (by similarity).	cytoplasm
Q6UR67	UREE_YERFR	cytoplasm (by similarity).	cytoplasm
Q6UR70	URE3_YERFR	cytoplasm (by similarity).	cytoplasm
Q6UR75	UREE_YERBE	cytoplasm (by similarity).	cytoplasm
Q6UR78	URE3_YERBE	cytoplasm (by similarity).	cytoplasm
Q6UR83	UREE_YERAL	cytoplasm (by similarity).	cytoplasm
Q6UR86	URE3_YERAL	cytoplasm (by similarity).	cytoplasm
Q6VAY5	DNAJ_PSEST	cytoplasm (probable).	cytoplasm
Q6W2B1	TAUB_RHISN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6WB51	PSTB_ALCFA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6WB63	PHNC_ALCFA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q6WB83	LLDD_ALCFA		cytoplasm
Q6WEG4	HRPA_PECCC	extracellular and secreted via type iii secretion system (by similarity).	extracellular

Q6XZ07	RECA_VIBNA	cytoplasm (by similarity).	cytoplasm
Q6YI30	RECF_SODGL	cytoplasm (by similarity).	cytoplasm
Q6ZZC6	MDH_MYXXA		cytoplasm
Q70FG9	PMRB_PECCC	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q70FH0	PMRA_PECCC	cytoplasm (potential).	cytoplasm
Q70GH6	GLMM_NEISU		cytoplasm
Q70I53	HDAH_ALCSD		cytoplasm
Q70JN9	SLDA1_GLUOX	secreted protein (potential).	cytoplasm
Q70JP0	SLDB1_GLUOX	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q714U9	FCS1_HAEIN		cytoplasm
Q725K9	PURA_DESVH	cytoplasm (by similarity).	cytoplasm
Q725M6	DNAJ_DESVH	cytoplasm (probable).	cytoplasm
Q725M7	RPOZ_DESVH		cytoplasm
Q725P0	ZRAP_DESVH		cytoplasm
Q725Q1	ILVD_DESVH		cytoplasm
Q725Q3	METE_DESVH		cytoplasm
Q725Q9	FMT_DESVH		cytoplasm
Q725S3	QUEA_DESVH	cytoplasm (potential).	cytoplasm
Q725T6	ATKA_DESVH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q725W6	Y3309_DESVH		cytoplasm
Q726D3	MURA_DESVH	cytoplasm (probable).	inner membrane
Q726D5	FPG_DESVH		cytoplasm
Q726E5	Y3208_DESVH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q726F4	HAM1_DESVH		extracellular
Q726K4	TOLB_DESVH	periplasm (potential).	periplasm
Q726N6	RUVX_DESVH	cytoplasm (potential).	cytoplasm
Q726S6	ACKA_DESVH	cytoplasm (by similarity).	cytoplasm
Q726X1	LEU3_DESVH	cytoplasm (by similarity).	cytoplasm
Q726X7	PSD_DESVH		cytoplasm
Q727C0	GPMA_DESVH		cytoplasm
Q727C6	RPOC_DESVH		cytoplasm
Q727C7	RPOB_DESVH		cytoplasm
Q727C9	RL10_DESVH		cytoplasm
Q727D0	RL1_DESVH		cytoplasm
Q727E3	RL31_DESVH		cytoplasm
Q727E5	Y2910_DESVH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q727F4	PYRB_DESVH		cytoplasm
Q727G3	Y2892_DESVH		inner membrane
Q727G4	NIKR_DESVH		cytoplasm
Q728D5	GLMU_DESVH	cytoplasm (by similarity).	cytoplasm
Q728G0	HTPG_DESVH	cytoplasm (by similarity).	cytoplasm
Q728R7	RL35_DESVH		cytoplasm
Q728R8	RL20_DESVH		cytoplasm
Q728T9	MRAZ_DESVH		inner membrane

Q728U5	MRAY_DESVH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q728U6	MURD_DESVH	cytoplasm (by similarity).	cytoplasm
Q728U8	MURG_DESVH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q728V0	MURB_DESVH	cytoplasm (probable).	inner membrane
Q729A4	PANC_DESVH		cytoplasm
Q729I2	LPXD_DESVH		cytoplasm
Q729L7	NDK_DESVH	cytoplasm (by similarity).	cytoplasm
Q729M2	HYP A1_DESVH		cytoplasm
Q729Q7	HYP A2_DESVH		cytoplasm
Q729T9	RRMJ_DESVH	cytoplasm (by similarity).	cytoplasm
Q729U1	RUVC_DESVH		cytoplasm
Q729U3	RUVA_DESVH		cytoplasm
Q729U4	THYX_DESVH		cytoplasm
Q729U6	DNAA_DESVH	cytoplasm (by similarity).	cytoplasm
Q729V4	GLGA_DESVH		cytoplasm
Q729V5	GLGB_DESVH		extracellular
Q72A55	SURE_DESVH	cytoplasm (potential).	periplasm
Q72A57	KTHY_DESVH		cytoplasm
Q72A64	Y2133_DESVH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q72AL6	CH60_DESVH	cytoplasm (by similarity).	cytoplasm
Q72AN9	PROA_DESVH	cytoplasm (by similarity).	cytoplasm
Q72AQ6	PHNC_DESVH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q72AR0	KAD_DESVH	cytoplasm (by similarity).	cytoplasm
Q72AR5	SYI_DESVH	cytoplasm (by similarity).	cytoplasm
Q72AT2	ACPS_DESVH	cytoplasm (by similarity).	cytoplasm
Q72AU4	RS20_DESVH		cytoplasm
Q72AV0	HEM3_DESVH		cytoplasm
Q72AV2	PPNK_DESVH	cytoplasm (by similarity).	cytoplasm
Q72AV7	Y1883_DESVH		inner membrane
Q72AW6	CLPB_DESVH	cytoplasm (probable).	cytoplasm
Q72AZ1	PIMT_DESVH	cytoplasm (by similarity).	cytoplasm
Q72B11	GIDA_DESVH	cytoplasm (by similarity).	cytoplasm
Q72B41	KSGA_DESVH		cytoplasm
Q72B46	RS21_DESVH		cytoplasm
Q72BE3	GLUQ_DESVH		cytoplasm
Q72BK4	Y1631_DESVH		cytoplasm
Q72BL6	GPMI_DESVH		cytoplasm
Q72BM6	DAPB_DESVH	cytoplasm (by similarity).	cytoplasm
Q72BN0	UVRB_DESVH	cytoplasm (by similarity).	cytoplasm
Q72BN2	LFTR_DESVH	cytoplasm (by similarity).	cytoplasm
Q72BN4	CLPS_DESVH		cytoplasm
Q72BN6	CRCB_DESVH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q72BQ7	HSLV_DESVH	cytoplasm (by similarity).	cytoplasm
Q72BQ8	ISPE_DESVH		cytoplasm
Q72BR0	RL25_DESVH		cytoplasm
Q72BR1	PTH_DESVH	cytoplasm (by similarity).	cytoplasm

Q72BV1	MIAA_DESVH		cytoplasm
Q72BV2	COAD_DESVH	cytoplasm (by similarity).	cytoplasm
Q72C17	HSLU_DESVH	cytoplasm (by similarity).	cytoplasm
Q72C18	ARGB_DESVH	cytoplasm (probable).	cytoplasm
Q72C23	HEM1_DESVH		cytoplasm
Q72C25	TILS_DESVH	cytoplasm (probable).	cytoplasm
Q72C30	ISPDF_DESVH		cytoplasm
Q72C44	RNC_DESVH	cytoplasm (by similarity).	cytoplasm
Q72CA6	ILVC_DESVH		cytoplasm
Q72CD5	EX7S_DESVH	cytoplasm (by similarity).	cytoplasm
Q72CD9	ISPG_DESVH		cytoplasm
Q72CE7	CLPX_DESVH		cytoplasm
Q72CE8	CLPP_DESVH	cytoplasm (by similarity).	cytoplasm
Q72CE9	TIG_DESVH		cytoplasm
Q72CF1	SELD_DESVH		cytoplasm
Q72CF4	RPOA_DESVH		cytoplasm
Q72CF5	RS4_DESVH		cytoplasm
Q72CF6	RS11_DESVH		cytoplasm
Q72CF7	RS13_DESVH		cytoplasm
Q72CG1	RL15_DESVH		cytoplasm
Q72CG3	RS5_DESVH		cytoplasm
Q72CG4	RL18_DESVH		cytoplasm
Q72CG6	RS8_DESVH		cytoplasm
Q72CG8	RL5_DESVH		cytoplasm
Q72CH1	RS17_DESVH		cytoplasm
Q72CH3	RL16_DESVH		cytoplasm
Q72CH4	RS3_DESVH		cytoplasm
Q72CH5	RL22_DESVH		cytoplasm
Q72CH6	RS19_DESVH		cytoplasm
Q72CH7	RL2_DESVH		cytoplasm
Q72CI1	RS10_DESVH		cytoplasm
Q72CI4	RS7_DESVH		cytoplasm
Q72CI5	RS12_DESVH		cytoplasm
Q72CK1	GLMM_DESVH		cytoplasm
Q72CM9	Y1254_DESVH		inner membrane
Q72CQ9	END4_DESVH		extracellular
Q72CS2	RL28_DESVH		cytoplasm
Q72CS4	RL32_DESVH		cytoplasm
Q72CS5	PLSX_DESVH		cytoplasm
Q72CT0	GLYA_DESVH	cytoplasm (by similarity).	cytoplasm
Q72D35	OTC_DESVH	cytoplasm (probable).	cytoplasm
Q72D36	ARLY_DESVH	cytoplasm (probable).	cytoplasm
Q72D46	PSTB_DESVH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q72D63	XGPT_DESVH	cytoplasm (by similarity).	cytoplasm
Q72D73	CBIO_DESVH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q72D86	GUAA_DESVH		cytoplasm
Q72DA0	HIS8_DESVH		cytoplasm
Q72DA1	KCY_DESVH	cytoplasm (by similarity).	cytoplasm
Q72DA4	UPP_DESVH		cytoplasm
Q72DG6	QUEF_DESVH	cytoplasm (probable).	cytoplasm
Q72DH1	RL9_DESVH		cytoplasm
Q72DH2	RS18_DESVH		cytoplasm

Q72DH3	RS6_DESVH		cytoplasm
Q72DH6	SYV_DESVH	cytoplasm (by similarity).	cytoplasm
Q72DI6	Y943_DESVH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q72DJ9	PROB_DESVH	cytoplasm (by similarity).	cytoplasm
Q72DK1	RL27_DESVH		cytoplasm
Q72DK5	Y924_DESVH		cytoplasm
Q72DL8	TRUA_DESVH		cytoplasm
Q72DM3	LIPB_DESVH	cytoplasm (potential).	cytoplasm
Q72DM9	KGUA_DESVH	cytoplasm (by similarity).	cytoplasm
Q72DN0	Y899_DESVH		inner membrane
Q72DN5	AROC_DESVH		cytoplasm
Q72DN7	AROK_DESVH	cytoplasm (probable).	cytoplasm
Q72DQ5	RS2_DESVH		cytoplasm
Q72DQ9	RRF_DESVH	cytoplasm (by similarity).	cytoplasm
Q72DR3	DXR_DESVH		cytoplasm
Q72DU2	RIMM_DESVH	cytoplasm (potential).	cytoplasm
Q72DU3	TRMD_DESVH	cytoplasm (potential).	cytoplasm
Q72DU4	RL19_DESVH		cytoplasm
Q72DU5	RNH2_DESVH	cytoplasm (potential).	extracellular
Q72DV1	SSRP_DESVH	cytoplasm (by similarity).	cytoplasm
Q72DW3	COBQ_DESVH		cytoplasm
Q72DW6	HRC_A_DESVH		cytoplasm
Q72DW7	GRPE_DESVH	cytoplasm (probable).	cytoplasm
Q72DW8	DNAK_DESVH		cytoplasm
Q72DX8	UVRC_DESVH	cytoplasm (by similarity).	cytoplasm
Q72E02	ATPA_DESVH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q72E03	ATPG_DESVH		inner membrane
Q72E05	ATPE_DESVH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q72E47	SYV_DESVH	cytoplasm (by similarity).	cytoplasm
Q72E76	LEPA_DESVH		cytoplasm
Q72EH1	SAHH_DESVH	cytoplasm (by similarity).	cytoplasm
Q72EN4	Y543_DESVH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q72EP8	CSRA_DESVH		cytoplasm
Q72EQ3	FLGI_DESVH	periplasm (by similarity).	periplasm
Q72EQ4	FLGH_DESVH	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q72EQ8	Y511_DESVH		inner membrane
Q72ER4	TRUB_DESVH		cytoplasm
Q72ER5	RS15_DESVH		cytoplasm
Q72ET7	HLDD_DESVH		cytoplasm
Q72EU7	TRPA_DESVH		cytoplasm
Q72EX7	ADEC_DESVH		cytoplasm
Q72F03	AMPA_DESVH	cytoplasm (by similarity).	cytoplasm
Q72F92	ENO_DESVH	cytoplasm (by similarity).	extracellular
Q72FU0	Y123_DESVH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q72G08	ISPH_DESVH		cytoplasm

Q72G45	LGT_DESVH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q73FM5	RPOZ_WOLPM		cytoplasm
Q73FP4	RIMM_WOLPM	cytoplasm (potential).	cytoplasm
Q73FQ1	ISPH_WOLPM		cytoplasm
Q73FR9	TILS_WOLPM	cytoplasm (probable).	cytoplasm
Q73FS2	KTHY_WOLPM		cytoplasm
Q73FT3	RNC_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73FU0	ATPG_WOLPM		inner membrane
Q73FX5	THYX_WOLPM		cytoplasm
Q73FX6	MURA_WOLPM	cytoplasm (probable).	inner membrane
Q73FY6	HEMH_WOLPM	cytoplasm (by similarity).	extracellular
Q73FY9	NDK_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73FZ1	RECR_WOLPM		cytoplasm
Q73G09	PDXH_WOLPM		cytoplasm
Q73G10	FPG_WOLPM		cytoplasm
Q73G18	RS20_WOLPM		cytoplasm
Q73G24	ISPDF_WOLPM		cytoplasm
Q73G26	APAG_WOLPM		inner membrane
Q73G32	Y1135_WOLPM		inner membrane
Q73G44	MDH_WOLPM		cytoplasm
Q73G50	SYI_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73G60	RNH2_WOLPM	cytoplasm (potential).	extracellular
Q73G61	MRAY_WOLPM	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q73G71	TRMB_WOLPM		cytoplasm
Q73G84	SURE_WOLPM	cytoplasm (potential).	periplasm
Q73G93	TRUA_WOLPM		cytoplasm
Q73GB0	PSD_WOLPM		cytoplasm
Q73GB4	Y1045_WOLPM	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q73GC3	GLYA_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73GD5	TRMD_WOLPM	cytoplasm (potential).	cytoplasm
Q73GD6	RL19_WOLPM		cytoplasm
Q73GE2	LIPB_WOLPM	cytoplasm (potential).	cytoplasm
Q73GE7	GIDA_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73GG3	DXR_WOLPM		cytoplasm
Q73GG7	RL32_WOLPM		cytoplasm
Q73GG8	PLSX_WOLPM		cytoplasm
Q73GG9	FABH_WOLPM	cytoplasm (probable).	cytoplasm
Q73GI2	CCME_WOLPM	bacterial cell membrane; single-pass type ii membrane protein (potential).	inner membrane
Q73GL7	DNAK_WOLPM		cytoplasm
Q73GN5	RS15_WOLPM		cytoplasm
Q73GP1	PYRB_WOLPM		cytoplasm
Q73GR4	GPMI_WOLPM		cytoplasm
Q73GR6	FMT_WOLPM		cytoplasm
Q73GR7	RL20_WOLPM		cytoplasm
Q73GR8	RL35_WOLPM		cytoplasm
Q73GT3	MURD_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73GW5	ACPS_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73GX9	GRPE_WOLPM	cytoplasm (probable).	cytoplasm

Q73GZ5	RL9_WOLPM		cytoplasm
Q73GZ6	RS18_WOLPM		cytoplasm
Q73H02	DAPA_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73H08	LGT_WOLPM	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q73H09	SSRP_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73H48	Y726_WOLPM		inner membrane
Q73H59	DAPD_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73H86	RS10_WOLPM		cytoplasm
Q73H89	RL2_WOLPM		cytoplasm
Q73H91	RL22_WOLPM		cytoplasm
Q73H92	RS3_WOLPM		cytoplasm
Q73H93	RL16_WOLPM		cytoplasm
Q73H94	RL29_WOLPM		cytoplasm
Q73H95	RS17_WOLPM		cytoplasm
Q73H98	RL5_WOLPM		cytoplasm
Q73HA0	RS8_WOLPM		cytoplasm
Q73HA2	RL18_WOLPM		cytoplasm
Q73HA6	KAD_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73HA9	RPOA_WOLPM		cytoplasm
Q73HB2	ATPA_WOLPM	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q73HK1	Y557_WOLPM		inner membrane
Q73HL4	MURB_WOLPM	cytoplasm (probable).	inner membrane
Q73HM1	RS2_WOLPM		cytoplasm
Q73HM7	COAD_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73HQ2	ENO_WOLPM	cytoplasm (by similarity).	extracellular
Q73HR8	LEPA_WOLPM		cytoplasm
Q73HS9	SYV_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73HV1	KGUA_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73HW7	SYI_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73HX8	CCMA_WOLPM	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q73HZ4	UBIE_WOLPM		cytoplasm
Q73HZ5	RL28_WOLPM		cytoplasm
Q73I13	DAPB_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73I23	ISPE_WOLPM		cytoplasm
Q73I41	Y341_WOLPM		cytoplasm
Q73I44	PURA_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73I58	TIG_WOLPM		cytoplasm
Q73I59	CLPP_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73I60	CLPX_WOLPM		cytoplasm
Q73I70	CH10_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73I71	CH60_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73I74	RNH_WOLPM	cytoplasm (potential).	extracellular
Q73IE4	CLPB_WOLPM	cytoplasm (probable).	cytoplasm
Q73IH1	GUAA_WOLPM		cytoplasm
Q73IH9	COAE_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73II8	PTH_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73II9	RL25_WOLPM		cytoplasm
Q73IK6	UVRC_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73IL4	RUVX_WOLPM	cytoplasm (potential).	cytoplasm
Q73IL5	RUVC_WOLPM		cytoplasm

Q73IM4	GLMU_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73IP1	ISPG_WOLPM		cytoplasm
Q73IQ0	SECB_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73IR3	KSGA_WOLPM		cytoplasm
Q73IS9	RRMJ_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73IU2	AMPA_WOLPM	cytoplasm (by similarity).	cytoplasm
Q73IV4	DNAJ_WOLPM	cytoplasm (probable).	cytoplasm
Q73IV6	TOLB_WOLPM	periplasm (potential).	periplasm
Q73IW9	RPOBC_WOLPM		cytoplasm
Q73IX8	RS7_WOLPM		cytoplasm
Q73IX9	RS12_WOLPM		cytoplasm
Q73IZ0	DNAA_WOLPM	cytoplasm (by similarity).	cytoplasm
Q746Q4	GIDA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q746Q5	GIDB_GEOSL		cytoplasm
Q746Y3	PPCK_GEOSL	cytoplasm (by similarity).	cytoplasm
Q746Z9	ISPD_GEOSL		cytoplasm
Q747A0	ISPF_GEOSL		extracellular
Q747B9	HSLO_GEOSL	cytoplasm (by similarity).	cytoplasm
Q747C1	KUP3_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q747C7	CH60_GEOSL	cytoplasm (by similarity).	cytoplasm
Q747C8	CH10_GEOSL	cytoplasm (by similarity).	cytoplasm
Q747F5	HEMH_GEOSL	cytoplasm (by similarity).	extracellular
Q747F9	PURA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q747I1	HEM3_GEOSL		cytoplasm
Q747I2	HEM1_GEOSL		cytoplasm
Q747I7	UVRC_GEOSL	cytoplasm (by similarity).	cytoplasm
Q747K3	UVRB_GEOSL	cytoplasm (by similarity).	cytoplasm
Q747K8	GLGA2_GEOSL		cytoplasm
Q747N0	RL27_GEOSL		cytoplasm
Q747Q3	PROB_GEOSL	cytoplasm (by similarity).	cytoplasm
Q747Q4	PROA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q747Q7	Y3208_GEOSL		inner membrane
Q747Q8	GPMI_GEOSL		cytoplasm
Q747W9	MOAA_GEOSL		cytoplasm
Q747X9	SYI_GEOSL	cytoplasm (by similarity).	cytoplasm
Q748A8	RL31_GEOSL		cytoplasm
Q748A9	THYX_GEOSL		cytoplasm
Q748B3	MURA_GEOSL	cytoplasm (probable).	inner membrane
Q748C9	MRAZ_GEOSL		inner membrane
Q748D3	MRAY_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q748D4	MURD_GEOSL	cytoplasm (by similarity).	cytoplasm
Q748D6	MURG_GEOSL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q748F4	FLGH_GEOSL	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q748F5	FLGI_GEOSL	periplasm (by similarity).	periplasm
Q748G1	CSRA_GEOSL		cytoplasm
Q748K0	CBIO2_GEOSL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q748M1	NIKR_GEOSL		cytoplasm

Q748X2	LEU3_GEOSL	cytoplasm (by similarity).	cytoplasm
Q748X6	ARGC_GEOSL	cytoplasm (probable).	cytoplasm
Q748Y3	RL1_GEOSL		cytoplasm
Q748Y4	RL10_GEOSL		cytoplasm
Q748Y6	RPOB_GEOSL		cytoplasm
Q748Y7	RS7_GEOSL		cytoplasm
Q748Z0	RS10_GEOSL		cytoplasm
Q748Z2	RS19_GEOSL		cytoplasm
Q748Z3	RL22_GEOSL		cytoplasm
Q748Z4	RS3_GEOSL		cytoplasm
Q748Z5	RL16_GEOSL		cytoplasm
Q748Z6	RL29_GEOSL		cytoplasm
Q748Z7	RS17_GEOSL		cytoplasm
Q748Z9	RL5_GEOSL		cytoplasm
Q749A1	RS8_GEOSL		cytoplasm
Q749A3	RL18_GEOSL		cytoplasm
Q749A4	RS5_GEOSL		cytoplasm
Q749A6	RL15_GEOSL		cytoplasm
Q749A8	KAD_GEOSL	cytoplasm (by similarity).	cytoplasm
Q749B0	RS13_GEOSL		cytoplasm
Q749B1	RS11_GEOSL		cytoplasm
Q749B2	RS4_GEOSL		cytoplasm
Q749B3	RPOA_GEOSL		cytoplasm
Q749C5	Y2818_GEOSL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q749E9	MSCL_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q749F2	Y2791_GEOSL		inner membrane
Q749N5	ACKA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q749X2	QUEA_GEOSL	cytoplasm (potential).	cytoplasm
Q749Y7	KCY_GEOSL	cytoplasm (by similarity).	cytoplasm
Q749Y8	ISPH_GEOSL		cytoplasm
Q74A44	GID_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74AA5	KUP2_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q74AB0	ATKA_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q74AC8	METX_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74AE2	LIPB_GEOSL	cytoplasm (potential).	cytoplasm
Q74AH4	TRPD_GEOSL		cytoplasm
Q74AI3	TRPA_GEOSL		cytoplasm
Q74AK4	KUP1_GEOSL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q74AM0	Y2333_GEOSL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q74AR6	ENO_GEOSL	cytoplasm (by similarity).	extracellular
Q74AR8	Y2284_GEOSL		inner membrane
Q74AT5	LPXD_GEOSL		cytoplasm
Q74AU2	LPXK_GEOSL		cytoplasm
Q74AW2	RPOZ_GEOSL		cytoplasm

Q74AW5	RL28_GEOSL		cytoplasm
Q74AX0	KTHY_GEOSL		cytoplasm
Q74AX1	RNC_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74AZ3	RS20_GEOSL		cytoplasm
Q74BF6	HLDE_GEOSL		cytoplasm
Q74BH6	PPNK_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74BJ6	SYV_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74BL4	AROC_GEOSL		cytoplasm
Q74BL5	AROK_GEOSL	cytoplasm (probable).	cytoplasm
Q74BL6	AROB_GEOSL	cytoplasm (probable).	cytoplasm
Q74BL9	AROQ_GEOSL		cytoplasm
Q74BP1	MIAA_GEOSL		cytoplasm
Q74BW1	RS2_GEOSL		cytoplasm
Q74BW4	DXR_GEOSL		cytoplasm
Q74BW7	ILVD_GEOSL		cytoplasm
Q74BW9	ILVC_GEOSL		cytoplasm
Q74BX0	PSD_GEOSL		cytoplasm
Q74BZ3	Y1884_GEOSL		cytoplasm
Q74C12	KSGA_GEOSL		cytoplasm
Q74C57	APGM_GEOSL		cytoplasm
Q74C65	TILS_GEOSL	cytoplasm (probable).	cytoplasm
Q74C70	GLMM_GEOSL		cytoplasm
Q74C71	ACPS_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74C79	RNPH_GEOSL		cytoplasm
Q74C80	HAM1_GEOSL		extracellular
Q74C81	TIG_GEOSL		cytoplasm
Q74C82	CLPP_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74C83	CLPX_GEOSL		cytoplasm
Q74CA8	EX7S_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74CC6	Y1748_GEOSL		cytoplasm
Q74CG4	SSRP_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74CG7	PANC_GEOSL		cytoplasm
Q74CG8	PANB_GEOSL		cytoplasm
Q74CI6	NRDR_GEOSL		cytoplasm
Q74CR0	GPMA_GEOSL		cytoplasm
Q74CR5	GLYA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74CR8	ACP_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74CS2	PLSX_GEOSL		cytoplasm
Q74CS3	RL32_GEOSL		cytoplasm
Q74CT0	RS15_GEOSL		cytoplasm
Q74CT2	RBFA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74CT6	Y1585_GEOSL		inner membrane
Q74CZ5	PIMT_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74CZ6	SURE_GEOSL	cytoplasm (potential).	periplasm
Q74D01	RL20_GEOSL		cytoplasm
Q74D02	RL35_GEOSL		cytoplasm
Q74D53	MDH_GEOSL		cytoplasm
Q74D60	ISPG_GEOSL		cytoplasm
Q74D67	Y1452_GEOSL		cytoplasm
Q74DB5	ACCA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74DI7	GLUQ_GEOSL		cytoplasm
Q74DI8	Y1328_GEOSL		inner membrane
Q74DK5	G6PI_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74DN5	CBIO1_GEOSL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q74DP5	PYRB_GEOSL		cytoplasm
Q74DS2	COAD_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74DU7	Y1218_GEOSL		cytoplasm
Q74E25	SYU_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74E54	NDK_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74E68	PSTB_GEOSL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q74E87	RUVA_GEOSL		cytoplasm
Q74E88	RUVC_GEOSL		cytoplasm
Q74ED9	GLGA1_GEOSL		cytoplasm
Q74EE6	Y1016_GEOSL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q74EG5	FPG_GEOSL		cytoplasm
Q74EM9	UPP_GEOSL		cytoplasm
Q74EU2	UBIE_GEOSL		cytoplasm
Q74F05	WRBA_GEOSL		cytoplasm
Q74FE1	RL9_GEOSL		cytoplasm
Q74FE3	RS18_GEOSL		cytoplasm
Q74FE4	RS6_GEOSL		cytoplasm
Q74FE6	PTH_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74FE7	RL25_GEOSL		cytoplasm
Q74FE9	ISPE_GEOSL		cytoplasm
Q74FF1	CLPB_GEOSL	cytoplasm (probable).	cytoplasm
Q74FG0	RNH2_GEOSL	cytoplasm (potential).	extracellular
Q74FG1	RL19_GEOSL		cytoplasm
Q74FG3	TRMD_GEOSL	cytoplasm (potential).	cytoplasm
Q74FG4	RIMM_GEOSL	cytoplasm (potential).	cytoplasm
Q74FK1	SELD_GEOSL		cytoplasm
Q74FL9	THIG_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74FM9	RECO_GEOSL		cytoplasm
Q74FS7	END4_GEOSL		extracellular
Q74FT1	DTD_GEOSL	cytoplasm (probable).	cytoplasm
Q74FT5	TRMB_GEOSL		cytoplasm
Q74FW8	Y484_GEOSL		inner membrane
Q74G02	Y450_GEOSL		cytoplasm
Q74G05	PRMA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74G62	RADC_GEOSL		cytoplasm
Q74G73	HYPB_GEOSL		cytoplasm
Q74GB4	AMPA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74GG6	DNAA_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74GH5	GLMU_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74GH6	GLMS_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74GS4	Y171_GEOSL		inner membrane
Q74GT5	DAPB_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74GT9	ARLY_GEOSL	cytoplasm (probable).	cytoplasm
Q74GU2	OTC_GEOSL	cytoplasm (probable).	cytoplasm
Q74GU4	ARGB_GEOSL	cytoplasm (probable).	cytoplasm
Q74GV1	CINA_GEOSL		inner membrane
Q74GV3	Y141_GEOSL		inner membrane
Q74GW4	FMT_GEOSL		cytoplasm
Q74GX9	ATPE_GEOSL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q74GY1	ATPG_GEOSL		inner membrane

Q74GY2	ATPA_GEOSL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q74GZ7	RECR_GEOSL		cytoplasm
Q74H58	DNAJ_GEOSL	cytoplasm (probable).	cytoplasm
Q74H59	DNAK_GEOSL		cytoplasm
Q74H60	GRPE_GEOSL	cytoplasm (probable).	cytoplasm
Q74H61	HRCA_GEOSL		cytoplasm
Q74H67	TOLB_GEOSL	periplasm (potential).	periplasm
Q74H90	RECF_GEOSL	cytoplasm (by similarity).	cytoplasm
Q74PY8	TYPH_YERPE		cytoplasm
Q74Q30	DJLA_YERPE	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q74R94	GPPA_YERPE		extracellular
Q74RF8	MALG_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q74RF9	MALF_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q74RZ6	AAS_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q74S44	Y1040_YERPE		cytoplasm
Q74TH2	SUFE_YERPE	cytoplasm (by similarity).	cytoplasm
Q74X11	CLPB_YERPE	cytoplasm (probable).	cytoplasm
Q74XD3	CUTA_YERPE	cytoplasm (by similarity).	periplasm
Q74Y45	BIOH_YERPE	cytoplasm (by similarity).	cytoplasm
Q75R59	NQRF_VIBAN	bacterial cell inner membrane (potential).	inner membrane
Q75R60	NQRE_VIBAN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q75R61	NQRD_VIBAN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q75R62	NQRC_VIBAN	bacterial cell inner membrane (potential).	inner membrane
Q75R63	NQRB_VIBAN	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q75R64	NQRA_VIBAN		inner membrane
Q75WD2	DNAJ_ACEAC	cytoplasm (probable).	cytoplasm
Q799A3	SUGE_KLEOX	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q79IF2	SUGE2_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q79K00	SUGE_SALCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q79RI9	TR96_PRORE		cytoplasm
Q79S39	TRSX_VIBCH		cytoplasm
Q79V15	GRPE_BRAJA	cytoplasm (probable).	cytoplasm

Q7A9L5	ADEC_ECO57		cytoplasm
Q7A9X4	XYLA_ECO57	cytoplasm (by similarity).	cytoplasm
Q7AAQ7	HLDE_ECO57		cytoplasm
Q7ABA6	HOSA_ECO57		cytoplasm
Q7ABI1	GRPE_ECO57	cytoplasm (probable).	cytoplasm
Q7ABK6	HMP_ECO57		cytoplasm
Q7AC24	ARNB_ECO57		cytoplasm
Q7ACM1	MDTC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7ADF8	YNIC_ECO57		cytoplasm
Q7ADI4	SUFS_ECO57	cytoplasm (by similarity).	cytoplasm
Q7ADI5	SUFE_ECO57	cytoplasm (by similarity).	cytoplasm
Q7AFA1	MDTH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7AFV7	CBPA_ECO57	nucleoid (by similarity).	inner membrane
Q7AH43	FBPC_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7AH91	BETB_ECO57		cytoplasm
Q7AHS5	DJLA_ECO57	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q7AHT0	FIXC_ECO57		periplasm
Q7AMH5	CLPB_SALTI	cytoplasm (probable).	cytoplasm
Q7ANK1	YAEH_SALTI		inner membrane
Q7BF87	ARNB_YERPS		cytoplasm
Q7BS31	EFE_PSECA		cytoplasm
Q7BS32	EFE_PSESG		cytoplasm
Q7BSX2	COPB_ECO57		cytoplasm
Q7BSX4	YFI3_ECO57		extracellular
Q7BZ80	RECF_SHIFL	cytoplasm (by similarity).	cytoplasm
Q7BZ90	ADEC_SHIFL		cytoplasm
Q7C093	ISPD_SHIFL		cytoplasm
Q7C0D0	GRPE_SHIFL	cytoplasm (probable).	cytoplasm
Q7C0F9	HMP_SHIFL		cytoplasm
Q7C1E6	SUFE_SHIFL	cytoplasm (by similarity).	cytoplasm
Q7C1M3	BTUD_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7C1M5	BTUC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7C254	CBPA_SHIFL	nucleoid (by similarity).	inner membrane
Q7C397	COAE_SHIFL	cytoplasm (by similarity).	cytoplasm
Q7C637	XYLA_SALTI	cytoplasm (by similarity).	cytoplasm
Q7CB38	YEIH_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7CEG6	CLPB_BRUSU	cytoplasm (probable).	cytoplasm
Q7CH40	GRPE_YERPE	cytoplasm (probable).	cytoplasm
Q7CLU6	RL22_XANCP		cytoplasm
Q7CNT2	URE21_BRUME	cytoplasm (by similarity).	cytoplasm
Q7CNT3	UREE_BRUME	cytoplasm (by similarity).	cytoplasm
Q7CP66	TYPH_SALTY		cytoplasm

Q7CP73	MDTM_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7CP92	ULAG_SALTY	cytoplasm (potential).	extracellular
Q7CP93	ULAR_SALTY	cytoplasm (potential).	cytoplasm
Q7CP98	SUGE_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7CPA2	CUTA_SALTY	cytoplasm (by similarity).	periplasm
Q7CPB2	YBJB_SALTY		cytoplasm
Q7CPE1	ATPA_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7CPF1	IBPA_SALTY	cytoplasm (by similarity).	cytoplasm
Q7CPH8	SECB_SALTY	cytoplasm (by similarity).	cytoplasm
Q7CPL4	RL16_SALTY		cytoplasm
Q7CPL6	RL18_SALTY		cytoplasm
Q7CPM7	AAEX_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7CPM8	AAEA_SALTY	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q7CPP9	GLMM_SALTY		cytoplasm
Q7CPR7	MUG_SALTY	cytoplasm (potential).	extracellular
Q7CPR9	HLDE_SALTY		cytoplasm
Q7CPV1	YGFY_SALTY		cytoplasm
Q7CPW1	MTFA_SALTY		cytoplasm
Q7CPZ4	GRPE_SALTY	cytoplasm (probable).	cytoplasm
Q7CQ01	CLPB_SALTY	cytoplasm (probable).	cytoplasm
Q7CQ05	GRCA_SALTY		cytoplasm
Q7CQ08	TADA_SALTY		cytoplasm
Q7CQ12	ISCA_SALTY		cytoplasm
Q7CQ21	HDA_SALTY		nucleoid
Q7CQ70	YEJL_SALTY		inner membrane
Q7CQ71	RL25_SALTY		cytoplasm
Q7CQB9	PGSA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7CQF3	KITH_SALTY	cytoplasm (by similarity).	cytoplasm
Q7CQK5	HNSB2_SALTY		cytoplasm
Q7CQK7	SYI_SALTY	cytoplasm (by similarity).	cytoplasm
Q7CQN5	SUFS_SALTY	cytoplasm (by similarity).	cytoplasm
Q7CQR9	YCDX_SALTY		cytoplasm
Q7CQS3	CBPM_SALTY		cytoplasm
Q7CQS8	TUSE_SALTY	cytoplasm (by similarity).	cytoplasm
Q7CQZ6	CITD1_SALTY	cytoplasm (by similarity).	cytoplasm
Q7CR41	ACPH_SALTY		cytoplasm
Q7CR64	YAEH_SALTY		inner membrane
Q7CR86	DJLA_SALTY	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q7CR91	CITD2_SALTY	cytoplasm (by similarity).	cytoplasm
Q7CU92	CLPB_AGRT5	cytoplasm (probable).	cytoplasm
Q7CUX4	ADEC2_AGRT5		cytoplasm
Q7CWK7	MDH_AGRT5		cytoplasm

Q7CXF0	ADEC1_AGRT5		cytoplasm
Q7D0Y8	Y671_AGRT5	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7DBF9	UDG_ECO57		cytoplasm
Q7DDI1	Y1088_NEIMB		cytoplasm
Q7DDL2	RRMJ_NEIMB	cytoplasm (by similarity).	cytoplasm
Q7DDM4	RS15_NEIMB		cytoplasm
Q7DDM9	GRPE_NEIMB	cytoplasm (probable).	cytoplasm
Q7DDT0	RL18_NEIMB		cytoplasm
Q7DDT3	RS17_NEIMB		cytoplasm
Q7DDT4	RL16_NEIMB		cytoplasm
Q7DDT5	RL22_NEIMB		cytoplasm
Q7M7H7	RNPA_PHOLL		extracellular
Q7M7H8	GSHB_PHOLL		cytoplasm
Q7M7I1	CH102_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7M7I2	CH602_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7M7I3	FABH2_VIBVY	cytoplasm (probable).	cytoplasm
Q7M7I6	CH101_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7M7I7	CH601_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7M7I8	MIAA_VIBVY		cytoplasm
Q7M7J0	TRMD_VIBVY	cytoplasm (potential).	cytoplasm
Q7M7J1	FABZ_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7M7J4	TRUA_VIBVY		cytoplasm
Q7M7J5	FABA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7M7J9	FABH1_VIBVY	cytoplasm (probable).	cytoplasm
Q7M7K0	DGTL1_VIBVY		cytoplasm
Q7M7K5	CCA_VIBVY		cytoplasm
Q7M7L1	RUVC_WOLSU		cytoplasm
Q7M7L6	DAPA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M7M2	DEF_WOLSU		cytoplasm
Q7M7M3	CLPP_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M7M4	TIG_WOLSU		cytoplasm
Q7M7P6	ASSY_WOLSU	cytoplasm (probable).	cytoplasm
Q7M7P9	HEMH_WOLSU	cytoplasm (by similarity).	extracellular
Q7M7Q5	COAE_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M7Q6	SPEE_WOLSU		cytoplasm
Q7M7Q9	ARGB_WOLSU	cytoplasm (probable).	cytoplasm
Q7M7R1	RS9_WOLSU		cytoplasm
Q7M7S6	LGT_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7M7T1	FLGH_WOLSU	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q7M7T9	GLMU_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M7U1	ARGC_WOLSU	cytoplasm (probable).	cytoplasm
Q7M7U7	RL25_WOLSU		cytoplasm
Q7M7U8	PTH_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M7V2	HIS7_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M7V7	UPPS_WOLSU		inner membrane
Q7M7V8	PURA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M7W8	ENGA_WOLSU		cytoplasm
Q7M7W9	GPMI_WOLSU		cytoplasm
Q7M7X0	MRAY_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q7M7X6	KHSE_WOLSU	cytoplasm (potential).	cytoplasm
Q7M7Y2	GATB_WOLSU		cytoplasm
Q7M7Y3	GPDA_WOLSU	cytoplasm (probable).	cytoplasm
Q7M7Y6	HIS8_WOLSU		cytoplasm
Q7M7Z0	DXS_WOLSU		cytoplasm
Q7M7Z3	NDK_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M7Z4	PLSX_WOLSU		cytoplasm
Q7M803	ACPS_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M806	UVRB_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M818	THIC_WOLSU		cytoplasm
Q7M840	RNC_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M841	AROC_WOLSU		cytoplasm
Q7M842	GATA_WOLSU		cytoplasm
Q7M847	GSA_WOLSU	cytoplasm (potential).	cytoplasm
Q7M851	ILVC_WOLSU		cytoplasm
Q7M877	TRPA_WOLSU		cytoplasm
Q7M878	PANB_WOLSU		cytoplasm
Q7M879	RUVB_WOLSU		cytoplasm
Q7M886	LEU3_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M887	LEUD_WOLSU		cytoplasm
Q7M8A5	KAD_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M8B4	END4_WOLSU		extracellular
Q7M8B5	OTC_WOLSU	cytoplasm (probable).	cytoplasm
Q7M8C4	HTPG_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M8D4	RL3_WOLSU		cytoplasm
Q7M8D5	RL4_WOLSU		cytoplasm
Q7M8D7	RL2_WOLSU		cytoplasm
Q7M8D8	RS19_WOLSU		cytoplasm
Q7M8D9	RL22_WOLSU		cytoplasm
Q7M8E0	RS3_WOLSU		cytoplasm
Q7M8E1	RL16_WOLSU		cytoplasm
Q7M8E2	RS17_WOLSU		cytoplasm
Q7M8E4	RL24_WOLSU		cytoplasm
Q7M8E5	RL5_WOLSU		cytoplasm
Q7M8E6	RS8_WOLSU		cytoplasm
Q7M8E8	RL18_WOLSU		cytoplasm
Q7M8E9	RS5_WOLSU		cytoplasm
Q7M8F4	RS13_WOLSU		cytoplasm
Q7M8F5	RS11_WOLSU		cytoplasm
Q7M8F6	RS4_WOLSU		cytoplasm
Q7M8F7	RPOA_WOLSU		cytoplasm
Q7M8F9	MURE_WOLSU	cytoplasm (probable).	cytoplasm
Q7M8G1	SURE_WOLSU	cytoplasm (potential).	periplasm
Q7M8H5	LEPA_WOLSU		cytoplasm
Q7M8H6	UPP_WOLSU		cytoplasm
Q7M8H7	SYV_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M8I1	AROQ_WOLSU		cytoplasm
Q7M8J0	KPRS_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M8K0	UVRC_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M8K2	GUAA_WOLSU		cytoplasm
Q7M8K6	FLGI_WOLSU	periplasm (by similarity).	periplasm
Q7M8K9	HISX_WOLSU		cytoplasm
Q7M8L2	HEM3_WOLSU		cytoplasm
Q7M8M4	PHNC_WOLSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q7M8N8	TGT_WOLSU		cytoplasm
Q7M8N9	AROB_WOLSU	cytoplasm (probable).	cytoplasm
Q7M8Q0	ENO_WOLSU	cytoplasm (by similarity).	extracellular
Q7M8Q1	RECA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M8U5	CLPX2_WOLSU		cytoplasm
Q7M8W6	AMPA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M8W7	TRPB2_WOLSU		cytoplasm
Q7M8Y6	ISPH_WOLSU		cytoplasm
Q7M8Z2	ISPG_WOLSU		cytoplasm
Q7M8Z4	PROA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M8Z5	RL9_WOLSU		cytoplasm
Q7M8Z6	HSLV_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M8Z9	UPPP_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7M915	Y1268_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7M933	GCH1_WOLSU		cytoplasm
Q7M936	RIBAB_WOLSU		cytoplasm
Q7M938	ARLY_WOLSU	cytoplasm (probable).	cytoplasm
Q7M962	NAPA_WOLSU	periplasm (by similarity).	periplasm
Q7M9A0	DCD_WOLSU		extracellular
Q7M9A7	MDH_WOLSU		cytoplasm
Q7M9B0	RS15_WOLSU		cytoplasm
Q7M9C0	CORA_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7M9C1	PGK_WOLSU	cytoplasm (potential).	cytoplasm
Q7M9F6	THIG_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9J0	SSRP_WOLSU	cytoplasm (potential).	cytoplasm
Q7M9J1	TRUB_WOLSU		cytoplasm
Q7M9J8	GIDB_WOLSU		cytoplasm
Q7M9K2	ECTB_WOLSU		cytoplasm
Q7M9K3	ECTA_WOLSU		cytoplasm
Q7M9L1	SELA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9L5	PURL_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9L7	RL20_WOLSU		cytoplasm
Q7M9L8	RL35_WOLSU		cytoplasm
Q7M9M2	GLMM_WOLSU		cytoplasm
Q7M9M3	RS20_WOLSU		cytoplasm
Q7M9M5	GIDA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9M7	DXR_WOLSU		cytoplasm
Q7M9N0	HIS1_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9N1	DUT_WOLSU		extracellular
Q7M9N9	THIE_WOLSU		cytoplasm
Q7M9R5	ADD_WOLSU		extracellular
Q7M9S1	TRPB1_WOLSU		cytoplasm
Q7M9T3	DNAJ_WOLSU	cytoplasm (probable).	cytoplasm
Q7M9U1	HLDE_WOLSU		cytoplasm
Q7M9U3	TRUA_WOLSU		cytoplasm
Q7M9W1	ACP_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9W3	ACCA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9W4	LEU1_WOLSU		cytoplasm
Q7M9W9	HIS4_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9X0	HIS5_WOLSU	cytoplasm (by similarity).	cytoplasm

Q7M9X2	COAD_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9X4	CLPB_WOLSU	cytoplasm (probable).	cytoplasm
Q7M9X5	PUR7_WOLSU		cytoplasm
Q7M9X7	PURQ_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7M9Y9	PYRB_WOLSU		cytoplasm
Q7M9Z1	Y575_WOLSU	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q7M9Z9	LEU2_WOLSU		cytoplasm
Q7MA00	UNG_WOLSU	cytoplasm (by similarity).	extracellular
Q7MA03	RL19_WOLSU		cytoplasm
Q7MA04	TRMD_WOLSU	cytoplasm (potential).	cytoplasm
Q7MA16	TOLB_WOLSU	periplasm (potential).	periplasm
Q7MA19	ATPG_WOLSU		inner membrane
Q7MA20	ATPA_WOLSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MA26	FMT_WOLSU		cytoplasm
Q7MA27	PROB_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MA29	RL27_WOLSU		cytoplasm
Q7MA34	GRPE_WOLSU	cytoplasm (probable).	cytoplasm
Q7MA35	DNAK_WOLSU		cytoplasm
Q7MA54	RS7_WOLSU		cytoplasm
Q7MA55	RS12_WOLSU		cytoplasm
Q7MA56	RPOBC_WOLSU		cytoplasm
Q7MA58	RL10_WOLSU		cytoplasm
Q7MA59	RL1_WOLSU		cytoplasm
Q7MA63	DAPB_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MA70	RUVA_WOLSU		cytoplasm
Q7MA71	DDL_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MA77	TPIS_WOLSU		cytoplasm
Q7MAA5	DCUP_WOLSU	cytoplasm (probable).	cytoplasm
Q7MAC1	MURA_WOLSU	cytoplasm (probable).	inner membrane
Q7MAC4	LFTR_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAC9	ARGJ_WOLSU	cytoplasm (probable).	cytoplasm
Q7MAD1	RL28_WOLSU		cytoplasm
Q7MAD4	MURG_WOLSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MAD7	PYRE_WOLSU		cytoplasm
Q7MAD8	RRF_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAE2	CH10_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAE3	CH60_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAE6	ARGD_WOLSU	cytoplasm (probable).	cytoplasm
Q7MAE7	DGTL1_WOLSU		cytoplasm
Q7MAE8	PYRF_WOLSU		cytoplasm
Q7MAE9	NUSB_WOLSU		cytoplasm
Q7MAF0	RISB_WOLSU		cytoplasm
Q7MAF1	KDSA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAH6	SYI_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAJ5	NADE_WOLSU		cytoplasm
Q7MAK0	RS2_WOLSU		cytoplasm
Q7MAK5	KGUA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAL0	SYI_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAN4	ILVD_WOLSU		cytoplasm
Q7MAP8	PDXJ_WOLSU	cytoplasm (probable).	cytoplasm
Q7MAQ8	HIS2_WOLSU	cytoplasm (by similarity).	cytoplasm

Q7MAR0	GLYA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAR3	LNT_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MAS1	HIS6_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAS2	FABZ_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MAS4	CLPX1_WOLSU		cytoplasm
Q7MAV0	Y004_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MAX1	TRME_PHOLL		cytoplasm
Q7MAX2	RNPH_PHOLL		cytoplasm
Q7MAX3	DUT_PHOLL		extracellular
Q7MAX5	CAPP_PHOLL		cytoplasm
Q7MAX6	TRMA_PHOLL		cytoplasm
Q7MAX8	MIAA_PHOLL		cytoplasm
Q7MAX9	PURA2_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MAY1	TRUB_PHOLL		cytoplasm
Q7MAZ1	PCP1_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MAZ6	CH10_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MAZ7	CH60_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MAZ9	CCA_PHOLL		cytoplasm
Q7MB01	TGT_PHOLL		cytoplasm
Q7MB17	TRUA_PHOLL		cytoplasm
Q7MB46	FABA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MB64	TRMD_PHOLL	cytoplasm (potential).	cytoplasm
Q7MB85	CYSH_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MBD4	RNPH_CHRVO		cytoplasm
Q7MBD6	TRMD_CHRVO	cytoplasm (potential).	cytoplasm
Q7MBE2	MIAA_CHRVO		cytoplasm
Q7MBE8	DUT_CHRVO		extracellular
Q7MBF4	CCA_CHRVO		cytoplasm
Q7MBF7	TRUB_CHRVO		cytoplasm
Q7MBG1	GLMU_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7MBG5	HEM1_WOLSU		cytoplasm
Q7MBG6	MQO_WOLSU		cytoplasm
Q7MBG9	STHA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MBT9	PDXH_VIBVY		cytoplasm
Q7MBZ3	UXAC_VIBVY		cytoplasm
Q7MBZ9	UXUA_VIBVY		cytoplasm
Q7MCE1	Y5446_VIBVY		inner membrane
Q7MCF1	Y5436_VIBVY		inner membrane
Q7MCN7	SY11_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MCQ2	DEF1_VIBVY		cytoplasm
Q7MCR1	TNAA_VIBVY		cytoplasm
Q7MCS5	RIMK_VIBVY		cytoplasm
Q7MD09	Y5227_VIBVY		inner membrane
Q7MD15	LUXP_VIBVY	periplasm (probable).	periplasm
Q7MD16	LUXQ_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q7MD31	NANK_VIBVY		cytoplasm
Q7MD32	NANE_VIBVY		cytoplasm
Q7MD44	NAPA_VIBVY	periplasm (by similarity).	periplasm
Q7MD57	GUAC_VIBVY		cytoplasm
Q7MDD4	TKT2_VIBVY		extracellular

Q7MDD5	TAL_VIBVY	cytoplasm (probable).	cytoplasm
Q7MDE5	GCH1_VIBVY		cytoplasm
Q7MDF0	CLCA_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q7MDF5	PEPT_VIBVY	cytoplasm (by similarity).	extracellular
Q7MDL6	ADD_VIBVY		extracellular
Q7MDN2	PHS_VIBVY		cytoplasm
Q7ME60	COBQ_VIBVY		cytoplasm
Q7ME63	MODC_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7ME78	Y4806_VIBVY		inner membrane
Q7ME85	DDL_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7ME92	Y4778_VIBVY		inner membrane
Q7ME99	Y4771_VIBVY		cytoplasm
Q7MED4	Y4736_VIBVY	periplasm (potential).	periplasm
Q7MEE8	FTHS_VIBVY		cytoplasm
Q7MEE9	GLGC2_VIBVY		cytoplasm
Q7MEH7	GLYA2_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MEH8	GCSH_VIBVY		cytoplasm
Q7MEH9	GCSP_VIBVY		cytoplasm
Q7MEL2	ACKA2_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MEL4	RNB_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MEP0	MGSA_VIBVY		cytoplasm
Q7MEW6	Y4554_VIBVY		inner membrane
Q7MEW9	GBPA_VIBVY	secreted protein (by similarity).	cytoplasm
Q7MF00	GLPB_VIBVY		cytoplasm
Q7MF05	Y4515_VIBVY		cytoplasm
Q7MF12	BETA_VIBVY		cytoplasm
Q7MF13	BETB_VIBVY		cytoplasm
Q7MF14	BETI_VIBVY		cytoplasm
Q7MF74	SPEE_VIBVY		cytoplasm
Q7MFB0	NADE_VIBVY		cytoplasm
Q7MFB3	PYRC_VIBVY		cytoplasm
Q7MFC1	MALG_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MFC2	MALF_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MFC4	MALK_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MFE8	PSTB2_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MFG6	DEOD2_VIBVY		cytoplasm
Q7MFH3	CBIO_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MFL5	TDH_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MFM6	CATA_VIBVY	periplasm (probable).	periplasm
Q7MFN1	Y4289_VIBVY		cytoplasm
Q7MFN7	Y4283_VIBVY		inner membrane

Q7MFR0	RIBB_VIBVY		cytoplasm
Q7MFU0	RUMB_VIBVY		cytoplasm
Q7MFY7	NORW_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MFY9	NORR_VIBVY		cytoplasm
Q7MG84	PCP_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MG90	GLGB_VIBVY		extracellular
Q7MG94	MALT_VIBVY		cytoplasm
Q7MGA2	ENGC2_VIBVY		cytoplasm
Q7MGC2	Y4047_VIBVY		inner membrane
Q7MGE1	NAGB_VIBVY		extracellular
Q7MGF9	Y4010_VIBVY		inner membrane
Q7MGG9	GIDA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MGH0	GIDB_VIBVY		cytoplasm
Q7MGH8	ATPA_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MGH9	ATPG_VIBVY		inner membrane
Q7MGI1	ATPE_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MGI2	GLMU_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MGI8	ILVD_VIBVY		cytoplasm
Q7MGK4	RSMB_VIBVY		cytoplasm
Q7MGK5	FMT_VIBVY		cytoplasm
Q7MGK6	DEF2_VIBVY		cytoplasm
Q7MGK9	SMG_VIBVY		cytoplasm
Q7MGL1	PURK_VIBVY		cytoplasm
Q7MGL2	PUR6_VIBVY		cytoplasm
Q7MGL4	HEM6_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MGL5	AROE_VIBVY		cytoplasm
Q7MGM2	CRCB_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MGM3	THIC_VIBVY		cytoplasm
Q7MGM7	THIG_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MGN5	Y3194_VIBVY		inner membrane
Q7MGP6	GPPA_VIBVY		extracellular
Q7MGP7	RHLB_VIBVY		cytoplasm
Q7MGQ8	MURB_VIBVY	cytoplasm (probable).	inner membrane
Q7MGQ9	COAA_VIBVY	cytoplasm (probable).	cytoplasm
Q7MGR5	RL1_VIBVY		cytoplasm
Q7MGR6	RL10_VIBVY		cytoplasm
Q7MGR7	RL7_VIBVY		cytoplasm
Q7MGR8	RPOB_VIBVY		cytoplasm
Q7MGS0	RPOC_VIBVY		cytoplasm
Q7MGS7	DCUP_VIBVY	cytoplasm (probable).	cytoplasm
Q7MGT4	PUR2_VIBVY		cytoplasm
Q7MGT5	PUR9_VIBVY		cytoplasm
Q7MGT8	FIS_VIBVY		cytoplasm
Q7MGU2	AROQ_VIBVY		cytoplasm
Q7MGU3	ACSA_VIBVY		cytoplasm
Q7MGV6	DSBD_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MGW6	K6PF_VIBVY	cytoplasm (by similarity).	cytoplasm

Q7MGX4	FRDD_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MGX5	FRDC_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MGX9	SYK3_VIBVY		cytoplasm
Q7MGY7	GPDA_VIBVY	cytoplasm (probable).	cytoplasm
Q7MGY8	SECB_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MGZ2	GPMI_VIBVY		cytoplasm
Q7MGZ5	PSD_VIBVY		cytoplasm
Q7MGZ6	ENGC1_VIBVY		cytoplasm
Q7MGZ7	ORN_VIBVY	cytoplasm (potential).	cytoplasm
Q7MH01	MUTL_VIBVY		cytoplasm
Q7MH02	HFQ_VIBVY		cytoplasm
Q7MH07	PURA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MH09	HMP_VIBVY		cytoplasm
Q7MH13	RLMB_VIBVY		cytoplasm
Q7MH14	GPH_VIBVY		cytoplasm
Q7MH19	ARGD_VIBVY	cytoplasm (probable).	cytoplasm
Q7MH24	Y3048_VIBVY		inner membrane
Q7MH33	SLYX_VIBVY		inner membrane
Q7MH37	TUSD_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MH38	TUSC_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MH40	RS12_VIBVY		cytoplasm
Q7MH41	RS7_VIBVY		cytoplasm
Q7MH47	TPIS_VIBVY		cytoplasm
Q7MH54	RRAA2_VIBVY		cytoplasm
Q7MH56	HSLU_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MH57	HSLV_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MH61	RL31_VIBVY		cytoplasm
Q7MH63	METJ_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MH68	CAPP_VIBVY		cytoplasm
Q7MH69	ARGE_VIBVY	cytoplasm (probable).	extracellular
Q7MH70	ARGC_VIBVY	cytoplasm (probable).	cytoplasm
Q7MH71	ARGB_VIBVY	cytoplasm (probable).	cytoplasm
Q7MH72	ASSY_VIBVY	cytoplasm (probable).	cytoplasm
Q7MH73	ARLY_VIBVY	cytoplasm (probable).	cytoplasm
Q7MH83	AROK_VIBVY	cytoplasm (probable).	cytoplasm
Q7MH84	AROB_VIBVY	cytoplasm (probable).	cytoplasm
Q7MH88	RS6_VIBVY		cytoplasm
Q7MH89	PRIB_VIBVY		cytoplasm
Q7MH90	RS18_VIBVY		cytoplasm
Q7MH91	RL9_VIBVY		cytoplasm
Q7MH97	G6PI_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHA5	CYSJ_VIBVY		cytoplasm
Q7MHA6	CYSI_VIBVY		cytoplasm
Q7MHA7	CYSH_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHB1	METH_VIBVY		cytoplasm
Q7MHB5	UVRA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHE0	Y2931_VIBVY	cytoplasm (potential).	inner membrane
Q7MHE6	RAPA_VIBVY		cytoplasm
Q7MHF0	PYRI_VIBVY		cytoplasm
Q7MHF1	PYRB_VIBVY		cytoplasm
Q7MHF2	OTC_VIBVY	cytoplasm (probable).	cytoplasm
Q7MHF3	ARCA_VIBVY	cytoplasm (by similarity).	extracellular

Q7MHG1	SYV_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHG4	AMPA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHH4	Y2897_VIBVY		inner membrane
Q7MHI4	FETP_VIBVY		cytoplasm
Q7MHI6	TRMB_VIBVY		cytoplasm
Q7MHI7	GLSA_VIBVY		extracellular
Q7MHJ0	HAM1_VIBVY		extracellular
Q7MHJ2	Y2877_VIBVY		inner membrane
Q7MHJ8	RUVX_VIBVY	cytoplasm (potential).	cytoplasm
Q7MHK0	Y2869_VIBVY		inner membrane
Q7MHK1	GSHB_VIBVY		cytoplasm
Q7MHK4	SPRT_VIBVY	cytoplasm (potential).	extracellular
Q7MHK7	TKT1_VIBVY		extracellular
Q7MHL1	PGK_VIBVY	cytoplasm (potential).	cytoplasm
Q7MHL5	ICIA_VIBVY		cytoplasm
Q7MHL9	RPIA_VIBVY		cytoplasm
Q7MHM2	Y2847_VIBVY		inner membrane
Q7MHM8	Y2841_VIBVY		cytoplasm
Q7MHN6	LEPA_VIBVY		cytoplasm
Q7MHN8	RNC_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHN9	ERA_VIBVY		cytoplasm
Q7MHP0	RECO_VIBVY		cytoplasm
Q7MHP1	PDXJ_VIBVY	cytoplasm (probable).	cytoplasm
Q7MHP2	ACPS_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHP7	RUMA_VIBVY		cytoplasm
Q7MHQ0	PYRG_VIBVY		cytoplasm
Q7MHQ1	ENO_VIBVY	cytoplasm (by similarity).	extracellular
Q7MHQ2	FTSB_VIBVY	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q7MHQ4	ISPD_VIBVY		cytoplasm
Q7MHQ5	ISPF_VIBVY		extracellular
Q7MHQ6	TRUD_VIBVY		cytoplasm
Q7MHQ7	SURE_VIBVY	cytoplasm (potential).	periplasm
Q7MHQ8	PIMT_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHR2	MUTS_VIBVY		cytoplasm
Q7MHR4	RECA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHR5	RECX_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHR8	CSRA_VIBVY		cytoplasm
Q7MHR9	OADG_VIBVY	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q7MHS5	GSH1_VIBVY		cytoplasm
Q7MHS7	LUXS_VIBVY		extracellular
Q7MHT1	RS16_VIBVY		cytoplasm
Q7MHT2	RIMM_VIBVY	cytoplasm (potential).	cytoplasm
Q7MHT3	RL19_VIBVY		cytoplasm
Q7MHT4	Y2784_VIBVY		cytoplasm
Q7MHT5	COAE_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MHV3	PANC_VIBVY		cytoplasm
Q7MHV4	PANB_VIBVY		cytoplasm
Q7MHV7	GLUQ_VIBVY		cytoplasm
Q7MHV9	SFSA_VIBVY		cytoplasm
Q7MHW8	Y2750_VIBVY		inner membrane

Q7MHY9	GSA_VIBVY	cytoplasm (potential).	cytoplasm
Q7MHZ4	SYI2_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MI01	RRMJ_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MI04	GLMM_VIBVY		cytoplasm
Q7MI07	Y2710_VIBVY		inner membrane
Q7MI10	RBFA_VIBVY	cytoplasm (potential).	cytoplasm
Q7MI11	TRUB_VIBVY		cytoplasm
Q7MI12	RS15_VIBVY		cytoplasm
Q7MI38	DEOC2_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MI39	TYPH_VIBVY		cytoplasm
Q7MI40	DEOB_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MI41	DEOD1_VIBVY		cytoplasm
Q7MI80	GAL1_VIBVY	cytoplasm (potential).	cytoplasm
Q7MI93	GLPK_VIBVY		cytoplasm
Q7MIA6	ARGA_VIBVY	cytoplasm (probable).	cytoplasm
Q7MIC7	NQRA_VIBVY		inner membrane
Q7MIC8	NQRB_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q7MIC9	NQRC_VIBVY	bacterial cell inner membrane (potential).	inner membrane
Q7MID0	NQRD_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MID1	NQRE_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MID2	NQRF_VIBVY	bacterial cell inner membrane (potential).	inner membrane
Q7MID6	DPO4_VIBVY	cytoplasm (probable).	cytoplasm
Q7MIE3	Y2574_VIBVY		inner membrane
Q7MIE7	RL31B_VIBVY		cytoplasm
Q7MIF1	Y2566_VIBVY		inner membrane
Q7MIF7	Y2560_VIBVY		inner membrane
Q7MIF8	GLND_VIBVY		cytoplasm
Q7MIG0	RS2_VIBVY		cytoplasm
Q7MIG2	PYRH_VIBVY		cytoplasm
Q7MIG3	RRF_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MIG4	UPPS_VIBVY		inner membrane
Q7MIG6	DXR_VIBVY		cytoplasm
Q7MIH0	LPXD_VIBVY		cytoplasm
Q7MIH1	LPXA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MIH2	LPXB_VIBVY		cytoplasm
Q7MIH3	RNH2_VIBVY	cytoplasm (potential).	extracellular
Q7MIH6	ACCA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MIH8	TILS_VIBVY	cytoplasm (probable).	cytoplasm
Q7MII1	Y2535_VIBVY	cytoplasm (potential).	inner membrane
Q7MII6	RNH_VIBVY	cytoplasm (potential).	extracellular
Q7MIK1	PUR5_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MIK2	UPP_VIBVY		cytoplasm
Q7MIL2	DAPA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MIR0	CCMA_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q7MIR4	CCME_VIBVY	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q7MIS4	FADI_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MIS5	FADJ_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MIS9	Y2436_VIBVY		inner membrane
Q7MIT1	AROC_VIBVY		cytoplasm
Q7MIT5	Y2430_VIBVY		cytoplasm
Q7MIT6	PDXB_VIBVY		cytoplasm
Q7MIV3	Y2410_VIBVY		inner membrane
Q7MIV4	RECR_VIBVY		cytoplasm
Q7MIV9	AQPZ_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MIW1	Y2402_VIBVY		inner membrane
Q7MIZ4	Y2369_VIBVY		inner membrane
Q7MJ01	LOLD_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MJ07	MSBA_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MJ08	LPXK_VIBVY		cytoplasm
Q7MJ10	KDSB_VIBVY	cytoplasm (potential).	cytoplasm
Q7MJ13	Y2350_VIBVY		inner membrane
Q7MJ16	Y2347_VIBVY		inner membrane
Q7MJ36	Y2327_VIBVY		inner membrane
Q7MJ40	CLPS_VIBVY		cytoplasm
Q7MJ43	ATE_VIBVY	cytoplasm (potential).	cytoplasm
Q7MJ45	AROA_VIBVY	cytoplasm (probable).	cytoplasm
Q7MJ49	GLGC1_VIBVY		cytoplasm
Q7MJ50	GLGA_VIBVY		cytoplasm
Q7MJ58	TORD_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MJ62	MUKF_VIBVY	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q7MJ75	RUVC_VIBVY		cytoplasm
Q7MJ77	RUVA_VIBVY		cytoplasm
Q7MJ78	RUVB_VIBVY		cytoplasm
Q7MJ87	TOLB_VIBVY	periplasm (potential).	periplasm
Q7MJ90	NADA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MJC0	MAO12_VIBVY		cytoplasm
Q7MJI2	FUCI_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MJL8	Y2143_VIBVY		cytoplasm
Q7MJM6	METE_VIBVY		cytoplasm
Q7MJM8	ACKA1_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MJM9	Y2132_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MJN7	DSBB_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MJP0	FADR_VIBVY	cytoplasm (potential).	cytoplasm

Q7MJT4	Y2076_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MK12	ACPD_VIBVY		extracellular
Q7MK21	LIFO_VIBVY	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
Q7MK24	SPEA_VIBVY		cytoplasm
Q7MK41	PNCB_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MK44	IHFA_VIBVY		cytoplasm
Q7MK46	TPMT_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MK48	CDD_VIBVY		cytoplasm
Q7MK58	HUTH_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MK59	HUTU_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MK61	HUTI_VIBVY	cytoplasm (potential).	cytoplasm
Q7MK67	RL35_VIBVY		cytoplasm
Q7MK68	RL20_VIBVY		cytoplasm
Q7MKP8	NORM_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MKR2	MOBA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MKT7	NPD_VIBVY	cytoplasm (probable).	cytoplasm
Q7MKW4	Y1664_VIBVY		inner membrane
Q7MKX3	PYRD_VIBVY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MKY1	PROQ_VIBVY	cytoplasm (potential).	inner membrane
Q7MKZ1	Y1636_VIBVY		inner membrane
Q7MLD5	META_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLE6	BTUD_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MLE7	BTUC_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MLE8	ASTE_VIBVY		periplasm
Q7MLF2	COBS_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MLF3	COBT_VIBVY		cytoplasm
Q7MLF6	Y1471_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MLF9	PUR7_VIBVY		cytoplasm
Q7MLG3	MAO11_VIBVY		cytoplasm
Q7MLH6	SERC_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLI9	Y1438_VIBVY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7MLJ0	Y1437_VIBVY		cytoplasm
Q7MLK7	RL25_VIBVY		cytoplasm
Q7MLM1	HCP_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLQ2	TORA_VIBVY	periplasm (by similarity).	periplasm
Q7MLR4	KITH_VIBVY	cytoplasm (potential).	cytoplasm
Q7MLR7	LPXH_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLR9	Y1358_VIBVY		inner membrane

Q7MLS0	HIS2_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLS1	HIS61_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLS2	HIS4_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLS3	HIS51_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLS4	HIS7_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLS5	HIS8_VIBVY		cytoplasm
Q7MLS6	HISX_VIBVY		cytoplasm
Q7MLS7	HIS1_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLT4	TRMU_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLT5	Y1342_VIBVY		inner membrane
Q7MLU5	HTPX_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q7MLU7	BIOD_VIBVY		cytoplasm
Q7MLV4	LOLA_VIBVY	periplasm (by similarity).	periplasm
Q7MLX2	PYRF_VIBVY		cytoplasm
Q7MLX5	IHFB_VIBVY		cytoplasm
Q7MLX7	KCY_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MLY4	DNAE2_VIBVY	cytoplasm (probable).	cytoplasm
Q7MLZ6	KTHY_VIBVY		cytoplasm
Q7MLZ9	ACP_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MM01	PLSX_VIBVY		cytoplasm
Q7MM02	RL32_VIBVY		cytoplasm
Q7MM04	Y1269_VIBVY	cytoplasm (potential).	inner membrane
Q7MM12	URK_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MM27	UBIG_VIBVY		cytoplasm
Q7MM39	UVRC_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MM44	EFPL_VIBVY		cytoplasm
Q7MM45	Y1228_VIBVY		inner membrane
Q7MM54	TRPD_VIBVY		cytoplasm
Q7MM56	TRPB_VIBVY		cytoplasm
Q7MM57	TRPA_VIBVY		cytoplasm
Q7MM60	ISPZ_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MM73	MOAC_VIBVY		cytoplasm
Q7MM75	MOAA_VIBVY		cytoplasm
Q7MM77	LUXU_VIBVY		inner membrane
Q7MM78	LUXO_VIBVY		cytoplasm
Q7MM80	UVRB_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MM81	RNFA_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MM84	RNFD_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q7MM85	RNFG_VIBVY	bacterial cell inner membrane (potential).	inner membrane
Q7MM86	RNFE_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MM91	RNT_VIBVY		cytoplasm
Q7MMA6	Y1166_VIBVY		inner membrane
Q7MMC4	MSRB_VIBVY		cytoplasm

Q7MMF0	Y1122_VIBVY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7MMF9	Y1113_VIBVY		cytoplasm
Q7MMG6	CLPX_VIBVY		cytoplasm
Q7MMG7	CLPP_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MMG8	TIG_VIBVY		cytoplasm
Q7MMI2	Y1089_VIBVY		inner membrane
Q7MML0	MINC_VIBVY		inner membrane
Q7MML2	MINE_VIBVY		inner membrane
Q7MMN3	SUCC_VIBVY		cytoplasm
Q7MMR4	HEMH_VIBVY	cytoplasm (by similarity).	extracellular
Q7MMR5	KAD_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MMR7	HTPG_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MMT3	Y984_VIBVY		inner membrane
Q7MMT6	ZIPA_VIBVY	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q7MMV0	FLGI_VIBVY	periplasm (by similarity).	periplasm
Q7MMV1	FLGH_VIBVY	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q7MMW9	Y948_VIBVY		inner membrane
Q7MMY2	KDSA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MMY7	HEM1_VIBVY		cytoplasm
Q7MMY8	LOLB_VIBVY	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q7MMZ0	ISPE_VIBVY		cytoplasm
Q7MMZ1	KPRS_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MMZ2	PTH_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MN02	Y915_VIBVY		inner membrane
Q7MN04	LNT_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MN10	Y907_VIBVY		inner membrane
Q7MN15	Y902_VIBVY		inner membrane
Q7MN16	LIPB_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MN17	LIPA_VIBVY	cytoplasm (potential).	cytoplasm
Q7MN19	GLYA1_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MN30	QUEF_VIBVY	cytoplasm (probable).	cytoplasm
Q7MN36	MTFA_VIBVY		cytoplasm
Q7MN44	THII_VIBVY	cytoplasm (potential).	cytoplasm
Q7MN47	EX7S_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MN49	DXS_VIBVY		cytoplasm
Q7MN52	NUSB_VIBVY		cytoplasm
Q7MN53	RISB_VIBVY		cytoplasm
Q7MN57	NRDR_VIBVY		cytoplasm
Q7MN58	PROA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MN59	PROB_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MN61	Y856_VIBVY		cytoplasm
Q7MN62	XGPT_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MN84	DNAJ_VIBVY	cytoplasm (probable).	cytoplasm
Q7MN85	DNAK_VIBVY		cytoplasm
Q7MN92	GRPE_VIBVY	cytoplasm (probable).	cytoplasm
Q7MN93	PPNK_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MN97	Y820_VIBVY		inner membrane
Q7MN99	SSRP_VIBVY	cytoplasm (potential).	cytoplasm

Q7MND7	DADA_VIBVY		cytoplasm
Q7MNE1	GUAA_VIBVY		cytoplasm
Q7MNE3	EX7L_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MNE7	ENGA_VIBVY		cytoplasm
Q7MNF1	ISPG_VIBVY		cytoplasm
Q7MNF4	NDK_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MNF5	PEPB_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MNF8	HSCA_VIBVY		cytoplasm
Q7MNF9	HSCB_VIBVY		inner membrane
Q7MNG2	ISCS_VIBVY		cytoplasm
Q7MNH1	TGT_VIBVY		cytoplasm
Q7MNH2	QUEA_VIBVY	cytoplasm (potential).	cytoplasm
Q7MNH9	CUTC_VIBVY	cytoplasm (probable).	cytoplasm
Q7MNI7	PSTB1_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MNJ0	PPK_VIBVY		cytoplasm
Q7MNJ5	RDGC_VIBVY	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q7MNK1	CLPB_VIBVY	cytoplasm (probable).	cytoplasm
Q7MNK7	Y708_VIBVY		inner membrane
Q7MNK8	TRPR_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MNM5	ISPH_VIBVY		cytoplasm
Q7MNM7	LSPA_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q7MNM8	SYI_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MNN1	RS20_VIBVY		cytoplasm
Q7MNN6	TYSY_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MNN7	LGT_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MNP0	NUDH_VIBVY		cytoplasm
Q7MNP1	MUTH_VIBVY	cytoplasm (by similarity).	extracellular
Q7MNP0	XERD_VIBVY	cytoplasm (by similarity).	nucleoid
Q7MNP6	Y659_VIBVY		inner membrane
Q7MNR0	UNG_VIBVY	cytoplasm (by similarity).	extracellular
Q7MNR1	END4_VIBVY		extracellular
Q7MNR2	GRCA_VIBVY		cytoplasm
Q7MNR4	KHSE_VIBVY	cytoplasm (potential).	cytoplasm
Q7MNR7	RRAA1_VIBVY		cytoplasm
Q7MNT2	BTUF_VIBVY	periplasm (by similarity).	periplasm
Q7MNU0	CARB_VIBVY		cytoplasm
Q7MNU1	CARA_VIBVY		cytoplasm
Q7MNU2	DAPB_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MNU6	LPXC_VIBVY		cytoplasm
Q7MNV0	MURC_VIBVY	cytoplasm (probable).	cytoplasm
Q7MNV1	MURG_VIBVY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MNV3	MURD_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MNV4	MRAY_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MNV6	MURE_VIBVY	cytoplasm (probable).	cytoplasm

Q7MNV9	MRAW_VIBVY		cytoplasm
Q7MNW2	Y603_VIBVY		inner membrane
Q7MNX0	RS9_VIBVY		cytoplasm
Q7MNY3	HLDE_VIBVY		cytoplasm
Q7MNY4	GLNE_VIBVY		cytoplasm
Q7MNZ4	UPPP_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MNZ7	Y567_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MP00	RS21_VIBVY		cytoplasm
Q7MP37	DEOC1_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MP60	MTLD_VIBVY		cytoplasm
Q7MP62	GLMS_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MP77	LEU1_VIBVY		cytoplasm
Q7MP78	LEU3_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MP79	LEU2_VIBVY		cytoplasm
Q7MP80	LEUD_VIBVY		cytoplasm
Q7MP82	DJLA_VIBVY	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q7MP83	OSTA_VIBVY	bacterial cell outer membrane (by similarity).	outer membrane
Q7MP85	PDXA_VIBVY	cytoplasm (probable).	cytoplasm
Q7MP86	KSGA_VIBVY		cytoplasm
Q7MP87	APAG_VIBVY		inner membrane
Q7MP88	APAH_VIBVY		extracellular
Q7MP93	RL27_VIBVY		cytoplasm
Q7MP97	MDH_VIBVY		cytoplasm
Q7MP98	ARGR_VIBVY	cytoplasm (probable).	cytoplasm
Q7MPA3	MURA_VIBVY	cytoplasm (probable).	inner membrane
Q7MPB9	Y445_VIBVY		cytoplasm
Q7MPC4	Y440_VIBVY		cytoplasm
Q7MPD2	IPYR_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MPF0	CYSC_VIBVY		cytoplasm
Q7MPF2	CYSN_VIBVY		cytoplasm
Q7MPF3	CYSD_VIBVY		cytoplasm
Q7MPG4	RPOA_VIBVY		cytoplasm
Q7MPG5	RS4_VIBVY		cytoplasm
Q7MPG6	RS11_VIBVY		cytoplasm
Q7MPG7	RS13_VIBVY		cytoplasm
Q7MPG9	RL15_VIBVY		cytoplasm
Q7MPH1	RS5_VIBVY		cytoplasm
Q7MPH2	RL18_VIBVY		cytoplasm
Q7MPH4	RS8_VIBVY		cytoplasm
Q7MPH6	RL5_VIBVY		cytoplasm
Q7MPH7	RL24_VIBVY		cytoplasm
Q7MPH9	RS17_VIBVY		cytoplasm
Q7MPI0	RL29_VIBVY		cytoplasm
Q7MPI1	RL16_VIBVY		cytoplasm
Q7MPI2	RS3_VIBVY		cytoplasm
Q7MPI3	RL22_VIBVY		cytoplasm
Q7MPI4	RS19_VIBVY		cytoplasm
Q7MPI5	RL2_VIBVY		cytoplasm
Q7MPI7	RL4_VIBVY		cytoplasm

Q7MPI8	RL3_VIBVY		cytoplasm
Q7MPI9	RS10_VIBVY		cytoplasm
Q7MPK7	HIS63_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MPK8	HIS53_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MPN6	HLDD_VIBVY		cytoplasm
Q7MPP3	HIS62_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MPP4	HIS52_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MPR5	KDKA_VIBVY	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q7MPS0	COAD_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MPS3	FPG_VIBVY		cytoplasm
Q7MPS5	RL33_VIBVY		cytoplasm
Q7MPS6	RL28_VIBVY		cytoplasm
Q7MPS7	RADC_VIBVY		cytoplasm
Q7MPS9	COABC_VIBVY		cytoplasm
Q7MPT0	SLMA_VIBVY	nucleoid-associated (by similarity).	cytoplasm
Q7MPT2	PYRE_VIBVY		cytoplasm
Q7MPT3	RNPH_VIBVY		cytoplasm
Q7MPW9	KGUA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MPX0	RPOZ_VIBVY		cytoplasm
Q7MPX6	GREB_VIBVY		cytoplasm
Q7MPY0	BIOH_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MPY4	Y227_VIBVY		inner membrane
Q7MQ02	HSLO_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MQ03	PPCK_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MQ06	DTD_VIBVY	cytoplasm (probable).	cytoplasm
Q7MQ07	RBN_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MQ19	Y191_VIBVY		inner membrane
Q7MQ23	ENGB_VIBVY		cytoplasm
Q7MQ29	TATB_VIBVY	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q7MQ30	TATA_VIBVY	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q7MQ31	UBIB_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MQ33	UBIE_VIBVY		cytoplasm
Q7MQ40	MURI_VIBVY		cytoplasm
Q7MQ42	BTUB_VIBVY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q7MQ79	TRMA_VIBVY		cytoplasm
Q7MQ83	STHA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MQ85	LEXA_VIBVY		cytoplasm
Q7MQ86	PLSB_VIBVY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MQ91	GLPE_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MQB9	XERC_VIBVY	cytoplasm (by similarity).	nucleoid
Q7MQC1	DAPF_VIBVY	cytoplasm (by similarity).	cytoplasm

Q7MQC4	CYAY_VIBVY		inner membrane
Q7MQC7	HEM3_VIBVY		cytoplasm
Q7MQD2	USPB_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MQD7	Y071_VIBVY		inner membrane
Q7MQH3	ILVC_VIBVY		cytoplasm
Q7MQH9	FADB_VIBVY		cytoplasm
Q7MQI0	FADA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MQI4	TUSA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MQJ5	RECF_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MQJ7	DNAA_VIBVY	cytoplasm (by similarity).	cytoplasm
Q7MQK4	RNPA_VIBVY		extracellular
Q7MQK5	OXAA_VIBVY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MQK6	TRME_VIBVY		cytoplasm
Q7MQK8	THYX_WOLSU		cytoplasm
Q7MQL4	Y2204_WOLSU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7MQP3	LUXS_WOLSU		extracellular
Q7MQS0	HISZ_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MQU0	Y2018_WOLSU		inner membrane
Q7MQU1	RBFA_WOLSU	cytoplasm (potential).	cytoplasm
Q7MQW9	ISPDF_WOLSU		cytoplasm
Q7MQZ8	RS6_WOLSU		cytoplasm
Q7MQZ9	RS18_WOLSU		cytoplasm
Q7MR19	HAM1_WOLSU		extracellular
Q7MR31	TILS_WOLSU	cytoplasm (probable).	cytoplasm
Q7MR39	Y1748_WOLSU		cytoplasm
Q7MR67	PPNK_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MRJ7	Y1287_WOLSU		inner membrane
Q7MRK6	LOLA_WOLSU	periplasm (by similarity).	periplasm
Q7MRQ2	CRCB_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MRV4	Y1016_WOLSU		inner membrane
Q7MRX1	Y973_WOLSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MS14	CSRA_WOLSU		cytoplasm
Q7MS24	ECTC_WOLSU		cytoplasm
Q7MS42	HYP_A_WOLSU		cytoplasm
Q7MSA5	RS21_WOLSU		cytoplasm
Q7MSD5	GATC_WOLSU		cytoplasm
Q7MSD7	Y555_WOLSU		inner membrane
Q7MSD9	RS16_WOLSU		cytoplasm
Q7MSF4	ATPE_WOLSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MSH1	RL31_WOLSU		cytoplasm
Q7MSI8	Y399_WOLSU		inner membrane
Q7MSL3	CLPS_WOLSU		cytoplasm
Q7MSQ0	RPOZ_WOLSU		cytoplasm
Q7MSS6	MRAW_WOLSU		cytoplasm

Q7MSS9	Y164_WOLSU		inner membrane
Q7MST2	Y156_WOLSU		inner membrane
Q7MST9	TRMB_WOLSU		cytoplasm
Q7MSU1	Y139_WOLSU		inner membrane
Q7MSY1	QUEF_WOLSU	cytoplasm (probable).	cytoplasm
Q7MSY2	DNAA_WOLSU	cytoplasm (by similarity).	cytoplasm
Q7MXZ3	Y4907_PHOLL		inner membrane
Q7MY02	SELA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MY03	FDHE_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MY11	FDHD_PHOLL	cytoplasm (potential).	cytoplasm
Q7MY25	PYRE_PHOLL		cytoplasm
Q7MY26	SLMA_PHOLL	nucleoid-associated (by similarity).	cytoplasm
Q7MY28	RADC_PHOLL		cytoplasm
Q7MY29	RL28_PHOLL		cytoplasm
Q7MY36	FPG_PHOLL		cytoplasm
Q7MY37	COAD_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MY46	HLDD_PHOLL		cytoplasm
Q7MY48	TDH_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MY53	SECB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MY54	GPDA_PHOLL	cytoplasm (probable).	cytoplasm
Q7MY63	GMHB2_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MYB3	TPIS_PHOLL		cytoplasm
Q7MYB6	GLPK_PHOLL		cytoplasm
Q7MYB9	RRAA_PHOLL		cytoplasm
Q7MYC1	HSLU_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MYC2	HSLV_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MYC6	RL31_PHOLL		cytoplasm
Q7MYC7	METJ_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MYD5	ARGE_PHOLL	cytoplasm (probable).	extracellular
Q7MYD6	ARGC_PHOLL	cytoplasm (probable).	cytoplasm
Q7MYD7	ARGB_PHOLL	cytoplasm (probable).	cytoplasm
Q7MYD8	ASSY_PHOLL	cytoplasm (probable).	cytoplasm
Q7MYD9	ARLY_PHOLL	cytoplasm (probable).	cytoplasm
Q7MYE3	BTUB_PHOLL	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q7MYE5	MURB_PHOLL	cytoplasm (probable).	inner membrane
Q7MYE7	COAA_PHOLL	cytoplasm (probable).	cytoplasm
Q7MYF1	RL3_PHOLL		cytoplasm
Q7MYF2	RL4_PHOLL		cytoplasm
Q7MYF4	RL2_PHOLL		cytoplasm
Q7MYF5	RS19_PHOLL		cytoplasm
Q7MYF6	RL22_PHOLL		cytoplasm
Q7MYF7	RS3_PHOLL		cytoplasm
Q7MYF8	RL16_PHOLL		cytoplasm
Q7MYF9	RL29_PHOLL		cytoplasm
Q7MYG0	RS17_PHOLL		cytoplasm
Q7MYG2	RL24_PHOLL		cytoplasm
Q7MYG3	RL5_PHOLL		cytoplasm
Q7MYG5	RS8_PHOLL		cytoplasm
Q7MYG7	RL18_PHOLL		cytoplasm
Q7MYG8	RS5_PHOLL		cytoplasm
Q7MYH0	RL15_PHOLL		cytoplasm
Q7MYH2	RS13_PHOLL		cytoplasm
Q7MYH3	RS11_PHOLL		cytoplasm

Q7MYH4	RS4_PHOLL		cytoplasm
Q7MYH5	RPOA_PHOLL		cytoplasm
Q7MYI0	RSMB_PHOLL		cytoplasm
Q7MYI1	FMT_PHOLL		cytoplasm
Q7MYI2	DEF_PHOLL		cytoplasm
Q7MYJ5	ILVD_PHOLL		cytoplasm
Q7MYK9	ILVC_PHOLL		cytoplasm
Q7MYL1	GPPA_PHOLL		extracellular
Q7MYL2	RHLB_PHOLL		cytoplasm
Q7MYM4	WECF_PHOLL	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q7MYM5	WZYE_PHOLL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7MYM6	WECG_PHOLL		cytoplasm
Q7MYN1	HEM3_PHOLL		cytoplasm
Q7MYN3	CYAY_PHOLL		inner membrane
Q7MYP0	CORA_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MYS6	PSD_PHOLL		cytoplasm
Q7MYS8	ORN_PHOLL	cytoplasm (potential).	cytoplasm
Q7MYU0	HFQ_PHOLL		cytoplasm
Q7MYU6	RLMB_PHOLL		cytoplasm
Q7MYU7	RS6_PHOLL		cytoplasm
Q7MYU8	PRIB_PHOLL		cytoplasm
Q7MYU9	RL9_PHOLL		cytoplasm
Q7MYW1	MSRA_PHOLL		cytoplasm
Q7MYW8	ARGR_PHOLL	cytoplasm (probable).	cytoplasm
Q7MYW9	MDH_PHOLL		cytoplasm
Q7MYX5	RL27_PHOLL		cytoplasm
Q7MYY0	RRMJ_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MYY3	GLMM_PHOLL		cytoplasm
Q7MYY5	Y4531_PHOLL		inner membrane
Q7MYY8	RBFA_PHOLL	cytoplasm (potential).	cytoplasm
Q7MYY9	RS15_PHOLL		cytoplasm
Q7MZ08	Y4503_PHOLL		cytoplasm
Q7MZ10	Y4501_PHOLL		inner membrane
Q7MZ14	PYRI_PHOLL		cytoplasm
Q7MZ16	PYRB_PHOLL		cytoplasm
Q7MZ18	OTC_PHOLL	cytoplasm (probable).	cytoplasm
Q7MZ25	SYV_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MZ27	AMPA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MZ74	METE_PHOLL		cytoplasm
Q7MZ76	PCP2_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MZ81	UBIE_PHOLL		cytoplasm
Q7MZ83	UBIB_PHOLL	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q7MZ91	FADA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MZ92	FADB_PHOLL		cytoplasm
Q7MZ97	META_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MZA0	ACEK_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7MZB4	G6PI_PHOLL	cytoplasm (by similarity).	cytoplasm

Q7MZB7	PLSB_PHOLL	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7MZB8	LEXA_PHOLL		cytoplasm
Q7MZI3	Y4304_PHOLL		inner membrane
Q7MZY0	SUGE_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7MZY6	SYK3_PHOLL		cytoplasm
Q7MZZ5	TUSA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N015	FIS_PHOLL		cytoplasm
Q7N030	AROQ_PHOLL		cytoplasm
Q7N035	Y4068_PHOLL	cytoplasm (potential).	inner membrane
Q7N042	Y4061_PHOLL		cytoplasm
Q7N053	Y4044_PHOLL		cytoplasm
Q7N068	MURA_PHOLL	cytoplasm (probable).	inner membrane
Q7N079	RS9_PHOLL		cytoplasm
Q7N087	MTGA_PHOLL	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q7N089	DIAA_PHOLL		cytoplasm
Q7N0A6	IDI2_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N0B7	Y3975_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N0B9	UPPP_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N0C2	GLNE_PHOLL		cytoplasm
Q7N0C3	HLDE_PHOLL		cytoplasm
Q7N0C8	RIBB_PHOLL		cytoplasm
Q7N0I2	ACPH_PHOLL		cytoplasm
Q7N0I3	QUEA_PHOLL	cytoplasm (potential).	cytoplasm
Q7N0I7	NRDR_PHOLL		cytoplasm
Q7N0I9	RISB_PHOLL		cytoplasm
Q7N0J0	NUSB_PHOLL		cytoplasm
Q7N0J7	DXS_PHOLL		cytoplasm
Q7N0J8	EX7S_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N0J9	THII_PHOLL	cytoplasm (potential).	cytoplasm
Q7N0K1	Y3881_PHOLL		inner membrane
Q7N0L2	TIG_PHOLL		cytoplasm
Q7N0L3	CLPP_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N0L4	CLPX_PHOLL		cytoplasm
Q7N0N3	CBIO_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7N0P1	Y3840_PHOLL		inner membrane
Q7N0P2	RECR_PHOLL		cytoplasm
Q7N0P4	HTPG_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N0P5	KAD_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N0P6	HEMH_PHOLL	cytoplasm (by similarity).	extracellular
Q7N0S3	LPXH_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N118	SPRT_PHOLL	cytoplasm (potential).	extracellular
Q7N121	SPEA_PHOLL		cytoplasm
Q7N122	SPEB_PHOLL		cytoplasm
Q7N124	MUTH_PHOLL	cytoplasm (by similarity).	extracellular

Q7N126	LEUD_PHOLL		cytoplasm
Q7N127	LEU2_PHOLL		cytoplasm
Q7N128	LEU3_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N129	LEU1_PHOLL		cytoplasm
Q7N138	MRAZ_PHOLL		inner membrane
Q7N139	MRAW_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N142	MURE_PHOLL	cytoplasm (probable).	cytoplasm
Q7N144	MRAY_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N145	MURD_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N147	MURG_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7N148	MURC_PHOLL	cytoplasm (probable).	cytoplasm
Q7N149	DDL_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N153	LPXC_PHOLL		cytoplasm
Q7N157	Y3643_PHOLL		cytoplasm
Q7N158	COAE_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N179	Y3616_PHOLL		inner membrane
Q7N183	ARGO_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N189	RPIA_PHOLL		cytoplasm
Q7N193	Y3602_PHOLL		inner membrane
Q7N197	GCST_PHOLL		cytoplasm
Q7N199	GCSP_PHOLL		cytoplasm
Q7N1C1	Y3555_PHOLL		cytoplasm
Q7N1G0	NORM_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N1U1	SSRP_PHOLL	cytoplasm (potential).	cytoplasm
Q7N1U3	Y3376_PHOLL		inner membrane
Q7N1U6	PPNK_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N1U7	GRPE_PHOLL	cytoplasm (probable).	cytoplasm
Q7N1U8	UNG_PHOLL	cytoplasm (by similarity).	extracellular
Q7N1V4	IDI1_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N1X3	LEPA_PHOLL		cytoplasm
Q7N1X5	RNC_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N1X7	RECO_PHOLL		cytoplasm
Q7N1X8	PDXJ_PHOLL	cytoplasm (probable).	cytoplasm
Q7N1X9	ACPS_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N215	HMP_PHOLL		cytoplasm
Q7N216	GLYA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N226	ISCA_PHOLL		cytoplasm
Q7N227	HSCB_PHOLL		inner membrane
Q7N228	HSCA_PHOLL		cytoplasm
Q7N231	PEPB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N287	FADI_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N288	FADJ_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N290	Y3198_PHOLL		cytoplasm
Q7N295	HUTU_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N296	HUTH_PHOLL	cytoplasm (potential).	cytoplasm
Q7N299	AROC_PHOLL		cytoplasm
Q7N2A3	Y3185_PHOLL		cytoplasm
Q7N2B2	PDXB_PHOLL		cytoplasm

Q7N2G7	ARGM_PHOLL		cytoplasm
Q7N2H1	ASTE_PHOLL		periplasm
Q7N2I1	ACKA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N2I2	Y3094_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N2I4	Y3092_PHOLL		cytoplasm
Q7N2M5	UBIG_PHOLL		cytoplasm
Q7N2S8	CBID_PHOLL		cytoplasm
Q7N2T7	COBQ_PHOLL		cytoplasm
Q7N2U4	COBT_PHOLL		cytoplasm
Q7N2V3	EUTC_PHOLL		cytoplasm
Q7N351	Y2871_PHOLL		inner membrane
Q7N352	RL25_PHOLL		cytoplasm
Q7N360	EFPL_PHOLL		cytoplasm
Q7N364	END4_PHOLL		extracellular
Q7N365	Y2856_PHOLL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7N376	Y2843_PHOLL		inner membrane
Q7N380	Y2839_PHOLL	cytoplasm (potential).	inner membrane
Q7N382	RL32_PHOLL		cytoplasm
Q7N383	PLSX_PHOLL		cytoplasm
Q7N384	FABH_PHOLL	cytoplasm (probable).	cytoplasm
Q7N391	KTHY_PHOLL		cytoplasm
Q7N397	NAGZ_PHOLL	cytoplasm (by similarity).	extracellular
Q7N3A6	LOLD_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7N3B4	Y2805_PHOLL		inner membrane
Q7N3D8	Y2779_PHOLL		cytoplasm
Q7N3E1	MDTC_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3E2	MDTB_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q7N3E3	MDTA_PHOLL	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q7N3F7	PUR5_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N3F8	UPP_PHOLL		cytoplasm
Q7N3G5	HDA_PHOLL		nucleoid
Q7N3H3	PUR7_PHOLL		cytoplasm
Q7N3K4	GUAA_PHOLL		cytoplasm
Q7N3L4	Y2701_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3L5	Y2700_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3M2	Y2693_PHOLL		inner membrane
Q7N3M3	YEBF_PHOLL	secreted protein (by similarity).	inner membrane
Q7N3N2	PROQ_PHOLL	cytoplasm (potential).	inner membrane
Q7N3P8	RL35_PHOLL		cytoplasm

Q7N3P9	RL20_PHOLL		cytoplasm
Q7N3Q3	BTUC_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3Q4	BTUD_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7N3Q5	ARNB_PHOLL		cytoplasm
Q7N3Q6	ARNC_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3Q7	ARNA_PHOLL		cytoplasm
Q7N3Q9	ARNT_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3R1	PBGE3_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3T4	Y2629_PHOLL		cytoplasm
Q7N3U5	SUFS_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N3U6	SUFE_PHOLL	cytoplasm (by similarity).	inner membrane
Q7N3V2	MDTK_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3W0	RNT_PHOLL		cytoplasm
Q7N3W5	PDXH_PHOLL		cytoplasm
Q7N3W6	SY11_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N3Y1	Y2581_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3Y5	ZNTB_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N3Z5	FADR_PHOLL	cytoplasm (potential).	cytoplasm
Q7N3Z6	DADA_PHOLL		cytoplasm
Q7N406	SELD_PHOLL		cytoplasm
Q7N451	Y2503_PHOLL		inner membrane
Q7N457	KITH_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N466	Y2488_PHOLL		inner membrane
Q7N475	Y2479_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N485	TRPA_PHOLL		cytoplasm
Q7N486	TRPB_PHOLL		cytoplasm
Q7N488	TRPD_PHOLL		cytoplasm
Q7N4C1	PYRF_PHOLL		cytoplasm
Q7N4F5	RNB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N4H5	ADD_PHOLL		extracellular
Q7N4H8	FUMC_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N4P7	XYLA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N4Y7	URE1_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N4Y8	URE2_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N4Y9	URE3_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N511	ACPD_PHOLL		extracellular
Q7N519	Y2141_PHOLL		inner membrane
Q7N522	MINC_PHOLL		inner membrane

Q7N547	RUVB_PHOLL		cytoplasm
Q7N548	RUVA_PHOLL		cytoplasm
Q7N549	RUVC_PHOLL		cytoplasm
Q7N550	Y2109_PHOLL		inner membrane
Q7N557	CUTC_PHOLL	cytoplasm (probable).	periplasm
Q7N562	Y2095_PHOLL	periplasm (potential).	periplasm
Q7N568	MDTH_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N574	LUXB_PHOLL		cytoplasm
Q7N575	LUXA_PHOLL		cytoplasm
Q7N576	LUXD_PHOLL		cytoplasm
Q7N577	LUXC_PHOLL		cytoplasm
Q7N583	KDSA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N587	HEM1_PHOLL		cytoplasm
Q7N588	LOLB_PHOLL	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q7N589	ISPE_PHOLL		cytoplasm
Q7N590	KPRS_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N5A1	PTH_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N5C1	AQPZ_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N5C7	UVRC_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N5C8	PGSA_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N5H9	SDHD_PHOLL		cytoplasm
Q7N5I5	WRBA_PHOLL		cytoplasm
Q7N5M6	FLGI_PHOLL	periplasm (by similarity).	periplasm
Q7N5M7	FLGH_PHOLL	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q7N5W3	Y1816_PHOLL		inner membrane
Q7N5Z1	TUSE_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N5Z7	MGSA_PHOLL		cytoplasm
Q7N603	Y1774_PHOLL		inner membrane
Q7N617	PYRD_PHOLL	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7N621	PNCB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6B9	MUKF_PHOLL	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q7N6C5	LPXK_PHOLL		cytoplasm
Q7N6C6	MSBA_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N6D4	KCY_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6D6	SERC_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6E9	LOLA_PHOLL	periplasm (by similarity).	periplasm
Q7N6F5	LFTR_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6F7	CLPS_PHOLL		cytoplasm
Q7N6G6	RUMB_PHOLL		cytoplasm
Q7N6I0	HIS1_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6I1	HIS8_PHOLL		cytoplasm
Q7N6I2	HIS7_PHOLL	cytoplasm (by similarity).	cytoplasm

Q7N6I3	HIS5_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6I4	HIS41_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6I5	HIS6_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6I6	HIS2_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6J3	DCD_PHOLL		extracellular
Q7N6K1	Y1549_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N6K3	CDD_PHOLL		cytoplasm
Q7N6K4	MAO1_PHOLL		cytoplasm
Q7N6K7	GCH1_PHOLL		cytoplasm
Q7N6Q1	UVRB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6R1	6PGL_PHOLL		cytoplasm
Q7N6R3	MODC_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7N6S0	GPMA_PHOLL		cytoplasm
Q7N6S3	NADA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6T6	TOLB_PHOLL	periplasm (potential).	periplasm
Q7N6W5	ATKA_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q7N6Y0	GLK_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N6Y5	ZIPA_PHOLL	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q7N6Z2	CYSA_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7N6Z9	HEM6_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N702	ENGA_PHOLL		cytoplasm
Q7N706	ISPG_PHOLL		cytoplasm
Q7N710	NDK_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N748	Y1310_PHOLL		inner membrane
Q7N750	LNT_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N760	Y1298_PHOLL		inner membrane
Q7N765	Y1293_PHOLL		inner membrane
Q7N766	LIPB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N767	LIPA_PHOLL	cytoplasm (potential).	cytoplasm
Q7N768	CRCB_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N772	NRDI_PHOLL		inner membrane
Q7N788	CLPB_PHOLL	cytoplasm (probable).	cytoplasm
Q7N798	RL19_PHOLL		cytoplasm
Q7N799	RIMM_PHOLL	cytoplasm (potential).	cytoplasm
Q7N7A0	RS16_PHOLL		cytoplasm
Q7N7A4	GSH1_PHOLL		cytoplasm
Q7N7A6	RECA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N7A9	AAS_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N7B0	AROL_PHOLL	cytoplasm (probable).	cytoplasm
Q7N7B1	PROA_PHOLL	cytoplasm (by similarity).	cytoplasm

Q7N7B2	PROB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N7B3	Y1242_PHOLL		cytoplasm
Q7N7B4	XGPT_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N7F7	GMHA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N7G6	Y1183_PHOLL		inner membrane
Q7N7G7	RUVX_PHOLL	cytoplasm (potential).	cytoplasm
Q7N7H2	HAM1_PHOLL		extracellular
Q7N7H7	GLSA_PHOLL		extracellular
Q7N7H9	TRMB_PHOLL		cytoplasm
Q7N7I1	FETP_PHOLL		cytoplasm
Q7N7I2	MLTC_PHOLL	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q7N7Z5	PGK_PHOLL	cytoplasm (potential).	cytoplasm
Q7N807	RNH_PHOLL	cytoplasm (potential).	extracellular
Q7N835	ENO_PHOLL	cytoplasm (by similarity).	extracellular
Q7N839	RUMA_PHOLL		cytoplasm
Q7N842	BTUF_PHOLL	periplasm (by similarity).	periplasm
Q7N845	GSA_PHOLL	cytoplasm (potential).	cytoplasm
Q7N864	SFSA_PHOLL		cytoplasm
Q7N866	GLUQ_PHOLL		cytoplasm
Q7N869	PANB_PHOLL		cytoplasm
Q7N870	PANC_PHOLL		cytoplasm
Q7N871	PAND_PHOLL		cytoplasm
Q7N892	SPEE_PHOLL		cytoplasm
Q7N893	SPED_PHOLL		cytoplasm
Q7N8B9	FBPC_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7N8C9	TNAA_PHOLL		cytoplasm
Q7N8D1	HIS42_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8D4	PURA1_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8K0	MUTS_PHOLL		cytoplasm
Q7N8K3	PIMT_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8K4	SURE_PHOLL	cytoplasm (potential).	periplasm
Q7N8K5	TRUD_PHOLL		cytoplasm
Q7N8K6	ISPF_PHOLL		extracellular
Q7N8K7	ISPD_PHOLL		cytoplasm
Q7N8L0	CYSN_PHOLL		cytoplasm
Q7N8L5	CYSI_PHOLL		cytoplasm
Q7N8L6	CYSJ_PHOLL		cytoplasm
Q7N8M0	Y699_PHOLL	cytoplasm (potential).	inner membrane
Q7N8M1	GMHB1_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8N0	TILS_PHOLL	cytoplasm (probable).	cytoplasm
Q7N8N1	ACCA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8N3	RNH2_PHOLL	cytoplasm (potential).	extracellular
Q7N8N4	LPXB_PHOLL		cytoplasm
Q7N8N6	FABZ_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8N7	LPXD_PHOLL		cytoplasm
Q7N8N8	SKP_PHOLL	periplasm (by similarity).	periplasm
Q7N8N9	YAET_PHOLL	bacterial cell outer membrane (by similarity).	outer membrane
Q7N8P2	UPPS_PHOLL		inner membrane
Q7N8P3	DXR_PHOLL		cytoplasm
Q7N8P4	RRF_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8P7	RS2_PHOLL		cytoplasm
Q7N8P9	GLND_PHOLL		cytoplasm

Q7N8Q0	DAPD_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8Q1	Y668_PHOLL		inner membrane
Q7N8Q7	QUEF_PHOLL	cytoplasm (probable).	cytoplasm
Q7N8R2	INH_PHOLL	periplasm (by similarity).	periplasm
Q7N8R3	PRTA_PHOLL	secreted protein (by similarity).	extracellular
Q7N8R4	MTFA_PHOLL		cytoplasm
Q7N8S4	ARGA_PHOLL	cytoplasm (probable).	cytoplasm
Q7N8U4	TYSY_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8U5	LGT_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N8U7	NUDH_PHOLL		cytoplasm
Q7N8V1	RAPA_PHOLL		cytoplasm
Q7N8V3	DJLA_PHOLL	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q7N8V4	OSTA_PHOLL	bacterial cell outer membrane (by similarity).	outer membrane
Q7N8V7	KSGA_PHOLL		cytoplasm
Q7N8V8	APAG_PHOLL		inner membrane
Q7N8V9	APAH_PHOLL		extracellular
Q7N8W2	CARA_PHOLL		cytoplasm
Q7N8W3	DAPB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8W9	ISPH_PHOLL		cytoplasm
Q7N8X2	SYI_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N8X4	RS20_PHOLL		cytoplasm
Q7N8Y3	DNAJ_PHOLL	cytoplasm (probable).	cytoplasm
Q7N8Y4	DNAK_PHOLL		cytoplasm
Q7N8Z1	TAL_PHOLL	cytoplasm (probable).	cytoplasm
Q7N8Z3	Y566_PHOLL		inner membrane
Q7N8Z5	KHSE_PHOLL	cytoplasm (potential).	cytoplasm
Q7N900	GPMB_PHOLL		cytoplasm
Q7N901	Y558_PHOLL		inner membrane
Q7N902	TRPR_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N930	DEOD_PHOLL		cytoplasm
Q7N931	DEOB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N932	DEOC_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N959	NFI_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N960	DCUP_PHOLL	cytoplasm (probable).	cytoplasm
Q7N961	NUDC_PHOLL		cytoplasm
Q7N963	THIC_PHOLL		cytoplasm
Q7N964	THIE_PHOLL		cytoplasm
Q7N966	THIG_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N976	MALT_PHOLL		cytoplasm
Q7N983	MALG_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N984	MALF_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7N986	MALK_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q7N987	LAMB_PHOLL	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q7N9A3	RPOC_PHOLL		cytoplasm
Q7N9A4	RPOB_PHOLL		cytoplasm
Q7N9A6	RL10_PHOLL		cytoplasm
Q7N9A7	RL1_PHOLL		cytoplasm
Q7N9B3	RS7_PHOLL		cytoplasm
Q7N9B4	RS12_PHOLL		cytoplasm
Q7N9B5	TUSB_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N9B6	TUSC_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N9B7	TUSD_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N9C0	SLYX_PHOLL		inner membrane
Q7N9E2	Y398_PHOLL		inner membrane
Q7N9E5	ARGD_PHOLL	cytoplasm (probable).	cytoplasm
Q7N9P2	KGUA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N9P3	RPOZ_PHOLL		cytoplasm
Q7N9S2	DTD_PHOLL	cytoplasm (probable).	cytoplasm
Q7N9T4	Y229_PHOLL		inner membrane
Q7N9U4	PSTB_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7N9V7	BIOH_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N9W0	SYI2_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N9W2	Y198_PHOLL		inner membrane
Q7N9W3	GLPE_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7N9X5	UXAC_PHOLL		cytoplasm
Q7N9Y1	UXUA_PHOLL		cytoplasm
Q7NA24	Y123_PHOLL		inner membrane
Q7NA33	CYNS_PHOLL		cytoplasm
Q7NA45	PPCK_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7NA46	HSLO_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7NA55	AROK_PHOLL	cytoplasm (probable).	cytoplasm
Q7NA56	AROB_PHOLL	cytoplasm (probable).	cytoplasm
Q7NA60	GPH_PHOLL		cytoplasm
Q7NA64	Y081_PHOLL		inner membrane
Q7NA72	ACTP_PHOLL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NA81	RAVA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7NA82	VIAA_PHOLL		cytoplasm
Q7NA83	ASNA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7NA86	GIDA_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7NA87	GIDB_PHOLL		cytoplasm
Q7NA92	ATPA_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7NA93	ATPG_PHOLL		inner membrane
Q7NA95	ATPE_PHOLL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7NA96	GLMU_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7NA97	GLMS_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7NA98	RL31B_PHOLL		cytoplasm
Q7NAD1	RECF_PHOLL	cytoplasm (by similarity).	cytoplasm
Q7NAD3	DNAA_PHOLL	cytoplasm (by similarity).	cytoplasm

Q7NPQ6	DARCY_CHRVO		cytoplasm
Q7NPS3	RNH2_CHRVO	cytoplasm (potential).	extracellular
Q7NPT6	RNPA_CHRVO		extracellular
Q7NPT7	Y4405_CHRVO		inner membrane
Q7NPT9	TRME_CHRVO		cytoplasm
Q7NPW5	BIOH_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NPY6	GATC_CHRVO		cytoplasm
Q7NPY7	GATA_CHRVO		cytoplasm
Q7NPY8	GATB_CHRVO		cytoplasm
Q7NPZ0	MRAZ_CHRVO		inner membrane
Q7NPZ1	MRAW_CHRVO		cytoplasm
Q7NPZ4	MURE_CHRVO	cytoplasm (probable).	cytoplasm
Q7NPZ6	MRAY_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NPZ7	MURD_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NPZ9	MURG_CHRVO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7NQ00	MURC_CHRVO	cytoplasm (probable).	cytoplasm
Q7NQ01	DDLB_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NQ20	Y4322_CHRVO		inner membrane
Q7NQ51	PROA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NQ65	GSHB_CHRVO		cytoplasm
Q7NQ72	SMG_CHRVO		cytoplasm
Q7NQ75	DEF_CHRVO		cytoplasm
Q7NQ76	FMT_CHRVO		cytoplasm
Q7NQ92	PYRE_CHRVO		cytoplasm
Q7NQB5	RUVC_CHRVO		cytoplasm
Q7NQB7	RUVA_CHRVO		cytoplasm
Q7NQC4	RIBB_CHRVO		cytoplasm
Q7NQC5	RUVB_CHRVO		cytoplasm
Q7NQC8	PROB_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NQE3	RL1_CHRVO		cytoplasm
Q7NQE4	RL10_CHRVO		cytoplasm
Q7NQE6	RPOB_CHRVO		cytoplasm
Q7NQE7	RPOC_CHRVO		cytoplasm
Q7NQE8	RS12_CHRVO		cytoplasm
Q7NQE9	RS7_CHRVO		cytoplasm
Q7NQF1	RS10_CHRVO		cytoplasm
Q7NQF2	RL3_CHRVO		cytoplasm
Q7NQF3	RL4_CHRVO		cytoplasm
Q7NQF5	RL2_CHRVO		cytoplasm
Q7NQF6	RS19_CHRVO		cytoplasm
Q7NQF7	RL22_CHRVO		cytoplasm
Q7NQF8	RS3_CHRVO		cytoplasm
Q7NQF9	RL16_CHRVO		cytoplasm
Q7NQG0	RL29_CHRVO		cytoplasm
Q7NQG1	RS17_CHRVO		cytoplasm
Q7NQG3	RL24_CHRVO		cytoplasm
Q7NQG4	RL5_CHRVO		cytoplasm
Q7NQG6	RS8_CHRVO		cytoplasm
Q7NQG8	RL18_CHRVO		cytoplasm
Q7NQG9	RS5_CHRVO		cytoplasm
Q7NQH1	RL15_CHRVO		cytoplasm
Q7NQH4	RS13_CHRVO		cytoplasm

Q7NQH5	RS11_CHRVO		cytoplasm
Q7NQH6	RS4_CHRVO		cytoplasm
Q7NQH7	RPOA_CHRVO		cytoplasm
Q7NQH9	GLSA_CHRVO		extracellular
Q7NQI3	LNT_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NQI5	Y4152_CHRVO		inner membrane
Q7NQG7	SPEE_CHRVO		cytoplasm
Q7NQS0	APAH_CHRVO		extracellular
Q7NQS5	FPG_CHRVO		cytoplasm
Q7NQS7	LOLB_CHRVO	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q7NQS8	ISPE_CHRVO		cytoplasm
Q7NQS9	KPRS_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NQT0	RL25_CHRVO		cytoplasm
Q7NQT1	PTH_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NQX1	CH602_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NR63	ARGB_CHRVO	cytoplasm (probable).	cytoplasm
Q7NR74	RUVX_CHRVO	cytoplasm (potential).	cytoplasm
Q7NR75	Y3909_CHRVO		inner membrane
Q7NRF7	COAE_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NRF8	Y3823_CHRVO		cytoplasm
Q7NRI3	RRMJ_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NRI6	GLMM_CHRVO		cytoplasm
Q7NRJ5	TRMB_CHRVO		cytoplasm
Q7NRJ9	ARCA_CHRVO	cytoplasm (by similarity).	extracellular
Q7NRK0	OTCC_CHRVO	cytoplasm (probable).	cytoplasm
Q7NRL1	KGUA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NRL2	RPOZ_CHRVO		cytoplasm
Q7NRL4	THIG_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NRL5	RS21_CHRVO		cytoplasm
Q7NRN0	QUEF_CHRVO	cytoplasm (probable).	cytoplasm
Q7NRN3	RS20_CHRVO		cytoplasm
Q7NRQ7	KTHY_CHRVO		cytoplasm
Q7NRS0	Y3710_CHRVO		cytoplasm
Q7NRS9	DEOC_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NRT0	TYPH_CHRVO		cytoplasm
Q7NRT1	DEOB_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NRT2	DEOD_CHRVO		cytoplasm
Q7NRT4	RS9_CHRVO		cytoplasm
Q7NRT5	ARGC_CHRVO	cytoplasm (probable).	cytoplasm
Q7NRU2	Y3688_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NRV1	SURE_CHRVO	cytoplasm (potential).	periplasm
Q7NRV5	RS16_CHRVO		cytoplasm
Q7NRV6	RIMM_CHRVO	cytoplasm (potential).	cytoplasm
Q7NRV7	RL19_CHRVO		cytoplasm
Q7NRW1	CLPS_CHRVO		cytoplasm
Q7NRW7	MUTS_CHRVO		cytoplasm
Q7NRY7	RS6_CHRVO		cytoplasm
Q7NRY9	RS18_CHRVO		cytoplasm
Q7NRZ0	RL9_CHRVO		cytoplasm
Q7NS06	UPP_CHRVO		cytoplasm
Q7NS12	PUR5_CHRVO	cytoplasm (by similarity).	cytoplasm

Q7NS23	METE_CHRVO		cytoplasm
Q7NS41	MTGA_CHRVO	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q7NS57	SYI_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NS59	ISPH_CHRVO		cytoplasm
Q7NS67	Y3559_CHRVO		inner membrane
Q7NS72	DCD_CHRVO		extracellular
Q7NS84	NDK_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NS88	ISPG_CHRVO		cytoplasm
Q7NS92	ENGA_CHRVO		cytoplasm
Q7NS93	HFQ_CHRVO		cytoplasm
Q7NS97	HISZ_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NS98	PURA2_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NSA6	ADC_CHRVO		cytoplasm
Q7NSB3	Y3513_CHRVO		inner membrane
Q7NSD8	HMP_CHRVO		cytoplasm
Q7NSE1	Y3485_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NSG1	GUAA_CHRVO		cytoplasm
Q7NSG2	SSRP_CHRVO	cytoplasm (potential).	cytoplasm
Q7NSG4	Y3462_CHRVO		inner membrane
Q7NSH0	RL28_CHRVO		cytoplasm
Q7NSJ1	GLUQ_CHRVO		cytoplasm
Q7NSJ3	GCST_CHRVO		cytoplasm
Q7NSJ5	GCSP_CHRVO		cytoplasm
Q7NSK2	PNCB_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NSK3	Y3420_CHRVO	cytoplasm (potential).	inner membrane
Q7NSK5	RL32_CHRVO		cytoplasm
Q7NSK6	PLSX_CHRVO		cytoplasm
Q7NSK7	FABH_CHRVO	cytoplasm (probable).	cytoplasm
Q7NSP5	MINC_CHRVO		inner membrane
Q7NSR4	FETP_CHRVO		cytoplasm
Q7NSR8	GPMI_CHRVO		cytoplasm
Q7NSS4	LPXK_CHRVO		cytoplasm
Q7NSS7	KAD_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NST4	Y3336_CHRVO		cytoplasm
Q7NST5	HPRK_CHRVO		cytoplasm
Q7NT31	CH601_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NT63	Y3198_CHRVO		inner membrane
Q7NT71	ACCA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NT72	TILS_CHRVO	cytoplasm (probable).	cytoplasm
Q7NT75	LPXH_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NT84	PCP_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NT87	Y3173_CHRVO		inner membrane
Q7NTA7	UVRB_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NTD5	Y3123_CHRVO		inner membrane
Q7NTF9	LIPA_CHRVO	cytoplasm (potential).	cytoplasm
Q7NTG0	LIPB_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NTG1	Y3095_CHRVO		inner membrane
Q7NTI2	SYI_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NTK7	KCY_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NTL2	PYRF_CHRVO		cytoplasm
Q7NTL6	HLDD_CHRVO		cytoplasm

Q7NTU0	LOLD_CHRVO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7NTY6	GLND_CHRVO		cytoplasm
Q7NTY9	AMPA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NTZ9	CAAL_CHRVO		cytoplasm
Q7NU21	FLGH2_CHRVO	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q7NU22	FLGI2_CHRVO	periplasm (by similarity).	periplasm
Q7NU26	ASTE_CHRVO		periplasm
Q7NU27	SPEA_CHRVO		cytoplasm
Q7NU39	AQPZ_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NU46	TAUB_CHRVO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7NUB6	LEU2_CHRVO		cytoplasm
Q7NUB8	LEUD2_CHRVO		cytoplasm
Q7NUC2	LEU3_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NUD6	TRUA_CHRVO		cytoplasm
Q7NUD8	TRPB_CHRVO		cytoplasm
Q7NUD9	TRPA_CHRVO		cytoplasm
Q7NUF2	KDSA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NUG7	KUP1_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NUI5	LIFO_CHRVO	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
Q7NUK5	DXS_CHRVO		cytoplasm
Q7NUK7	EX7S_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NUM8	SYV_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NUP9	SDHD_CHRVO		cytoplasm
Q7NUY8	TIG_CHRVO		cytoplasm
Q7NUY9	CLPP_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NUZ0	CLPX_CHRVO		cytoplasm
Q7NV65	HEMH_CHRVO	cytoplasm (by similarity).	extracellular
Q7NV72	DDLA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NVB8	KPTA_CHRVO		extracellular
Q7NVF2	NUSB_CHRVO		cytoplasm
Q7NVF3	RISB_CHRVO		cytoplasm
Q7NVH4	G6PI2_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NVH5	ORN_CHRVO	cytoplasm (potential).	cytoplasm
Q7NVH7	PHS_CHRVO		cytoplasm
Q7NVM0	PPNK_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NVN5	CYSN_CHRVO		cytoplasm
Q7NVP1	SERC_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NVY1	LPXB_CHRVO		cytoplasm
Q7NVY3	FABZ_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NVY4	LPXD_CHRVO		cytoplasm
Q7NVY8	DXR_CHRVO		cytoplasm
Q7NVZ0	UPPS_CHRVO		inner membrane
Q7NVZ1	RRF_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NVZ4	RS2_CHRVO		cytoplasm
Q7NW07	APAG_CHRVO		inner membrane

Q7NW10	GPH_CHRVO		cytoplasm
Q7NW17	TRPD_CHRVO		cytoplasm
Q7NW22	LEUD1_CHRVO		cytoplasm
Q7NW95	GLNE_CHRVO		cytoplasm
Q7NWB7	NAGZ_CHRVO	cytoplasm (by similarity).	extracellular
Q7NWB8	ACPS_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NWC0	PDXJ_CHRVO	cytoplasm (probable).	cytoplasm
Q7NWC1	RECO_CHRVO		cytoplasm
Q7NWC4	RNC_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NWC7	LEPA_CHRVO		cytoplasm
Q7NWE3	Y2047_CHRVO		inner membrane
Q7NWI9	HSLO_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NWI5	ASSY_CHRVO	cytoplasm (probable).	cytoplasm
Q7NWI6	OTC_CHRVO	cytoplasm (probable).	cytoplasm
Q7NWN7	CLPB_CHRVO	cytoplasm (probable).	cytoplasm
Q7NWN9	Y1942_CHRVO		cytoplasm
Q7NWP1	CRCB_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NWP9	RUMA_CHRVO		cytoplasm
Q7NWQ2	UPPP2_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NWR6	DADA_CHRVO		cytoplasm
Q7NWU9	CYNS_CHRVO		cytoplasm
Q7NWX3	CYSA2_CHRVO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7NX01	CYSA1_CHRVO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7NX17	PYRD_CHRVO	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7NX30	ATE_CHRVO	cytoplasm (potential).	cytoplasm
Q7NX31	LFTR_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NX34	DAPB_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NX61	Y1768_CHRVO		inner membrane
Q7NX81	RL31_CHRVO		cytoplasm
Q7NX89	Y1738_CHRVO		cytoplasm
Q7NX94	Y1733_CHRVO		inner membrane
Q7NXB9	FLGI1_CHRVO	periplasm (by similarity).	periplasm
Q7NXC0	FLGH1_CHRVO	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q7NXH5	TDH_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NXI1	DNAJ_CHRVO	cytoplasm (probable).	cytoplasm
Q7NXI3	DNAK_CHRVO		cytoplasm
Q7NXI4	GRPE_CHRVO	cytoplasm (probable).	cytoplasm
Q7NXI6	RRAA_CHRVO		cytoplasm
Q7NXI9	PAND_CHRVO		cytoplasm
Q7NXJ0	PANC_CHRVO		cytoplasm
Q7NXJ1	PANB_CHRVO		cytoplasm
Q7NXK2	FDHD_CHRVO	cytoplasm (potential).	cytoplasm
Q7NXL2	LOLA_CHRVO	periplasm (by similarity).	periplasm
Q7NXL4	RECR_CHRVO		cytoplasm
Q7NXL5	Y1611_CHRVO		inner membrane

Q7NXL8	HRCA_CHRVO		cytoplasm
Q7NXL9	RECA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NXM7	ATKA_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q7NXN3	NORM_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NXN4	MURB_CHRVO	cytoplasm (probable).	inner membrane
Q7NXQ0	COBQ_CHRVO		cytoplasm
Q7NXR0	CBID_CHRVO		cytoplasm
Q7NXU5	ACKA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NY11	RS15_CHRVO		cytoplasm
Q7NY12	RBFA_CHRVO	cytoplasm (potential).	cytoplasm
Q7NY15	Y1460_CHRVO		inner membrane
Q7NY61	CUTC_CHRVO	cytoplasm (probable).	cytoplasm
Q7NYB4	MSCL_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NYC3	RL20_CHRVO		cytoplasm
Q7NYC4	RL35_CHRVO		cytoplasm
Q7NYE7	UPPP1_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NYF6	HTPG_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NYG9	UVRC_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NYI7	NRDR_CHRVO		cytoplasm
Q7NYI8	GLYA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NYJ7	ILVD_CHRVO		cytoplasm
Q7NYJ8	LGT_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NYL4	RPIA_CHRVO		cytoplasm
Q7NYL5	ISPF_CHRVO		extracellular
Q7NYL6	ISPD_CHRVO		cytoplasm
Q7NYM3	DTD_CHRVO	cytoplasm (probable).	cytoplasm
Q7NYM4	Y1250_CHRVO		inner membrane
Q7NYT5	AROC_CHRVO		cytoplasm
Q7NYV9	TNAA_CHRVO		cytoplasm
Q7NYX3	RLMB_CHRVO		cytoplasm
Q7NYZ2	GCH1_CHRVO		cytoplasm
Q7NYZ3	GPDA_CHRVO	cytoplasm (probable).	cytoplasm
Q7NYZ5	SECB_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NZ00	DCUP_CHRVO	cytoplasm (probable).	cytoplasm
Q7NZ02	FUMC_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NZ10	NUDH_CHRVO		cytoplasm
Q7NZ19	COAD_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NZ31	HSCB_CHRVO		inner membrane
Q7NZ33	HSCA_CHRVO		cytoplasm
Q7NZ60	MDH_CHRVO		cytoplasm
Q7NZ91	UBIG_CHRVO		cytoplasm
Q7NZ95	TYSY_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NZD1	UBIB_CHRVO	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q7NZD3	UBIE_CHRVO		cytoplasm

Q7NZD4	QUEA_CHRVO	cytoplasm (potential).	cytoplasm
Q7NZD5	AROQ_CHRVO		cytoplasm
Q7NZF7	SAHH_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7NZI3	TPIS_CHRVO		cytoplasm
Q7NZI7	PSTB_CHRVO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7NZJ6	HAM1_CHRVO		extracellular
Q7NZZ2	RL27_CHRVO		cytoplasm
Q7NZU3	AROK_CHRVO	cytoplasm (probable).	cytoplasm
Q7NZU4	AROB_CHRVO	cytoplasm (probable).	cytoplasm
Q7NZU6	MSBA_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7NZY0	Y789_CHRVO		cytoplasm
Q7NZY3	METX_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P006	Y763_CHRVO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7P012	HEM6_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P071	PURA1_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P094	ATPE_CHRVO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7P096	ATPG_CHRVO		inner membrane
Q7P097	ATPA_CHRVO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7P0A5	GIDB_CHRVO		cytoplasm
Q7P0E6	HIS2_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P0E7	HIS3_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P0E9	HIS6_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P0F0	HIS4_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P0F1	HIS5_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P0F3	HIS7_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P0F4	HIS8_CHRVO		cytoplasm
Q7P0F6	HISX_CHRVO		cytoplasm
Q7P0F7	HIS1_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P0H2	LEU1_CHRVO		cytoplasm
Q7P0H6	PSD_CHRVO		cytoplasm
Q7P0H9	ILVC_CHRVO		cytoplasm
Q7P0J4	KUP2_CHRVO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7P0P9	Y517_CHRVO		inner membrane
Q7P0S3	COBT_CHRVO		cytoplasm
Q7P0S6	SPEB_CHRVO		cytoplasm
Q7P0T8	ARGJ_CHRVO	cytoplasm (probable).	cytoplasm
Q7P0V2	Y463_CHRVO		inner membrane
Q7P0W0	SLYX_CHRVO		inner membrane
Q7P0W5	DAPD_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P0X5	MURA_CHRVO	cytoplasm (probable).	inner membrane
Q7P112	HSLU_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P113	HSLV_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P144	PYRI_CHRVO		cytoplasm
Q7P145	PYRB_CHRVO		cytoplasm

Q7P188	HUTH_CHRVO	cytoplasm (potential).	cytoplasm
Q7P190	HUTU_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P1B9	NPD_CHRVO	cytoplasm (probable).	cytoplasm
Q7P1G2	GLPK_CHRVO		cytoplasm
Q7P1H8	THIC_CHRVO		cytoplasm
Q7P1I1	UNG_CHRVO	cytoplasm (by similarity).	extracellular
Q7P1M4	PGK_CHRVO	cytoplasm (potential).	cytoplasm
Q7P1P8	PUR7_CHRVO		cytoplasm
Q7P1R3	THIE_CHRVO		cytoplasm
Q7P1R4	G6PI1_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7P1T8	Y124_CHRVO	cytoplasm (potential).	inner membrane
Q7P1U1	KSGA_CHRVO		cytoplasm
Q7P1U7	ARLY_CHRVO	cytoplasm (probable).	cytoplasm
Q7P1V3	TOLB_CHRVO	periplasm (potential).	periplasm
Q7P1Y3	HEM1_CHRVO		cytoplasm
Q7P203	PDXH_CHRVO		cytoplasm
Q7P206	CAPP_CHRVO		cytoplasm
Q7P207	HEM3_CHRVO		cytoplasm
Q7P221	CYAY_CHRVO		inner membrane
Q7P259	DNAA_CHRVO	cytoplasm (by similarity).	cytoplasm
Q7PBU9	UPPS_RICSI		inner membrane
Q7SIF3	ARSS_ALCFA		cytoplasm
Q7SIF4	AOXB_ALCFA		periplasm
Q7TTP0	HMP_BORPE		cytoplasm
Q7TTP2	HMP_BORPA		cytoplasm
Q7U317	CH60_HELHP	cytoplasm (by similarity).	cytoplasm
Q7U318	CH10_HELHP	cytoplasm (by similarity).	cytoplasm
Q7U319	FABZ_HELHP	cytoplasm (by similarity).	cytoplasm
Q7U321	TRMD_HELHP	cytoplasm (potential).	cytoplasm
Q7U322	MURG_HELHP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7U326	Y693_HELHP		cytoplasm
Q7U328	TRUA_HELHP		cytoplasm
Q7U330	TRUB_HELHP		cytoplasm
Q7U331	TRMD_HAEDU	cytoplasm (potential).	cytoplasm
Q7U335	TRUA_HAEDU		cytoplasm
Q7U336	MURG_HAEDU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7U337	FABH_HAEDU	cytoplasm (probable).	cytoplasm
Q7U338	MIAA_HAEDU		cytoplasm
Q7U339	TRMA_HAEDU		cytoplasm
Q7U342	TRMU_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7U343	FABA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7U344	TRME_HAEDU		cytoplasm
Q7U345	TRMD_BLOFL	cytoplasm (potential).	cytoplasm
Q7U346	MURG_BLOFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7U347	MIAA_BLOFL		cytoplasm
Q7U348	CH60_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7U350	CCA_BLOFL		cytoplasm
Q7U351	OXAA_BLOFL	bacterial cell inner membrane; multi- pass membrane protein (by similarity).	inner membrane

Q7U352	FABA_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7U356	TRUA_BLOFL		cytoplasm
Q7U358	CCA_BORPE		cytoplasm
Q7U361	TRMD_BORPE	cytoplasm (potential).	cytoplasm
Q7U363	TRUA_BORPE		cytoplasm
Q7U365	HPRK_BORPE		cytoplasm
Q7U366	MIAA_BORPE		cytoplasm
Q7U367	CCA_BORPA		cytoplasm
Q7U370	HPRK_BORPA		cytoplasm
Q7U372	MIAA_BORPA		cytoplasm
Q7U374	TRMD_BORPA	cytoplasm (potential).	cytoplasm
Q7U376	TRUA_BORPA		cytoplasm
Q7U381	HPRK_BORBR		cytoplasm
Q7U383	MIAA_BORBR		cytoplasm
Q7U384	TRMD_BORBR	cytoplasm (potential).	cytoplasm
Q7U386	TRUA_BORBR		cytoplasm
Q7U391	CCA_BORBR		cytoplasm
Q7UAE7	RUVC_SHIFL		cytoplasm
Q7UAH4	SUFS_SHIFL	cytoplasm (by similarity).	cytoplasm
Q7UAH8	MDTK_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7UAV7	YHIQ_SHIFL		inner membrane
Q7UB34	ILVC_SHIFL		cytoplasm
Q7UB84	GLPK_SHIFL		cytoplasm
Q7UBC5	DUSA_SHIFL		cytoplasm
Q7UBC6	PLSB_SHIFL	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7UBD0	MALK_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7UBD3	RSMB_SHIFL		cytoplasm
Q7UBE9	AAEX_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7UBI7	GLNE_SHIFL		cytoplasm
Q7UBI8	HLDE_SHIFL		cytoplasm
Q7UBJ0	GLGS_SHIFL		cytoplasm
Q7UBJ3	ZUPT_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7UBK4	YQHA_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7UBP8	ARGO_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7UBT0	CYSH_SHIFL	cytoplasm (by similarity).	cytoplasm
Q7UBW5	CLPB_SHIFL	cytoplasm (probable).	cytoplasm
Q7UC29	CYSA_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7UC56	NUOG_SHIFL		cytoplasm

Q7UC63	ARNC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7UC91	DUSC_SHIFL		cytoplasm
Q7UCD5	HOKD_SHIFL		inner membrane
Q7UCG7	SOTB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7UCH6	Y1537_SHIFL	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q7UCI6	ASTE_SHIFL		periplasm
Q7UCL3	YEBN_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7UCT6	DADA_SHIFL		cytoplasm
Q7UCU1	MINC_SHIFL		inner membrane
Q7UCW4	NAGZ_SHIFL	cytoplasm (by similarity).	extracellular
Q7UDD0	LIPB_SHIFL	cytoplasm (by similarity).	cytoplasm
Q7UDF9	HOKE_SHIFL		inner membrane
Q7UDG0	HOKF_SHIFL		inner membrane
Q7UDL2	PROB_SHIFL	cytoplasm (by similarity).	cytoplasm
Q7UDQ6	TILS_SHIFL	cytoplasm (probable).	cytoplasm
Q7UDT5	RAPA_SHIFL		cytoplasm
Q7UDT6	DJLA_SHIFL	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q7UDT8	ISPH_SHIFL		cytoplasm
Q7UDU1	DNAJ_SHIFL	cytoplasm (probable).	cytoplasm
Q7VF06	DCUP_HELHP	cytoplasm (probable).	cytoplasm
Q7VF14	MURE_HELHP	cytoplasm (probable).	cytoplasm
Q7VF28	THIG_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VF43	HAM1_HELHP		extracellular
Q7VF52	PURL_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VF77	Y1800_HELHP		inner membrane
Q7VF98	GLMM_HELHP		cytoplasm
Q7VF99	RS20_HELHP		cytoplasm
Q7VFA6	KAD_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VFD3	RL31_HELHP		cytoplasm
Q7VFD7	HTPG_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VFE9	HEM3_HELHP		cytoplasm
Q7VFF1	HEM1_HELHP		cytoplasm
Q7VFF5	HISX_HELHP		cytoplasm
Q7VFF6	MQO_HELHP		cytoplasm
Q7VFF8	ARLY_HELHP	cytoplasm (probable).	cytoplasm
Q7VFG4	GCH1_HELHP		cytoplasm
Q7VFH6	COAE_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VFJ7	CORA_HELHP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VFL1	GLYA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VFP7	LNT_HELHP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VFP9	DAPA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VFAQ	PURQ_HELHP	cytoplasm (by similarity).	cytoplasm

Q7VFQ6	PUR7_HELHP		cytoplasm
Q7VFS1	Y1604_HELHP		inner membrane
Q7VFU3	ISPDF_HELHP		cytoplasm
Q7VFU4	THIC_HELHP		cytoplasm
Q7VFX4	MDH_HELHP		cytoplasm
Q7VFW0	CLPS_HELHP		cytoplasm
Q7VFX7	UVRB_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VFX6	ENGA_HELHP		cytoplasm
Q7VFX9	KPRS_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VFX3	HLDE_HELHP		cytoplasm
Q7VG06	DNAJ_HELHP	cytoplasm (probable).	cytoplasm
Q7VG29	PTH_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VG30	RL25_HELHP		cytoplasm
Q7VG37	QUEA_HELHP	cytoplasm (potential).	cytoplasm
Q7VG38	GIDB_HELHP		cytoplasm
Q7VG78	GUAA_HELHP		cytoplasm
Q7VGA7	TRPB_HELHP		cytoplasm
Q7VGA8	TRPA_HELHP		cytoplasm
Q7VGB3	FLIE_HELHP		flagellar
Q7VGB9	RPOA_HELHP		cytoplasm
Q7VGC0	RS4_HELHP		cytoplasm
Q7VGC1	RS11_HELHP		cytoplasm
Q7VGC2	RS13_HELHP		cytoplasm
Q7VGC7	RS5_HELHP		cytoplasm
Q7VGC8	RL18_HELHP		cytoplasm
Q7VGD0	RS8_HELHP		cytoplasm
Q7VGD2	RL5_HELHP		cytoplasm
Q7VGD3	RL24_HELHP		cytoplasm
Q7VGD5	RS17_HELHP		cytoplasm
Q7VGD6	RL29_HELHP		cytoplasm
Q7VGD7	RL16_HELHP		cytoplasm
Q7VGD8	RS3_HELHP		cytoplasm
Q7VGD9	RL22_HELHP		cytoplasm
Q7VGE0	RS19_HELHP		cytoplasm
Q7VGE1	RL2_HELHP		cytoplasm
Q7VGE3	RL4_HELHP		cytoplasm
Q7VGE4	RL3_HELHP		cytoplasm
Q7VGE5	RS10_HELHP		cytoplasm
Q7VGF0	AMPA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VGF9	KTHY_HELHP		cytoplasm
Q7VGG0	COAD_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VGJ6	HIS7_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VGK6	TPIS_HELHP		cytoplasm
Q7VGL3	ACKA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VGM5	PPNK_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VGN1	RL28_HELHP		cytoplasm
Q7VGP1	MRAW_HELHP		cytoplasm
Q7VGR5	TILS_HELHP	cytoplasm (probable).	cytoplasm
Q7VGS1	UNG_HELHP	cytoplasm (by similarity).	extracellular
Q7VGT8	Y1232_HELHP		inner membrane
Q7VGU1	RUVX_HELHP	cytoplasm (potential).	cytoplasm
Q7VGU9	ASSY_HELHP	cytoplasm (probable).	cytoplasm
Q7VGW6	ILVC_HELHP		cytoplasm
Q7VGX8	SYI_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VGY5	CLPX_HELHP		cytoplasm
Q7VGZ1	HIS6_HELHP	cytoplasm (by similarity).	cytoplasm

Q7VGZ3	KSGA_HELHP		cytoplasm
Q7VGZ8	GPMI_HELHP		cytoplasm
Q7VGZ9	MRAY_HELHP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VH06	Y1161_HELHP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7VH30	LEU1_HELHP		cytoplasm
Q7VH31	LEU2_HELHP		cytoplasm
Q7VH32	LEUD_HELHP		cytoplasm
Q7VH33	LEU3_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VH56	RUVC_HELHP		cytoplasm
Q7VH57	DNAA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VH68	LPXD_HELHP		cytoplasm
Q7VH74	UPPS_HELHP		inner membrane
Q7VH95	RS2_HELHP		cytoplasm
Q7VHA5	Y1062_HELHP		inner membrane
Q7VHF4	KHSE_HELHP	cytoplasm (potential).	cytoplasm
Q7VHF9	NADE_HELHP		cytoplasm
Q7VHH1	HEMH_HELHP	cytoplasm (by similarity).	extracellular
Q7VHJ0	OTC_HELHP	cytoplasm (probable).	cytoplasm
Q7VHK3	GSA_HELHP	cytoplasm (potential).	cytoplasm
Q7VHL1	HISZ_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VHL2	PURA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VHM1	AROQ_HELHP		cytoplasm
Q7VHM7	RS16_HELHP		cytoplasm
Q7VHN0	RL19_HELHP		cytoplasm
Q7VHN2	SYV_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VHP1	Y923_HELHP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VHR4	FLGH_HELHP	bacterial cell outer membrane (probable).	inner membrane
Q7VHS5	CARA_HELHP		cytoplasm
Q7VHT1	AROB_HELHP	cytoplasm (probable).	cytoplasm
Q7VHV1	PDXJ_HELHP	cytoplasm (probable).	cytoplasm
Q7VHW3	ILVD_HELHP		cytoplasm
Q7VHX2	TGT_HELHP		cytoplasm
Q7VHY0	RPOZ_HELHP		cytoplasm
Q7VHY1	SYV_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VHY5	HIS4_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VHY7	PRMA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VI03	HYPV_HELHP		cytoplasm
Q7VI04	ISPG_HELHP		cytoplasm
Q7VI05	PROA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VI13	GATB_HELHP		cytoplasm
Q7VI15	GPDA_HELHP	cytoplasm (probable).	cytoplasm
Q7VI24	TRMB_HELHP		cytoplasm
Q7VI31	RS21_HELHP		cytoplasm
Q7VI68	SELA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VI80	ACP_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VI82	ACCA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VIA1	KGUA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VIA7	GATA_HELHP		cytoplasm
Q7VIA8	AROC_HELHP		cytoplasm

Q7VIA9	RNC_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VIB6	GLMU_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VIC6	PLSX_HELHP		cytoplasm
Q7VIC9	NDK_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VIE2	GRPE_HELHP	cytoplasm (probable).	cytoplasm
Q7VIE3	DNAK_HELHP		cytoplasm
Q7VIH2	RECA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VIH4	ENO_HELHP	cytoplasm (by similarity).	extracellular
Q7VIH7	AROK_HELHP	cytoplasm (probable).	cytoplasm
Q7VIJ3	HIS8_HELHP		cytoplasm
Q7VIJ7	DXS_HELHP		cytoplasm
Q7VIM8	MURA_HELHP	cytoplasm (probable).	inner membrane
Q7VIN2	NPD_HELHP	cytoplasm (probable).	cytoplasm
Q7VIN5	DEF_HELHP		cytoplasm
Q7VIN7	CLPP_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VIN8	TIG_HELHP		cytoplasm
Q7VIR3	RS15_HELHP		cytoplasm
Q7VIR8	RS6_HELHP		cytoplasm
Q7VIT0	DXR_HELHP		cytoplasm
Q7VIT3	PYRB_HELHP		cytoplasm
Q7VIU0	RUVA_HELHP		cytoplasm
Q7VIU4	LGT_HELHP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VIU8	RUVB_HELHP		cytoplasm
Q7VIV5	RS9_HELHP		cytoplasm
Q7VIW7	DAPB_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VIW9	HIS5_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VJ02	HIS2_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VJ06	RL20_HELHP		cytoplasm
Q7VJ07	RL35_HELHP		cytoplasm
Q7VJ10	RRF_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VJ12	MURB_HELHP	cytoplasm (probable).	inner membrane
Q7VJ16	TOLB_HELHP	periplasm (potential).	periplasm
Q7VJ20	ATPE_HELHP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VJ22	ATPG_HELHP		inner membrane
Q7VJ23	ATPA_HELHP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VJ31	PROB_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VJ36	UREE_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VJ78	RL11_HELHP		cytoplasm
Q7VJ79	RL1_HELHP		cytoplasm
Q7VJ80	RL10_HELHP		cytoplasm
Q7VJ82	RPOBC_HELHP		cytoplasm
Q7VJ83	RS12_HELHP		cytoplasm
Q7VJ84	RS7_HELHP		cytoplasm
Q7VJ91	NIKR_HELHP		cytoplasm
Q7VJA8	SURE_HELHP	cytoplasm (potential).	periplasm
Q7VJB1	Y332_HELHP		inner membrane
Q7VJB2	PAND_HELHP		cytoplasm
Q7VJB6	PGK_HELHP	cytoplasm (potential).	cytoplasm
Q7VJC7	LOLA_HELHP	periplasm (by similarity).	periplasm
Q7VJD2	RL9_HELHP		cytoplasm

Q7VJD3	HSLV_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VJN5	RIBB_HELHP		cytoplasm
Q7VJQ0	DCD_HELHP		extracellular
Q7VJR4	ARGB_HELHP	cytoplasm (probable).	cytoplasm
Q7VJR7	LUXS_HELHP		extracellular
Q7VJT5	NAPA_HELHP	periplasm (by similarity).	periplasm
Q7VJU0	DUT_HELHP		extracellular
Q7VJU4	HIS1_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VJV1	FLGI_HELHP	periplasm (by similarity).	periplasm
Q7VJV5	ISPH_HELHP		cytoplasm
Q7VJW2	DDL_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VJW9	CSRA_HELHP		cytoplasm
Q7VJX1	SSRP_HELHP	cytoplasm (potential).	cytoplasm
Q7VJX7	RNPA_HELHP		extracellular
Q7VJY0	OXAA_HELHP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VJY3	CLPB_HELHP	cytoplasm (probable).	cytoplasm
Q7VJZ1	LEPA_HELHP		cytoplasm
Q7VK48	GCH2_HELHP		extracellular
Q7VK53	KDSA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VK54	RISB_HELHP		cytoplasm
Q7VK55	NUSB_HELHP		cytoplasm
Q7VK73	UVRC_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VK78	Y014_HELHP	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q7VK79	GIDA_HELHP	cytoplasm (by similarity).	cytoplasm
Q7VK83	RL27_HELHP		cytoplasm
Q7VK98	FMT_HAEDU		cytoplasm
Q7VKA0	MSCL_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VKA1	MUTS_HAEDU		cytoplasm
Q7VKA3	Y2022_HAEDU		inner membrane
Q7VKA8	Y2015_HAEDU		inner membrane
Q7VKB0	LIPB_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKB1	LIPA_HAEDU	cytoplasm (potential).	cytoplasm
Q7VKB3	HSLU_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKB4	HSLV_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKB6	FETP_HAEDU		cytoplasm
Q7VKB7	MLTC_HAEDU	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q7VKC2	SELA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKC4	RSMB_HAEDU		cytoplasm
Q7VKC6	RS20_HAEDU		cytoplasm
Q7VKC9	FUMC_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKD1	RS10_HAEDU		cytoplasm
Q7VKD2	RL3_HAEDU		cytoplasm
Q7VKD3	RL4_HAEDU		cytoplasm
Q7VKD5	RL2_HAEDU		cytoplasm
Q7VKD6	RL22_HAEDU		cytoplasm
Q7VKD7	RS3_HAEDU		cytoplasm
Q7VKD8	RL16_HAEDU		cytoplasm
Q7VKD9	RL29_HAEDU		cytoplasm
Q7VKE0	RS17_HAEDU		cytoplasm
Q7VKE4	RL24_HAEDU		cytoplasm

Q7VKE5	RL5_HAEDU		cytoplasm
Q7VKE7	RS8_HAEDU		cytoplasm
Q7VKE9	RL18_HAEDU		cytoplasm
Q7VKF0	RS5_HAEDU		cytoplasm
Q7VKF1	RL30_HAEDU		cytoplasm
Q7VKF2	RL15_HAEDU		cytoplasm
Q7VKF5	RS13_HAEDU		cytoplasm
Q7VKF6	RS11_HAEDU		cytoplasm
Q7VKF7	RS4_HAEDU		cytoplasm
Q7VKF8	RPOA_HAEDU		cytoplasm
Q7VKG0	RS16_HAEDU		cytoplasm
Q7VKG1	RIMM_HAEDU	cytoplasm (potential).	cytoplasm
Q7VKG2	RL19_HAEDU		cytoplasm
Q7VKG8	XERC_HAEDU	cytoplasm (by similarity).	nucleoid
Q7VKH0	LPXH_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKH4	KTHY_HAEDU		cytoplasm
Q7VKH6	ACP_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKH9	RL31B_HAEDU		cytoplasm
Q7VKI2	RPOZ_HAEDU		cytoplasm
Q7VKI8	Y1916_HAEDU		cytoplasm
Q7VKJ0	ASNA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKJ3	MURI_HAEDU		cytoplasm
Q7VKJ6	TUSD_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKJ9	SLYX_HAEDU		inner membrane
Q7VKK4	GLMS_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKK8	HLDD_HAEDU		cytoplasm
Q7VKK9	DEF_HAEDU		cytoplasm
Q7VKL4	RL10_HAEDU		cytoplasm
Q7VKL5	RL7_HAEDU		cytoplasm
Q7VKL7	RPOB_HAEDU		cytoplasm
Q7VKL8	RPOC_HAEDU		cytoplasm
Q7VKN0	NANE_HAEDU		cytoplasm
Q7VKN1	NAGB_HAEDU		extracellular
Q7VKP2	XGPT_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKP3	KGUA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKP5	Y1828_HAEDU		inner membrane
Q7VKP7	MODC_HAEDU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VKR0	UVRC_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKR5	PURA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKR6	THII_HAEDU	cytoplasm (potential).	cytoplasm
Q7VKR7	Y1805_HAEDU	cytoplasm (potential).	inner membrane
Q7VKR8	PPNK_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VKS1	RL35_HAEDU		cytoplasm
Q7VKS2	RL20_HAEDU		cytoplasm
Q7VKS6	RUVX_HAEDU	cytoplasm (potential).	cytoplasm
Q7VKS7	Y1794_HAEDU		inner membrane
Q7VKU3	TOLB_HAEDU	periplasm (potential).	periplasm
Q7VKU4	DJLA_HAEDU	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q7VKU5	DUSC_HAEDU		cytoplasm
Q7VKU9	RUMA_HAEDU		cytoplasm
Q7VKV0	RAPA_HAEDU		cytoplasm
Q7VKV3	PYRE_HAEDU		cytoplasm

Q7VKV4	RUVA_HAEDU		cytoplasm
Q7VKV5	RUVB_HAEDU		cytoplasm
Q7VKW2	UBIG_HAEDU		cytoplasm
Q7VKX3	RS15_HAEDU		cytoplasm
Q7VKX4	PPID_HAEDU	bacterial cell membrane; single-pass type ii membrane protein (potential).	periplasm
Q7VKY2	DSBB_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VKY8	Y1715_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VKZ1	ISPZ_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VKZ4	GLNE_HAEDU		cytoplasm
Q7VKZ5	Y1708_HAEDU		inner membrane
Q7VKZ7	Y1706_HAEDU		inner membrane
Q7VKZ9	OSTA_HAEDU	bacterial cell outer membrane (by similarity).	outer membrane
Q7VL00	GREA_HAEDU		cytoplasm
Q7VL12	BIOD2_HAEDU		cytoplasm
Q7VL13	Y1679_HAEDU		inner membrane
Q7VL21	GMHB_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VL22	QUEF_HAEDU	cytoplasm (probable).	cytoplasm
Q7VL26	DCD_HAEDU		extracellular
Q7VL27	URK_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VL28	GPMA_HAEDU		cytoplasm
Q7VL35	Y1650_HAEDU		inner membrane
Q7VL52	MSBA_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VL53	LOLB_HAEDU	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q7VL54	ISPE_HAEDU		cytoplasm
Q7VL55	KPRS_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VL56	PYRD_HAEDU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VL73	LEPA_HAEDU		cytoplasm
Q7VL75	RNC_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VL76	ERA_HAEDU		cytoplasm
Q7VL77	RECO_HAEDU		cytoplasm
Q7VL79	RS2_HAEDU		cytoplasm
Q7VL82	PYRH_HAEDU		cytoplasm
Q7VL83	RRF_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VL86	PDXS_HAEDU		extracellular
Q7VL87	PDXT_HAEDU		cytoplasm
Q7VL94	MUKF_HAEDU	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q7VLE3	Y1515_HAEDU		inner membrane
Q7VLE6	GLMU_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLE9	GUAA_HAEDU		cytoplasm
Q7VLF2	RRMJ_HAEDU	cytoplasm (by similarity).	cytoplasm

Q7VLF7	RS9_HAEDU		cytoplasm
Q7VLG4	IHFA_HAEDU		cytoplasm
Q7VLG6	BIOD1_HAEDU		cytoplasm
Q7VLH2	KITH_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLI0	Y1463_HAEDU		inner membrane
Q7VLI3	RBFA_HAEDU	cytoplasm (potential).	cytoplasm
Q7VLI5	ACKA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLK1	USPA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLK4	SSPA_HAEDU		cytoplasm
Q7VLL3	UVRB_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLM6	DAPB_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLM7	MTFA_HAEDU		cytoplasm
Q7VLN1	MOAA_HAEDU		cytoplasm
Q7VLN2	MOAC_HAEDU		cytoplasm
Q7VLN4	MOAE_HAEDU		cytoplasm
Q7VLN5	Y1387_HAEDU		inner membrane
Q7VLN8	ARGA_HAEDU	cytoplasm (probable).	cytoplasm
Q7VLN9	AROA_HAEDU	cytoplasm (probable).	cytoplasm
Q7VLP0	SERC_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLP5	Y1377_HAEDU		inner membrane
Q7VLP8	NRDR_HAEDU		cytoplasm
Q7VLQ4	TGT_HAEDU		cytoplasm
Q7VLR0	Y1362_HAEDU		cytoplasm
Q7VLR3	Y1359_HAEDU		inner membrane
Q7VLR5	PYRF_HAEDU		cytoplasm
Q7VLR8	IHFB_HAEDU		cytoplasm
Q7VLS0	KCY_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLS2	Y1349_HAEDU		inner membrane
Q7VLS3	HAM1_HAEDU		extracellular
Q7VLT3	PPCK_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLT4	FTSB_HAEDU	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q7VLT5	ISPD_HAEDU		cytoplasm
Q7VLU0	EX7S_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLW2	UVRA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLW6	FTHS_HAEDU		cytoplasm
Q7VLX2	AROC_HAEDU		cytoplasm
Q7VLX8	RNB_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLX9	APAH_HAEDU		extracellular
Q7VLY1	QUEA_HAEDU	cytoplasm (potential).	cytoplasm
Q7VLY6	RL25_HAEDU		cytoplasm
Q7VLZ2	CITD_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VLZ5	RIMK_HAEDU		cytoplasm
Q7VLZ6	LOLA_HAEDU	periplasm (by similarity).	periplasm
Q7VM15	RNH_HAEDU	cytoplasm (potential).	extracellular
Q7VM20	UPPS_HAEDU		inner membrane
Q7VM24	LPXD_HAEDU		cytoplasm
Q7VM25	FABZ_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VM26	LPXA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VM27	DXR_HAEDU		cytoplasm
Q7VM29	ENGA_HAEDU		cytoplasm
Q7VM30	HLDE_HAEDU		cytoplasm
Q7VM31	PYRR_HAEDU		cytoplasm

Q7VM33	KSGA_HAEDU		cytoplasm
Q7VM34	UPP_HAEDU		cytoplasm
Q7VM42	GREB_HAEDU		cytoplasm
Q7VM44	RISB_HAEDU		cytoplasm
Q7VM49	GLPB_HAEDU		cytoplasm
Q7VM56	LNT_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VM59	Y1141_HAEDU		cytoplasm
Q7VM62	Y1137_HAEDU		inner membrane
Q7VM64	SSRP_HAEDU	cytoplasm (potential).	cytoplasm
Q7VM69	Y1129_HAEDU		cytoplasm
Q7VM70	COAE_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VM84	FDHE_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VM87	DAPA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VM90	GCH1_HAEDU		cytoplasm
Q7VMA4	HSCA_HAEDU		cytoplasm
Q7VMA6	HSCB_HAEDU		inner membrane
Q7VMA9	ISCS_HAEDU		cytoplasm
Q7VMB5	NORM_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VMB7	GRPE_HAEDU	cytoplasm (probable).	cytoplasm
Q7VMC0	UNG_HAEDU	cytoplasm (by similarity).	extracellular
Q7VMC2	GRCA_HAEDU		cytoplasm
Q7VMC4	END4_HAEDU		extracellular
Q7VMD0	NDK_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMD5	RL9_HAEDU		cytoplasm
Q7VMD6	PRIB_HAEDU		cytoplasm
Q7VMD7	RS6_HAEDU		cytoplasm
Q7VME2	ISPG_HAEDU		cytoplasm
Q7VME5	ARLY_HAEDU	cytoplasm (probable).	cytoplasm
Q7VMF1	ENG_C_HAEDU		cytoplasm
Q7VMF2	RNH2_HAEDU	cytoplasm (potential).	extracellular
Q7VMG8	PROQ_HAEDU	cytoplasm (potential).	inner membrane
Q7VMH2	TYSY_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMI1	PTH_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMI2	ENG_D_HAEDU		cytoplasm
Q7VMJ1	Y986_HAEDU		inner membrane
Q7VMJ6	CDD_HAEDU		cytoplasm
Q7VMJ8	Y977_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VMS5	ARGD_HAEDU	cytoplasm (probable).	cytoplasm
Q7VMS6	ARGB_HAEDU	cytoplasm (probable).	cytoplasm
Q7VMS8	DEOD_HAEDU		cytoplasm
Q7VMS9	DEOC_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMU2	SPRT_HAEDU	cytoplasm (potential).	extracellular
Q7VMV0	PGK_HAEDU	cytoplasm (potential).	cytoplasm
Q7VMV4	LOLD_HAEDU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VMV6	KDSA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMW1	DNAA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMW3	RECF_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMW5	LPXB_HAEDU		cytoplasm

Q7VMW7	HSLO_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMX4	6PGD_HAEDU		cytoplasm
Q7VMX6	ZIPA_HAEDU	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q7VMY0	KAD_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMY1	MURC_HAEDU	cytoplasm (probable).	cytoplasm
Q7VMY2	DDL_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VMY6	LPXC_HAEDU		cytoplasm
Q7VMZ4	DSBD_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VMZ9	Y802_HAEDU		inner membrane
Q7VN08	CCME_HAEDU	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q7VN12	CCMA_HAEDU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VN15	Y778_HAEDU		inner membrane
Q7VN18	TRMB_HAEDU		cytoplasm
Q7VN19	PLSX_HAEDU		cytoplasm
Q7VN20	RL32_HAEDU		cytoplasm
Q7VN27	TPIS_HAEDU		cytoplasm
Q7VN28	SURE_HAEDU	cytoplasm (potential).	periplasm
Q7VN34	RNPA_HAEDU		extracellular
Q7VN41	DTD_HAEDU	cytoplasm (probable).	cytoplasm
Q7VN43	MUTL_HAEDU		cytoplasm
Q7VN44	HFQ_HAEDU		cytoplasm
Q7VN48	SLMA_HAEDU	nucleoid-associated (by similarity).	cytoplasm
Q7VN49	DUT_HAEDU		extracellular
Q7VN51	RADC_HAEDU		cytoplasm
Q7VN52	RL28_HAEDU		cytoplasm
Q7VN53	RL33_HAEDU		cytoplasm
Q7VN59	CORA_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VN62	UBIB_HAEDU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q7VN63	TATA_HAEDU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q7VN64	TATB_HAEDU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q7VN67	FPG_HAEDU		cytoplasm
Q7VN81	SYI_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VN90	METJ_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VN93	SYV_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VN98	TIG_HAEDU		cytoplasm
Q7VN99	SECB_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VNA0	GPDA_HAEDU	cytoplasm (probable).	cytoplasm
Q7VNA7	PSD_HAEDU		cytoplasm

Q7VNB5	Y640_HAEDU		inner membrane
Q7VNC2	AROQ_HAEDU		cytoplasm
Q7VNC4	DAPD_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VNC9	ALR_HAEDU		cytoplasm
Q7VND8	RNT_HAEDU		cytoplasm
Q7VNF3	Y596_HAEDU		inner membrane
Q7VNF5	RUVC_HAEDU		cytoplasm
Q7VNG0	SELD_HAEDU		cytoplasm
Q7VNH1	CLPB_HAEDU	cytoplasm (probable).	cytoplasm
Q7VNH9	AMPA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VNI5	PLSB_HAEDU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VNI6	LEXA_HAEDU		cytoplasm
Q7VNI7	MUTH_HAEDU	cytoplasm (by similarity).	extracellular
Q7VNI8	NAGZ_HAEDU	cytoplasm (by similarity).	extracellular
Q7VNM5	RUMB_HAEDU		cytoplasm
Q7VNM6	ENO_HAEDU	cytoplasm (by similarity).	extracellular
Q7VNN7	COAD_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VNN8	ORN_HAEDU	cytoplasm (potential).	cytoplasm
Q7VNP1	FIS_HAEDU		cytoplasm
Q7VNP2	DUSB_HAEDU		cytoplasm
Q7VNP7	DXS_HAEDU		cytoplasm
Q7VNP2	NUSB_HAEDU		cytoplasm
Q7VNR4	AROK_HAEDU	cytoplasm (probable).	cytoplasm
Q7VNR5	AROB_HAEDU	cytoplasm (probable).	cytoplasm
Q7VNR9	G6PI_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VNS2	AROE_HAEDU		cytoplasm
Q7VNS6	RECX_HAEDU	cytoplasm (potential).	cytoplasm
Q7VNS7	RECA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VNT1	RNFE_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VNT3	RNFD_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q7VNT4	RNFC_HAEDU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	cytoplasm
Q7VNT6	RNFA_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VNU4	NQRF_HAEDU	bacterial cell inner membrane (potential).	inner membrane
Q7VNU5	NQRE_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q7VNU6	NQRD_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q7VNU7	NQRC_HAEDU	bacterial cell inner membrane (potential).	inner membrane
Q7VNU8	NQRB_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q7VNU9	NQRA_HAEDU		inner membrane

Q7VNV0	Y378_HAEDU		inner membrane
Q7VNV1	ADD_HAEDU		extracellular
Q7VNV2	DUSA_HAEDU		cytoplasm
Q7VNV5	PYRG_HAEDU		cytoplasm
Q7VNV8	LUXS_HAEDU		extracellular
Q7VNW8	Y358_HAEDU		inner membrane
Q7VNX2	HTPX_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q7VNY0	ENGB_HAEDU		cytoplasm
Q7VNY2	TUSA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VNY6	KDSB_HAEDU	cytoplasm (potential).	cytoplasm
Q7VNZ0	GLPE_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VNZ1	Y326_HAEDU		inner membrane
Q7VP02	TAL_HAEDU	cytoplasm (probable).	cytoplasm
Q7VP10	RNPH_HAEDU		cytoplasm
Q7VP34	SYI_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VP36	RLMB_HAEDU		cytoplasm
Q7VP41	MDH_HAEDU		cytoplasm
Q7VP42	ARGR_HAEDU	cytoplasm (probable).	cytoplasm
Q7VP51	MURA_HAEDU	cytoplasm (probable).	inner membrane
Q7VP56	MURD_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VP57	MRAY_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VP59	MURE_HAEDU	cytoplasm (probable).	cytoplasm
Q7VP62	MRAW_HAEDU		cytoplasm
Q7VP66	CARA_HAEDU		cytoplasm
Q7VP67	CARB_HAEDU		cytoplasm
Q7VP68	OTC_HAEDU	cytoplasm (probable).	cytoplasm
Q7VP78	CLPP_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VP79	CLPX_HAEDU		cytoplasm
Q7VP80	LPXK_HAEDU		cytoplasm
Q7VP83	PDXH_HAEDU		cytoplasm
Q7VP85	RIMI_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VP91	RL27_HAEDU		cytoplasm
Q7VP94	GLMM_HAEDU		cytoplasm
Q7VP95	RPIA_HAEDU		cytoplasm
Q7VPB1	Y178_HAEDU		inner membrane
Q7VPB5	LGT_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VPB7	NUDH_HAEDU		cytoplasm
Q7VPC0	IPYR_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VPC3	MQO_HAEDU		cytoplasm
Q7VPI9	LLDD_HAEDU		cytoplasm
Q7VPJ2	MURB_HAEDU	cytoplasm (probable).	inner membrane
Q7VPJ7	NAPA_HAEDU	periplasm (by similarity).	periplasm
Q7VPK3	Y065_HAEDU		cytoplasm
Q7VPK4	ISPH_HAEDU		cytoplasm
Q7VPK5	LSPA_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q7VPK9	COAA_HAEDU	cytoplasm (probable).	cytoplasm
Q7VPL3	ACCA_HAEDU	cytoplasm (by similarity).	cytoplasm

Q7VPM2	OXAA_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VPM9	SYK3_HAEDU		cytoplasm
Q7VPN0	DAPF_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VPN2	CRCB_HAEDU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VPN5	PRMA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VPN7	TILS_HAEDU	cytoplasm (probable).	cytoplasm
Q7VPN8	XERD_HAEDU	cytoplasm (by similarity).	nucleoid
Q7VPN9	ATPE_HAEDU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VPP1	ATPG_HAEDU		inner membrane
Q7VPP2	ATPA_HAEDU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VPP8	GIDB_HAEDU		cytoplasm
Q7VPP9	GIDA_HAEDU	cytoplasm (by similarity).	cytoplasm
Q7VQB1	TGT_BLOFL		cytoplasm
Q7VQB2	QUEA_BLOFL	cytoplasm (potential).	cytoplasm
Q7VQC0	DEF_BLOFL		cytoplasm
Q7VQC1	FMT_BLOFL		cytoplasm
Q7VQC3	RPOA_BLOFL		cytoplasm
Q7VQC4	RS4_BLOFL		cytoplasm
Q7VQC5	RS11_BLOFL		cytoplasm
Q7VQC6	RS13_BLOFL		cytoplasm
Q7VQD1	RS5_BLOFL		cytoplasm
Q7VQD2	RL18_BLOFL		cytoplasm
Q7VQD4	RS8_BLOFL		cytoplasm
Q7VQD6	RL5_BLOFL		cytoplasm
Q7VQD7	RL24_BLOFL		cytoplasm
Q7VQD9	RS17_BLOFL		cytoplasm
Q7VQE1	RL16_BLOFL		cytoplasm
Q7VQE2	RS3_BLOFL		cytoplasm
Q7VQE3	RL22_BLOFL		cytoplasm
Q7VQE4	RS19_BLOFL		cytoplasm
Q7VQE5	RL2_BLOFL		cytoplasm
Q7VQE7	RL4_BLOFL		cytoplasm
Q7VQE8	RL3_BLOFL		cytoplasm
Q7VQE9	RS10_BLOFL		cytoplasm
Q7VQF2	MURB_BLOFL	cytoplasm (probable).	inner membrane
Q7VQF3	CLPB_BLOFL	cytoplasm (probable).	cytoplasm
Q7VQF8	RL19_BLOFL		cytoplasm
Q7VQF9	RIMM_BLOFL	cytoplasm (potential).	cytoplasm
Q7VQG0	RS16_BLOFL		cytoplasm
Q7VQG2	CSRA_BLOFL		cytoplasm
Q7VQH1	CYSI_BLOFL		cytoplasm
Q7VQH2	CYSJ_BLOFL		cytoplasm
Q7VQH3	ENO_BLOFL	cytoplasm (by similarity).	extracellular
Q7VQI7	MURC_BLOFL	cytoplasm (probable).	cytoplasm
Q7VQI9	MURD_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQJ0	MRAY_BLOFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q7VQJ2	MURE_BLOFL	cytoplasm (probable).	cytoplasm
Q7VQJ5	MRAW_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQJ6	LEU1_BLOFL		cytoplasm
Q7VQJ7	LEU3_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQJ8	LEU2_BLOFL		cytoplasm
Q7VQJ9	LEUD_BLOFL		cytoplasm
Q7VQK3	KSGA_BLOFL		cytoplasm
Q7VQK8	DAPB_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQL0	SYI_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQL2	RS20_BLOFL		cytoplasm
Q7VQL3	DNAJ_BLOFL	cytoplasm (probable).	cytoplasm
Q7VQL4	DNAK_BLOFL		cytoplasm
Q7VQL6	KHSE_BLOFL	cytoplasm (potential).	cytoplasm
Q7VQM1	RS15_BLOFL		cytoplasm
Q7VQM2	RBFA_BLOFL	cytoplasm (potential).	cytoplasm
Q7VQM5	GLMM_BLOFL		cytoplasm
Q7VQM8	RRMJ_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQN1	RL27_BLOFL		cytoplasm
Q7VQN7	RL9_BLOFL		cytoplasm
Q7VQN8	RS18_BLOFL		cytoplasm
Q7VQN9	RS6_BLOFL		cytoplasm
Q7VQP0	RLMB_BLOFL		cytoplasm
Q7VQP1	PURA_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQP7	ORN_BLOFL	cytoplasm (potential).	cytoplasm
Q7VQP8	PSD_BLOFL		cytoplasm
Q7VQQ4	RIBB_BLOFL		cytoplasm
Q7VQQ6	HLDE_BLOFL		cytoplasm
Q7VQQ8	Y060_BLOFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VQR0	RS21_BLOFL		cytoplasm
Q7VQR6	RS9_BLOFL		cytoplasm
Q7VQS0	MURA_BLOFL	cytoplasm (probable).	inner membrane
Q7VQT0	AMPA_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQT2	SYV_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQT3	OTC_BLOFL	cytoplasm (probable).	cytoplasm
Q7VQV1	RNPA_BLOFL		extracellular
Q7VQV2	Y1013_BLOFL		inner membrane
Q7VQV3	TRME_BLOFL		cytoplasm
Q7VQV4	GLMU_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQV5	ATPE_BLOFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VQV7	ATPG_BLOFL		inner membrane
Q7VQV8	ATPA_BLOFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VQW3	GIDA_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQW4	HIS2_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQW5	HIS6_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQW6	HIS4_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQW7	HIS5_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQW8	HIS7_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQW9	HIS8_BLOFL		cytoplasm
Q7VQX0	HISX_BLOFL		cytoplasm
Q7VQX1	HIS1_BLOFL	cytoplasm (by similarity).	cytoplasm

Q7VQX3	Y460_BLOFL		inner membrane
Q7VQX5	MDTH_BLOFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VQZ1	MINC_BLOFL		inner membrane
Q7VQZ6	KITH_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VQZ9	TRPA_BLOFL		cytoplasm
Q7VR00	TRPB_BLOFL		cytoplasm
Q7VR05	GCH2_BLOFL		extracellular
Q7VR12	PGSA_BLOFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VR16	RL32_BLOFL		cytoplasm
Q7VR17	PLSX_BLOFL		cytoplasm
Q7VR18	FABH_BLOFL	cytoplasm (probable).	cytoplasm
Q7VR21	ACP_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VR23	KTHY_BLOFL		cytoplasm
Q7VR29	LOLD_BLOFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VR38	LOLA_BLOFL	periplasm (by similarity).	periplasm
Q7VR40	SERC_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VR42	KCY_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VR44	MSBA_BLOFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VR45	LPXK_BLOFL		cytoplasm
Q7VR50	FUMC_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VR52	SYX_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VR53	PDXH_BLOFL		cytoplasm
Q7VR54	SLYA_BLOFL		cytoplasm
Q7VR55	RNT_BLOFL		cytoplasm
Q7VR65	LPLA_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VR68	RL20_BLOFL		cytoplasm
Q7VR69	RL35_BLOFL		cytoplasm
Q7VR76	KPRS_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VR80	GPMA_BLOFL		cytoplasm
Q7VR81	6PGL_BLOFL		cytoplasm
Q7VR84	TOLB_BLOFL	periplasm (potential).	periplasm
Q7VRA3	Y316_BLOFL		inner membrane
Q7VRA5	LNT_BLOFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VRB5	LPXH_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRB6	KAD_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRC5	AROQ_BLOFL		cytoplasm
Q7VRC9	TILS_BLOFL	cytoplasm (probable).	cytoplasm
Q7VRD0	ACCA_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRD2	RNH2_BLOFL	cytoplasm (potential).	extracellular
Q7VRD3	LPXB_BLOFL		cytoplasm
Q7VRD5	FABZ_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRD6	LPXD_BLOFL		cytoplasm
Q7VRD7	SKP_BLOFL	periplasm (by similarity).	periplasm
Q7VRE1	UPPS_BLOFL		inner membrane
Q7VRE2	DXR_BLOFL		cytoplasm
Q7VRE3	RRF_BLOFL	cytoplasm (by similarity).	cytoplasm

Q7VRE6	RS2_BLOFL		cytoplasm
Q7VRE8	DAPD_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRF2	LGT_BLOFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VRF3	NUDH_BLOFL		cytoplasm
Q7VRG1	RPIA_BLOFL		cytoplasm
Q7VRG3	PGK_BLOFL	cytoplasm (potential).	cytoplasm
Q7VRG4	SPEB_BLOFL		cytoplasm
Q7VRG6	Y251_BLOFL		inner membrane
Q7VRG7	RUVX_BLOFL	cytoplasm (potential).	cytoplasm
Q7VRG9	FETP_BLOFL		cytoplasm
Q7VRH0	CLPX_BLOFL		cytoplasm
Q7VRH1	CLPP_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRH9	DXS_BLOFL		cytoplasm
Q7VRI1	NUSB_BLOFL		cytoplasm
Q7VRI2	RISB_BLOFL		cytoplasm
Q7VRI4	G6PI_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRI8	METE_BLOFL		cytoplasm
Q7VRJ1	UBIE_BLOFL		cytoplasm
Q7VRJ7	KGUA_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRK0	DUT_BLOFL		extracellular
Q7VRK1	RL28_BLOFL		cytoplasm
Q7VRK7	GPDA_BLOFL	cytoplasm (probable).	cytoplasm
Q7VRL0	TPIS_BLOFL		cytoplasm
Q7VRL2	RL31_BLOFL		cytoplasm
Q7VRL8	ILVD_BLOFL		cytoplasm
Q7VRM0	ILVC_BLOFL		cytoplasm
Q7VRM4	HEM3_BLOFL		cytoplasm
Q7VRM7	CORA_BLOFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VRN1	Y573_BLOFL		inner membrane
Q7VRN2	AROK_BLOFL	cytoplasm (probable).	cytoplasm
Q7VRN3	AROB_BLOFL	cytoplasm (probable).	cytoplasm
Q7VRN7	RS12_BLOFL		cytoplasm
Q7VRN8	RS7_BLOFL		cytoplasm
Q7VRP3	RL11_BLOFL		cytoplasm
Q7VRP4	RL1_BLOFL		cytoplasm
Q7VRP5	RL10_BLOFL		cytoplasm
Q7VRP7	RPOB_BLOFL		cytoplasm
Q7VRP8	RPOC_BLOFL		cytoplasm
Q7VRQ5	PPNK_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRQ6	GRPE_BLOFL	cytoplasm (probable).	cytoplasm
Q7VRQ7	UNG_BLOFL	cytoplasm (by similarity).	extracellular
Q7VRQ8	LEPA_BLOFL		cytoplasm
Q7VRR0	RNC_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRR1	PDXJ_BLOFL	cytoplasm (probable).	cytoplasm
Q7VRR2	ACPS_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRR4	GLYA_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRS0	MQO_BLOFL		cytoplasm
Q7VRS2	GUAA_BLOFL		cytoplasm
Q7VRS4	URE3_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRS5	URE2_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRS6	URE1_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRS9	UPP_BLOFL		cytoplasm

Q7VRT1	DAPA_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRT4	TAL_BLOFL	cytoplasm (probable).	cytoplasm
Q7VRU6	AROC_BLOFL		cytoplasm
Q7VRU9	PDXB_BLOFL		cytoplasm
Q7VRV7	NUOG_BLOFL		cytoplasm
Q7VRX1	DDL_BLOFL	cytoplasm (by similarity).	cytoplasm
Q7VRX2	RL25_BLOFL		cytoplasm
Q7VRX3	GCH1_BLOFL		cytoplasm
Q7VRY0	LEXA_BORPE		cytoplasm
Q7VRY3	HEM1_BORPE		cytoplasm
Q7VRZ3	GLMS_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VS43	GPMA_BORPE		cytoplasm
Q7VS46	SECB_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VS47	GPDA_BORPE	cytoplasm (probable).	cytoplasm
Q7VS51	CAAL_BORPE		cytoplasm
Q7VS76	ASPD1_BORPE		cytoplasm
Q7VS88	DEF2_BORPE		cytoplasm
Q7VS89	FMT_BORPE		cytoplasm
Q7VSA0	OAT_BORPE		cytoplasm
Q7VSB4	Y521_BORPE		extracellular
Q7VSD7	Y494_BORPE		inner membrane
Q7VSD8	RNPA_BORPE		extracellular
Q7VSE0	DNAA_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VSE8	Y480_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VSF4	RIBB_BORPE		cytoplasm
Q7VSG7	SMG_BORPE		cytoplasm
Q7VSH3	ARGD2_BORPE	cytoplasm (probable).	cytoplasm
Q7VSM2	ARGB_BORPE	cytoplasm (probable).	cytoplasm
Q7VSN1	GATC_BORPE		cytoplasm
Q7VSN2	GATA_BORPE		cytoplasm
Q7VSN3	GATB_BORPE		cytoplasm
Q7VSN4	PYRE_BORPE		cytoplasm
Q7VSR5	TRME_BORPE		cytoplasm
Q7VSU7	GLPK_BORPE		cytoplasm
Q7VSV5	COAE_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VSW3	ARGJ_BORPE	cytoplasm (probable).	cytoplasm
Q7VSY4	HIS2_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VSY5	HIS3_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VSY6	HIS6_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VSY7	HIS4_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VSY8	HIS5_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VSY9	HIS7_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VSZ0	HIS82_BORPE		cytoplasm
Q7VSZ1	HISX_BORPE		cytoplasm
Q7VSZ2	HIS1_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VSZ3	MURA_BORPE	cytoplasm (probable).	inner membrane
Q7VT27	GLMU_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VT61	TRMB_BORPE		cytoplasm
Q7VT92	DGTL1_BORPE		cytoplasm
Q7VT93	AROB_BORPE	cytoplasm (probable).	cytoplasm
Q7VT94	AROK_BORPE	cytoplasm (probable).	cytoplasm
Q7VT96	CYAY_BORPE		inner membrane
Q7VTA7	RS11_BORPE		cytoplasm
Q7VTA8	RS13_BORPE		cytoplasm

Q7VTB2	RL15_BORPE		cytoplasm
Q7VTB4	RS5_BORPE		cytoplasm
Q7VTB5	RL18_BORPE		cytoplasm
Q7VTB7	RS8_BORPE		cytoplasm
Q7VTB9	RL5_BORPE		cytoplasm
Q7VTC0	RL24_BORPE		cytoplasm
Q7VTC4	RS17_BORPE		cytoplasm
Q7VTC6	RL16_BORPE		cytoplasm
Q7VTC7	RS3_BORPE		cytoplasm
Q7VTC8	RL22_BORPE		cytoplasm
Q7VTC9	RS19_BORPE		cytoplasm
Q7VTD0	RL2_BORPE		cytoplasm
Q7VTD2	RL4_BORPE		cytoplasm
Q7VTD3	RL3_BORPE		cytoplasm
Q7VTD4	RS10_BORPE		cytoplasm
Q7VTD6	RS7_BORPE		cytoplasm
Q7VTD7	RS12_BORPE		cytoplasm
Q7VTE5	THIG_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VTF0	TRPA_BORPE		cytoplasm
Q7VTF1	TRPB_BORPE		cytoplasm
Q7VTF7	PNCB_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VTJ7	ARGD1_BORPE	cytoplasm (probable).	cytoplasm
Q7VTJ8	OTC_BORPE	cytoplasm (probable).	cytoplasm
Q7VTJ9	ASSY_BORPE	cytoplasm (probable).	cytoplasm
Q7VTK0	KUP_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VTK4	RLMB_BORPE		cytoplasm
Q7VTN3	NUSB_BORPE		cytoplasm
Q7VTN4	RISB_BORPE		cytoplasm
Q7VTP4	COAD_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VTR8	ATE_BORPE	cytoplasm (potential).	cytoplasm
Q7VTR9	PYRD_BORPE	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VTT6	RUVB_BORPE		cytoplasm
Q7VTT9	RUVA_BORPE		cytoplasm
Q7VTU0	RUVC_BORPE		cytoplasm
Q7VTZ7	NUDH_BORPE		cytoplasm
Q7VU03	TOLB_BORPE	periplasm (potential).	periplasm
Q7VU11	KSGA_BORPE		cytoplasm
Q7VU17	PHS_BORPE		cytoplasm
Q7VU41	DCUP_BORPE	cytoplasm (probable).	cytoplasm
Q7VU43	ATPE_BORPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VU45	ATPG_BORPE		inner membrane
Q7VU46	ATPA_BORPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VU61	APAG_BORPE		inner membrane
Q7VU66	TRPD_BORPE		cytoplasm
Q7VU72	PDXH_BORPE		cytoplasm
Q7VU93	MINC_BORPE		inner membrane
Q7VUD0	URE3_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUD2	URE2_BORPE	cytoplasm (by similarity).	cytoplasm

Q7VUD3	URE1_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUD4	UREE_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUE3	TYSY_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUF4	G6PI_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUG2	HGD_BORPE		cytoplasm
Q7VUG7	FPG_BORPE		cytoplasm
Q7VUG9	LOLB_BORPE	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q7VUH0	ISPE_BORPE		cytoplasm
Q7VUH1	KPRS_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUH2	RL25_BORPE		cytoplasm
Q7VUH3	PTH_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUJ5	MODC_BORPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VUJ9	HSLU_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUK0	HSLV_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUL8	SAHH_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUM8	UNG_BORPE	cytoplasm (by similarity).	extracellular
Q7VUN5	LGT_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VUN6	ILVD2_BORPE		cytoplasm
Q7VUP5	MRAZ_BORPE		inner membrane
Q7VUP6	MRAW_BORPE		cytoplasm
Q7VUQ0	MRAY_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VUQ1	MURD_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUQ3	MURG_BORPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VUQ4	MURC_BORPE	cytoplasm (probable).	cytoplasm
Q7VUQ5	DDL_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUS7	AROQ_BORPE		cytoplasm
Q7VUV5	Y2965_BORPE		inner membrane
Q7VUV9	RS9_BORPE		cytoplasm
Q7VUW0	ARGC_BORPE	cytoplasm (probable).	cytoplasm
Q7VUW5	SYV_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUW7	GLYA_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VUW8	NRDR_BORPE		cytoplasm
Q7VUZ0	Y2925_BORPE		inner membrane
Q7VUZ9	Y2916_BORPE		inner membrane
Q7VV53	PANB_BORPE		inner membrane
Q7VV78	Y2808_BORPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7VV85	EX7S_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VV87	DXS_BORPE		cytoplasm
Q7VV88	Y2797_BORPE		inner membrane
Q7VV89	RS6_BORPE		cytoplasm
Q7VV91	RS18_BORPE		cytoplasm
Q7VV92	RL9_BORPE		cytoplasm
Q7VVA6	RS20_BORPE		cytoplasm
Q7VVB2	LPXK_BORPE		cytoplasm
Q7VVC1	CLPS_BORPE		cytoplasm

Q7VVG6	ARLY_BORPE	cytoplasm (probable).	cytoplasm
Q7VVH7	ORN_BORPE	cytoplasm (potential).	cytoplasm
Q7VVM4	GUAA_BORPE		cytoplasm
Q7VVR1	DUT_BORPE		extracellular
Q7VVR2	RL35_BORPE		cytoplasm
Q7VVR3	RL20_BORPE		cytoplasm
Q7VVU3	METE_BORPE		cytoplasm
Q7VVU9	HEM3_BORPE		cytoplasm
Q7VVX6	PPNK_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VVX7	HRC_A_BORPE		cytoplasm
Q7VVX8	HEMH_BORPE	cytoplasm (by similarity).	extracellular
Q7VVY0	GRPE_BORPE	cytoplasm (probable).	cytoplasm
Q7VVY2	DNAK_BORPE		cytoplasm
Q7VVY3	DNAJ_BORPE	cytoplasm (probable).	cytoplasm
Q7VVZ9	ATKA_BORPE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q7VW06	LOLA_BORPE	periplasm (by similarity).	periplasm
Q7VW21	Y2452_BORPE		inner membrane
Q7VW25	Y2447_BORPE	cytoplasm (potential).	inner membrane
Q7VW27	RL32_BORPE		cytoplasm
Q7VW28	PLSX_BORPE		cytoplasm
Q7VW29	FABH_BORPE	cytoplasm (probable).	cytoplasm
Q7VW32	ACP_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VW39	RNC_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VW48	AMPA_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VW77	KDSA_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VW79	ENO_BORPE	cytoplasm (by similarity).	extracellular
Q7VW97	MDH_BORPE		cytoplasm
Q7VWC1	RPIA_BORPE		cytoplasm
Q7VWC4	FETP_BORPE		cytoplasm
Q7VWD8	MSBA_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VWE3	Y2314_BORPE	cytoplasm (potential).	inner membrane
Q7VWE4	Y2313_BORPE		inner membrane
Q7VWE6	NADD_BORPE		cytoplasm
Q7VWE7	HEM6_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VWE9	Y2308_BORPE		inner membrane
Q7VWK6	SYV_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VWK7	NDK_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VWL0	ISPG_BORPE		cytoplasm
Q7VWL4	ENGA_BORPE		cytoplasm
Q7VWL5	HIS81_BORPE		cytoplasm
Q7VWL6	HFQ_BORPE		cytoplasm
Q7VWM0	HISZ_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VWM1	PURA_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VWM3	RS21_BORPE		cytoplasm
Q7VWV5	QUEF_BORPE	cytoplasm (probable).	cytoplasm
Q7VWV7	UVRC_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VWV8	NAGZ_BORPE	cytoplasm (by similarity).	extracellular
Q7VWV9	ACPS_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VWW0	PDXJ_BORPE	cytoplasm (probable).	cytoplasm
Q7VWY1	RL28_BORPE		cytoplasm
Q7VWZ0	PROA_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VWZ5	Y2036_BORPE		inner membrane

Q7VX46	NPD_BORPE	cytoplasm (probable).	cytoplasm
Q7VX83	ASPD2_BORPE		cytoplasm
Q7VX92	TILS_BORPE	cytoplasm (probable).	cytoplasm
Q7VX94	ACCA_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VX99	LPXH_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VXA0	UPPP_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VXB5	Y1875_BORPE		inner membrane
Q7VXD2	KTHY_BORPE		cytoplasm
Q7VXD7	RIMM_BORPE	cytoplasm (potential).	cytoplasm
Q7VXD8	RS16_BORPE		cytoplasm
Q7VXF8	PAND_BORPE		cytoplasm
Q7VXG7	HSCA_BORPE		cytoplasm
Q7VXG8	HSCB_BORPE		inner membrane
Q7VXH4	UVRB_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VXI6	CLPX_BORPE		cytoplasm
Q7VXI7	CLPP_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VXI8	TIG_BORPE		cytoplasm
Q7VXK3	SYI_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VXL5	Y1738_BORPE		cytoplasm
Q7VXM6	RUMA_BORPE		cytoplasm
Q7VXN2	SURE_BORPE	cytoplasm (potential).	periplasm
Q7VXN3	Y1718_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VXX8	HAM1_BORPE		extracellular
Q7VXY3	RNPB_BORPE		cytoplasm
Q7VXZ3	KGUA_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VXZ4	RPOZ_BORPE		cytoplasm
Q7VY01	MUTS_BORPE		cytoplasm
Q7VY12	RECR_BORPE		cytoplasm
Q7VY13	Y1550_BORPE		inner membrane
Q7VY42	PUR7_BORPE		cytoplasm
Q7VY62	GSHB_BORPE		cytoplasm
Q7VY73	LEU3_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VY74	LEUD2_BORPE		cytoplasm
Q7VY75	LEU2_BORPE		cytoplasm
Q7VY92	AROC_BORPE		cytoplasm
Q7VY99	TAL_BORPE	cytoplasm (probable).	cytoplasm
Q7VYA8	SSRP_BORPE	cytoplasm (potential).	cytoplasm
Q7VYB4	Y1435_BORPE		cytoplasm
Q7VYB6	RNH2_BORPE	cytoplasm (potential).	extracellular
Q7VYB7	LPXB_BORPE		cytoplasm
Q7VYB9	FABZ_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VYC0	LPXD_BORPE		cytoplasm
Q7VYC4	DXR_BORPE		cytoplasm
Q7VYC6	UPPS_BORPE		inner membrane
Q7VYC7	RRF_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VYD0	RS2_BORPE		cytoplasm
Q7VYD2	GLND_BORPE		cytoplasm
Q7VYG2	FLGI_BORPE	periplasm (by similarity).	periplasm
Q7VYG3	FLGH_BORPE	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q7VYI1	LEUD1_BORPE		cytoplasm
Q7VYK0	GLGB_BORPE		extracellular

Q7VYM4	PSD_BORPE		cytoplasm
Q7VYQ0	GLNE_BORPE		cytoplasm
Q7VYR1	RBFA_BORPE	cytoplasm (potential).	cytoplasm
Q7VYR4	Y1245_BORPE		inner membrane
Q7VYS2	ISPH_BORPE		cytoplasm
Q7VYS4	RADC_BORPE		cytoplasm
Q7VYS7	RL31B_BORPE		cytoplasm
Q7VYU0	CRCB_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7VYV4	Y1203_BORPE		inner membrane
Q7VYV6	CLPB_BORPE	cytoplasm (probable).	cytoplasm
Q7VZ31	LOLD_BORPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VZ47	Y1090_BORPE		inner membrane
Q7VZ56	RRMJ_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VZ59	GLMM_BORPE		cytoplasm
Q7VZ66	PSTB_BORPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VZ79	UPP_BORPE		cytoplasm
Q7VZ81	QUEA_BORPE	cytoplasm (potential).	cytoplasm
Q7VZ88	Y1039_BORPE		inner membrane
Q7VZA1	FLHD_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VZB8	PGK_BORPE	cytoplasm (potential).	cytoplasm
Q7VZE0	RL19_BORPE		cytoplasm
Q7VZE5	CYSA_BORPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7VZF5	HLDD_BORPE		cytoplasm
Q7VZG1	KCY_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VZG4	SERC_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VZG7	UBIG_BORPE		cytoplasm
Q7VZM5	TPMT_BORPE	cytoplasm (by similarity).	cytoplasm
Q7VZN1	ISPF_BORPE		extracellular
Q7VZN2	ISPD_BORPE		cytoplasm
Q7VZT5	TPIS_BORPE		cytoplasm
Q7VZU1	RS15_BORPE		cytoplasm
Q7VZU4	ILVC_BORPE		cytoplasm
Q7VZX5	RL27_BORPE		cytoplasm
Q7VZX7	PROB_BORPE	cytoplasm (by similarity).	cytoplasm
Q7W014	Y690_BORPE		cytoplasm
Q7W039	MTGA_BORPE	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q7W041	APAH_BORPE		extracellular
Q7W044	PYRB_BORPE		cytoplasm
Q7W045	RUVX_BORPE	cytoplasm (potential).	cytoplasm
Q7W046	Y319_BORPE		inner membrane
Q7W049	THIE_BORPE		cytoplasm
Q7W066	53DR_BORPE		extracellular
Q7W069	ILVD1_BORPE		cytoplasm
Q7W079	MSCL_BORPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q7W0A2	FUMC_BORPE	cytoplasm (by similarity).	cytoplasm
Q7W0A7	PUR5_BORPE	cytoplasm (by similarity).	cytoplasm
Q7W0D0	CAPP_BORPE		cytoplasm
Q7W0D7	THIC_BORPE		cytoplasm
Q7W0E3	GCSP_BORPE		cytoplasm
Q7W0E5	GCST_BORPE		cytoplasm
Q7W0F2	DCD_BORPE		extracellular
Q7W0F8	Y179_BORPE		inner membrane
Q7W0H1	UBIE_BORPE		cytoplasm
Q7W0K8	LIPA_BORPE	cytoplasm (potential).	cytoplasm
Q7W0K9	LIPB_BORPE	cytoplasm (by similarity).	cytoplasm
Q7W0L0	Y104_BORPE		inner membrane
Q7W0M8	HTPG_BORPE	cytoplasm (by similarity).	cytoplasm
Q7W0P6	METX_BORPE	cytoplasm (by similarity).	cytoplasm
Q7W0Q0	DEF1_BORPE		cytoplasm
Q7W0R8	RPOC_BORPE		cytoplasm
Q7W0R9	RPOB_BORPE		cytoplasm
Q7W0S1	RL10_BORPE		cytoplasm
Q7W0S2	RL1_BORPE		cytoplasm
Q7W0S3	RL11_BORPE		cytoplasm
Q7W0S9	GIDB_BORPE		cytoplasm
Q7W0T0	GIDA_BORPE	cytoplasm (by similarity).	cytoplasm
Q7W0T5	LEXA_BORPA		cytoplasm
Q7W134	CH60_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W142	NUSB_BORPA		cytoplasm
Q7W143	RISB_BORPA		cytoplasm
Q7W148	PHNC_BORPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W154	COAD_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W179	PTH_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W180	RL25_BORPA		cytoplasm
Q7W181	KPRS_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W182	ISPE_BORPA		cytoplasm
Q7W183	LOLB_BORPA	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q7W185	FPG_BORPA		cytoplasm
Q7W189	HGD_BORPA		cytoplasm
Q7W197	G6PI_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W1A9	TYSY_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W1C4	GCSP_BORPA		cytoplasm
Q7W1C6	GCST_BORPA		cytoplasm
Q7W1D3	DCD_BORPA		extracellular
Q7W1D9	Y754_BORPA		inner membrane
Q7W1E4	OAT_BORPA		cytoplasm
Q7W1F4	MODC_BORPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W1I6	GLYA1_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W1P1	RL27_BORPA		cytoplasm
Q7W1P3	PROB_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W1Q0	THIC_BORPA		cytoplasm
Q7W1Q6	GPMA_BORPA		cytoplasm
Q7W1Q9	SECB_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W1R0	GPDA_BORPA	cytoplasm (probable).	cytoplasm
Q7W1R5	CAAL_BORPA		cytoplasm

Q7W1T9	ASPD1_BORPA		cytoplasm
Q7W1V2	FMT_BORPA		cytoplasm
Q7W1V3	DEF1_BORPA		cytoplasm
Q7W1Y7	UNG_BORPA	cytoplasm (by similarity).	extracellular
Q7W1Z7	SAHH_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W215	HSLV_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W216	HSLU_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W222	LIPA_BORPA	cytoplasm (potential).	cytoplasm
Q7W223	LIPB_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W224	Y168_BORPA		inner membrane
Q7W2B5	DGTL1_BORPA		cytoplasm
Q7W2B6	AROB_BORPA	cytoplasm (probable).	cytoplasm
Q7W2B7	AROK_BORPA	cytoplasm (probable).	cytoplasm
Q7W2B9	CYAY_BORPA		inner membrane
Q7W2D1	RS11_BORPA		cytoplasm
Q7W2D2	RS13_BORPA		cytoplasm
Q7W2D6	RL15_BORPA		cytoplasm
Q7W2D8	RS5_BORPA		cytoplasm
Q7W2D9	RL18_BORPA		cytoplasm
Q7W2E1	RS8_BORPA		cytoplasm
Q7W2E3	RL5_BORPA		cytoplasm
Q7W2E4	RL24_BORPA		cytoplasm
Q7W2E7	RS17_BORPA		cytoplasm
Q7W2E9	RL16_BORPA		cytoplasm
Q7W2F0	RS3_BORPA		cytoplasm
Q7W2F1	RL22_BORPA		cytoplasm
Q7W2F2	RS19_BORPA		cytoplasm
Q7W2F3	RL2_BORPA		cytoplasm
Q7W2F5	RL4_BORPA		cytoplasm
Q7W2F6	RL3_BORPA		cytoplasm
Q7W2F7	RS10_BORPA		cytoplasm
Q7W2F9	RS7_BORPA		cytoplasm
Q7W2G0	RS12_BORPA		cytoplasm
Q7W2G8	RPOC_BORPA		cytoplasm
Q7W2G9	RPOB_BORPA		cytoplasm
Q7W2H1	RL10_BORPA		cytoplasm
Q7W2H2	RL1_BORPA		cytoplasm
Q7W2H3	RL11_BORPA		cytoplasm
Q7W2I0	GIDB_BORPA		cytoplasm
Q7W2I1	GIDA_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W2J0	TRME_BORPA		cytoplasm
Q7W2K1	OXAA_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W2K2	Y4404_BORPA		inner membrane
Q7W2K3	RNPA_BORPA		extracellular
Q7W2K5	DNAA_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W2M1	RIBB_BORPA		cytoplasm
Q7W2N4	GLUQ_BORPA		cytoplasm
Q7W2N5	SMG_BORPA		cytoplasm
Q7W2N9	ARGD2_BORPA	cytoplasm (probable).	cytoplasm
Q7W2P4	ACEK_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W2W9	MSCL_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W2X7	HIS2_BORPA	cytoplasm (by similarity).	cytoplasm

Q7W2X8	HIS3_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W2X9	HIS6_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W2Y0	HIS4_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W2Y1	HIS5_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W2Y2	HIS7_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W2Y3	HIS82_BORPA		cytoplasm
Q7W2Y4	HISX_BORPA		cytoplasm
Q7W2Y5	HIS1_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W2Y6	MURA_BORPA	cytoplasm (probable).	inner membrane
Q7W321	GLMU_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W334	GLMS_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W388	TRPD_BORPA		cytoplasm
Q7W393	APAG_BORPA		inner membrane
Q7W3A8	ATPA_BORPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W3A9	ATPG_BORPA		inner membrane
Q7W3B1	ATPE_BORPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W3B3	DCUP_BORPA	cytoplasm (probable).	cytoplasm
Q7W3E7	PANB_BORPA		inner membrane
Q7W3F2	GSH1_BORPA		cytoplasm
Q7W3F8	METX_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W3H5	PYRE_BORPA		cytoplasm
Q7W3H6	GATB_BORPA		cytoplasm
Q7W3H7	GATA_BORPA		cytoplasm
Q7W3H8	GATC_BORPA		cytoplasm
Q7W3I6	ARGB_BORPA	cytoplasm (probable).	cytoplasm
Q7W3J5	Y4038_BORPA		cytoplasm
Q7W3M5	Y4006_BORPA		inner membrane
Q7W3N6	UBIE_BORPA		cytoplasm
Q7W3P5	THIG_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W3R1	GLPK_BORPA		cytoplasm
Q7W3R8	COAE_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W3S6	ARGJ_BORPA	cytoplasm (probable).	cytoplasm
Q7W3U2	THIE_BORPA		cytoplasm
Q7W3U5	Y3932_BORPA		inner membrane
Q7W3U6	RUVX_BORPA	cytoplasm (potential).	cytoplasm
Q7W3U7	PYRB_BORPA		cytoplasm
Q7W3V0	APAH_BORPA		extracellular
Q7W3V2	MTGA_BORPA	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q7W3V9	AROQ_BORPA		cytoplasm
Q7W3W2	PRMA_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W3Y8	Y3887_BORPA		inner membrane
Q7W3Z2	RS9_BORPA		cytoplasm
Q7W3Z3	ARGC_BORPA	cytoplasm (probable).	cytoplasm
Q7W3Z8	SYX_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W400	GLYA2_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W401	NRDR_BORPA		cytoplasm
Q7W417	URE1_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W418	UREE_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W476	TOLB_BORPA	periplasm (potential).	periplasm
Q7W482	NUDH_BORPA		cytoplasm

Q7W496	LGT_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W497	ILVD2_BORPA		cytoplasm
Q7W4A6	MRAZ_BORPA		inner membrane
Q7W4A7	MRAW_BORPA		cytoplasm
Q7W4B1	MRAY_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W4B2	MURD_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W4B4	MURG_BORPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W4B5	MURC_BORPA	cytoplasm (probable).	cytoplasm
Q7W4B6	DDL_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W4D4	Y3732_BORPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7W4J0	PHS_BORPA		cytoplasm
Q7W4J6	KSGA_BORPA		cytoplasm
Q7W4K0	DEF2_BORPA		cytoplasm
Q7W4N4	PUR5_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W4N9	FUMC_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W4Q1	PDXH_BORPA		cytoplasm
Q7W4S9	MINC_BORPA		inner membrane
Q7W4T6	RUVB_BORPA		cytoplasm
Q7W4T9	RUVA_BORPA		cytoplasm
Q7W4U0	RUVC_BORPA		cytoplasm
Q7W4W6	PYRD_BORPA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W4W7	ATE_BORPA	cytoplasm (potential).	cytoplasm
Q7W4W8	LFTR_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W509	MURB_BORPA	cytoplasm (probable).	inner membrane
Q7W510	DAPB_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W513	PPNK_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W514	HRCB_BORPA		cytoplasm
Q7W515	HEMH_BORPA	cytoplasm (by similarity).	extracellular
Q7W517	GRPE_BORPA	cytoplasm (probable).	cytoplasm
Q7W519	DNAK_BORPA		cytoplasm
Q7W520	DNAJ_BORPA	cytoplasm (probable).	cytoplasm
Q7W536	ATKA_BORPA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q7W543	LOLA_BORPA	periplasm (by similarity).	periplasm
Q7W561	Y3440_BORPA		inner membrane
Q7W566	ILVC_BORPA		cytoplasm
Q7W569	RS15_BORPA		cytoplasm
Q7W575	TPIS_BORPA		cytoplasm
Q7W5C9	ISPD_BORPA		cytoplasm
Q7W5D0	ISPF_BORPA		extracellular
Q7W5G3	PNCB_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W5G8	TRPB_BORPA		cytoplasm
Q7W5G9	TRPA_BORPA		cytoplasm

Q7W5H7	Y3314_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W5I0	Y3311_BORPA	cytoplasm (potential).	inner membrane
Q7W5I2	RL32_BORPA		cytoplasm
Q7W5I3	PLSX_BORPA		cytoplasm
Q7W5I4	FABH_BORPA	cytoplasm (probable).	cytoplasm
Q7W5I7	ACP_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W5J4	LEPA_BORPA		cytoplasm
Q7W5J6	RNC_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W5K6	AMPA_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W5N7	KDSA_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W5N9	ENO_BORPA	cytoplasm (by similarity).	extracellular
Q7W5Q8	MDH_BORPA		cytoplasm
Q7W5V2	Y3185_BORPA		cytoplasm
Q7W5Z6	UBIG_BORPA		cytoplasm
Q7W5Z9	SERC_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W603	KCY_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W609	HLDD_BORPA		cytoplasm
Q7W641	DADA_BORPA		cytoplasm
Q7W669	Y3058_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W670	SURE_BORPA	cytoplasm (potential).	periplasm
Q7W676	RUMA_BORPA		cytoplasm
Q7W690	Y3031_BORPA		inner membrane
Q7W6B1	RPOZ_BORPA		cytoplasm
Q7W6B2	KGUA_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W6C1	RNPH_BORPA		cytoplasm
Q7W6C6	HAM1_BORPA		extracellular
Q7W6I5	PSD_BORPA		cytoplasm
Q7W6L4	GLGB_BORPA		extracellular
Q7W6L5	GLGA_BORPA		cytoplasm
Q7W6N6	RS16_BORPA		cytoplasm
Q7W6N7	RIMM_BORPA	cytoplasm (potential).	cytoplasm
Q7W6P3	SYV_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W6P4	NDK_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W6P6	ISPG_BORPA		cytoplasm
Q7W6Q0	ENGA_BORPA		cytoplasm
Q7W6Q1	HIS81_BORPA		cytoplasm
Q7W6Q2	HFQ_BORPA		cytoplasm
Q7W6Q6	HISZ_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W6Q7	PURA_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W6Q9	RS21_BORPA		cytoplasm
Q7W733	NAPA_BORPA	periplasm (by similarity).	periplasm
Q7W736	CCMA_BORPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W740	CCME_BORPA	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q7W749	LEUD2_BORPA		cytoplasm
Q7W785	HEM3_BORPA		cytoplasm
Q7W791	METE_BORPA		cytoplasm
Q7W7C8	RL20_BORPA		cytoplasm

Q7W7C9	RL35_BORPA		cytoplasm
Q7W7E8	CLPS_BORPA		cytoplasm
Q7W7F7	LPXK_BORPA		cytoplasm
Q7W7G0	KAD_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W7G3	RS20_BORPA		cytoplasm
Q7W7H6	ARGD1_BORPA	cytoplasm (probable).	cytoplasm
Q7W7H7	OTC_BORPA	cytoplasm (probable).	cytoplasm
Q7W7H8	ASSY_BORPA	cytoplasm (probable).	cytoplasm
Q7W7H9	KUP_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W7I7	RLMB_BORPA		cytoplasm
Q7W7P5	RL9_BORPA		cytoplasm
Q7W7P6	RS18_BORPA		cytoplasm
Q7W7P8	RS6_BORPA		cytoplasm
Q7W7P9	Y2465_BORPA		inner membrane
Q7W7Q0	DXS_BORPA		cytoplasm
Q7W7Q2	EX7S_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W7T8	Y2422_BORPA		inner membrane
Q7W7U0	HEM6_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W7U1	NADD_BORPA		cytoplasm
Q7W7U3	Y2417_BORPA		inner membrane
Q7W7U4	Y2416_BORPA	cytoplasm (potential).	inner membrane
Q7W850	ASPD2_BORPA		cytoplasm
Q7W859	TILS_BORPA	cytoplasm (probable).	cytoplasm
Q7W860	ACCA_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W865	LPXH_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W866	UPPP_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W880	MUTS_BORPA		cytoplasm
Q7W8H4	TPMT_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W8J2	Y2141_BORPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7W8Q6	PSTB_BORPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W8R3	GLMM_BORPA		cytoplasm
Q7W8R6	RRMJ_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W8T0	LOLD_BORPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W8U9	HSCA_BORPA		cytoplasm
Q7W8V0	HSCB_BORPA		inner membrane
Q7W8V6	UVRB_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W8X1	CLPX_BORPA		cytoplasm
Q7W8X2	CLPP_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W8X3	TIG_BORPA		cytoplasm
Q7W8Y2	DAPD_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W8Z2	SYI_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W8Z8	DUT_BORPA		extracellular
Q7W910	GSHB_BORPA		cytoplasm
Q7W917	AQPZ_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q7W929	LEU3_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W930	LEUD1_BORPA		cytoplasm
Q7W931	LEU2_BORPA		cytoplasm
Q7W950	AROC_BORPA		cytoplasm
Q7W954	Y1919_BORPA		inner membrane
Q7W972	PAND_BORPA		cytoplasm
Q7W977	ECTD_BORPA		cytoplasm
Q7W978	ECTC_BORPA		cytoplasm
Q7W979	ECTB_BORPA		cytoplasm
Q7W980	ECTA_BORPA		cytoplasm
Q7W993	GLNE_BORPA		cytoplasm
Q7W9A4	RBFA_BORPA	cytoplasm (potential).	cytoplasm
Q7W9A7	Y1860_BORPA		inner membrane
Q7W9B4	ISPH_BORPA		cytoplasm
Q7W9B6	RADC_BORPA		cytoplasm
Q7W9B8	RL31B_BORPA		cytoplasm
Q7W9D1	CRCB_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W9E3	Y1817_BORPA		inner membrane
Q7W9E6	CLPB_BORPA	cytoplasm (probable).	cytoplasm
Q7W9J1	QUEF_BORPA	cytoplasm (probable).	cytoplasm
Q7W9J5	NAGZ_BORPA	cytoplasm (by similarity).	extracellular
Q7W9J6	ACPS_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W9J7	PDXJ_BORPA	cytoplasm (probable).	cytoplasm
Q7W9L9	RL28_BORPA		cytoplasm
Q7W9M7	PROA_BORPA	cytoplasm (by similarity).	cytoplasm
Q7W9N2	Y1723_BORPA		inner membrane
Q7W9N7	MSBA_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7W9Q2	FETP_BORPA		cytoplasm
Q7W9Q5	RPIA_BORPA		cytoplasm
Q7W9T4	ORN_BORPA	cytoplasm (potential).	cytoplasm
Q7W9T9	RL19_BORPA		cytoplasm
Q7W9U5	CYSA_BORPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7W9W1	DNAE2_BORPA	cytoplasm (probable).	cytoplasm
Q7WA16	Y1579_BORPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7WA31	KTHY_BORPA		cytoplasm
Q7WA41	SSRP_BORPA	cytoplasm (potential).	cytoplasm
Q7WA44	Y1543_BORPA		cytoplasm
Q7WA46	RNH2_BORPA	cytoplasm (potential).	extracellular
Q7WA47	LPXB_BORPA		cytoplasm
Q7WA49	FABZ_BORPA	cytoplasm (by similarity).	cytoplasm
Q7WA50	LPXD_BORPA		cytoplasm
Q7WA54	DXR_BORPA		cytoplasm
Q7WA56	UPPS_BORPA		inner membrane
Q7WA57	RRF_BORPA	cytoplasm (by similarity).	cytoplasm
Q7WA60	RS2_BORPA		cytoplasm
Q7WA62	GLND_BORPA		cytoplasm
Q7WA91	FLGI_BORPA	periplasm (by similarity).	periplasm

Q7WA92	FLGH_BORPA	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q7WAB3	FLHD_BORPA	cytoplasm (by similarity).	cytoplasm
Q7WAE4	ARLY_BORPA	cytoplasm (probable).	cytoplasm
Q7WAK8	PPCK_BORPA	cytoplasm (by similarity).	cytoplasm
Q7WAK9	NORM_BORPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WAV6	GUAA_BORPA		cytoplasm
Q7WAY2	TAL_BORPA	cytoplasm (probable).	cytoplasm
Q7WAY7	RECR_BORPA		cytoplasm
Q7WAY8	Y1222_BORPA		inner membrane
Q7WB15	PUR7_BORPA		cytoplasm
Q7WB43	PGK_BORPA	cytoplasm (potential).	cytoplasm
Q7WB58	UPP_BORPA		cytoplasm
Q7WB60	QUEA_BORPA	cytoplasm (potential).	cytoplasm
Q7WB68	Y1139_BORPA		inner membrane
Q7WBA1	FDHD_BORPA	cytoplasm (potential).	cytoplasm
Q7WBC1	Y1084_BORPA		inner membrane
Q7WBD0	Y1075_BORPA		inner membrane
Q7WBF2	Y1051_BORPA		inner membrane
Q7WBG5	ADD_BORPA		extracellular
Q7WC23	TRMB_BORPA		cytoplasm
Q7WC32	HTPG_BORPA	cytoplasm (by similarity).	cytoplasm
Q7WC84	Y444_BORPA		extracellular
Q7WC96	53DR_BORPA		extracellular
Q7WC98	ILVD1_BORPA		cytoplasm
Q7WCA9	CAPP_BORPA		cytoplasm
Q7WCE4	HEM1_BORPA		cytoplasm
Q7WCK0	LEXA_BORBR		cytoplasm
Q7WCL8	ATKA_BORBR	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q7WCM5	LOLA_BORBR	periplasm (by similarity).	periplasm
Q7WCP1	Y3890_BORBR		inner membrane
Q7WCP6	ILVC_BORBR		cytoplasm
Q7WCP9	RS15_BORBR		cytoplasm
Q7WCQ5	TPIS_BORBR		cytoplasm
Q7WCW3	ISPD_BORBR		cytoplasm
Q7WCW4	ISPF_BORBR		extracellular
Q7WCZ8	PNCB_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WD04	TRPB_BORBR		cytoplasm
Q7WD05	TRPA_BORBR		cytoplasm
Q7WD13	Y3765_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WD16	Y3762_BORBR	cytoplasm (potential).	inner membrane
Q7WD18	RL32_BORBR		cytoplasm
Q7WD19	PLSX_BORBR		cytoplasm
Q7WD20	FABH_BORBR	cytoplasm (probable).	cytoplasm
Q7WD23	ACP_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WD30	LEPA_BORBR		cytoplasm
Q7WD32	RNC_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WD42	AMPA_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WD73	KDSA_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WD75	ENO_BORBR	cytoplasm (by similarity).	extracellular

Q7WD94	MDH_BORBR		cytoplasm
Q7WDI4	TRME_BORBR		cytoplasm
Q7WDJ6	Y4992_BORBR		inner membrane
Q7WDJ7	RNPA_BORBR		extracellular
Q7WDJ9	DNAA_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WDL7	RIBB_BORBR		cytoplasm
Q7WDN0	GLUQ_BORBR		cytoplasm
Q7WDN1	SMG_BORBR		cytoplasm
Q7WDN7	ARGD2_BORBR	cytoplasm (probable).	cytoplasm
Q7WDP2	ACEK_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WDW9	MSCL_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WDX7	HIS2_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WDX8	HIS3_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WDX9	HIS6_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WDY0	HIS4_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WDY1	HIS5_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WDY2	HIS7_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WDY3	HIS82_BORBR		cytoplasm
Q7WDY4	HISX_BORBR		cytoplasm
Q7WDY5	HIS1_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WDY6	MURA_BORBR	cytoplasm (probable).	inner membrane
Q7WE21	GLMU_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WE36	GLMS_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WE57	ASPD3_BORBR		cytoplasm
Q7WE96	SELU_BORBR		cytoplasm
Q7WE97	SELD_BORBR		cytoplasm
Q7WEA9	Y4725_BORBR		inner membrane
Q7WEK7	TRPD_BORBR		cytoplasm
Q7WEL1	APAG_BORBR		inner membrane
Q7WEM7	ATPA_BORBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WEM8	ATPG_BORBR		inner membrane
Q7WEN0	ATPE_BORBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WEN2	DCUP_BORBR	cytoplasm (probable).	cytoplasm
Q7WER7	PANB_BORBR		inner membrane
Q7WES2	GSH1_BORBR		cytoplasm
Q7WES8	METX_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WEU9	PYRE_BORBR		cytoplasm
Q7WEV0	GATB_BORBR		cytoplasm
Q7WEV1	GATA_BORBR		cytoplasm
Q7WEV2	GATC_BORBR		cytoplasm
Q7WEW1	ARGB_BORBR	cytoplasm (probable).	cytoplasm
Q7WEX0	Y4511_BORBR		cytoplasm
Q7WF01	Y4479_BORBR		inner membrane
Q7WF12	UBIE_BORBR		cytoplasm
Q7WF21	THIG_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WF38	GLPK_BORBR		cytoplasm
Q7WF46	COAE_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WF54	ARGJ_BORBR	cytoplasm (probable).	cytoplasm
Q7WF72	THIE_BORBR		cytoplasm
Q7WF75	Y4405_BORBR		inner membrane

Q7WF76	RUVX_BORBR	cytoplasm (potential).	cytoplasm
Q7WF77	PYRB_BORBR		cytoplasm
Q7WF80	APAH_BORBR		extracellular
Q7WF82	MTGA_BORBR	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q7WF89	AROQ_BORBR		cytoplasm
Q7WF92	PRMA_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WFC0	Y4360_BORBR		inner membrane
Q7WFC4	RS9_BORBR		cytoplasm
Q7WFC5	ARGC_BORBR	cytoplasm (probable).	cytoplasm
Q7WFD0	SYT_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WFD2	GLYA2_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WFD3	NRDR_BORBR		cytoplasm
Q7WFF2	UREE_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WFN4	TOLB_BORBR	periplasm (potential).	periplasm
Q7WFP0	NUDH_BORBR		cytoplasm
Q7WFAQ4	LGT_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WFAQ5	ILVD3_BORBR		cytoplasm
Q7WFR4	MRAZ_BORBR		inner membrane
Q7WFR5	MRAW_BORBR		cytoplasm
Q7WFR9	MRAY_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WFS0	MURD_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WFS2	MURG_BORBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WFS3	MURC_BORBR	cytoplasm (probable).	cytoplasm
Q7WFS4	DDL_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WFU2	Y4178_BORBR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7WG14	PHS_BORBR		cytoplasm
Q7WG20	KSGA_BORBR		cytoplasm
Q7WG25	DEF2_BORBR		cytoplasm
Q7WG60	PUR5_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WG65	FUMC_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WG77	PDXH_BORBR		cytoplasm
Q7WGA5	MINC_BORBR		inner membrane
Q7WGB3	RUVB_BORBR		cytoplasm
Q7WGB6	RUVA_BORBR		cytoplasm
Q7WGB7	RUVC_BORBR		cytoplasm
Q7WGE4	PYRD_BORBR	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WGE5	ATE_BORBR	cytoplasm (potential).	cytoplasm
Q7WGE6	LFTR_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WGH4	MURB_BORBR	cytoplasm (probable).	inner membrane
Q7WGH5	DAPB_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WGH8	PPNK_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WGH9	HRCA_BORBR		cytoplasm
Q7WGI0	HEMH_BORBR	cytoplasm (by similarity).	extracellular
Q7WGI2	GRPE_BORBR	cytoplasm (probable).	cytoplasm

Q7WGI4	DNAK_BORBR		cytoplasm
Q7WGI5	DNAJ_BORBR	cytoplasm (probable).	cytoplasm
Q7WVK3	Y3586_BORBR		cytoplasm
Q7WGT9	UBIG_BORBR		cytoplasm
Q7WGU2	SERC_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WGU9	HLDD_BORBR		cytoplasm
Q7WGW1	CYSA_BORBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WGW8	RL19_BORBR		cytoplasm
Q7WGX3	ORN_BORBR	cytoplasm (potential).	cytoplasm
Q7WH03	RPIA_BORBR		cytoplasm
Q7WH06	FETP_BORBR		cytoplasm
Q7WH20	MSBA_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WH25	Y3385_BORBR		inner membrane
Q7WH30	PROA_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WH39	RL28_BORBR		cytoplasm
Q7WH63	PDXJ_BORBR	cytoplasm (probable).	cytoplasm
Q7WH64	ACPS_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WH65	NAGZ_BORBR	cytoplasm (by similarity).	extracellular
Q7WH69	QUEF_BORBR	cytoplasm (probable).	cytoplasm
Q7WHB6	CLPB_BORBR	cytoplasm (probable).	cytoplasm
Q7WHB9	Y3289_BORBR		inner membrane
Q7WHD3	CRCB_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WHE7	RL31B_BORBR		cytoplasm
Q7WHF0	RADC_BORBR		cytoplasm
Q7WHF2	ISPH_BORBR		cytoplasm
Q7WHG0	Y3248_BORBR		inner membrane
Q7WHG3	RBFA_BORBR	cytoplasm (potential).	cytoplasm
Q7WHH4	GLNE_BORBR		cytoplasm
Q7WHI7	ECTA_BORBR		cytoplasm
Q7WHI8	ECTB_BORBR		cytoplasm
Q7WHI9	ECTC_BORBR		cytoplasm
Q7WHJ0	ECTD_BORBR		cytoplasm
Q7WHJ6	PAND_BORBR		cytoplasm
Q7WHL9	RS16_BORBR		cytoplasm
Q7WHM0	RIMM_BORBR	cytoplasm (potential).	cytoplasm
Q7WHM6	SYV_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WHM7	NDK_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WHN0	ISPG_BORBR		cytoplasm
Q7WHN4	ENGA_BORBR		cytoplasm
Q7WHN5	HIS81_BORBR		cytoplasm
Q7WHN6	HFQ_BORBR		cytoplasm
Q7WHP0	HISZ_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WHP1	PURA_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WHP3	RS21_BORBR		cytoplasm
Q7WHW5	HMP_BORBR		cytoplasm
Q7WHW8	DNAE2_BORBR	cytoplasm (probable).	cytoplasm
Q7WI07	DADA_BORBR		cytoplasm
Q7WI35	Y3021_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q7WI36	SURE_BORBR	cytoplasm (potential).	periplasm
Q7WI42	RUMA_BORBR		cytoplasm
Q7WI59	Y2997_BORBR		inner membrane
Q7WI79	RPOZ_BORBR		cytoplasm
Q7WI80	KGUA_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WI90	RNPH_BORBR		cytoplasm
Q7WI95	HAM1_BORBR		extracellular
Q7WIF7	PSD_BORBR		cytoplasm
Q7WII7	GLGB_BORBR		extracellular
Q7WIJ0	GLGA_BORBR		cytoplasm
Q7WIN3	LEUD2_BORBR		cytoplasm
Q7WIP4	CCME_BORBR	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q7WIP8	CCMA_BORBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WIQ1	NAPA_BORBR	periplasm (by similarity).	periplasm
Q7WJ43	Y2657_BORBR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7WJ55	HUTG_BORBR		cytoplasm
Q7WJ63	KTHY_BORBR		cytoplasm
Q7WJ73	SSRP_BORBR	cytoplasm (potential).	cytoplasm
Q7WJ78	Y2621_BORBR		cytoplasm
Q7WJ80	RNH2_BORBR	cytoplasm (potential).	extracellular
Q7WJ81	LPXB_BORBR		cytoplasm
Q7WJ83	FABZ_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WJ84	LPXD_BORBR		cytoplasm
Q7WJ88	DXR_BORBR		cytoplasm
Q7WJ90	UPPS_BORBR		inner membrane
Q7WJ91	RRF_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WJ94	RS2_BORBR		cytoplasm
Q7WJ96	GLND_BORBR		cytoplasm
Q7WJD1	FLGI_BORBR	periplasm (by similarity).	periplasm
Q7WJD2	FLGH_BORBR	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q7WJI7	ARLY_BORBR	cytoplasm (probable).	cytoplasm
Q7WJP7	ILVD2_BORBR		cytoplasm
Q7WJQ9	PPCK_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WJR0	NORM_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WK13	GUAA_BORBR		cytoplasm
Q7WK40	LOLD_BORBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WK59	HSCA_BORBR		cytoplasm
Q7WK60	HSCB_BORBR		inner membrane
Q7WK66	UVRB_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WK82	CLPX_BORBR		cytoplasm
Q7WK83	CLPP_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WK84	TIG_BORBR		cytoplasm
Q7WKD5	SYI_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WKE3	DUT_BORBR		extracellular

Q7WKF5	GSHB_BORBR		cytoplasm
Q7WKG2	AQPZ_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WKH4	LEU3_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WKH5	LEUD1_BORBR		cytoplasm
Q7WKH6	LEU2_BORBR		cytoplasm
Q7WKJ5	AROC_BORBR		cytoplasm
Q7WKJ9	Y2107_BORBR		inner membrane
Q7WKM1	HEM3_BORBR		cytoplasm
Q7WKM7	METE_BORBR		cytoplasm
Q7WKR6	RL20_BORBR		cytoplasm
Q7WKR7	RL35_BORBR		cytoplasm
Q7WKT7	CLPS_BORBR		cytoplasm
Q7WKU5	LPXK_BORBR		cytoplasm
Q7WKU8	KAD_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WKV2	RS20_BORBR		cytoplasm
Q7WKW5	ARGD1_BORBR	cytoplasm (probable).	cytoplasm
Q7WKW6	OTC_BORBR	cytoplasm (probable).	cytoplasm
Q7WKW7	ASSY_BORBR	cytoplasm (probable).	cytoplasm
Q7WKW8	KUP_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WKX5	RLMB_BORBR		cytoplasm
Q7WL32	RL9_BORBR		cytoplasm
Q7WL33	RS18_BORBR		cytoplasm
Q7WL35	RS6_BORBR		cytoplasm
Q7WL36	Y1913_BORBR		inner membrane
Q7WL37	DXS_BORBR		cytoplasm
Q7WL39	EX7S_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WL78	Y1871_BORBR		inner membrane
Q7WL80	HEM6_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WL81	NADD_BORBR		cytoplasm
Q7WL83	Y1866_BORBR		inner membrane
Q7WL84	Y1865_BORBR	cytoplasm (potential).	inner membrane
Q7WLE5	NPD_BORBR	cytoplasm (probable).	cytoplasm
Q7WLJ8	ASPD2_BORBR		cytoplasm
Q7WLK7	TILS_BORBR	cytoplasm (probable).	cytoplasm
Q7WLK8	ACCA_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WLL3	LPXH_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WLL4	UPPP_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WLT5	MUTS_BORBR		cytoplasm
Q7WM36	TPMT_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WM55	Y1538_BORBR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q7WMC3	PSTB_BORBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WMD0	GLMM_BORBR		cytoplasm
Q7WMD3	RRMJ_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WME6	TAL_BORBR	cytoplasm (probable).	cytoplasm
Q7WMF1	RECR_BORBR		cytoplasm
Q7WMF2	Y1439_BORBR		inner membrane

Q7WMI1	PUR7_BORBR		cytoplasm
Q7WMK9	PGK_BORBR	cytoplasm (potential).	cytoplasm
Q7WMM5	UPP_BORBR		cytoplasm
Q7WMM7	QUEA_BORBR	cytoplasm (potential).	cytoplasm
Q7WMN5	Y1355_BORBR		inner membrane
Q7WMN7	LNT_BORBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WMS0	FDHD_BORBR	cytoplasm (potential).	cytoplasm
Q7WMU0	Y1300_BORBR		inner membrane
Q7WMU9	Y1291_BORBR		inner membrane
Q7WMX3	Y1267_BORBR		inner membrane
Q7WMY6	ADD_BORBR		extracellular
Q7WNK2	RBLL_BORBR		cytoplasm
Q7WNS4	CH60_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WNT2	NUSB_BORBR		cytoplasm
Q7WNT3	RISB_BORBR		cytoplasm
Q7WNT8	PHNC_BORBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WNU4	COAD_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WNY2	PTH_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WNY3	RL25_BORBR		cytoplasm
Q7WNY4	KPRS_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WNY5	ISPE_BORBR		cytoplasm
Q7WNY6	LOLB_BORBR	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q7WNZ3	HGD_BORBR		cytoplasm
Q7WP01	G6PI_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WP13	TYSY_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WP29	GCSP_BORBR		cytoplasm
Q7WP31	GCST_BORBR		cytoplasm
Q7WP40	DCD_BORBR		extracellular
Q7WP46	Y839_BORBR		inner membrane
Q7WP51	OAT_BORBR		cytoplasm
Q7WP62	MODC_BORBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WPH6	GLYA1_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WQ22	TRMB_BORBR		cytoplasm
Q7WQ31	HTPG_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WQ88	Y445_BORBR		extracellular
Q7WQA0	53DR_BORBR		extracellular
Q7WQA2	ILVD1_BORBR		cytoplasm
Q7WQB3	CAPP_BORBR		cytoplasm
Q7WQF0	HEM1_BORBR		cytoplasm
Q7WQL7	RL27_BORBR		cytoplasm
Q7WQL9	PROB_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WQM6	THIC_BORBR		cytoplasm
Q7WQN2	GPMA_BORBR		cytoplasm
Q7WQN5	SECB_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WQN6	GPDA_BORBR	cytoplasm (probable).	cytoplasm
Q7WQP1	CAAL_BORBR		cytoplasm
Q7WQR6	ASPD1_BORBR		cytoplasm
Q7WQS8	FMT_BORBR		cytoplasm
Q7WQS9	DEF1_BORBR		cytoplasm

Q7WQW5	UNG_BORBR	cytoplasm (by similarity).	extracellular
Q7WQX5	SAHH_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WQZ3	HSLV_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WQZ4	HSLU_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WR00	LIPA_BORBR	cytoplasm (potential).	cytoplasm
Q7WR01	LIPB_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WR02	Y170_BORBR		inner membrane
Q7WR83	DGTL1_BORBR		cytoplasm
Q7WR84	AROB_BORBR	cytoplasm (probable).	cytoplasm
Q7WR85	AROK_BORBR	cytoplasm (probable).	cytoplasm
Q7WR87	CYAY_BORBR		inner membrane
Q7WR99	RS11_BORBR		cytoplasm
Q7WRA0	RS13_BORBR		cytoplasm
Q7WRA4	RL15_BORBR		cytoplasm
Q7WRA6	RS5_BORBR		cytoplasm
Q7WRA7	RL18_BORBR		cytoplasm
Q7WRA9	RS8_BORBR		cytoplasm
Q7WRB1	RL5_BORBR		cytoplasm
Q7WRB2	RL24_BORBR		cytoplasm
Q7WRB6	RS17_BORBR		cytoplasm
Q7WRB8	RL16_BORBR		cytoplasm
Q7WRB9	RS3_BORBR		cytoplasm
Q7WRC0	RL22_BORBR		cytoplasm
Q7WRC1	RS19_BORBR		cytoplasm
Q7WRC2	RL2_BORBR		cytoplasm
Q7WRC4	RL4_BORBR		cytoplasm
Q7WRC5	RL3_BORBR		cytoplasm
Q7WRC6	RS10_BORBR		cytoplasm
Q7WRC8	RS7_BORBR		cytoplasm
Q7WRC9	RS12_BORBR		cytoplasm
Q7WRD8	RPOC_BORBR		cytoplasm
Q7WRD9	RPOB_BORBR		cytoplasm
Q7WRE1	RL10_BORBR		cytoplasm
Q7WRE2	RL1_BORBR		cytoplasm
Q7WRE3	RL11_BORBR		cytoplasm
Q7WRF0	GIDB_BORBR		cytoplasm
Q7WRF1	GIDA_BORBR	cytoplasm (by similarity).	cytoplasm
Q7WS85	MDH_SALPA		cytoplasm
Q7WTJ2	DMPP_ACICA		cytoplasm
Q7WTJ6	DMPL_ACICA		cytoplasm
Q7WTR3	MDTK_ERWAM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WUM8	HMP_RHIME		cytoplasm
Q7WXB8	CCME_RALEU	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q7WXC1	CCMA_RALEU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q7WXK8	KUP_RALEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q7WZ32	CH603_METCA	cytoplasm (by similarity).	cytoplasm
Q7WZE4	RPOZ_SHEVI		cytoplasm

Q7WZE5	KGUA_SHEVI	cytoplasm (by similarity).	cytoplasm
Q7X222	TPIS_KLEPN		cytoplasm
Q7X2C7	BCHE_RHOGE		cytoplasm
Q7X2N3	SYV_SPHEL	cytoplasm (by similarity).	cytoplasm
Q7X2N6	SSRP_SPHEL	cytoplasm (by similarity).	cytoplasm
Q7X3P1	SPEB_PROMI		cytoplasm
Q7X3X5	MDH_MORS2		cytoplasm
Q7X4S1	XCBA_NEIME		cytoplasm
Q7X5H2	IDI_CITFR	cytoplasm (by similarity).	cytoplasm
Q7ZAI9	XERC_VIBVU	cytoplasm (by similarity).	nucleoid
Q7ZAJ0	XERD_VIBVU	cytoplasm (by similarity).	nucleoid
Q7ZAJ5	XERC_SHEON	cytoplasm (by similarity).	nucleoid
Q7ZAJ8	XERD_SHEON	cytoplasm (by similarity).	nucleoid
Q7ZAL9	XERC_SHIFL	cytoplasm (by similarity).	nucleoid
Q7ZAM0	XERD_SHIFL	cytoplasm (by similarity).	nucleoid
Q7ZAN6	XERD_BRUSU	cytoplasm (by similarity).	nucleoid
Q7ZAN7	XERC_BRUSU	cytoplasm (by similarity).	nucleoid
Q81ZK5	Y2358_NITEU		cytoplasm
Q81ZU6	GLGB_NITEU		extracellular
Q81ZZ2	UBIG_NITEU		cytoplasm
Q81ZZ5	METX_NITEU	cytoplasm (by similarity).	cytoplasm
Q81ZZ7	GLNE_NITEU		cytoplasm
Q81ZZ8	UBIE_NITEU		cytoplasm
Q81ZZ9	PRMA_NITEU	cytoplasm (by similarity).	cytoplasm
Q820A2	RRMJ_NITEU	cytoplasm (by similarity).	cytoplasm
Q820A4	LGT_NITEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q820B4	LNT_COXBU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q820B5	UBIG_COXBU		cytoplasm
Q820C5	UBIG_SHIFL		cytoplasm
Q820C8	LNT_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q820C9	LNT_NITEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q820D2	HIS5_NITEU	cytoplasm (by similarity).	cytoplasm
Q820H8	LEPA_NITEU		cytoplasm
Q820I0	RNC_NITEU	cytoplasm (by similarity).	cytoplasm
Q820I1	ACPS_NITEU	cytoplasm (by similarity).	cytoplasm
Q820I7	PROA_NITEU	cytoplasm (by similarity).	cytoplasm
Q820J1	GATA_NITEU		cytoplasm
Q820J5	DGTL1_NITEU		cytoplasm
Q820J6	MIAA_NITEU		cytoplasm
Q820J7	FMT_NITEU		cytoplasm
Q820K1	PYRE_NITEU		cytoplasm
Q820K4	TRMD_NITEU	cytoplasm (potential).	cytoplasm
Q820M0	LEU1_NITEU		cytoplasm
Q820M4	PROB_NITEU	cytoplasm (by similarity).	cytoplasm
Q820M6	SYI_NITEU	cytoplasm (by similarity).	cytoplasm
Q820N1	ARGB_NITEU	cytoplasm (probable).	cytoplasm
Q820N3	COAD_NITEU	cytoplasm (by similarity).	cytoplasm
Q820N6	UVRC_NITEU	cytoplasm (by similarity).	cytoplasm

Q820P2	TRUB_NITEU		cytoplasm
Q820P8	TRUA_NITEU		cytoplasm
Q820Q0	HIS6_NITEU	cytoplasm (by similarity).	cytoplasm
Q820Q1	DCD_NITEU		extracellular
Q820Q8	RL18_NITEU		cytoplasm
Q820Q9	RL5_NITEU		cytoplasm
Q820R0	RL24_NITEU		cytoplasm
Q820R1	RS3_NITEU		cytoplasm
Q820S0	SERC_NITEU	cytoplasm (by similarity).	cytoplasm
Q820S3	RUVA_NITEU		cytoplasm
Q820S4	ATPE_NITEU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q820V9	CCA_COXBU		cytoplasm
Q820W3	MIAA_COXBU		cytoplasm
Q820W7	FABZ_COXBU	cytoplasm (by similarity).	cytoplasm
Q820X0	FABH_COXBU	cytoplasm (probable).	cytoplasm
Q820X1	TRMD_COXBU	cytoplasm (potential).	cytoplasm
Q820X3	MURG_COXBU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q820Z9	TRMA_SHIFL		cytoplasm
Q821A6	CCA_SHIFL		cytoplasm
Q821A9	PTFB1_SHIFL	cytoplasm (potential).	inner membrane
Q821B1	ECOT_SHIFL	periplasm (by similarity).	periplasm
Q821B6	CITD_SHIFL	cytoplasm (by similarity).	cytoplasm
Q821C1	GSA_SHIFL	cytoplasm (potential).	cytoplasm
Q821C2	LEU2_SHIFL		cytoplasm
Q82S07	NUSB_NITEU		cytoplasm
Q82S08	RISB_NITEU		cytoplasm
Q82S12	FPG_NITEU		cytoplasm
Q82S78	GIDA_NITEU	cytoplasm (by similarity).	cytoplasm
Q82S79	GIDB_NITEU		cytoplasm
Q82S90	DAPD_NITEU	cytoplasm (by similarity).	cytoplasm
Q82S98	SSB_NITEU		cytoplasm
Q82SA7	Y2439_NITEU		cytoplasm
Q82SA8	OPGH_NITEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q82SA9	OPGG_NITEU	periplasm (by similarity).	periplasm
Q82SD3	FLHD_NITEU	cytoplasm (by similarity).	cytoplasm
Q82SD8	CLPB_NITEU	cytoplasm (probable).	cytoplasm
Q82SG6	Y2372_NITEU		cytoplasm
Q82SI5	ACSA_NITEU		cytoplasm
Q82SI6	RS20_NITEU		cytoplasm
Q82SI9	Y2333_NITEU		inner membrane
Q82SJ7	PDXJ_NITEU	cytoplasm (probable).	cytoplasm
Q82SJ8	NAGZ_NITEU	cytoplasm (by similarity).	extracellular
Q82SL8	BIOH_NITEU	cytoplasm (by similarity).	cytoplasm
Q82SM5	FUMC_NITEU	cytoplasm (by similarity).	cytoplasm
Q82SM6	QUEF_NITEU	cytoplasm (probable).	cytoplasm
Q82SP3	GLGA_NITEU		cytoplasm
Q82SP4	G6PI_NITEU	cytoplasm (by similarity).	cytoplasm
Q82SP6	HSLU_NITEU	cytoplasm (by similarity).	cytoplasm
Q82SP7	HSLV_NITEU	cytoplasm (by similarity).	cytoplasm
Q82SQ2	RPOZ_NITEU		cytoplasm

Q82SQ3	KGUA_NITEU	cytoplasm (by similarity).	cytoplasm
Q82SQ9	Y2248_NITEU		inner membrane
Q82ST3	PYRD_NITEU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q82ST4	ATE_NITEU	cytoplasm (potential).	cytoplasm
Q82ST5	LFTR_NITEU	cytoplasm (by similarity).	cytoplasm
Q82SU1	ARGJ_NITEU	cytoplasm (probable).	cytoplasm
Q82SU4	SECB_NITEU	cytoplasm (by similarity).	cytoplasm
Q82SU6	GPDA_NITEU	cytoplasm (probable).	cytoplasm
Q82SV5	Y2194_NITEU		inner membrane
Q82SW5	KTHY_NITEU		cytoplasm
Q82T00	RS4B_NITEU		cytoplasm
Q82T07	PHK_NITEU		cytoplasm
Q82T49	FLIE_NITEU		flagellar
Q82T56	GATB_NITEU		cytoplasm
Q82T57	GATC_NITEU		cytoplasm
Q82T68	RS12_NITEU		cytoplasm
Q82T69	RS7_NITEU		cytoplasm
Q82T71	RL11_NITEU		cytoplasm
Q82T72	RL1_NITEU		cytoplasm
Q82T73	RL10_NITEU		cytoplasm
Q82T75	RPOB_NITEU		cytoplasm
Q82T76	RPOC_NITEU		cytoplasm
Q82T88	GLGC_NITEU		cytoplasm
Q82TB9	AROB_NITEU	cytoplasm (probable).	cytoplasm
Q82TC0	AROK_NITEU	cytoplasm (probable).	cytoplasm
Q82TC8	DEF2_NITEU		cytoplasm
Q82TD1	SMG_NITEU		cytoplasm
Q82TD5	KCY_NITEU	cytoplasm (by similarity).	cytoplasm
Q82TD8	PYRF_NITEU		cytoplasm
Q82TF3	HPPA_NITEU	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q82TF6	RECA_NITEU	cytoplasm (by similarity).	cytoplasm
Q82TG6	RBL_NITEU		extracellular
Q82TH3	HEM1_NITEU		cytoplasm
Q82TK9	AROC_NITEU		cytoplasm
Q82TL0	HEM6_NITEU	cytoplasm (by similarity).	cytoplasm
Q82TM7	Y1857_NITEU		cytoplasm
Q82TN0	ARLY_NITEU	cytoplasm (probable).	cytoplasm
Q82TN2	MURA_NITEU	cytoplasm (probable).	inner membrane
Q82TN5	Y1849_NITEU		cytoplasm
Q82TP9	MINC_NITEU		inner membrane
Q82TQ2	LOLB_NITEU	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q82TQ3	ISPE_NITEU		cytoplasm
Q82TQ4	KPRS_NITEU	cytoplasm (by similarity).	cytoplasm
Q82TQ5	RL25_NITEU		cytoplasm
Q82TQ6	PTH_NITEU	cytoplasm (by similarity).	cytoplasm
Q82TU0	GPMA2_NITEU		cytoplasm
Q82TU1	TPIS_NITEU		cytoplasm
Q82TV8	HTPG_NITEU	cytoplasm (by similarity).	cytoplasm
Q82TW4	DEF1_NITEU		cytoplasm
Q82TX6	RPIA_NITEU		cytoplasm
Q82TY3	CLPS_NITEU		cytoplasm

Q82TZ6	RS2_NITEU		cytoplasm
Q82TZ8	RRF_NITEU	cytoplasm (by similarity).	cytoplasm
Q82TZ9	UPPS_NITEU		inner membrane
Q82U01	DXR_NITEU		cytoplasm
Q82U05	FABZ_NITEU	cytoplasm (by similarity).	cytoplasm
Q82U06	RNH2_NITEU	cytoplasm (potential).	extracellular
Q82U08	MUTS_NITEU		cytoplasm
Q82U09	CRCB_NITEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q82U36	RL19_NITEU		cytoplasm
Q82U37	RIMM_NITEU	cytoplasm (potential).	cytoplasm
Q82U38	RS16_NITEU		cytoplasm
Q82U41	Y1668_NITEU		inner membrane
Q82U42	RUVX_NITEU	cytoplasm (potential).	cytoplasm
Q82U44	PYRB_NITEU		cytoplasm
Q82U60	FABH_NITEU	cytoplasm (probable).	cytoplasm
Q82U61	PLSX_NITEU		cytoplasm
Q82U62	RL32_NITEU		cytoplasm
Q82U64	Y1642_NITEU	cytoplasm (potential).	inner membrane
Q82U73	MTGA_NITEU	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q82U82	CCA_NITEU		cytoplasm
Q82UC1	APAG_NITEU		inner membrane
Q82UJ5	LIPA_NITEU	cytoplasm (potential).	cytoplasm
Q82UJ6	LIPB_NITEU	cytoplasm (by similarity).	cytoplasm
Q82UJ7	Y1487_NITEU		inner membrane
Q82UK1	RS9_NITEU		cytoplasm
Q82UK2	ARGC_NITEU	cytoplasm (probable).	cytoplasm
Q82UK6	PPNK_NITEU	cytoplasm (by similarity).	cytoplasm
Q82UK7	HRCA_NITEU		cytoplasm
Q82UK8	HEMH_NITEU	cytoplasm (by similarity).	extracellular
Q82UL8	RL28_NITEU		cytoplasm
Q82UL9	RADC_NITEU		cytoplasm
Q82UM1	DUT_NITEU		extracellular
Q82UP3	ARGD_NITEU	cytoplasm (probable).	cytoplasm
Q82UP4	OTC_NITEU	cytoplasm (probable).	cytoplasm
Q82UP5	ASSY_NITEU	cytoplasm (probable).	cytoplasm
Q82UP6	METE_NITEU		cytoplasm
Q82UP9	GLYA_NITEU	cytoplasm (by similarity).	cytoplasm
Q82UQ0	NRDR_NITEU		cytoplasm
Q82UQ1	SYU_NITEU	cytoplasm (by similarity).	cytoplasm
Q82UQ7	THIE_NITEU		cytoplasm
Q82UR9	ISPD_NITEU		cytoplasm
Q82US7	ISPF_NITEU		extracellular
Q82UZ3	ILVC_NITEU		cytoplasm
Q82UZ4	PSD_NITEU		cytoplasm
Q82UZ9	NUDH_NITEU		cytoplasm
Q82V16	GSHB_NITEU		cytoplasm
Q82V19	RL27_NITEU		cytoplasm
Q82V23	HFQ_NITEU		cytoplasm
Q82V28	HISZ_NITEU	cytoplasm (by similarity).	cytoplasm
Q82V29	PURA_NITEU	cytoplasm (by similarity).	cytoplasm
Q82VD1	Y1163_NITEU		inner membrane
Q82VD3	DXS_NITEU		cytoplasm

Q82VD5	EX7S_NITEU	cytoplasm (by similarity).	cytoplasm
Q82VF0	TGT_NITEU		cytoplasm
Q82VF1	QUEA_NITEU	cytoplasm (potential).	cytoplasm
Q82VL9	LOLD_NITEU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q82VM2	LOLA_NITEU	periplasm (by similarity).	periplasm
Q82VN1	RL31_NITEU		cytoplasm
Q82VN4	Y1032_NITEU		inner membrane
Q82VP4	TILS_NITEU	cytoplasm (probable).	cytoplasm
Q82VP5	ACCA_NITEU	cytoplasm (by similarity).	cytoplasm
Q82VR4	PSTB_NITEU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q82VS0	DDL_NITEU	cytoplasm (by similarity).	cytoplasm
Q82VS1	MURB_NITEU	cytoplasm (probable).	inner membrane
Q82VS2	MURC_NITEU	cytoplasm (probable).	cytoplasm
Q82VS3	MURG_NITEU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q82VS5	MURD_NITEU	cytoplasm (by similarity).	cytoplasm
Q82VS6	MRAY_NITEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q82VS8	MURE_NITEU	cytoplasm (probable).	cytoplasm
Q82VT1	MRAW_NITEU		cytoplasm
Q82VT2	MRAZ_NITEU		inner membrane
Q82VV3	RL35_NITEU		cytoplasm
Q82VV4	RL20_NITEU		cytoplasm
Q82VV9	SURE_NITEU	cytoplasm (potential).	periplasm
Q82W15	KSGA_NITEU		cytoplasm
Q82W26	HISX_NITEU		cytoplasm
Q82W27	HIS1_NITEU	cytoplasm (by similarity).	cytoplasm
Q82W32	PUR7_NITEU		cytoplasm
Q82WA9	UVRB_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WB9	MDH_NITEU		cytoplasm
Q82WC5	CCME_NITEU	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q82WC8	CCMA_NITEU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q82WC9	RBFA_NITEU	cytoplasm (potential).	cytoplasm
Q82WD1	Y759_NITEU		inner membrane
Q82WE9	SELU_NITEU		cytoplasm
Q82WF6	Y724_NITEU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q82WG1	Y719_NITEU		inner membrane
Q82WG7	Y711_NITEU		inner membrane
Q82WI1	TRPA_NITEU		cytoplasm
Q82WI2	TRPB_NITEU		cytoplasm
Q82WI6	LEU3_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WI7	LEUD_NITEU		cytoplasm
Q82WI9	LEU2_NITEU		cytoplasm

Q82WL1	SAHH_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WL9	AROQ_NITEU		cytoplasm
Q82WM1	ISPH_NITEU		cytoplasm
Q82WM3	HIS81_NITEU		cytoplasm
Q82WM4	HIS7_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WM5	HIS4_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WM6	HIS3_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WM7	HIS2_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WP9	DAPB_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WQ0	APAH_NITEU		extracellular
Q82WQ3	GCSPB_NITEU		cytoplasm
Q82WQ4	GCSPA_NITEU		cytoplasm
Q82WQ5	GCSH_NITEU		cytoplasm
Q82WQ6	GCST_NITEU		cytoplasm
Q82WR3	Y599_NITEU		inner membrane
Q82WR4	COAE_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WS2	HEM3_NITEU		cytoplasm
Q82WS3	CAPP_NITEU		cytoplasm
Q82WT5	CYSA_NITEU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q82WU3	TYSY_NITEU	cytoplasm (by similarity).	cytoplasm
Q82WX6	GLMM_NITEU		cytoplasm
Q82WZ5	RLUD_NITEU		cytoplasm
Q82X50	DCUP_NITEU	cytoplasm (probable).	cytoplasm
Q82X51	SYV_NITEU	cytoplasm (by similarity).	cytoplasm
Q82X54	AMPA_NITEU	cytoplasm (by similarity).	cytoplasm
Q82X61	Y434_NITEU		inner membrane
Q82X65	SSRP_NITEU	cytoplasm (potential).	cytoplasm
Q82X69	RPOA_NITEU		cytoplasm
Q82X70	RS4A_NITEU		cytoplasm
Q82X71	RS11_NITEU		cytoplasm
Q82X73	RL15_NITEU		cytoplasm
Q82X75	RS5_NITEU		cytoplasm
Q82X77	RS8_NITEU		cytoplasm
Q82X80	RS17_NITEU		cytoplasm
Q82X81	RL29_NITEU		cytoplasm
Q82X82	RL16_NITEU		cytoplasm
Q82X83	RL22_NITEU		cytoplasm
Q82X84	RS19_NITEU		cytoplasm
Q82X85	RL2_NITEU		cytoplasm
Q82X87	RL4_NITEU		cytoplasm
Q82X88	RL3_NITEU		cytoplasm
Q82X89	RS10_NITEU		cytoplasm
Q82X93	Y395_NITEU		inner membrane
Q82X99	RNPA_NITEU		extracellular
Q82XA0	Y388_NITEU		inner membrane
Q82XA1	TRME_NITEU		cytoplasm
Q82XC7	Y356_NITEU	cytoplasm (potential).	inner membrane
Q82XD1	RLMB_NITEU		cytoplasm
Q82XD4	SPEE_NITEU		cytoplasm
Q82XE0	HIS82_NITEU		cytoplasm
Q82XE8	PGK_NITEU	cytoplasm (potential).	cytoplasm
Q82XF2	FETP_NITEU		cytoplasm
Q82XG5	FLGI_NITEU	periplasm (by similarity).	periplasm

Q82XG6	FLGH_NITEU	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q82XI7	THIG_NITEU	cytoplasm (by similarity).	cytoplasm
Q82XJ3	HAM1_NITEU		extracellular
Q82XJ4	RNPH_NITEU		cytoplasm
Q82XN3	Y224_NITEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q82XN9	TOLB_NITEU	periplasm (potential).	periplasm
Q82XP4	RUVB_NITEU		cytoplasm
Q82XP5	RUVC_NITEU		cytoplasm
Q82XP6	Y210_NITEU		inner membrane
Q82XP7	GLMU_NITEU	cytoplasm (by similarity).	cytoplasm
Q82XP9	ATPG_NITEU		inner membrane
Q82XQ0	ATPA_NITEU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q82XQ6	RS6_NITEU		cytoplasm
Q82XQ7	PRIB_NITEU		cytoplasm
Q82XQ8	RS18_NITEU		cytoplasm
Q82XQ9	RL9_NITEU		cytoplasm
Q82XS4	GPMA1_NITEU		cytoplasm
Q82XS9	RS15_NITEU		cytoplasm
Q82XU6	ENGA_NITEU		cytoplasm
Q82XV0	ISPG_NITEU		cytoplasm
Q82XV5	NDK_NITEU	cytoplasm (by similarity).	cytoplasm
Q82XW4	Y131_NITEU		cytoplasm
Q82XY7	ILVD_NITEU		cytoplasm
Q82XZ6	GUAA_NITEU		cytoplasm
Q82Y02	PUR5_NITEU	cytoplasm (by similarity).	cytoplasm
Q82Y12	ENGC_NITEU		cytoplasm
Q82Y13	PHS_NITEU		cytoplasm
Q82Y15	ORN_NITEU	cytoplasm (potential).	cytoplasm
Q82Y17	PANC_NITEU		cytoplasm
Q82Y18	PANB_NITEU		cytoplasm
Q82Y30	HPRK_NITEU		cytoplasm
Q82Y49	UPPP_NITEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q82Y50	THIC_NITEU		cytoplasm
Q82Y56	CLPX_NITEU		cytoplasm
Q82Y57	CLPP_NITEU	cytoplasm (by similarity).	cytoplasm
Q82Y58	TIG_NITEU		cytoplasm
Q82Y60	CH60_NITEU	cytoplasm (by similarity).	cytoplasm
Q82Y61	CH10_NITEU	cytoplasm (by similarity).	cytoplasm
Q82Y74	TRPD_NITEU		cytoplasm
Q82Y84	DNAA_NITEU	cytoplasm (by similarity).	cytoplasm
Q83A16	PYRB_COXBU		cytoplasm
Q83A17	RUVX_COXBU	cytoplasm (potential).	cytoplasm
Q83A18	Y2093_COXBU		inner membrane
Q83A19	PPCK_COXBU	cytoplasm (by similarity).	cytoplasm
Q83A37	HEM3_COXBU		cytoplasm
Q83A42	KITH_COXBU	cytoplasm (by similarity).	cytoplasm
Q83A62	METE_COXBU		cytoplasm
Q83A77	SAHH_COXBU	cytoplasm (by similarity).	cytoplasm
Q83A90	UBIE_COXBU		cytoplasm

Q83A94	HSLU_COXBU	cytoplasm (by similarity).	cytoplasm
Q83A95	HSLV_COXBU	cytoplasm (by similarity).	cytoplasm
Q83AA6	SMG_COXBU		cytoplasm
Q83AA8	FMT_COXBU		cytoplasm
Q83AB7	APAH_COXBU		extracellular
Q83AC2	KSGA_COXBU		cytoplasm
Q83AF3	GLMU_COXBU	cytoplasm (by similarity).	cytoplasm
Q83AF4	ATPE_COXBU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83AF6	ATPG_COXBU		inner membrane
Q83AF7	ATPA_COXBU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83AJ2	AROB_COXBU	cytoplasm (probable).	cytoplasm
Q83AJ3	AROK_COXBU	cytoplasm (probable).	cytoplasm
Q83AK6	DEF2_COXBU		cytoplasm
Q83AK9	Y1876_COXBU		inner membrane
Q83AL0	GSHB_COXBU		cytoplasm
Q83AP0	PTH_COXBU	cytoplasm (by similarity).	cytoplasm
Q83AP1	RL25_COXBU		cytoplasm
Q83AQ1	KPRS_COXBU	cytoplasm (by similarity).	cytoplasm
Q83AQ2	LOLB_COXBU	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q83AQ4	PSD_COXBU		cytoplasm
Q83AR8	PDXB_COXBU		cytoplasm
Q83AU6	PGK_COXBU	cytoplasm (potential).	cytoplasm
Q83AX9	RS9_COXBU		cytoplasm
Q83AY5	Y1742_COXBU		inner membrane
Q83AZ0	PUR5_COXBU	cytoplasm (by similarity).	cytoplasm
Q83AZ6	HEM6_COXBU	cytoplasm (by similarity).	cytoplasm
Q83B06	GCST_COXBU		cytoplasm
Q83B07	GCSH_COXBU		cytoplasm
Q83B08	GCSPA_COXBU		cytoplasm
Q83B09	GCSPB_COXBU		cytoplasm
Q83B31	DCD_COXBU		extracellular
Q83B43	KDSA_COXBU	cytoplasm (by similarity).	cytoplasm
Q83B44	ENO_COXBU	cytoplasm (by similarity).	extracellular
Q83B60	HLDE_COXBU		cytoplasm
Q83BB0	RIMK_COXBU		cytoplasm
Q83BB9	RS21_COXBU		cytoplasm
Q83BD3	RBN_COXBU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83BE0	RUVB_COXBU		cytoplasm
Q83BE2	RUVA_COXBU		cytoplasm
Q83BE3	RUVC_COXBU		cytoplasm
Q83BE4	Y1566_COXBU		inner membrane
Q83BF8	NUDH_COXBU		cytoplasm
Q83BG0	LGT_COXBU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83BG2	TYSY_COXBU	cytoplasm (by similarity).	cytoplasm
Q83BH2	GPMI_COXBU		cytoplasm
Q83BI9	SECB_COXBU	cytoplasm (by similarity).	cytoplasm
Q83BJ0	GPDA_COXBU	cytoplasm (probable).	cytoplasm

Q83BJ8	ACCA_COXBU	cytoplasm (by similarity).	cytoplasm
Q83BJ9	TILS_COXBU	cytoplasm (probable).	cytoplasm
Q83BK3	LEPA_COXBU		cytoplasm
Q83BL1	PDXJ_COXBU	cytoplasm (probable).	cytoplasm
Q83BL5	LPXH_COXBU	cytoplasm (by similarity).	cytoplasm
Q83BM8	GATB_COXBU		cytoplasm
Q83BM9	GATA_COXBU		cytoplasm
Q83BQ3	TPIS_COXBU		cytoplasm
Q83BR9	Y1434_COXBU		inner membrane
Q83BS2	RBFA_COXBU	cytoplasm (potential).	cytoplasm
Q83BT3	GLYA_COXBU	cytoplasm (by similarity).	cytoplasm
Q83BT4	NRDR_COXBU		cytoplasm
Q83BT5	NUSB_COXBU		cytoplasm
Q83BV4	RRF_COXBU	cytoplasm (by similarity).	cytoplasm
Q83BV5	UPPS_COXBU		inner membrane
Q83BY4	RRMJ_COXBU	cytoplasm (by similarity).	cytoplasm
Q83BY7	GLMM_COXBU		cytoplasm
Q83BZ6	GUAA_COXBU		cytoplasm
Q83BZ8	RNH2_COXBU	cytoplasm (potential).	extracellular
Q83BZ9	DDL_COXBU	cytoplasm (by similarity).	cytoplasm
Q83C12	RL35_COXBU		cytoplasm
Q83C13	RL20_COXBU		cytoplasm
Q83C29	SSRP_COXBU	cytoplasm (potential).	cytoplasm
Q83C31	Y1303_COXBU		inner membrane
Q83C38	PPNK_COXBU	cytoplasm (by similarity).	cytoplasm
Q83C41	GRPE_COXBU	cytoplasm (probable).	cytoplasm
Q83C63	LIPA_COXBU	cytoplasm (potential).	cytoplasm
Q83C64	LIPB_COXBU	cytoplasm (by similarity).	cytoplasm
Q83C71	NDK_COXBU	cytoplasm (by similarity).	cytoplasm
Q83C83	ENGA_COXBU		cytoplasm
Q83C87	MDH_COXBU		cytoplasm
Q83C89	Y1239_COXBU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83C93	ORN_COXBU	cytoplasm (potential).	cytoplasm
Q83CA8	PUR7_COXBU		cytoplasm
Q83CC6	QUEA_COXBU	cytoplasm (potential).	cytoplasm
Q83CD5	UVRC_COXBU	cytoplasm (by similarity).	cytoplasm
Q83CD9	THII_COXBU	cytoplasm (potential).	cytoplasm
Q83CG2	TRPA_COXBU		cytoplasm
Q83CJ1	CBPM_COXBU		cytoplasm
Q83CJ2	CBPA_COXBU	nucleoid (by similarity).	inner membrane
Q83CL8	FUMC_COXBU	cytoplasm (by similarity).	cytoplasm
Q83CQ2	MUTS_COXBU		cytoplasm
Q83CQ4	RECA_COXBU	cytoplasm (by similarity).	cytoplasm
Q83CQ8	CSRA2_COXBU		cytoplasm
Q83CV2	LOLD_COXBU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83CV4	PURA_COXBU	cytoplasm (by similarity).	cytoplasm
Q83CV9	DEF1_COXBU		cytoplasm
Q83CW4	UNG_COXBU	cytoplasm (by similarity).	extracellular
Q83CW6	RLMB_COXBU		cytoplasm
Q83CY0	PYRD_COXBU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane

Q83D06	FETP_COXBU		cytoplasm
Q83D14	GLPK_COXBU		cytoplasm
Q83D18	PDXH_COXBU		cytoplasm
Q83D39	RL31_COXBU		cytoplasm
Q83D55	BDBC_COXBU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q83D67	AROC_COXBU		cytoplasm
Q83D70	Y870_COXBU		cytoplasm
Q83D73	RL9_COXBU		cytoplasm
Q83D75	RS18_COXBU		cytoplasm
Q83D76	RS6_COXBU		cytoplasm
Q83D83	LPXK_COXBU		cytoplasm
Q83D84	MSBA_COXBU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83D86	MGSA_COXBU		cytoplasm
Q83D88	RS15_COXBU		cytoplasm
Q83D91	G6PI_COXBU	cytoplasm (by similarity).	cytoplasm
Q83DD0	SYV_COXBU	cytoplasm (by similarity).	cytoplasm
Q83DE3	GCH1_COXBU		cytoplasm
Q83DI0	MURA_COXBU	cytoplasm (probable).	inner membrane
Q83DJ1	CLPX_COXBU		cytoplasm
Q83DJ2	CLPP_COXBU	cytoplasm (by similarity).	cytoplasm
Q83DJ3	TIG_COXBU		cytoplasm
Q83DN1	DAPD_COXBU	cytoplasm (by similarity).	cytoplasm
Q83DP0	RECR_COXBU		cytoplasm
Q83DP1	Y656_COXBU		inner membrane
Q83DP8	RISB_COXBU		cytoplasm
Q83DS5	LPXB_COXBU		cytoplasm
Q83DT0	LPXD_COXBU		cytoplasm
Q83DV5	GLPE_COXBU	cytoplasm (by similarity).	cytoplasm
Q83DX5	Y567_COXBU		inner membrane
Q83E06	PYRF_COXBU		cytoplasm
Q83E10	KCY_COXBU	cytoplasm (by similarity).	cytoplasm
Q83E12	SERC_COXBU	cytoplasm (by similarity).	cytoplasm
Q83E18	UVRB_COXBU	cytoplasm (by similarity).	cytoplasm
Q83E35	KTHY_COXBU		cytoplasm
Q83E40	PLSX_COXBU		cytoplasm
Q83E41	RL32_COXBU		cytoplasm
Q83E63	EX7S_COXBU	cytoplasm (by similarity).	cytoplasm
Q83E75	KAD_COXBU	cytoplasm (by similarity).	cytoplasm
Q83E83	RS16_COXBU		cytoplasm
Q83E84	RIMM_COXBU	cytoplasm (potential).	cytoplasm
Q83E85	RL19_COXBU		cytoplasm
Q83EC9	SYI_COXBU	cytoplasm (by similarity).	cytoplasm
Q83ED6	RS20_COXBU		cytoplasm
Q83ED9	RL27_COXBU		cytoplasm
Q83EI7	THIG_COXBU	cytoplasm (by similarity).	cytoplasm
Q83EJ0	THIC_COXBU		cytoplasm
Q83EL0	HTPG_COXBU	cytoplasm (by similarity).	cytoplasm
Q83EL6	RPOZ_COXBU		cytoplasm
Q83EL7	KGUA_COXBU	cytoplasm (by similarity).	cytoplasm
Q83EL9	RNPH_COXBU		cytoplasm
Q83EM4	RL28_COXBU		cytoplasm
Q83EM7	COAD_COXBU	cytoplasm (by similarity).	cytoplasm

Q83EP0	DCUP_COXBU	cytoplasm (probable).	cytoplasm
Q83EP4	SSB_COXBU		cytoplasm
Q83EP5	Y270_COXBU		cytoplasm
Q83EQ2	RPOA_COXBU		cytoplasm
Q83EQ3	RS4_COXBU		cytoplasm
Q83EQ4	RS11_COXBU		cytoplasm
Q83EQ7	RL15_COXBU		cytoplasm
Q83EQ9	RS5_COXBU		cytoplasm
Q83ER0	RL18_COXBU		cytoplasm
Q83ER2	RS8_COXBU		cytoplasm
Q83ER4	RL5_COXBU		cytoplasm
Q83ER5	RL24_COXBU		cytoplasm
Q83ER7	RS17_COXBU		cytoplasm
Q83ER8	RL29_COXBU		cytoplasm
Q83ER9	RL16_COXBU		cytoplasm
Q83ES0	RS19_COXBU		cytoplasm
Q83ES1	RL2_COXBU		cytoplasm
Q83ES3	RL4_COXBU		cytoplasm
Q83ES4	RL3_COXBU		cytoplasm
Q83ES5	RS10_COXBU		cytoplasm
Q83ES8	RS7_COXBU		cytoplasm
Q83ES9	RS12_COXBU		cytoplasm
Q83ET0	RPOC_COXBU		cytoplasm
Q83ET2	RL10_COXBU		cytoplasm
Q83ET3	RL1_COXBU		cytoplasm
Q83ET4	RL11_COXBU		cytoplasm
Q83EV9	COAA_COXBU	cytoplasm (probable).	cytoplasm
Q83EX7	SYX_COXBU	cytoplasm (by similarity).	cytoplasm
Q83F01	COAE_COXBU	cytoplasm (by similarity).	cytoplasm
Q83F02	QUEF_COXBU	cytoplasm (probable).	cytoplasm
Q83F03	Y150_COXBU		inner membrane
Q83F11	LPXC_COXBU		cytoplasm
Q83F16	MURB_COXBU	cytoplasm (probable).	inner membrane
Q83F17	MURC_COXBU	cytoplasm (probable).	cytoplasm
Q83F20	MURD_COXBU	cytoplasm (by similarity).	cytoplasm
Q83F26	MRAY_COXBU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83F28	MURE_COXBU	cytoplasm (probable).	cytoplasm
Q83F35	MRAW_COXBU		cytoplasm
Q83F36	MRAZ_COXBU		inner membrane
Q83F37	Y114_COXBU		inner membrane
Q83F39	TDH_COXBU	cytoplasm (by similarity).	cytoplasm
Q83F55	CLPB_COXBU	cytoplasm (probable).	cytoplasm
Q83F59	TOLB_COXBU	periplasm (potential).	periplasm
Q83FA3	HAM1_COXBU		extracellular
Q83FA4	HEMH_COXBU	cytoplasm (by similarity).	extracellular
Q83FB4	RPIA_COXBU		cytoplasm
Q83FB6	CSRA1_COXBU		cytoplasm
Q83FD6	RECF_COXBU	cytoplasm (by similarity).	cytoplasm
Q83FD8	DNAA_COXBU	cytoplasm (by similarity).	cytoplasm
Q83IH7	OPGB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83II8	ULAE_SHIFL		cytoplasm

Q83II9	ULAA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83IJ0	ULAG_SHIFL	cytoplasm (potential).	extracellular
Q83IJ9	ORN_SHIFL	cytoplasm (potential).	cytoplasm
Q83IK0	ENGC_SHIFL		cytoplasm
Q83IL8	PYRI_SHIFL		cytoplasm
Q83IM2	OTC_SHIFL	cytoplasm (probable).	cytoplasm
Q83IM5	SYV_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83IQ8	MDTO_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83IR6	RLUF_SHIFL		cytoplasm
Q83IS3	NUDC_SHIFL		cytoplasm
Q83IS4	BTUB_SHIFL	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q83IS7	CAPP_SHIFL		cytoplasm
Q83IU3	RHAB_SHIFL		cytoplasm
Q83IU4	RHAA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83IV7	RDOA_SHIFL	cytoplasm (probable).	cytoplasm
Q83IW0	METE_SHIFL		cytoplasm
Q83IW8	YIGA_SHIFL		inner membrane
Q83IX2	CYAY_SHIFL		inner membrane
Q83IX5	WECG_SHIFL		cytoplasm
Q83IX6	WZYE_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q83IY2	VIAA_SHIFL		nucleoid
Q83IY3	GLMU_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83IY4	GLMS_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83J28	SELA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83J60	DCTR_SHIFL		cytoplasm
Q83J77	NIKE_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83J78	NIKD_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83J89	GLGX_SHIFL		extracellular
Q83J92	GNTX_SHIFL		cytoplasm
Q83J96	PPCK_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83JB1	FRLD_SHIFL		cytoplasm
Q83JB2	FRLC_SHIFL		cytoplasm
Q83JC0	TUSD_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83JC1	TUSC_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83JC2	TUSB_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83JD2	YRDB_SHIFL		cytoplasm
Q83JE1	YHDE_SHIFL	cytoplasm (potential).	inner membrane
Q83JE2	AAEB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83JF4	KDSD_SHIFL		cytoplasm
Q83JG5	YHBQ_SHIFL		cytoplasm
Q83JL2	YGIQ_SHIFL		cytoplasm
Q83JS0	HAM1_SHIFL		extracellular

Q83JS1	YGGU_SHIFL		inner membrane
Q83JS4	RUVX_SHIFL	cytoplasm (potential).	cytoplasm
Q83JS6	SPRT_SHIFL	cytoplasm (potential).	extracellular
Q83JU1	GCST_SHIFL		cytoplasm
Q83JU2	YGFY_SHIFL		cytoplasm
Q83JV7	AAS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83JW6	MTFA_SHIFL		cytoplasm
Q83JW9	QUEF_SHIFL	cytoplasm (probable).	cytoplasm
Q83JX8	CYSN_SHIFL		cytoplasm
Q83JY2	SURE_SHIFL	cytoplasm (potential).	periplasm
Q83JY3	PIMT_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83JZ0	YGAD_SHIFL		inner membrane
Q83JZ4	LUXS_SHIFL		extracellular
Q83K04	CSID_SHIFL		cytoplasm
Q83K21	GRCA_SHIFL		cytoplasm
Q83K24	ACPS_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83K43	ISPG_SHIFL		cytoplasm
Q83K84	SYE_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83K86	GLK_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83K95	FADI_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83KB1	YFBV_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83KB5	YFBJ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83KB6	ARNT_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83KD5	CCMA_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83KD7	YEJL_SHIFL		inner membrane
Q83KD8	NDPA_SHIFL	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q83KE6	EFPL_SHIFL		cytoplasm
Q83KF1	RIHB_SHIFL		extracellular
Q83KF5	MDTQ_SHIFL	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
Q83KH2	THIM_SHIFL		cytoplasm
Q83KI3	YEGP_SHIFL		extracellular
Q83KI4	MDTC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83KI5	MDTA_SHIFL	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q83KI8	YEGH_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q83KJ5	HIS5_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83KJ6	HIS8_SHIFL		cytoplasm
Q83KJ7	HIS1_SHIFL	cytoplasm (by similarity).	cytoplasm

Q83KJ8	YEFM_SHIFL		cytoplasm
Q83KK3	YEEN_SHIFL		inner membrane
Q83KM2	YEDZ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83KN1	DSRB_SHIFL		inner membrane
Q83KR4	RUVA_SHIFL		cytoplasm
Q83KR5	RUVB_SHIFL		cytoplasm
Q83KS1	YEBF_SHIFL	secreted protein (by similarity).	inner membrane
Q83KS5	ZNTB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83KW2	SUFB_SHIFL		inner membrane
Q83KY0	SYI_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83KZ1	MANA_SHIFL	cytoplasm (probable).	cytoplasm
Q83L10	YNFA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83L33	YDIU_SHIFL		inner membrane
Q83L66	MSRB_SHIFL		cytoplasm
Q83L75	YOA_H_SHIFL		inner membrane
Q83L78	YOB_D_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83LB2	YCJF_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83LC9	YCIC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83LD1	YCIU_SHIFL		inner membrane
Q83LD8	ISPE_SHIFL		cytoplasm
Q83LE1	PTH_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83LF6	RLUE_SHIFL		cytoplasm
Q83LF8	YCFC_SHIFL		inner membrane
Q83LH9	THIK_SHIFL		cytoplasm
Q83LI5	YCEQ_SHIFL		cytoplasm
Q83LI8	MDTH_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83LI9	YCEI_SHIFL	periplasm (potential).	periplasm
Q83LK1	YCDX_SHIFL		cytoplasm
Q83LP0	MSBA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83LP1	KCY_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83LP3	SERC_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83LP7	6PGL_SHIFL		cytoplasm
Q83LS2	YBJQ_SHIFL		inner membrane
Q83LU7	DING_SHIFL		cytoplasm
Q83LX0	CRCB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83LY1	RIHA_SHIFL		extracellular
Q83LZ7	END8_SHIFL		cytoplasm

Q83M01	GAL1_SHIFL	cytoplasm (potential).	cytoplasm
Q83M07	USPG_SHIFL		cytoplasm
Q83M14	CAAL_SHIFL		cytoplasm
Q83M28	LPXH_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83M36	YBAP_SHIFL		inner membrane
Q83M66	AROL_SHIFL	cytoplasm (probable).	cytoplasm
Q83M67	YAII_SHIFL		inner membrane
Q83M81	YAFA_SHIFL		cytoplasm
Q83MA0	TAUB_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83MD3	DXR_SHIFL		cytoplasm
Q83MD4	GLND_SHIFL		cytoplasm
Q83MD6	DGTP_SHIFL		cytoplasm
Q83MD7	BTUF_SHIFL	periplasm (by similarity).	periplasm
Q83ME2	GLUQ_SHIFL		cytoplasm
Q83ME5	PANB_SHIFL		cytoplasm
Q83MF0	SPEE_SHIFL		cytoplasm
Q83MF4	YACF_SHIFL		inner membrane
Q83MF7	DDLB_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83MF8	MURC_SHIFL	cytoplasm (probable).	cytoplasm
Q83MF9	MURD_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83MG0	MURE_SHIFL	cytoplasm (probable).	cytoplasm
Q83MG1	MRAZ_SHIFL		inner membrane
Q83MG5	ARAB_SHIFL		cytoplasm
Q83MG8	KSGA_SHIFL		cytoplasm
Q83MG9	CAIC_SHIFL		cytoplasm
Q83MH0	RIHC_SHIFL		extracellular
Q83MH2	SYI_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83MH5	DNAK_SHIFL		cytoplasm
Q83MH9	KHSE_SHIFL	cytoplasm (potential).	cytoplasm
Q83MI1	STHA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83MJ0	NFI_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83MJ2	RBN_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83MJ8	TRUB_SHIFL		cytoplasm
Q83MJ9	IDI_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83MK8	SDHD_SHIFL		cytoplasm
Q83ML6	MAO1_SHIFL		cytoplasm
Q83ML8	FUMC_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83MM7	TOLB_SHIFL	periplasm (by similarity).	inner membrane
Q83MN1	RNH2_SHIFL	cytoplasm (potential).	extracellular
Q83MN4	MURG_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83P00	DEOD_SHIFL		cytoplasm
Q83P01	TYPH_SHIFL		cytoplasm
Q83P02	DEOC_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83P26	ULAF_SHIFL		cytoplasm
Q83P27	ULAD_SHIFL		cytoplasm
Q83P28	ULAC_SHIFL	cytoplasm (probable).	cytoplasm
Q83P29	ULAB_SHIFL	cytoplasm (potential).	cytoplasm
Q83P30	ULAR_SHIFL	cytoplasm (potential).	cytoplasm
Q83P33	PURA_SHIFL	cytoplasm (by similarity).	cytoplasm

Q83P38	FRDC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83P43	CUTA_SHIFL	cytoplasm (by similarity).	periplasm
Q83P53	YJGA_SHIFL		cytoplasm
Q83P64	AMPA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83P86	MDTN_SHIFL	bacterial cell inner membrane; single-pass type ii membrane protein (potential).	inner membrane
Q83P87	MDTP_SHIFL	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
Q83P94	ACTP_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83P97	PHNC_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83PB5	PEPE_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PB7	DCUP_SHIFL	cytoplasm (probable).	cytoplasm
Q83PB8	THIC_SHIFL		cytoplasm
Q83PB9	THIE_SHIFL		cytoplasm
Q83PC0	THIG_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PC2	RL1_SHIFL		cytoplasm
Q83PC3	RL11_SHIFL		cytoplasm
Q83PC5	COAA_SHIFL	cytoplasm (probable).	cytoplasm
Q83PC7	MURB_SHIFL	cytoplasm (probable).	inner membrane
Q83PD4	RL31_SHIFL		cytoplasm
Q83PD6	FIEF_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83PD8	RHAT_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83PD9	RHAR_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PE0	RHAS_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PE1	RHAD_SHIFL	cytoplasm (by similarity).	extracellular
Q83PE5	FDHD_SHIFL	cytoplasm (potential).	cytoplasm
Q83PE8	FDHE_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PF0	YIHT_SHIFL		cytoplasm
Q83PG1	FADB_SHIFL		cytoplasm
Q83PG2	FADA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PH4	HEM3_SHIFL		cytoplasm
Q83PH8	WECF_SHIFL	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q83PI3	GPPA_SHIFL		extracellular
Q83PI6	ILVD_SHIFL		cytoplasm
Q83PJ2	KUP_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83PJ3	RAVA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PJ4	ASNA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PJ6	GIDA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PJ7	GIDB_SHIFL		cytoplasm
Q83PL3	TRME_SHIFL		cytoplasm

Q83PL6	MDTL_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83PM8	RPOZ_SHIFL		cytoplasm
Q83PN2	RNPH_SHIFL		cytoplasm
Q83PN3	DUT_SHIFL		extracellular
Q83PP2	HLDD_SHIFL		cytoplasm
Q83PP7	LLDD_SHIFL		cytoplasm
Q83PQ0	MTLD_SHIFL		cytoplasm
Q83PQ6	DLGD_SHIFL	cytoplasm (potential).	cytoplasm
Q83PR0	GADX_SHIFL		cytoplasm
Q83PR1	DCEA_SHIFL		cytoplasm
Q83PS8	TREF_SHIFL	cytoplasm (by similarity).	extracellular
Q83PV3	GLGB_SHIFL		extracellular
Q83PV4	GLGA_SHIFL		cytoplasm
Q83PW0	BIOH_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83PW3	FEOB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83PX0	AROB_SHIFL	cytoplasm (probable).	cytoplasm
Q83PX1	GPH_SHIFL		cytoplasm
Q83PX4	FRLR_SHIFL		cytoplasm
Q83PX9	KEFG_SHIFL		cytoplasm
Q83PY0	KEFB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83PY1	SLYX_SHIFL		inner membrane
Q83PY4	RL4_SHIFL		cytoplasm
Q83PY5	RL22_SHIFL		cytoplasm
Q83PY6	RL16_SHIFL		cytoplasm
Q83PY7	RL29_SHIFL		cytoplasm
Q83PY8	RL24_SHIFL		cytoplasm
Q83PY9	RL5_SHIFL		cytoplasm
Q83PZ0	FMT_SHIFL		cytoplasm
Q83PZ1	DEF_SHIFL		cytoplasm
Q83PZ5	DUSB_SHIFL		cytoplasm
Q83Q03	AAEA_SHIFL	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q83Q04	MDH_SHIFL		cytoplasm
Q83Q07	RS9_SHIFL		cytoplasm
Q83Q08	SSPB_SHIFL		cytoplasm
Q83Q15	GLMM_SHIFL		cytoplasm
Q83Q35	ALX_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83Q83	MLTC_SHIFL	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q83Q91	GSHB_SHIFL		cytoplasm
Q83Q93	SPEA_SHIFL	periplasm (by similarity).	cytoplasm
Q83QA2	GCSP_SHIFL		cytoplasm
Q83QB2	KDUI_SHIFL		cytoplasm
Q83QC1	MUTH_SHIFL	cytoplasm (by similarity).	extracellular
Q83QC3	PTRA_SHIFL	periplasm (by similarity).	extracellular

Q83QD0	SDAC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83QD5	RUMA_SHIFL		cytoplasm
Q83QD9	CYSJ_SHIFL		cytoplasm
Q83QE0	CYSI_SHIFL		cytoplasm
Q83QE3	TRUD_SHIFL		cytoplasm
Q83QE9	MUTS_SHIFL		cytoplasm
Q83QI6	RECO_SHIFL		cytoplasm
Q83QK3	HSCB_SHIFL		inner membrane
Q83QK4	HSCA_SHIFL		cytoplasm
Q83QK5	PEPB_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83QL1	GUAA_SHIFL		cytoplasm
Q83QL4	PUR5_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83QM8	TALA_SHIFL	cytoplasm (probable).	cytoplasm
Q83QN1	HEM6_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83QN4	YFEU_SHIFL		cytoplasm
Q83QN8	CYSZ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83QN9	ZIPA_SHIFL	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q83QP2	PTFC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83QP3	PTFAX_SHIFL	cytoplasm (probable).	cytoplasm
Q83QQ0	FADJ_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83QQ8	YFCK_SHIFL		cytoplasm
Q83QR1	PDXB_SHIFL		cytoplasm
Q83QR3	TRUA_SHIFL		cytoplasm
Q83QS1	YFBU_SHIFL		inner membrane
Q83QS4	YFBR_SHIFL		cytoplasm
Q83QT8	ARNA_SHIFL		cytoplasm
Q83QT9	ARNB_SHIFL		cytoplasm
Q83QV0	NAPA_SHIFL	periplasm (by similarity).	periplasm
Q83QV3	CCMB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83QW6	END4_SHIFL		extracellular
Q83QZ5	DCD_SHIFL		extracellular
Q83R03	HIS6_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83R04	HIS4_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83R05	HIS7_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83R13	COBS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83R14	COBT_SHIFL		cytoplasm
Q83R35	FLIE_SHIFL		flagellar
Q83R47	PGSA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83R81	ACPD_SHIFL		extracellular
Q83R85	OPGD_SHIFL	periplasm (by similarity).	periplasm
Q83RC0	ADD_SHIFL		extracellular

Q83RD1	MDTI_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83RD2	ASR_SHIFL		inner membrane
Q83RE1	UXAB_SHIFL		cytoplasm
Q83RE2	GLSA2_SHIFL		extracellular
Q83RE8	YDFG_SHIFL		cytoplasm
Q83RF3	YDEP_SHIFL		periplasm
Q83RF4	YDEO_SHIFL		cytoplasm
Q83RG1	RL20_SHIFL		cytoplasm
Q83RG5	NADE_SHIFL		cytoplasm
Q83RI9	CSPC_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83RJ2	PROQ_SHIFL	cytoplasm (potential).	inner membrane
Q83RK8	YCJQ_SHIFL		cytoplasm
Q83RM1	PYRF_SHIFL		cytoplasm
Q83RM8	CLS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83RN5	NARH_SHIFL	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q83RP1	HEM1_SHIFL		cytoplasm
Q83RP6	TREA_SHIFL	periplasm (by similarity).	periplasm
Q83RQ1	CVRA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q83RR0	PHOP_SHIFL	cytoplasm (probable).	cytoplasm
Q83RR1	PHOQ_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83RR7	POTC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83RR8	NPD_SHIFL	cytoplasm (probable).	cytoplasm
Q83RS0	LOLD_SHIFL	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q83RS8	FABH_SHIFL	cytoplasm (probable).	cytoplasm
Q83RS9	PLSX_SHIFL		cytoplasm
Q83RT9	MTOX_SHIFL		cytoplasm
Q83RU3	OPGG_SHIFL	periplasm (by similarity).	periplasm
Q83RW9	MBHS_SHIFL	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	periplasm
Q83RY7	LPXK_SHIFL		cytoplasm
Q83RZ5	YCAD_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83RZ7	DMSB_SHIFL		periplasm
Q83S05	HCP_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83S14	RUMB_SHIFL		cytoplasm
Q83S18	RIMK_SHIFL		cytoplasm
Q83S20	YBJL_SHIFL	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q83S36	YBHG_SHIFL	periplasm (potential).	inner membrane

Q83S42	UVRB_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83S79	FUR_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83S80	FLAV_SHIFL		cytoplasm
Q83S83	ATKA_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q83S86	YBGL_SHIFL		inner membrane
Q83SA2	ZITB_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83SB6	ENTS_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83SD5	SELU_SHIFL		cytoplasm
Q83SE1	GLSA1_SHIFL		extracellular
Q83SE5	HEMH_SHIFL	cytoplasm (by similarity).	extracellular
Q83SE6	HTPG_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83SG1	THII_SHIFL	cytoplasm (potential).	cytoplasm
Q83SG2	DXS_SHIFL		cytoplasm
Q83SG7	ACPH_SHIFL		cytoplasm
Q83SH9	PROA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83SL3	LPXB_SHIFL		cytoplasm
Q83SL4	RS2_SHIFL		cytoplasm
Q83SL9	SFSA_SHIFL		cytoplasm
Q83SM0	PANC_SHIFL		cytoplasm
Q83SM9	GUAC_SHIFL		cytoplasm
Q83SN7	MRAW_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83SP0	LEU1_SHIFL		cytoplasm
Q83SP1	LEU3_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83SQ0	OSTA_SHIFL	bacterial cell outer membrane (by similarity).	outer membrane
Q83SQ2	APAH_SHIFL		extracellular
Q83SQ3	KEFC_SHIFL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83SQ4	KEFF_SHIFL		cytoplasm
Q83SQ6	FIXX_SHIFL		periplasm
Q83SQ7	FIXC_SHIFL		periplasm
Q83SQ8	CAIE_SHIFL		cytoplasm
Q83SQ9	DAPB_SHIFL	cytoplasm (by similarity).	cytoplasm
Q83SR1	RS20_SHIFL		cytoplasm
Q83ST5	FEOB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q83SW5	YGIQ_SALTI		cytoplasm
Q83U59	SACB3_NEIMA		cytoplasm
Q83WI8	CH601_VIBHA	cytoplasm (by similarity).	cytoplasm
Q83WI9	CH101_VIBHA	cytoplasm (by similarity).	cytoplasm
Q83WJ5	TRUA_BURCE		cytoplasm
Q83WL3	RPP_XYLFT		nucleoid
Q83WS0	FTHS_METEX		cytoplasm
Q83WU4	YPNA_THIRO		cytoplasm
Q83Z42	OPGH_XANCV	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q845U5	HIS2_BURCE	cytoplasm (by similarity).	cytoplasm

Q845U6	HIS3_BURCE	cytoplasm (by similarity).	cytoplasm
Q845U7	HIS6_BURCE	cytoplasm (by similarity).	cytoplasm
Q845U8	HIS4_BURCE	cytoplasm (by similarity).	cytoplasm
Q845U9	HIS5_BURCE	cytoplasm (by similarity).	cytoplasm
Q845V1	HIS7_BURCE	cytoplasm (by similarity).	cytoplasm
Q845V2	HIS8_BURCE		cytoplasm
Q845V3	HISX_BURCE		cytoplasm
Q845V4	HIS1_BURCE	cytoplasm (by similarity).	cytoplasm
Q845W3	LEU3_BURCE	cytoplasm (by similarity).	cytoplasm
Q845X9	TRPD_BURCE		cytoplasm
Q847G1	SSBD_PSEPU		cytoplasm
Q848I3	DCTA_PSECL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q848I4	G6PI_PSEFL	cytoplasm (by similarity).	cytoplasm
Q848I5	PAND_PSEFL		cytoplasm
Q848R7	LCBA2_AERHY		cytoplasm
Q849P2	TRPB_PSESH		cytoplasm
Q849Q9	SEPA_PSEPU	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
Q849R0	SEPB_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q849R1	SEPC_PSEPU	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
Q84BK9	LCBA1_AERHY		cytoplasm
Q84BL4	RL20_VIBAN		cytoplasm
Q84CH1	CPS1A_ACTSU		cytoplasm
Q84CP9	DAPB_PSEFL	cytoplasm (by similarity).	cytoplasm
Q84CZ9	SACB4_NEIMA		cytoplasm
Q84D00	SACB5_NEIMA		cytoplasm
Q84F69	INH_PHOLU	periplasm (by similarity).	periplasm
Q84F70	PRTA1_PHOLU	secreted protein (by similarity).	extracellular
Q84F94	HSLV_MYXXA	cytoplasm (by similarity).	cytoplasm
Q84FF7	TOLB_MYXXA	periplasm (potential).	periplasm
Q84FY8	MDH_METEX		cytoplasm
Q84GJ9	TRPB_NEIGO		cytoplasm
Q84HD6	AMYS_NEIME	secreted protein (by similarity).	cytoplasm
Q84I51	HIS8_BUCSC		cytoplasm
Q84I52	HIS8_BUCMH		cytoplasm
Q84I53	HIS8_BUCDN		cytoplasm
Q879S2	RNPA_XYLFT		extracellular
Q879S3	OXAA_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q879S5	TRME_XYLFT		cytoplasm
Q879Y6	RHLB_XYLFT		cytoplasm
Q879Z0	UNG_XYLFT	cytoplasm (by similarity).	extracellular
Q879Z5	NORM_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q879Z7	OPGD_XYLFT	periplasm (by similarity).	periplasm
Q87A18	HEM1_XYLFT		cytoplasm

Q87A20	LOLB_XYLFT	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q87A21	ISPE_XYLFT		cytoplasm
Q87A22	KPRS_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87A23	RL25_XYLFT		cytoplasm
Q87A24	PTH_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87A27	NUSG_XYLFT		cytoplasm
Q87A28	RL11_XYLFT		cytoplasm
Q87A29	RL1_XYLFT		cytoplasm
Q87A30	RL10_XYLFT		cytoplasm
Q87A31	RL7_XYLFT		cytoplasm
Q87A32	RPOB_XYLFT		cytoplasm
Q87A33	RPOC_XYLFT		cytoplasm
Q87A34	RS7_XYLFT		cytoplasm
Q87A36	HTPX_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q87A39	Y1992_XYLFT		inner membrane
Q87A46	Y1985_XYLFT		inner membrane
Q87A51	PHS_XYLFT		cytoplasm
Q87A52	Y1979_XYLFT		inner membrane
Q87A63	PPK_XYLFT		cytoplasm
Q87A66	LPXH_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87A73	ISPG_XYLFT		cytoplasm
Q87A76	MURB_XYLFT	cytoplasm (probable).	inner membrane
Q87A77	PYRD_XYLFT	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q87A84	RS18_XYLFT		cytoplasm
Q87A85	RL9_XYLFT		cytoplasm
Q87A87	ZIPA_XYLFT	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q87A92	MTNA_XYLFT		cytoplasm
Q87A98	SUCC_XYLFT		cytoplasm
Q87AA7	COAE_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87AB0	RIMK_XYLFT		cytoplasm
Q87AB4	RL20_XYLFT		cytoplasm
Q87AB7	IHFA_XYLFT		cytoplasm
Q87AC0	MTFA_XYLFT		cytoplasm
Q87AC5	EX7L_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87AC9	MUTL_XYLFT		cytoplasm
Q87AD1	DCD_XYLFT		extracellular
Q87AE8	THIG_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87AE9	TRMB_XYLFT		cytoplasm
Q87AF1	MRAZ_XYLFT		inner membrane
Q87AF2	MRAW_XYLFT		cytoplasm
Q87AF5	MURE_XYLFT	cytoplasm (probable).	cytoplasm
Q87AF7	MRAY_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87AF9	MURG_XYLFT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87AG0	MURC_XYLFT	cytoplasm (probable).	cytoplasm
Q87AG1	DDL_XYLFT	cytoplasm (by similarity).	cytoplasm

Q87AG5	LPXC_XYLFT		cytoplasm
Q87AH8	PGK_XYLFT	cytoplasm (potential).	cytoplasm
Q87AI0	ALF1_XYLFT		cytoplasm
Q87AI9	OSTA_XYLFT	bacterial cell outer membrane (by similarity).	outer membrane
Q87AJ1	PDXA_XYLFT	cytoplasm (probable).	cytoplasm
Q87AJ9	MSRB_XYLFT		cytoplasm
Q87AK0	DADA_XYLFT		cytoplasm
Q87AK1	ALR_XYLFT		cytoplasm
Q87AK6	SURE_XYLFT	cytoplasm (potential).	periplasm
Q87AP2	Y1781_XYLFT		inner membrane
Q87AQ6	TOP1_XYLFT		cytoplasm
Q87AQ7	SMG_XYLFT		cytoplasm
Q87AR0	FMT_XYLFT		cytoplasm
Q87AR1	RSMB_XYLFT		cytoplasm
Q87AR6	Y1756_XYLFT	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q87AR7	RLUD_XYLFT		cytoplasm
Q87AR8	Y1754_XYLFT		cytoplasm
Q87AS0	MQO_XYLFT		cytoplasm
Q87AS2	GLYA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87AS3	NRDR_XYLFT		cytoplasm
Q87AS7	RISB_XYLFT		cytoplasm
Q87AS8	NUSB_XYLFT		cytoplasm
Q87AT3	DAPA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87AT6	UVRB_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87AV1	Y1712_XYLFT		inner membrane
Q87AV3	DSBB_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87AV9	C1332_XYLFT		cytoplasm
Q87AW7	PBPA_XYLFT	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q87AX5	C1331_XYLFT		cytoplasm
Q87AX6	THIE_XYLFT		cytoplasm
Q87AX8	CLPB_XYLFT	cytoplasm (probable).	cytoplasm
Q87AY2	DUSA_XYLFT		cytoplasm
Q87AY7	NUDH_XYLFT		cytoplasm
Q87AZ9	RMUC_XYLFT		cytoplasm
Q87B16	RS21_XYLFT		cytoplasm
Q87B33	PURA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87B35	NDK_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87B41	ENGA_XYLFT		cytoplasm
Q87B70	Y1588_XYLFT		cytoplasm
Q87B72	Y1586_XYLFT		inner membrane
Q87B73	ENGC_XYLFT		cytoplasm
Q87B79	TATB_XYLFT	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q87B80	TATA_XYLFT	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q87B82	HEMH_XYLFT	cytoplasm (by similarity).	extracellular

Q87B84	RDGC_XYLFT	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q87B86	FABA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87B89	KTHY_XYLFT		cytoplasm
Q87B94	PUR5_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BA5	Y1553_XYLFT		cytoplasm
Q87BC0	CH60_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BC5	AROE_XYLFT		cytoplasm
Q87BE2	TILS_XYLFT	cytoplasm (probable).	cytoplasm
Q87BE3	EX7S_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BG0	BIOD_XYLFT		cytoplasm
Q87BG1	SUHB_XYLFT		cytoplasm
Q87BG5	UBIG_XYLFT		cytoplasm
Q87BG6	GPH_XYLFT		cytoplasm
Q87BG9	METX_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BH3	DSBE_XYLFT	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q87BH5	CCME_XYLFT	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q87BH8	CCMA_XYLFT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87BI2	RLUB_XYLFT		cytoplasm
Q87BJ6	KCY_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BJ8	IHFB_XYLFT		cytoplasm
Q87BK6	GUAA_XYLFT		cytoplasm
Q87BK9	UVRA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BL1	RL27_XYLFT		cytoplasm
Q87BL5	SYI_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BL6	LSPA_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q87BP7	QUEF_XYLFT	cytoplasm (probable).	cytoplasm
Q87BP9	LEU2_XYLFT		cytoplasm
Q87BQ0	LEUD_XYLFT		cytoplasm
Q87BQ1	LEU3_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BR3	RUMA_XYLFT		cytoplasm
Q87BR5	NAGZ_XYLFT	cytoplasm (by similarity).	extracellular
Q87BR7	MTNP_XYLFT		cytoplasm
Q87BS0	SSRP_XYLFT	cytoplasm (potential).	cytoplasm
Q87BS2	Y1376_XYLFT		inner membrane
Q87BS5	RECN_XYLFT		cytoplasm
Q87BS6	HRCA_XYLFT		cytoplasm
Q87BS7	GRPE_XYLFT	cytoplasm (probable).	cytoplasm
Q87BS8	DNAK_XYLFT		cytoplasm
Q87BS9	DNAJ_XYLFT	cytoplasm (probable).	cytoplasm
Q87BT3	LGT_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87BT4	TYSY_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BU0	SERC_XYLFT	cytoplasm (by similarity).	cytoplasm

Q87BU2	AROA_XYLFT	cytoplasm (probable).	cytoplasm
Q87BV3	Y1345_XYLFT		extracellular
Q87BW3	GSA_XYLFT	cytoplasm (potential).	cytoplasm
Q87BX2	UVRC_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87BY1	PARB_XYLFT		cytoplasm
Q87BY8	METE_XYLFT		cytoplasm
Q87BZ2	GLPK_XYLFT		cytoplasm
Q87BZ9	Y1297_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87C00	ACSA_XYLFT		cytoplasm
Q87C03	DXS_XYLFT		cytoplasm
Q87C04	RECO_XYLFT		cytoplasm
Q87C05	ERA_XYLFT		cytoplasm
Q87C06	RNC_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87C09	LEPA_XYLFT		cytoplasm
Q87C20	Y1276_XYLFT		inner membrane
Q87C22	PYRB_XYLFT		cytoplasm
Q87C24	KHSE_XYLFT	cytoplasm (potential).	cytoplasm
Q87C28	HIS1_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87C29	HISX_XYLFT		cytoplasm
Q87C30	HIS8_XYLFT		cytoplasm
Q87C31	HIS7_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87C32	HIS5_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87C33	HIS4_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87C34	HIS6_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87C35	HIS2_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87C43	EFPL_XYLFT		cytoplasm
Q87C45	PRMA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87C62	NADD_XYLFT		cytoplasm
Q87C76	RNH_XYLFT	cytoplasm (potential).	extracellular
Q87C81	KDKA_XYLFT	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q87C83	APAH_XYLFT		extracellular
Q87C84	APAG_XYLFT		inner membrane
Q87C85	KSGA_XYLFT		cytoplasm
Q87C86	RNT_XYLFT		cytoplasm
Q87C87	PHOU_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87C88	PSTB_XYLFT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87C89	PSTA_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87C90	PSTC_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87C91	PSTS_XYLFT	periplasm (potential).	periplasm
Q87CA2	Y1186_XYLFT		cytoplasm
Q87CB7	SLYX_XYLFT		inner membrane
Q87CC5	OPGH_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87CC9	HEM3_XYLFT		cytoplasm

Q87CI7	MTGA_XYLFT	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q87CI8	MUTS_XYLFT		cytoplasm
Q87CK2	SECB_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87CK3	GPDA_XYLFT	cytoplasm (probable).	cytoplasm
Q87CK7	Y1058_XYLFT		inner membrane
Q87CK8	RECR_XYLFT		cytoplasm
Q87CL6	RL32_XYLFT		cytoplasm
Q87CL7	FABH_XYLFT	cytoplasm (probable).	cytoplasm
Q87CL8	LEU1_XYLFT		cytoplasm
Q87CM2	ILVC_XYLFT		cytoplasm
Q87CN1	UBIB_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87CN3	Y1032_XYLFT		cytoplasm
Q87CN8	UPPP_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87CU3	Y970_XYLFT		cytoplasm
Q87CY6	THIC_XYLFT		cytoplasm
Q87CZ0	NFI_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87CZ1	GPMA_XYLFT		cytoplasm
Q87CZ5	TOLB_XYLFT	periplasm (potential).	periplasm
Q87D00	RUVB_XYLFT		cytoplasm
Q87D01	KUP_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87D02	RUVA_XYLFT		cytoplasm
Q87D03	RUVC_XYLFT		cytoplasm
Q87D04	Y885_XYLFT		inner membrane
Q87D06	FETP_XYLFT		cytoplasm
Q87D16	TRPR_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87D18	NADA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87D19	NADB_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87D24	GIDB_XYLFT		cytoplasm
Q87D27	MSRA_XYLFT		cytoplasm
Q87D42	GSHB_XYLFT		cytoplasm
Q87D58	PUR9_XYLFT		cytoplasm
Q87D59	PUR2_XYLFT		cytoplasm
Q87D63	GCH1_XYLFT		cytoplasm
Q87D65	RLMB_XYLFT		cytoplasm
Q87D75	FISL_XYLFT		cytoplasm
Q87D84	Y802_XYLFT		inner membrane
Q87D89	RPIA_XYLFT		cytoplasm
Q87DA0	PPNK_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DA5	Y781_XYLFT		inner membrane
Q87DB3	GIDA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DC2	FUMC_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DD1	SPED_XYLFT		cytoplasm
Q87DD4	RS9_XYLFT		cytoplasm
Q87DD6	RL31B_XYLFT		cytoplasm
Q87DF7	HEMK_XYLFT		cytoplasm
Q87DF8	PIP_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DG2	HAM1_XYLFT		extracellular
Q87DG3	RNPH_XYLFT		cytoplasm

Q87DG5	KGUA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DG6	RPOZ_XYLFT		cytoplasm
Q87DG7	CYSNC_XYLFT		cytoplasm
Q87DG8	CYSD_XYLFT		cytoplasm
Q87DH1	CYSH_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DH6	Y709_XYLFT		extracellular
Q87DI1	UBIE_XYLFT		cytoplasm
Q87DI2	XERC_XYLFT	cytoplasm (by similarity).	nucleoid
Q87DI4	DAPF_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DJ2	CSD_XYLFT		cytoplasm
Q87DJ6	GLMM_XYLFT		cytoplasm
Q87DK2	Y680_XYLFT		cytoplasm
Q87DK8	CRCB_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87DL0	LOLA_XYLFT	periplasm (by similarity).	periplasm
Q87DL5	LFTR_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DL8	CLPS_XYLFT		cytoplasm
Q87DM0	TRMU_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DM1	Y661_XYLFT		inner membrane
Q87DM6	ENGB_XYLFT		cytoplasm
Q87DM8	ARGD_XYLFT	cytoplasm (probable).	cytoplasm
Q87DN0	XERD_XYLFT	cytoplasm (by similarity).	nucleoid
Q87DN8	MURA_XYLFT	cytoplasm (probable).	inner membrane
Q87DP7	HPRK_XYLFT		cytoplasm
Q87DP8	Y634_XYLFT		cytoplasm
Q87DQ0	PTHP_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DQ5	SSB_XYLFT		cytoplasm
Q87DR1	GCSP_XYLFT		cytoplasm
Q87DR8	TRPA_XYLFT		cytoplasm
Q87DR9	TRPB_XYLFT		cytoplasm
Q87DS0	TRPF_XYLFT		cytoplasm
Q87DS1	TRUA_XYLFT		cytoplasm
Q87DS4	AROC_XYLFT		cytoplasm
Q87DS5	Y606_XYLFT		cytoplasm
Q87DS7	PSD_XYLFT		cytoplasm
Q87DS9	CCA_XYLFT		cytoplasm
Q87DT3	BIOH_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DT7	RPOD_XYLFT		cytoplasm
Q87DT9	CYSA_XYLFT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87DU4	CUTC_XYLFT	cytoplasm (probable).	periplasm
Q87DU7	PDXH_XYLFT		cytoplasm
Q87DU8	AROK_XYLFT	cytoplasm (probable).	cytoplasm
Q87DU9	AROB_XYLFT	cytoplasm (probable).	cytoplasm
Q87DV0	DCUP_XYLFT	cytoplasm (probable).	cytoplasm
Q87DW2	MINC_XYLFT		inner membrane
Q87DW7	QUEA_XYLFT	cytoplasm (potential).	cytoplasm
Q87DY3	ISPF_XYLFT		extracellular
Q87DY4	ISPD_XYLFT		cytoplasm
Q87DY5	FTSB_XYLFT	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane

Q87DY6	ENO_XYLFT	cytoplasm (by similarity).	extracellular
Q87DY7	KDSA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DY8	PYRG_XYLFT		cytoplasm
Q87DZ7	Y532_XYLFT		inner membrane
Q87DZ8	LIPB_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87DZ9	LIPA_XYLFT	cytoplasm (potential).	cytoplasm
Q87E04	Y525_XYLFT		cytoplasm
Q87E07	ORN_XYLFT	cytoplasm (potential).	cytoplasm
Q87E11	Y518_XYLFT		cytoplasm
Q87E35	MDH_XYLFT		cytoplasm
Q87E48	DBH_XYLFT		cytoplasm
Q87E50	CLPX_XYLFT		cytoplasm
Q87E51	CLPP_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87E52	TIG_XYLFT		cytoplasm
Q87E55	LIFO_XYLFT	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
Q87E60	RS4_XYLFT		cytoplasm
Q87E61	RS13_XYLFT		cytoplasm
Q87E65	RS5_XYLFT		cytoplasm
Q87E66	RL18_XYLFT		cytoplasm
Q87E68	RS8_XYLFT		cytoplasm
Q87E69	RS14_XYLFT		cytoplasm
Q87E70	RL5_XYLFT		cytoplasm
Q87E71	RL24_XYLFT		cytoplasm
Q87E73	RS17_XYLFT		cytoplasm
Q87E74	RL29_XYLFT		cytoplasm
Q87E75	RL16_XYLFT		cytoplasm
Q87E76	RS3_XYLFT		cytoplasm
Q87E77	RL22_XYLFT		cytoplasm
Q87E78	RS19_XYLFT		cytoplasm
Q87E79	RL2_XYLFT		cytoplasm
Q87E81	RL4_XYLFT		cytoplasm
Q87E82	RL3_XYLFT		cytoplasm
Q87E88	ATPA_XYLFT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87E89	ATPG_XYLFT		inner membrane
Q87E91	ATPE_XYLFT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87E93	GLMU_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87E99	Y419_XYLFT		inner membrane
Q87EA3	Y415_XYLFT	cytoplasm (potential).	inner membrane
Q87EA7	MURD_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EB7	GREA_XYLFT		cytoplasm
Q87EB8	CARB_XYLFT		cytoplasm
Q87EB9	CARA_XYLFT		cytoplasm
Q87EC0	DAPB_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EC4	PNCB_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EE9	LPXK_XYLFT		cytoplasm
Q87EF0	MSBA_XYLFT	bacterial cell inner membrane; multi- pass membrane protein (by similarity).	inner membrane

Q87EF4	LOLD_XYLFT	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87EF5	LOLC_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87EF7	Y354_XYLFT		cytoplasm
Q87EG6	GLK_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EG7	6PGL_XYLFT		extracellular
Q87EH1	PYRH_XYLFT		cytoplasm
Q87EH6	RRF_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EH7	UPPS_XYLFT		inner membrane
Q87EH9	DXR_XYLFT		cytoplasm
Q87EI0	Y327_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q87EI2	LPXD_XYLFT		cytoplasm
Q87EI3	FABZ_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EI4	LPXA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EI5	LPXB_XYLFT		cytoplasm
Q87EI6	RNH2_XYLFT	cytoplasm (potential).	extracellular
Q87EI8	SAHH_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EJ1	PLSB_XYLFT	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q87EJ8	ATE_XYLFT	cytoplasm (potential).	cytoplasm
Q87EJ9	MNTH_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q87EK9	PROA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EL0	PROB_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EL1	ARGC_XYLFT	cytoplasm (probable).	cytoplasm
Q87EL2	ARGB_XYLFT	cytoplasm (probable).	cytoplasm
Q87EL4	AOTC_XYLFT	cytoplasm (probable).	cytoplasm
Q87EM1	Y282_XYLFT		inner membrane
Q87EM8	COAD_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EN0	HTPG_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EN2	DCTA_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87EQ7	TPIS_XYLFT		cytoplasm
Q87ES6	KAD_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EV1	RS15_XYLFT		cytoplasm
Q87EV2	TRUB_XYLFT		cytoplasm
Q87EV3	RBFA_XYLFT	cytoplasm (potential).	cytoplasm
Q87EV6	Y192_XYLFT		inner membrane
Q87EV7	G6PI_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EV8	PAND_XYLFT		cytoplasm
Q87EV9	PANC_XYLFT		cytoplasm
Q87EW0	PANB_XYLFT		cytoplasm
Q87EW6	TGT_XYLFT		cytoplasm
Q87EX2	TRPC_XYLFT		cytoplasm
Q87EX3	TRPD_XYLFT		cytoplasm
Q87EX9	PUR7_XYLFT		cytoplasm
Q87EY0	DPO3A_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87EY1	ACCA_XYLFT	cytoplasm (by similarity).	cytoplasm

Q87EZ6	GCST_XYLFT		cytoplasm
Q87EZ7	GCSH_XYLFT		cytoplasm
Q87F06	SYX_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87F16	PYRE_XYLFT		cytoplasm
Q87F19	DUT_XYLFT		extracellular
Q87F21	RADC_XYLFT		cytoplasm
Q87F25	SPEA_XYLFT		cytoplasm
Q87F26	SPEE_XYLFT		cytoplasm
Q87F28	GLMS_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87F32	AMPA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87F36	SYV_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87F44	RECA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87F45	LEXA_XYLFT		cytoplasm
Q87F51	DAPD_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87F53	RL19_XYLFT		cytoplasm
Q87F54	TRMD_XYLFT	cytoplasm (potential).	cytoplasm
Q87F55	RIMM_XYLFT	cytoplasm (potential).	cytoplasm
Q87F56	RS16_XYLFT		cytoplasm
Q87F63	ILVD_XYLFT		cytoplasm
Q87F66	RRMJ_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87F70	MIAA_XYLFT		cytoplasm
Q87F71	HFQ_XYLFT		cytoplasm
Q87F88	PDXJ_XYLFT	cytoplasm (probable).	cytoplasm
Q87F90	HMP_XYLFT		cytoplasm
Q87F94	AROQ_XYLFT		cytoplasm
Q87FA1	MSCL_XYLFT	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87FA3	PYRF_XYLFT		cytoplasm
Q87FB2	HEM6_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87FC4	RECF_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87FC6	DNAA_XYLFT	cytoplasm (by similarity).	cytoplasm
Q87FD6	Y5743_VIBPA		inner membrane
Q87FE9	PDXH_VIBPA		cytoplasm
Q87FH3	UXAC_VIBPA		cytoplasm
Q87FH9	UXUA_VIBPA		cytoplasm
Q87FK5	ARAB_VIBPA		cytoplasm
Q87FP9	ENGC2_VIBPA		cytoplasm
Q87FQ5	MALT_VIBPA		cytoplasm
Q87FR0	GLGB_VIBPA		extracellular
Q87FR1	MTGA_VIBPA	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q87FT0	GBPA_VIBPA	secreted protein (by similarity).	cytoplasm
Q87FT1	Y5597_VIBPA		inner membrane
Q87FU4	Y5584_VIBPA		inner membrane
Q87FV4	MDTL_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87FZ3	Y5527_VIBPA		cytoplasm
Q87G18	LLDD_VIBPA		cytoplasm
Q87G35	CBIO_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87G42	DEOD2_VIBPA		cytoplasm

Q87G59	PSTB2_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87GA3	Y5414_VIBPA		inner membrane
Q87GB5	MALK_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87GB7	MALF_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87GB8	MALG_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87GM3	ADD2_VIBPA		extracellular
Q87GR5	RIBB_VIBPA		cytoplasm
Q87GT9	Y5226_VIBPA		inner membrane
Q87GU3	Y5222_VIBPA		cytoplasm
Q87GU4	LUXP_VIBPA	periplasm (probable).	periplasm
Q87GU5	LUXQ_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q87GW6	NAPA_VIBPA	periplasm (by similarity).	periplasm
Q87GX2	PEPT_VIBPA	cytoplasm (by similarity).	extracellular
Q87GY4	TKT2_VIBPA		extracellular
Q87GY5	TAL_VIBPA	cytoplasm (probable).	cytoplasm
Q87GZ6	GCH1_VIBPA		cytoplasm
Q87GZ9	CLCA_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q87H06	GUAC_VIBPA		cytoplasm
Q87H51	BETI_VIBPA		cytoplasm
Q87H52	BETB_VIBPA		cytoplasm
Q87HG4	ALR2_VIBPA		cytoplasm
Q87HJ2	FABH2_VIBPA	cytoplasm (probable).	cytoplasm
Q87HN1	COBQ_VIBPA		cytoplasm
Q87HN4	MODC_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87HP7	Y4916_VIBPA		inner membrane
Q87HS0	DDL_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87HS9	Y4877_VIBPA		inner membrane
Q87HT6	Y4870_VIBPA		cytoplasm
Q87HV6	Y4850_VIBPA	periplasm (potential).	periplasm
Q87HX2	FTHS_VIBPA		cytoplasm
Q87HX3	GLGC2_VIBPA		cytoplasm
Q87I03	GLYA2_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87I04	GCSH_VIBPA		cytoplasm
Q87I05	GCSP_VIBPA		cytoplasm
Q87I22	DEF2_VIBPA		cytoplasm
Q87ID3	MGSA_VIBPA		cytoplasm
Q87IJ5	ACKA2_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87IJ9	RNB_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87IL9	PCP_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87IM1	TRPB2_VIBPA		cytoplasm
Q87IM9	PHS_VIBPA		cytoplasm
Q87IV1	Y4505_VIBPA		cytoplasm

Q87J41	NADE_VIBPA		cytoplasm
Q87J46	PYRC_VIBPA		cytoplasm
Q87J80	Y4373_VIBPA		cytoplasm
Q87J85	SYI1_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87JD1	Y4321_VIBPA		inner membrane
Q87JE4	PNCB_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87JE8	CATA_VIBPA	periplasm (probable).	periplasm
Q87JG6	CH602_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87JG7	CH102_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87JH5	IDI2_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87JI2	FLG11_VIBPA	periplasm (by similarity).	periplasm
Q87JI3	FLGH2_VIBPA	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q87JP5	CRYD_VIBPA		cytoplasm
Q87JQ6	TNAA_VIBPA		cytoplasm
Q87JS5	RIMK_VIBPA		cytoplasm
Q87JS8	SPEA_VIBPA		cytoplasm
Q87JV8	Y4140_VIBPA		inner membrane
Q87JW7	THIM_VIBPA		cytoplasm
Q87JW8	THIE_VIBPA		cytoplasm
Q87JZ5	Y4103_VIBPA		inner membrane
Q87K41	Y4057_VIBPA		inner membrane
Q87K60	NAGB_VIBPA		extracellular
Q87K77	Y4021_VIBPA		cytoplasm
Q87K87	Y4011_VIBPA		inner membrane
Q87K98	GIDA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87K99	GIDB_VIBPA		cytoplasm
Q87KA4	ATPF_VIBPA		inner membrane
Q87KA6	ATPA_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87KA7	ATPG_VIBPA		inner membrane
Q87KA9	ATPE_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87KB0	GLMU_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87KB6	ILVD_VIBPA		cytoplasm
Q87KD3	RSMB_VIBPA		cytoplasm
Q87KD4	FMT_VIBPA		cytoplasm
Q87KD5	DEF1_VIBPA		cytoplasm
Q87KD8	SMG_VIBPA		cytoplasm
Q87KE0	PURK_VIBPA		cytoplasm
Q87KE1	PUR6_VIBPA		cytoplasm
Q87KE3	HEM6_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87KE4	AROE_VIBPA		cytoplasm
Q87KE9	CRCB_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87KF0	THIC_VIBPA		cytoplasm
Q87KF4	THIG_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87KG2	Y3015_VIBPA		inner membrane
Q87KH4	GPPA_VIBPA		extracellular
Q87KH5	RHLB_VIBPA		cytoplasm
Q87KI9	HEM3_VIBPA		cytoplasm
Q87KJ1	CYAY_VIBPA		inner membrane
Q87KJ4	DAPF_VIBPA	cytoplasm (by similarity).	cytoplasm

Q87KM5	GLPE_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87KN0	PLSB_VIBPA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q87KN2	LEXA_VIBPA		cytoplasm
Q87KN5	STHA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87KN8	TRMA_VIBPA		cytoplasm
Q87KN9	BTUB_VIBPA	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q87KP1	MURI_VIBPA		cytoplasm
Q87KP5	MURB_VIBPA	cytoplasm (probable).	inner membrane
Q87KP9	NUSG_VIBPA		cytoplasm
Q87KQ0	RL11_VIBPA		cytoplasm
Q87KQ1	RL1_VIBPA		cytoplasm
Q87KQ2	RL10_VIBPA		cytoplasm
Q87KQ3	RL7_VIBPA		cytoplasm
Q87KQ4	RPOB_VIBPA		cytoplasm
Q87KQ5	RPOC_VIBPA		cytoplasm
Q87KQ7	NUDC_VIBPA		cytoplasm
Q87KR0	DCUP_VIBPA	cytoplasm (probable).	cytoplasm
Q87KS8	PUR2_VIBPA		cytoplasm
Q87KT0	PUR9_VIBPA		cytoplasm
Q87KU1	DUSB_VIBPA		cytoplasm
Q87KU2	PRMA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87KU6	AROQ_VIBPA		cytoplasm
Q87KU7	ACSA_VIBPA		cytoplasm
Q87KV8	CVRA_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q87KW0	DSBD_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87KX0	K6PF_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87KX3	CH101_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87KY1	FRDD_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87KY2	FRDC_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87KY6	SYK3_VIBPA		cytoplasm
Q87KZ2	GPDA_VIBPA	cytoplasm (probable).	cytoplasm
Q87KZ3	SECB_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87KZ5	GPMI_VIBPA		cytoplasm
Q87KZ9	PSD_VIBPA		cytoplasm
Q87L00	ENGC1_VIBPA		cytoplasm
Q87L01	ORN_VIBPA	cytoplasm (potential).	cytoplasm
Q87L05	MUTL_VIBPA		cytoplasm
Q87L06	MIAA_VIBPA		cytoplasm
Q87L07	HFQ_VIBPA		cytoplasm
Q87L11	RLMB_VIBPA		cytoplasm
Q87L12	GPH_VIBPA		cytoplasm
Q87L20	ARGD_VIBPA	cytoplasm (probable).	cytoplasm
Q87L25	Y2791_VIBPA		inner membrane
Q87L36	SLYX_VIBPA		inner membrane

Q87L40	TUSD_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87L41	TUSC_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87L43	RS12_VIBPA		cytoplasm
Q87L44	RS7_VIBPA		cytoplasm
Q87L49	METJ_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87L54	CAPP_VIBPA		cytoplasm
Q87L55	ARGC_VIBPA	cytoplasm (probable).	cytoplasm
Q87L56	ARGB_VIBPA	cytoplasm (probable).	cytoplasm
Q87L67	AROK_VIBPA	cytoplasm (probable).	cytoplasm
Q87L68	AROB_VIBPA	cytoplasm (probable).	cytoplasm
Q87L72	RS6_VIBPA		cytoplasm
Q87L73	PRIB_VIBPA		cytoplasm
Q87L75	RL9_VIBPA		cytoplasm
Q87L78	ALR1_VIBPA		cytoplasm
Q87L81	G6PI_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87L85	DUSA_VIBPA		cytoplasm
Q87L90	CYSJ_VIBPA		cytoplasm
Q87L91	CYSI_VIBPA		cytoplasm
Q87L92	CYSH_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87L95	METH_VIBPA		cytoplasm
Q87LA0	UVRA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87LA3	SSB_VIBPA		cytoplasm
Q87LC4	Y2688_VIBPA	cytoplasm (potential).	inner membrane
Q87LD0	RAPA_VIBPA		cytoplasm
Q87LD3	RLUA_VIBPA		cytoplasm
Q87LD5	Y2677_VIBPA		cytoplasm
Q87LD9	Y2673_VIBPA		cytoplasm
Q87LF4	MURA_VIBPA	cytoplasm (probable).	inner membrane
Q87LF7	PYRI_VIBPA		cytoplasm
Q87LF8	PYRB_VIBPA		cytoplasm
Q87LF9	OTC_VIBPA	cytoplasm (probable).	cytoplasm
Q87LG0	ARCA_VIBPA	cytoplasm (potential).	extracellular
Q87LG6	SYV_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87LG8	AMPA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87LH9	Y2633_VIBPA		inner membrane
Q87LI5	FETP_VIBPA		cytoplasm
Q87LI7	TRMB_VIBPA		cytoplasm
Q87LI9	GLSA_VIBPA		extracellular
Q87LJ1	HAM1_VIBPA		extracellular
Q87LJ3	Y2619_VIBPA		inner membrane
Q87LJ9	RUVX_VIBPA	cytoplasm (potential).	cytoplasm
Q87LK0	Y2612_VIBPA		inner membrane
Q87LK1	GSHB_VIBPA		cytoplasm
Q87LK4	SPRT_VIBPA	cytoplasm (potential).	extracellular
Q87LK8	TKT1_VIBPA		extracellular
Q87LL1	PGK_VIBPA	cytoplasm (potential).	cytoplasm
Q87LL6	ICIA_VIBPA		cytoplasm
Q87LL9	RPIA_VIBPA		cytoplasm
Q87LM3	Y2588_VIBPA		inner membrane
Q87LM9	Y2582_VIBPA		cytoplasm
Q87LN7	LEPA_VIBPA		cytoplasm
Q87LN9	RNC_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87LP0	ERA_VIBPA		cytoplasm
Q87LP1	RECO_VIBPA		cytoplasm
Q87LP2	PDXJ_VIBPA	cytoplasm (probable).	cytoplasm
Q87LP3	ACPS_VIBPA	cytoplasm (by similarity).	cytoplasm

Q87LP5	RUMA_VIBPA		cytoplasm
Q87LP9	PYRG_VIBPA		cytoplasm
Q87LQ0	ENO_VIBPA	cytoplasm (by similarity).	extracellular
Q87LQ1	FTSB_VIBPA	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q87LQ2	ISPD1_VIBPA		cytoplasm
Q87LQ3	ISPF_VIBPA		extracellular
Q87LQ4	TRUD_VIBPA		cytoplasm
Q87LQ5	SURE_VIBPA	cytoplasm (potential).	periplasm
Q87LQ6	PIMT_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87LQ9	MUTS_VIBPA		cytoplasm
Q87LR1	RECA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87LR2	RECX_VIBPA	cytoplasm (potential).	cytoplasm
Q87LR5	CSRA_VIBPA		cytoplasm
Q87LR6	OADG_VIBPA	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q87LS2	GSH1_VIBPA		cytoplasm
Q87LS4	LUXS_VIBPA		extracellular
Q87LS8	RS16_VIBPA		cytoplasm
Q87LS9	RIMM_VIBPA	cytoplasm (potential).	cytoplasm
Q87LT0	TRMD_VIBPA	cytoplasm (potential).	cytoplasm
Q87LT1	RL19_VIBPA		cytoplasm
Q87LT2	Y2529_VIBPA		cytoplasm
Q87LT3	Y2528_VIBPA		inner membrane
Q87LT4	COAE_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87LV1	PANC_VIBPA		cytoplasm
Q87LV2	PANB_VIBPA		cytoplasm
Q87LV6	GLUQ_VIBPA		cytoplasm
Q87LV9	SFSA_VIBPA		cytoplasm
Q87LW4	Y2494_VIBPA		inner membrane
Q87LY3	GSA_VIBPA	cytoplasm (potential).	cytoplasm
Q87LY8	SYI2_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87LZ1	OMPU_VIBPA	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q87LZ2	GREA_VIBPA		cytoplasm
Q87LZ4	RRMJ_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87LZ7	GLMM_VIBPA		cytoplasm
Q87M00	Y2458_VIBPA		inner membrane
Q87M03	RBFA_VIBPA	cytoplasm (potential).	cytoplasm
Q87M04	TRUB_VIBPA		cytoplasm
Q87M05	RS15_VIBPA		cytoplasm
Q87M22	DEOC_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87M23	TYPH_VIBPA		cytoplasm
Q87M24	DEOB_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87M25	DEOD1_VIBPA		cytoplasm
Q87M60	GAL1_VIBPA	cytoplasm (potential).	cytoplasm
Q87M72	GLPK_VIBPA		cytoplasm
Q87M87	ARGA_VIBPA	cytoplasm (probable).	cytoplasm
Q87MA6	NQRA_VIBPA		inner membrane

Q87MA7	NQRB_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q87MA8	NQRC_VIBPA	bacterial cell inner membrane (potential).	inner membrane
Q87MA9	NQRD_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q87MB0	NQRE_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q87MB4	DPO4_VIBPA	cytoplasm (probable).	cytoplasm
Q87MC1	Y2335_VIBPA		inner membrane
Q87MC5	RL31B_VIBPA		cytoplasm
Q87MC8	Y2328_VIBPA		inner membrane
Q87MD4	TRUC_VIBPA		cytoplasm
Q87MD5	Y2321_VIBPA		inner membrane
Q87MD6	GLND_VIBPA		cytoplasm
Q87MD8	RS2_VIBPA		cytoplasm
Q87ME0	PYRH_VIBPA		cytoplasm
Q87ME1	UPPS_VIBPA		inner membrane
Q87ME3	DXR_VIBPA		cytoplasm
Q87ME7	LPXD_VIBPA		cytoplasm
Q87ME8	FABZ_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87ME9	LPXA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87MF0	LPXB_VIBPA		cytoplasm
Q87MF1	RNH2_VIBPA	cytoplasm (potential).	extracellular
Q87MF3	ACCA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87MF4	TILS_VIBPA	cytoplasm (probable).	cytoplasm
Q87MF7	Y2298_VIBPA	cytoplasm (potential).	inner membrane
Q87MG2	RNH_VIBPA	cytoplasm (potential).	extracellular
Q87MG7	GMHA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87MH0	PUR5_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87MH1	UPP_VIBPA		cytoplasm
Q87MK8	CCMA_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87ML2	CCME_VIBPA	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q87MM2	FADI_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87MM3	FADJ_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87MM7	Y2204_VIBPA		inner membrane
Q87MM9	AROC_VIBPA		cytoplasm
Q87MN6	Y2195_VIBPA		cytoplasm
Q87MN8	PDXB_VIBPA		cytoplasm
Q87MP1	TRUA_VIBPA		cytoplasm
Q87MQ3	Y2178_VIBPA		inner membrane
Q87MQ4	RECR_VIBPA		cytoplasm
Q87MQ5	AQPZ_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87MR2	Y2169_VIBPA		inner membrane
Q87MS5	MSRB_VIBPA		cytoplasm
Q87MV2	Y2129_VIBPA		inner membrane

Q87MV3	NDPA_VIBPA	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q87MW9	RNFE_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87MX0	RNFG_VIBPA	bacterial cell inner membrane (potential).	inner membrane
Q87MX1	RNFD_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q87MX4	RNFA_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87MX6	UVRB_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87MX7	LUXO_VIBPA		cytoplasm
Q87MX8	LUXU_VIBPA		inner membrane
Q87MY0	MOAA_VIBPA		cytoplasm
Q87MY2	MOAC_VIBPA		cytoplasm
Q87MZ4	ACKA1_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87MZ5	Y2081_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87N01	DUSC_VIBPA		cytoplasm
Q87N03	DSBB_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87N05	FADR_VIBPA	cytoplasm (potential).	cytoplasm
Q87N09	URK_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87N15	RLUC_VIBPA		cytoplasm
Q87N16	Y2060_VIBPA	cytoplasm (potential).	inner membrane
Q87N18	RL32_VIBPA		cytoplasm
Q87N19	PLSX_VIBPA		cytoplasm
Q87N20	FABH1_VIBPA	cytoplasm (probable).	cytoplasm
Q87N26	KTHY_VIBPA		cytoplasm
Q87N39	DNAE2_VIBPA	cytoplasm (probable).	cytoplasm
Q87N44	KCY_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87N46	IHFB_VIBPA		cytoplasm
Q87N49	PYRF_VIBPA		cytoplasm
Q87N95	Y1980_VIBPA		cytoplasm
Q87NA1	METE_VIBPA		cytoplasm
Q87NA5	ISPZ_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87NB7	RLUB_VIBPA		cytoplasm
Q87NB9	Y1949_VIBPA		inner membrane
Q87NC0	EFPL_VIBPA		cytoplasm
Q87NC4	UVRC_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87ND5	UBIG_VIBPA		cytoplasm
Q87NE7	GCH2_VIBPA		extracellular
Q87NJ8	Y1870_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87NU5	AGUA_VIBPA		extracellular
Q87NW7	META_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87NZ6	ECTA_VIBPA		cytoplasm

Q87NZ7	ECTB_VIBPA		cytoplasm
Q87NZ8	ECTC_VIBPA		cytoplasm
Q87PB0	PROQ_VIBPA	cytoplasm (potential).	inner membrane
Q87PB7	PYRD_VIBPA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q87PC5	FABA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87PC8	Y1589_VIBPA		inner membrane
Q87PH8	NPD_VIBPA	cytoplasm (probable).	cytoplasm
Q87PK5	MOBA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87PM1	Y1481_VIBPA		cytoplasm
Q87PP9	ACPD_VIBPA		extracellular
Q87Q30	ISPD2_VIBPA		cytoplasm
Q87Q38	BTUD_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87Q39	BTUC_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87Q40	ASTE_VIBPA		periplasm
Q87Q45	COBS_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87Q46	COBT_VIBPA		cytoplasm
Q87Q50	Y1300_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87Q52	CDD_VIBPA		cytoplasm
Q87Q54	TPMT_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87Q56	IHFA_VIBPA		cytoplasm
Q87Q67	Y1283_VIBPA		inner membrane
Q87Q69	RL35_VIBPA		cytoplasm
Q87Q74	HUTI_VIBPA	cytoplasm (potential).	cytoplasm
Q87Q75	HUTG_VIBPA		cytoplasm
Q87Q76	HUTU_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87Q77	HUTH_VIBPA	cytoplasm (potential).	cytoplasm
Q87Q87	PUR7_VIBPA		cytoplasm
Q87Q92	MAO1_VIBPA		cytoplasm
Q87QA2	SDHD_VIBPA		cytoplasm
Q87QA3	SERC_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QB8	Y1232_VIBPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q87QB9	Y1231_VIBPA		cytoplasm
Q87QD4	RSUA_VIBPA		cytoplasm
Q87QD9	RL25_VIBPA		cytoplasm
Q87QG0	HCP_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QI7	TORA_VIBPA	periplasm (by similarity).	periplasm
Q87QJ8	KITH_VIBPA	cytoplasm (potential).	cytoplasm
Q87QK1	LPXH_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QK4	Y1145_VIBPA		inner membrane
Q87QK5	HIS2_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QK6	HIS6_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QK7	HIS4_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QK8	HIS5_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QK9	HIS7_VIBPA	cytoplasm (by similarity).	cytoplasm

Q87QL0	HIS8_VIBPA		cytoplasm
Q87QL1	HISX_VIBPA		cytoplasm
Q87QL2	HIS1_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QL9	TRMU_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QM0	Y1129_VIBPA		inner membrane
Q87QN1	HTPX_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q87QN3	BIOD_VIBPA		cytoplasm
Q87QP3	LOLA_VIBPA	periplasm (by similarity).	periplasm
Q87QT6	NADA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QT9	TOLB_VIBPA	periplasm (potential).	periplasm
Q87QU7	RUVB_VIBPA		cytoplasm
Q87QU8	RUVA_VIBPA		cytoplasm
Q87QV1	RUVC_VIBPA		cytoplasm
Q87QW4	MUKF_VIBPA	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q87QW8	TORD_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QX5	GLGA_VIBPA		cytoplasm
Q87QX6	GLGC1_VIBPA		cytoplasm
Q87QX9	AROA_VIBPA	cytoplasm (probable).	cytoplasm
Q87QY1	LFTR_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87QY2	ATE_VIBPA	cytoplasm (potential).	cytoplasm
Q87QY5	CLPS_VIBPA		cytoplasm
Q87QY8	RLUE_VIBPA		cytoplasm
Q87QY9	Y1009_VIBPA		inner membrane
Q87R08	Y990_VIBPA		inner membrane
Q87R12	Y986_VIBPA		inner membrane
Q87R14	KDSB_VIBPA	cytoplasm (potential).	cytoplasm
Q87R15	LPXK_VIBPA		cytoplasm
Q87R16	MSBA_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87R20	LOLD_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87R29	Y969_VIBPA		inner membrane
Q87R62	Y936_VIBPA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q87R72	Y926_VIBPA		cytoplasm
Q87R73	DGTL1_VIBPA		cytoplasm
Q87R79	CLPX_VIBPA		cytoplasm
Q87R80	CLPP_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87R81	TIG_VIBPA		cytoplasm
Q87R88	Y909_VIBPA		inner membrane
Q87RC3	MINC_VIBPA		inner membrane
Q87RC5	MINE_VIBPA		inner membrane
Q87RE8	SUCC_VIBPA		cytoplasm
Q87RH3	HEMH_VIBPA	cytoplasm (by similarity).	extracellular
Q87RH4	KAD_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RH5	HTPG_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RJ5	ZIPA_VIBPA	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane

Q87RJ6	CYSZ_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87RL7	Y761_VIBPA		inner membrane
Q87RN0	KDSA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RN5	HEM1_VIBPA		cytoplasm
Q87RN6	LOLB_VIBPA	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q87RN7	ISPE_VIBPA		cytoplasm
Q87RN8	KPRS_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RN9	PTH_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RP6	Y731_VIBPA		inner membrane
Q87RP8	LNT_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87RQ4	Y723_VIBPA		inner membrane
Q87RQ9	Y718_VIBPA		inner membrane
Q87RR0	LIPB_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RR1	LIPA_VIBPA	cytoplasm (potential).	cytoplasm
Q87RR2	GLYA1_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RS6	QUEF_VIBPA	cytoplasm (probable).	cytoplasm
Q87RT2	MTFA_VIBPA		cytoplasm
Q87RT6	THII_VIBPA	cytoplasm (potential).	cytoplasm
Q87RT8	EX7S_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RU0	DXS_VIBPA		cytoplasm
Q87RU3	NUSB_VIBPA		cytoplasm
Q87RU4	RISB_VIBPA		cytoplasm
Q87RU8	NRDR_VIBPA		cytoplasm
Q87RU9	PROA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RV0	PROB_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RV2	Y674_VIBPA		inner membrane
Q87RV3	XGPT_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RX2	DNAJ_VIBPA	cytoplasm (probable).	cytoplasm
Q87RX3	DNAK_VIBPA		cytoplasm
Q87RX5	GRPE_VIBPA	cytoplasm (probable).	cytoplasm
Q87RX6	PPNK_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87RY0	Y646_VIBPA		inner membrane
Q87RY2	SSRP_VIBPA	cytoplasm (potential).	cytoplasm
Q87S07	GUAA_VIBPA		cytoplasm
Q87S09	EX7L_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87S12	ENGA_VIBPA		cytoplasm
Q87S16	ISPG_VIBPA		cytoplasm
Q87S20	NDK_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87S21	PEPB_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87S24	HSCA_VIBPA		cytoplasm
Q87S25	HSCB_VIBPA		inner membrane
Q87S28	ISCS_VIBPA		cytoplasm
Q87S36	TGT_VIBPA		cytoplasm
Q87S37	QUEA_VIBPA	cytoplasm (potential).	cytoplasm
Q87S45	CUTC_VIBPA	cytoplasm (probable).	periplasm
Q87S48	PSTB1_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87S51	PPK_VIBPA		cytoplasm

Q87S56	RDGC_VIBPA	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q87S63	CLPB_VIBPA	cytoplasm (probable).	cytoplasm
Q87S65	RLUD_VIBPA		cytoplasm
Q87S70	Y554_VIBPA		inner membrane
Q87S71	TRPR_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87S81	Y543_VIBPA		cytoplasm
Q87S86	Y538_VIBPA		cytoplasm
Q87S87	ISPH_VIBPA		cytoplasm
Q87S89	LSPA_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q87S90	SYI_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87S93	RS20_VIBPA		cytoplasm
Q87SA0	TYSY_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87SA1	LGT_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87SA4	NUDH_VIBPA		cytoplasm
Q87SA6	MUTH_VIBPA	cytoplasm (by similarity).	extracellular
Q87SC0	Y504_VIBPA		inner membrane
Q87SC4	UNG_VIBPA	cytoplasm (by similarity).	extracellular
Q87SC6	END4_VIBPA		extracellular
Q87SC7	GRCA_VIBPA		cytoplasm
Q87SC9	KHSE_VIBPA	cytoplasm (potential).	cytoplasm
Q87SD2	RRAA1_VIBPA		cytoplasm
Q87SE7	BTUF_VIBPA	periplasm (by similarity).	periplasm
Q87SF3	CARB_VIBPA		cytoplasm
Q87SF4	CARA_VIBPA		cytoplasm
Q87SF5	DAPB_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87SF9	LPXC_VIBPA		cytoplasm
Q87SG3	MURC_VIBPA	cytoplasm (probable).	cytoplasm
Q87SG4	MURG_VIBPA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87SG6	MURD_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87SG7	MRAY_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87SG9	MURE_VIBPA	cytoplasm (probable).	cytoplasm
Q87SH5	Y448_VIBPA		inner membrane
Q87SI4	RS9_VIBPA		cytoplasm
Q87SJ9	HLDE_VIBPA		cytoplasm
Q87SK0	GLNE_VIBPA		cytoplasm
Q87SK9	CCA_VIBPA		cytoplasm
Q87SL0	UPPP_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87SL3	Y410_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87SQ3	MTLD_VIBPA		cytoplasm
Q87SR3	GLMS_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87SS7	LEU1_VIBPA		cytoplasm
Q87SS8	LEU3_VIBPA	cytoplasm (by similarity).	cytoplasm

Q87SS9	LEU2_VIBPA		cytoplasm
Q87ST0	LEUD_VIBPA		cytoplasm
Q87ST2	DJLA_VIBPA	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q87ST3	OSTA_VIBPA	bacterial cell outer membrane (by similarity).	outer membrane
Q87ST5	PDXA_VIBPA	cytoplasm (probable).	cytoplasm
Q87ST6	KSGA_VIBPA		cytoplasm
Q87ST7	APAG_VIBPA		inner membrane
Q87ST8	APAH_VIBPA		extracellular
Q87SU3	RL27_VIBPA		cytoplasm
Q87SU7	MDH_VIBPA		cytoplasm
Q87SU8	ARGR_VIBPA	cytoplasm (probable).	cytoplasm
Q87SW1	IPYR_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87SW6	MSRA_VIBPA		cytoplasm
Q87SX6	CYSC_VIBPA		cytoplasm
Q87SX9	CYSN_VIBPA		cytoplasm
Q87SY0	CYSD_VIBPA		cytoplasm
Q87SZ0	RPOA_VIBPA		cytoplasm
Q87SZ1	RS4_VIBPA		cytoplasm
Q87SZ2	RS13_VIBPA		cytoplasm
Q87SZ4	RL15_VIBPA		cytoplasm
Q87SZ6	RS5_VIBPA		cytoplasm
Q87SZ7	RL18_VIBPA		cytoplasm
Q87SZ9	RS8_VIBPA		cytoplasm
Q87T01	RL5_VIBPA		cytoplasm
Q87T02	RL24_VIBPA		cytoplasm
Q87T04	RS17_VIBPA		cytoplasm
Q87T05	RL29_VIBPA		cytoplasm
Q87T06	RL16_VIBPA		cytoplasm
Q87T07	RS3_VIBPA		cytoplasm
Q87T08	RL22_VIBPA		cytoplasm
Q87T09	RS19_VIBPA		cytoplasm
Q87T10	RL2_VIBPA		cytoplasm
Q87T12	RL4_VIBPA		cytoplasm
Q87T13	RL3_VIBPA		cytoplasm
Q87T15	RL31_VIBPA		cytoplasm
Q87T20	HSLV_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87T21	HSLU_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87T23	RRAA2_VIBPA		cytoplasm
Q87T31	TPIS_VIBPA		cytoplasm
Q87T56	HLDD_VIBPA		cytoplasm
Q87T74	KDKA_VIBPA	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q87T80	COAD_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87T81	FPG_VIBPA		cytoplasm
Q87T84	RL33_VIBPA		cytoplasm
Q87T85	RL28_VIBPA		cytoplasm
Q87T86	RADC_VIBPA		cytoplasm
Q87T89	COABC_VIBPA		cytoplasm
Q87T90	SLMA_VIBPA	nucleoid-associated (by similarity).	cytoplasm
Q87T92	PYRE_VIBPA		cytoplasm
Q87T93	RNPH_VIBPA		cytoplasm

Q87TA9	KGUA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87TB0	RPOZ_VIBPA		cytoplasm
Q87TB8	GREB_VIBPA		cytoplasm
Q87TC2	BIOH_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87TC4	Y146_VIBPA		inner membrane
Q87TE0	HSLO_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87TE1	PPCK_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87TE4	DTD_VIBPA	cytoplasm (probable).	cytoplasm
Q87TE5	RBN_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87TF3	ADD1_VIBPA		extracellular
Q87TF6	Y113_VIBPA		inner membrane
Q87TG0	ENGB_VIBPA		cytoplasm
Q87TH0	TATB_VIBPA	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q87TH1	TATA_VIBPA	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q87TH2	UBIB_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87TH4	UBIE_VIBPA		cytoplasm
Q87TJ1	USPB_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87TJ6	Y073_VIBPA		inner membrane
Q87TN4	ILVC_VIBPA		cytoplasm
Q87TN9	FADB_VIBPA		cytoplasm
Q87TP0	FADA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87TP4	TUSA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87TQ5	RECF_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87TQ7	DNAA_VIBPA	cytoplasm (by similarity).	cytoplasm
Q87TR4	RNPA_VIBPA		extracellular
Q87TR5	OXAA_VIBPA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87TR6	TRME_VIBPA		cytoplasm
Q87TR9	RNPA_PSESM		extracellular
Q87TS0	Y5613_PSESM		inner membrane
Q87TS1	OXAA_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87TS2	TRME_PSESM		cytoplasm
Q87TS3	GIDA_PSESM	cytoplasm (by similarity).	cytoplasm
Q87TS4	GIDB_PSESM		cytoplasm
Q87TT2	ATPA_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87TT3	ATPG_PSESM		inner membrane
Q87TT5	ATPE_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87TT6	GLMU_PSESM	cytoplasm (by similarity).	cytoplasm
Q87TT8	GLMS_PSESM	cytoplasm (by similarity).	cytoplasm

Q87TX5	OPGD_PSESM	periplasm (by similarity).	periplasm
Q87TY7	CLS_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87U31	PSTB2_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87UB2	AGUA_PSESM		extracellular
Q87UC6	Y5379_PSESM		inner membrane
Q87UC7	Y5378_PSESM		inner membrane
Q87UF5	FETP_PSESM		cytoplasm
Q87UG0	HIS7_PSESM	cytoplasm (by similarity).	cytoplasm
Q87UG1	HIS5_PSESM	cytoplasm (by similarity).	cytoplasm
Q87UG3	HIS4_PSESM	cytoplasm (by similarity).	cytoplasm
Q87UG4	HIS6_PSESM	cytoplasm (by similarity).	cytoplasm
Q87UH3	SECB_PSESM	cytoplasm (by similarity).	cytoplasm
Q87UH7	TAUB_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87UK7	RPIA_PSESM		cytoplasm
Q87UL1	NUDH_PSESM		cytoplasm
Q87UL3	LGT_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87UL4	TYSY_PSESM	cytoplasm (by similarity).	cytoplasm
Q87UL9	HUTI_PSESM	cytoplasm (potential).	cytoplasm
Q87UM1	HUTH_PSESM	cytoplasm (potential).	cytoplasm
Q87UM6	HUTU_PSESM	cytoplasm (by similarity).	cytoplasm
Q87UP6	PPK_PSESM		cytoplasm
Q87US2	Y5224_PSESM		inner membrane
Q87UX9	DTD_PSESM	cytoplasm (probable).	cytoplasm
Q87UY0	OPGG_PSESM	periplasm (by similarity).	periplasm
Q87UY1	OPGH_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87UY8	HIS2_PSESM	cytoplasm (by similarity).	cytoplasm
Q87UY9	HIS3_PSESM	cytoplasm (by similarity).	cytoplasm
Q87UZ0	UBIB_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87UZ2	UBIE_PSESM		cytoplasm
Q87V00	HSLU_PSESM	cytoplasm (by similarity).	cytoplasm
Q87V01	HSLV_PSESM	cytoplasm (by similarity).	cytoplasm
Q87V05	RL31_PSESM		cytoplasm
Q87V14	AROK_PSESM	cytoplasm (probable).	cytoplasm
Q87V15	AROB_PSESM	cytoplasm (probable).	cytoplasm
Q87V23	DCUP_PSESM	cytoplasm (probable).	cytoplasm
Q87V56	MDCC_PSESM	cytoplasm (by similarity).	cytoplasm
Q87V59	MDCG_PSESM		cytoplasm
Q87V70	Y5071_PSESM	periplasm (potential).	periplasm
Q87V73	SAHH_PSESM	cytoplasm (by similarity).	cytoplasm
Q87V83	ILVD_PSESM		cytoplasm
Q87V88	HAM1_PSESM		extracellular
Q87V90	METX_PSESM	cytoplasm (by similarity).	cytoplasm
Q87V99	PYRB_PSESM		cytoplasm
Q87VA1	RUVX_PSESM	cytoplasm (potential).	cytoplasm

Q87VA2	Y5037_PSESM		inner membrane
Q87VA4	GSHB_PSESM		cytoplasm
Q87VB1	Y5028_PSESM		inner membrane
Q87VB6	CVRA_PSESM	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q87VD5	GLNE_PSESM		cytoplasm
Q87VF3	MSBA_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87VF4	HLDE_PSESM		cytoplasm
Q87VG1	THIC_PSESM		cytoplasm
Q87VI5	ENGC_PSESM		cytoplasm
Q87VI6	ORN_PSESM	cytoplasm (potential).	cytoplasm
Q87VJ2	MUTL_PSESM		cytoplasm
Q87VJ3	MIAA_PSESM		cytoplasm
Q87VJ4	HFQ_PSESM		cytoplasm
Q87VJ8	HISZ_PSESM	cytoplasm (by similarity).	cytoplasm
Q87VJ9	PURA_PSESM	cytoplasm (by similarity).	cytoplasm
Q87VK2	RLMB_PSESM		cytoplasm
Q87VK3	RS6_PSESM		cytoplasm
Q87VK4	RS18_PSESM		cytoplasm
Q87VK6	RL9_PSESM		cytoplasm
Q87VK8	Y4928_PSESM		cytoplasm
Q87VL4	NADE_PSESM		cytoplasm
Q87VM7	UREE2_PSESM	cytoplasm (by similarity).	cytoplasm
Q87VN8	CBPA_PSESM	nucleoid (by similarity).	inner membrane
Q87VP0	URE1_PSESM	cytoplasm (by similarity).	cytoplasm
Q87VP1	URE2_PSESM	cytoplasm (by similarity).	cytoplasm
Q87VP4	URE3_PSESM	cytoplasm (by similarity).	cytoplasm
Q87VQ6	CBID_PSESM		cytoplasm
Q87VR8	PUR2_PSESM		cytoplasm
Q87VR9	PUR9_PSESM		cytoplasm
Q87VS1	DUSB_PSESM		cytoplasm
Q87VS3	PRMA_PSESM	cytoplasm (by similarity).	cytoplasm
Q87VS6	AROQ_PSESM		cytoplasm
Q87VU3	SPEA_PSESM		cytoplasm
Q87VV6	PROA_PSESM	cytoplasm (by similarity).	cytoplasm
Q87VV7	NADD_PSESM		cytoplasm
Q87VV9	Y4826_PSESM		inner membrane
Q87VW5	Y4820_PSESM		inner membrane
Q87VW6	LIPB_PSESM	cytoplasm (by similarity).	cytoplasm
Q87VW7	LIPA_PSESM	cytoplasm (potential).	cytoplasm
Q87VX7	LNT_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87VX9	Y4806_PSESM		inner membrane
Q87VY5	GSA_PSESM	cytoplasm (potential).	cytoplasm
Q87VY6	THIE_PSESM		cytoplasm
Q87VZ5	KPTA_PSESM		cytoplasm
Q87W79	NPD4_PSESM	cytoplasm (probable).	cytoplasm
Q87WA0	TRMA_PSESM		cytoplasm
Q87WB2	MSCL_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87WC1	GLYA2_PSESM	cytoplasm (by similarity).	cytoplasm

Q87WD5	ENO2_PSESM	cytoplasm (by similarity).	extracellular
Q87WD6	IPYR2_PSESM	cytoplasm (by similarity).	cytoplasm
Q87WG7	Y4583_PSESM	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q87WJ7	KDGD_PSESM		cytoplasm
Q87WN1	SSRP_PSESM	cytoplasm (potential).	cytoplasm
Q87WN4	Y4512_PSESM		inner membrane
Q87WN9	GRPE_PSESM	cytoplasm (probable).	cytoplasm
Q87WP0	DNAK_PSESM		cytoplasm
Q87WP1	DNAJ_PSESM	cytoplasm (probable).	cytoplasm
Q87WP2	DAPB_PSESM	cytoplasm (by similarity).	cytoplasm
Q87WP3	CARA_PSESM		cytoplasm
Q87WP4	CARB_PSESM		cytoplasm
Q87WP5	GREA_PSESM		cytoplasm
Q87WP7	RRMJ_PSESM	cytoplasm (by similarity).	cytoplasm
Q87WQ0	GLMM_PSESM		cytoplasm
Q87WQ1	TPIS_PSESM		cytoplasm
Q87WQ3	Y4492_PSESM		inner membrane
Q87WQ6	RBFA_PSESM	cytoplasm (potential).	cytoplasm
Q87WQ7	RS15_PSESM		cytoplasm
Q87WR8	GATB_PSESM		cytoplasm
Q87WR9	GATA_PSESM		cytoplasm
Q87WS0	GATC_PSESM		cytoplasm
Q87WS4	Y4469_PSESM	cytoplasm (potential).	inner membrane
Q87WS9	Y4464_PSESM		inner membrane
Q87WT7	Y4456_PSESM		cytoplasm
Q87WV2	MURA_PSESM	cytoplasm (probable).	inner membrane
Q87WV4	HIS1_PSESM	cytoplasm (by similarity).	cytoplasm
Q87WV5	HISX_PSESM		cytoplasm
Q87WV6	HIS8_PSESM		cytoplasm
Q87WW0	CYSD_PSESM		cytoplasm
Q87WW8	RS9_PSESM		cytoplasm
Q87WX3	Y4420_PSESM		inner membrane
Q87WX6	MRAZ_PSESM		inner membrane
Q87WX7	MRAW_PSESM		cytoplasm
Q87WY0	MURE_PSESM	cytoplasm (probable).	cytoplasm
Q87WY2	MRAY_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87WY3	MURD_PSESM	cytoplasm (by similarity).	cytoplasm
Q87WY5	MURG_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87WY6	MURC_PSESM	cytoplasm (probable).	cytoplasm
Q87WY7	DDLB_PSESM	cytoplasm (by similarity).	cytoplasm
Q87WZ1	LPXC_PSESM		cytoplasm
Q87WZ4	ARGJ_PSESM	cytoplasm (probable).	cytoplasm
Q87X00	Y4393_PSESM		inner membrane
Q87X13	CH10_PSESM	cytoplasm (by similarity).	cytoplasm
Q87X14	CH60_PSESM	cytoplasm (by similarity).	cytoplasm
Q87XE2	UNG_PSESM	cytoplasm (by similarity).	extracellular
Q87XF1	Y4227_PSESM		cytoplasm
Q87XF8	LEPA_PSESM		cytoplasm
Q87XG1	RNC_PSESM	cytoplasm (by similarity).	cytoplasm
Q87XG2	ERA_PSESM		cytoplasm

Q87XG3	RECO_PSESM		cytoplasm
Q87XG4	PDXJ_PSESM	cytoplasm (probable).	cytoplasm
Q87XJ5	RL31B_PSESM		cytoplasm
Q87XJ6	DDLA_PSESM	cytoplasm (by similarity).	cytoplasm
Q87XJ9	METE_PSESM		cytoplasm
Q87XL0	GLPK_PSESM		cytoplasm
Q87XL4	OTC_PSESM	cytoplasm (probable).	cytoplasm
Q87XM0	RNT_PSESM		cytoplasm
Q87XM1	PYRC_PSESM		cytoplasm
Q87XM3	ASSY_PSESM	cytoplasm (probable).	cytoplasm
Q87XN2	DCD_PSESM		extracellular
Q87XP9	Y4127_PSESM		inner membrane
Q87XR0	PDXH_PSESM		cytoplasm
Q87XS2	RAPA_PSESM		cytoplasm
Q87XW6	MUTS_PSESM		cytoplasm
Q87XY7	RECA_PSESM	cytoplasm (by similarity).	cytoplasm
Q87XY8	RECX_PSESM	cytoplasm (potential).	cytoplasm
Q87Y19	KUP_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87Y22	DPO4_PSESM	cytoplasm (probable).	cytoplasm
Q87Y27	SLYX_PSESM		inner membrane
Q87Y32	Y3980_PSESM		inner membrane
Q87Y33	RUVC_PSESM		cytoplasm
Q87Y34	RUVA_PSESM		cytoplasm
Q87Y35	RUVB_PSESM		cytoplasm
Q87Y40	TOLB_PSESM	periplasm (potential).	periplasm
Q87Y50	NADA_PSESM	cytoplasm (by similarity).	cytoplasm
Q87Y59	PUR7_PSESM		cytoplasm
Q87Y79	MAO1_PSESM		cytoplasm
Q87Y85	Y3918_PSESM		inner membrane
Q87YA4	RDGC_PSESM	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q87YC5	APEB_PSESM		extracellular
Q87YC6	MINE_PSESM		inner membrane
Q87YC8	MINC_PSESM		inner membrane
Q87YF5	LPXK_PSESM		cytoplasm
Q87YF7	KDSB_PSESM	cytoplasm (potential).	cytoplasm
Q87YF8	MURB_PSESM	cytoplasm (probable).	inner membrane
Q87YG2	Y3837_PSESM	cytoplasm (potential).	inner membrane
Q87YG4	RL32_PSESM		cytoplasm
Q87YG5	PLSX_PSESM		cytoplasm
Q87YH1	KTHY_PSESM		cytoplasm
Q87YI1	TRPF_PSESM		cytoplasm
Q87YK2	PPNK_PSESM	cytoplasm (by similarity).	cytoplasm
Q87YP0	Y3755_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87YP9	LPXH_PSESM	cytoplasm (by similarity).	cytoplasm
Q87YR5	TIG_PSESM		cytoplasm
Q87YR6	CLPP_PSESM	cytoplasm (by similarity).	cytoplasm
Q87YR7	CLPX_PSESM		cytoplasm
Q87YT0	RNH_PSESM	cytoplasm (potential).	extracellular
Q87YW7	1A1D_PSESM		cytoplasm
Q87YX3	ALLA_PSESM		extracellular

Q87YX4	ALLC_PSESM		extracellular
Q87YY5	ZIPA_PSESM	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q87YZ4	Y3645_PSESM		inner membrane
Q87Z00	RECR_PSESM		cytoplasm
Q87Z03	CCMA_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q87Z07	CCME_PSESM	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q87Z65	CSRA3_PSESM		cytoplasm
Q87Z72	MASZ2_PSESM	cytoplasm (by similarity).	cytoplasm
Q87Z79	HGD_PSESM		cytoplasm
Q87ZB2	FADB_PSESM		cytoplasm
Q87ZB3	FADA_PSESM	cytoplasm (by similarity).	cytoplasm
Q87ZB9	LEXA1_PSESM		cytoplasm
Q87ZC3	MTNP_PSESM		cytoplasm
Q87ZG3	SSUD_PSESM		cytoplasm
Q87ZQ4	NUOG_PSESM		cytoplasm
Q87ZR5	Y3359_PSESM		inner membrane
Q87ZR6	TRMU_PSESM	cytoplasm (by similarity).	cytoplasm
Q87ZS0	CLPS_PSESM		cytoplasm
Q87ZS3	ATE_PSESM	cytoplasm (potential).	cytoplasm
Q87ZS4	LFTR_PSESM	cytoplasm (by similarity).	cytoplasm
Q87ZS6	LOLA_PSESM	periplasm (by similarity).	periplasm
Q87ZS8	CRCB_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q87ZU3	INH_PSESM	periplasm (by similarity).	periplasm
Q880A6	PSTB1_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q880J5	Y3162_PSESM		inner membrane
Q880L3	UPPP_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q880M9	GLGA_PSESM		cytoplasm
Q880X7	UVRC_PSESM	cytoplasm (by similarity).	cytoplasm
Q880Z4	XYLA_PSESM	cytoplasm (by similarity).	cytoplasm
Q881C1	MODC_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q881F7	Y2936_PSESM		inner membrane
Q881H6	Y2917_PSESM		inner membrane
Q881T7	DNAE2_PSESM	cytoplasm (probable).	cytoplasm
Q881U0	LEXA2_PSESM		cytoplasm
Q881X0	GLGB_PSESM		extracellular
Q881Z4	Y2735_PSESM		inner membrane
Q882A9	NUDC_PSESM		cytoplasm
Q882G0	RIBAB_PSESM		cytoplasm
Q882K4	NPD3_PSESM	cytoplasm (probable).	cytoplasm
Q882K8	ARGD2_PSESM	cytoplasm (probable).	cytoplasm
Q882P7	Y2578_PSESM		inner membrane

Q882S0	PHNC2_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q882V6	MOAA_PSESM		cytoplasm
Q882Y4	GLSA_PSESM		extracellular
Q883F1	UREE1_PSESM	cytoplasm (by similarity).	cytoplasm
Q883F3	URE23_PSESM	cytoplasm (by similarity).	cytoplasm
Q883H6	IHFA_PSESM		cytoplasm
Q883K6	MOBA_PSESM	cytoplasm (by similarity).	cytoplasm
Q883P1	PYRD_PSESM	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q883Q6	RRAA_PSESM		cytoplasm
Q883Q9	Y2291_PSESM		cytoplasm
Q883T5	GREB_PSESM		cytoplasm
Q883V4	ATKC_PSESM	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q883V6	ATKA_PSESM	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q883Y4	GPDA_PSESM	cytoplasm (probable).	cytoplasm
Q883Y6	FABA_PSESM	cytoplasm (by similarity).	cytoplasm
Q883Y9	HTPG_PSESM	cytoplasm (by similarity).	cytoplasm
Q884C0	LEU3_PSESM	cytoplasm (by similarity).	cytoplasm
Q884C1	LEUD_PSESM		cytoplasm
Q884C2	LEU2_PSESM		cytoplasm
Q884C6	DUSC_PSESM		cytoplasm
Q884C9	UVRB_PSESM	cytoplasm (by similarity).	cytoplasm
Q884G7	DUSA_PSESM		cytoplasm
Q884H4	TAL_PSESM	cytoplasm (probable).	cytoplasm
Q884I1	QUEF_PSESM	cytoplasm (probable).	cytoplasm
Q884I3	LOLD_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q884I6	STHA_PSESM	cytoplasm (by similarity).	cytoplasm
Q884N3	SPEE_PSESM		cytoplasm
Q884P5	AROC_PSESM		cytoplasm
Q884Q3	GCH12_PSESM		cytoplasm
Q884R6	Y2022_PSESM		inner membrane
Q884R9	PDXB_PSESM		cytoplasm
Q884S0	TUSA_PSESM	cytoplasm (by similarity).	cytoplasm
Q884S1	MTFA_PSESM		cytoplasm
Q884X9	FLIE_PSESM		flagellar
Q884Z4	FLGI_PSESM	periplasm (by similarity).	periplasm
Q884Z5	FLGH_PSESM	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q885A6	CYNS_PSESM		cytoplasm
Q885J3	RISB2_PSESM		cytoplasm
Q885J4	ASTE_PSESM		periplasm
Q885K0	ARGD1_PSESM	cytoplasm (probable).	cytoplasm
Q885K7	ACSA_PSESM		cytoplasm
Q885L1	PHS_PSESM		cytoplasm
Q885M2	ISPZ_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q885Q1	MSRB_PSESM		cytoplasm
Q885Q3	HTPX_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q885Q4	TPMT_PSESM	cytoplasm (by similarity).	cytoplasm
Q885T0	IHFB_PSESM		cytoplasm
Q885T2	KCY_PSESM	cytoplasm (by similarity).	cytoplasm
Q885T5	SERC_PSESM	cytoplasm (by similarity).	cytoplasm
Q885T7	MTNA_PSESM		cytoplasm
Q885T9	UBIG_PSESM		cytoplasm
Q885U7	Y1734_PSESM		inner membrane
Q885V0	FUMC_PSESM	cytoplasm (by similarity).	cytoplasm
Q885W4	COBS_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q885W6	COBT_PSESM		cytoplasm
Q885W8	COBQ_PSESM		cytoplasm
Q885X7	NPD2_PSESM	cytoplasm (probable).	cytoplasm
Q885Y1	PUR5_PSESM	cytoplasm (by similarity).	cytoplasm
Q885Y8	RUMA_PSESM		cytoplasm
Q885Z8	SPRT_PSESM	cytoplasm (potential).	extracellular
Q885Z9	DCTA_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q886E0	Y1639_PSESM		cytoplasm
Q886E9	Y1630_PSESM		inner membrane
Q886F0	CSRA1_PSESM		cytoplasm
Q886F1	Y1628_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q886I1	DEF2_PSESM		cytoplasm
Q886I3	Y1596_PSESM		periplasm
Q886L4	PIMT_PSESM	cytoplasm (by similarity).	cytoplasm
Q886L5	SURE_PSESM	cytoplasm (potential).	periplasm
Q886L6	TRUD_PSESM		cytoplasm
Q886L7	ISPF_PSESM		extracellular
Q886M1	ISPD_PSESM		cytoplasm
Q886M3	ENO1_PSESM	cytoplasm (by similarity).	extracellular
Q886M4	KDSA_PSESM	cytoplasm (by similarity).	cytoplasm
Q886M5	PYRG_PSESM		cytoplasm
Q886M6	TILS_PSESM	cytoplasm (probable).	cytoplasm
Q886M7	ACCA_PSESM	cytoplasm (by similarity).	cytoplasm
Q886M9	RNH2_PSESM	cytoplasm (potential).	extracellular
Q886N0	LPXB_PSESM		cytoplasm
Q886N1	LPXA_PSESM	cytoplasm (by similarity).	cytoplasm
Q886N2	FABZ_PSESM	cytoplasm (by similarity).	cytoplasm
Q886N3	LPXD_PSESM		cytoplasm
Q886N7	DXR_PSESM		cytoplasm
Q886N9	UPPS_PSESM		inner membrane
Q886P0	RRF_PSESM	cytoplasm (by similarity).	cytoplasm
Q886P1	PYRH_PSESM		cytoplasm
Q886P3	RS2_PSESM		cytoplasm
Q886P5	GLND_PSESM		cytoplasm
Q886Q7	PLSB_PSESM	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane

Q886R2	Y1515_PSESM		inner membrane
Q886R8	KAD_PSESM	cytoplasm (by similarity).	cytoplasm
Q886R9	CAPP_PSESM		cytoplasm
Q886U9	RL19_PSESM		cytoplasm
Q886V0	TRMD_PSESM	cytoplasm (potential).	cytoplasm
Q886V1	RIMM_PSESM	cytoplasm (potential).	cytoplasm
Q886V2	RS16_PSESM		cytoplasm
Q886X5	GUAA_PSESM		cytoplasm
Q886X9	EX7L_PSESM	cytoplasm (by similarity).	cytoplasm
Q886Y1	LEU1_PSESM		cytoplasm
Q886Y6	ENGA_PSESM		cytoplasm
Q886Z0	ISPG_PSESM		cytoplasm
Q886Z4	NDK_PSESM	cytoplasm (by similarity).	cytoplasm
Q886Z7	HSCA_PSESM		cytoplasm
Q886Z8	HSCB_PSESM		inner membrane
Q887B0	TGT_PSESM		cytoplasm
Q887B5	HRCN_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q887B7	HRCQA_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	extracellular
Q887D9	AMIF_PSESM		cytoplasm
Q887H2	DSBB_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q887L4	GCSH1_PSESM		cytoplasm
Q887L5	GCSP_PSESM		cytoplasm
Q887M0	AMPA_PSESM	cytoplasm (by similarity).	cytoplasm
Q887M3	SYV_PSESM	cytoplasm (by similarity).	cytoplasm
Q887M6	NDPA_PSESM	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q887N8	RHLB_PSESM		cytoplasm
Q887P4	MOAC_PSESM		cytoplasm
Q887P7	Y1244_PSESM		inner membrane
Q887P8	ALGD_PSESM		cytoplasm
Q887P9	ALG8_PSESM	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q887Q0	ALG44_PSESM		inner membrane
Q887Q1	ALGK_PSESM	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q887Q2	ALGE_PSESM	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
Q887Q3	ALGG_PSESM	periplasm (probable).	extracellular
Q887Q4	ALGX_PSESM	periplasm (potential).	periplasm
Q887Q5	ALGL_PSESM	periplasm (potential).	periplasm
Q887Q6	ALGI_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q887Q7	ALGJ_PSESM	bacterial cell inner membrane; peripheral membrane protein; periplasmic side (by similarity).	periplasm

Q887Q8	ALGF_PSESM	periplasm (by similarity).	periplasm
Q887Q9	ALGA_PSESM		cytoplasm
Q887V1	GCH11_PSESM		cytoplasm
Q887V6	CYSZ_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q887Z4	MQO_PSESM		cytoplasm
Q888A0	UPP_PSESM		cytoplasm
Q888A2	HEMH_PSESM	cytoplasm (by similarity).	extracellular
Q888B8	MURI_PSESM		cytoplasm
Q888C2	HEM1_PSESM		cytoplasm
Q888C4	LOLB_PSESM	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q888C5	ISPE_PSESM		cytoplasm
Q888C6	KPRS_PSESM	cytoplasm (by similarity).	cytoplasm
Q888C7	RL25_PSESM		cytoplasm
Q888C8	PTH_PSESM	cytoplasm (by similarity).	cytoplasm
Q888L8	SOTB_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q888N1	Y986_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q888N2	Y985_PSESM		cytoplasm
Q888N4	ILVC_PSESM		cytoplasm
Q888P6	SFSA_PSESM		cytoplasm
Q888P9	GLUQ_PSESM		cytoplasm
Q888Q5	PANB_PSESM		cytoplasm
Q888Q6	PANC_PSESM		cytoplasm
Q888Q7	G6PI_PSESM	cytoplasm (by similarity).	cytoplasm
Q888U3	COAE_PSESM	cytoplasm (by similarity).	cytoplasm
Q888U4	Y922_PSESM		cytoplasm
Q889C2	CLPB_PSESM	cytoplasm (probable).	cytoplasm
Q889E1	ISPH_PSESM		cytoplasm
Q889E3	LSPA_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q889E4	SYI_PSESM	cytoplasm (by similarity).	cytoplasm
Q889E8	RS20_PSESM		cytoplasm
Q889F0	PROB_PSESM	cytoplasm (by similarity).	cytoplasm
Q889F2	RL27_PSESM		cytoplasm
Q889F6	SELU_PSESM		cytoplasm
Q889F7	SELD_PSESM		cytoplasm
Q889G0	PHNC1_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q889J2	ADD_PSESM		extracellular
Q889M3	EUTC_PSESM		cytoplasm
Q889M7	IPYR1_PSESM	cytoplasm (by similarity).	cytoplasm
Q889P9	EX7S_PSESM	cytoplasm (by similarity).	cytoplasm
Q889Q1	DXS_PSESM		cytoplasm
Q889Q3	GCH2_PSESM		extracellular
Q889Q5	NUSB_PSESM		cytoplasm
Q889Q6	RISB1_PSESM		cytoplasm
Q889R0	NRDR_PSESM		cytoplasm
Q889U1	SSB_PSESM		cytoplasm

Q889U6	RPOA_PSESM		cytoplasm
Q889U7	RS4_PSESM		cytoplasm
Q889U8	RS11_PSESM		cytoplasm
Q889U9	RS13_PSESM		cytoplasm
Q889V2	RL15_PSESM		cytoplasm
Q889V4	RS5_PSESM		cytoplasm
Q889V5	RL18_PSESM		cytoplasm
Q889V7	RS8_PSESM		cytoplasm
Q889V9	RL5_PSESM		cytoplasm
Q889W0	RL24_PSESM		cytoplasm
Q889W2	RS17_PSESM		cytoplasm
Q889W3	RL29_PSESM		cytoplasm
Q889W4	RL16_PSESM		cytoplasm
Q889W5	RS3_PSESM		cytoplasm
Q889W6	RL22_PSESM		cytoplasm
Q889W7	RS19_PSESM		cytoplasm
Q889W8	RL2_PSESM		cytoplasm
Q889X0	RL4_PSESM		cytoplasm
Q889X1	RL3_PSESM		cytoplasm
Q889X2	RS10_PSESM		cytoplasm
Q889X5	RS7_PSESM		cytoplasm
Q889X6	RS12_PSESM		cytoplasm
Q889X7	RPOC_PSESM		cytoplasm
Q889X8	RPOB_PSESM		cytoplasm
Q889X9	RL7_PSESM		cytoplasm
Q889Y0	RL10_PSESM		cytoplasm
Q889Y1	RL1_PSESM		cytoplasm
Q889Y2	RL11_PSESM		cytoplasm
Q889Y8	SYX_PSESM	cytoplasm (by similarity).	cytoplasm
Q889Z3	ARGC_PSESM	cytoplasm (probable).	cytoplasm
Q889Z9	SPED_PSESM		cytoplasm
Q88A03	TRPC_PSESM		cytoplasm
Q88A04	TRPD_PSESM		cytoplasm
Q88A13	NPD1_PSESM	cytoplasm (probable).	cytoplasm
Q88A30	GPH_PSESM		cytoplasm
Q88A45	PDXA_PSESM	cytoplasm (probable).	cytoplasm
Q88A46	KSGA_PSESM		cytoplasm
Q88A47	APAG_PSESM		inner membrane
Q88A48	APAH_PSESM		extracellular
Q88A49	GLPE_PSESM	cytoplasm (by similarity).	cytoplasm
Q88A51	Y546_PSESM		inner membrane
Q88A53	CCA_PSESM		cytoplasm
Q88A56	Y541_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88A58	RS21_PSESM		cytoplasm
Q88A79	PQQF_PSESM		extracellular
Q88A81	PQQB_PSESM		cytoplasm
Q88A82	PQQC_PSESM		cytoplasm
Q88A83	PQQD_PSESM		cytoplasm
Q88A84	PQQE_PSESM		cytoplasm
Q88AB2	MASZ1_PSESM	cytoplasm (by similarity).	cytoplasm
Q88AD1	GLYA1_PSESM	cytoplasm (by similarity).	cytoplasm
Q88AE7	BETA_PSESM		cytoplasm
Q88AE9	BETB_PSESM		cytoplasm
Q88AF0	BETI_PSESM		cytoplasm

Q88AF5	TRMB_PSESM		cytoplasm
Q88AF6	THIG_PSESM	cytoplasm (by similarity).	cytoplasm
Q88AF9	MTGA_PSESM	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q88AH3	COAD_PSESM	cytoplasm (by similarity).	cytoplasm
Q88AH6	FPG_PSESM		cytoplasm
Q88AI5	MSRA_PSESM		cytoplasm
Q88AK3	PGK_PSESM	cytoplasm (potential).	cytoplasm
Q88AM9	THII_PSESM	cytoplasm (potential).	cytoplasm
Q88AP4	ENGB_PSESM		cytoplasm
Q88AQ2	ALGB_PSESM		cytoplasm
Q88AR1	GSH1_PSESM		cytoplasm
Q88AR2	ARGA_PSESM	cytoplasm (probable).	cytoplasm
Q88AR9	GCSH2_PSESM		cytoplasm
Q88AS0	GCST_PSESM		cytoplasm
Q88AS5	CYSA_PSESM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88AZ4	PPCK_PSESM	cytoplasm (by similarity).	cytoplasm
Q88AZ5	HSLO_PSESM	cytoplasm (by similarity).	cytoplasm
Q88AZ9	RIMK_PSESM		cytoplasm
Q88B06	CYAY_PSESM		inner membrane
Q88B09	DAPF_PSESM	cytoplasm (by similarity).	cytoplasm
Q88B41	RSMB_PSESM		cytoplasm
Q88B42	FMT_PSESM		cytoplasm
Q88B43	DEF1_PSESM		cytoplasm
Q88B49	HEM6_PSESM	cytoplasm (by similarity).	cytoplasm
Q88B60	TRPA_PSESM		cytoplasm
Q88B61	TRPB_PSESM		cytoplasm
Q88B91	HEM3_PSESM		cytoplasm
Q88B94	ARLY_PSESM	cytoplasm (probable).	cytoplasm
Q88BA2	NORM_PSESM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88BB6	DADA_PSESM		cytoplasm
Q88BC7	RL33_PSESM		cytoplasm
Q88BC8	RL28_PSESM		cytoplasm
Q88BD1	RADC_PSESM		cytoplasm
Q88BD3	DUT_PSESM		extracellular
Q88BD4	ALGC_PSESM		cytoplasm
Q88BD5	ARGB_PSESM	cytoplasm (probable).	cytoplasm
Q88BD7	PYRE_PSESM		cytoplasm
Q88BE0	RNPH_PSESM		cytoplasm
Q88BE2	KGUA_PSESM	cytoplasm (by similarity).	cytoplasm
Q88BE3	RPOZ_PSESM		cytoplasm
Q88BK1	RECF_PSESM	cytoplasm (by similarity).	cytoplasm
Q88BK3	DNAA_PSESM	cytoplasm (by similarity).	cytoplasm
Q88BX2	ATPA_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88BX3	ATPG_PSEPK		inner membrane
Q88BX5	ATPE_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88BX6	GLMU_PSEPK	cytoplasm (by similarity).	cytoplasm

Q88BX8	GLMS_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88BY5	NPD_PSEPK	cytoplasm (probable).	cytoplasm
Q88C19	CLS_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88C57	PSTB2_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88C82	RPOZ_PSEPK		cytoplasm
Q88C87	KGUA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88C89	RNPH_PSEPK		cytoplasm
Q88C92	PYRE_PSEPK		cytoplasm
Q88C93	ALGC_PSEPK		cytoplasm
Q88C95	DUT_PSEPK		extracellular
Q88C97	RADC_PSEPK		cytoplasm
Q88C99	RL28_PSEPK		cytoplasm
Q88CA0	RL33_PSEPK		cytoplasm
Q88CB1	DADA2_PSEPK		cytoplasm
Q88CB9	NORM_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88CF1	XERC_PSEPK	cytoplasm (by similarity).	nucleoid
Q88CF3	DAPF_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88CF6	CYAY_PSEPK		inner membrane
Q88CG0	Y5221_PSEPK		inner membrane
Q88CG4	PPK_PSEPK		cytoplasm
Q88CI0	Y5201_PSEPK		inner membrane
Q88CI7	GCST_PSEPK		cytoplasm
Q88CI8	GCSH2_PSEPK		cytoplasm
Q88CI9	GCSP2_PSEPK		cytoplasm
Q88CL2	CYSA_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88CN0	RPIA_PSEPK		cytoplasm
Q88CN4	NUDH_PSEPK		cytoplasm
Q88CN8	LGT_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88CN9	TYSY_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88CQ2	ILVD_PSEPK		cytoplasm
Q88CQ5	FPG_PSEPK		cytoplasm
Q88CQ7	COAD_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88CS3	MTGA_PSEPK	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q88CS6	THIG_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88CS7	TRMB_PSEPK		cytoplasm
Q88CT0	HAM1_PSEPK		extracellular
Q88CT3	METX_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88CU3	RL31_PSEPK		cytoplasm
Q88CV1	AROK_PSEPK	cytoplasm (probable).	cytoplasm
Q88CV2	AROB_PSEPK	cytoplasm (probable).	cytoplasm
Q88CV6	DCUP_PSEPK	cytoplasm (probable).	cytoplasm
Q88CW2	Y5068_PSEPK		inner membrane

Q88CW4	CVRA_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q88CW6	BETA_PSEPK		cytoplasm
Q88CW7	BETB_PSEPK		cytoplasm
Q88CX4	GPMI_PSEPK		cytoplasm
Q88CX7	SECB_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88CY4	THII_PSEPK	cytoplasm (potential).	cytoplasm
Q88CZ6	HUTU_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88CZ7	HUTH_PSEPK	cytoplasm (potential).	cytoplasm
Q88CZ9	HUTI_PSEPK	cytoplasm (potential).	cytoplasm
Q88D02	DTD_PSEPK	cytoplasm (probable).	cytoplasm
Q88D03	OPGG_PSEPK	periplasm (by similarity).	periplasm
Q88D04	OPGH_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88D14	HIS2_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88D15	HIS3_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88D17	UBIE_PSEPK		cytoplasm
Q88D27	HSLU_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88D28	HSLV_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88D30	PYRB_PSEPK		cytoplasm
Q88D32	RUVX_PSEPK	cytoplasm (potential).	cytoplasm
Q88D33	Y4995_PSEPK		inner membrane
Q88D35	GSHB_PSEPK		cytoplasm
Q88D47	Y4981_PSEPK	periplasm (potential).	periplasm
Q88D64	PGK_PSEPK	cytoplasm (potential).	cytoplasm
Q88D92	MSBA_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88D93	HLDE_PSEPK		cytoplasm
Q88DA5	THIC_PSEPK		cytoplasm
Q88DB9	PSD_PSEPK		cytoplasm
Q88DC4	ENGC_PSEPK		cytoplasm
Q88DC5	ORN_PSEPK	cytoplasm (potential).	cytoplasm
Q88DD1	MUTL_PSEPK		cytoplasm
Q88DD2	MIAA_PSEPK		cytoplasm
Q88DD3	HFQ_PSEPK		cytoplasm
Q88DD7	HISZ_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88DD8	PURA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88DE7	RLMB_PSEPK		cytoplasm
Q88DE8	RS6_PSEPK		cytoplasm
Q88DE9	RS18_PSEPK		cytoplasm
Q88DF1	RL9_PSEPK		cytoplasm
Q88DF3	Y4872_PSEPK		cytoplasm
Q88DF6	NADE_PSEPK		cytoplasm
Q88DF7	PNCB_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88DH7	CBPA_PSEPK	nucleoid (by similarity).	inner membrane
Q88DH8	CBPM_PSEPK		cytoplasm
Q88DJ4	CBID_PSEPK		cytoplasm
Q88DK2	PUR2_PSEPK		cytoplasm
Q88DK3	PUR9_PSEPK		cytoplasm
Q88DK5	DUSB_PSEPK		cytoplasm
Q88DK7	PRMA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88DL4	PROA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88DL5	NADD_PSEPK		cytoplasm

Q88DL7	Y4808_PSEPK		inner membrane
Q88DM3	Y4802_PSEPK		inner membrane
Q88DM4	LIPB_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88DM5	LIPA_PSEPK	cytoplasm (potential).	cytoplasm
Q88DN4	LNT_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88DN6	Y4788_PSEPK		inner membrane
Q88DP0	GSA_PSEPK	cytoplasm (potential).	cytoplasm
Q88DP1	THIE_PSEPK		cytoplasm
Q88DT3	LLDD_PSEPK		cytoplasm
Q88DT6	SSRP_PSEPK	cytoplasm (potential).	cytoplasm
Q88DU1	GRPE_PSEPK	cytoplasm (probable).	cytoplasm
Q88DU2	DNAK_PSEPK		cytoplasm
Q88DU3	DNAJ_PSEPK	cytoplasm (probable).	cytoplasm
Q88DU4	DAPB_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88DU5	CARA_PSEPK		cytoplasm
Q88DU6	CARB_PSEPK		cytoplasm
Q88DU7	GREA_PSEPK		cytoplasm
Q88DV0	RRMJ_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88DV3	GLMM_PSEPK		cytoplasm
Q88DV4	TPIS_PSEPK		cytoplasm
Q88DV5	Y4714_PSEPK		inner membrane
Q88DV8	RBFA_PSEPK	cytoplasm (potential).	cytoplasm
Q88DV9	RS15_PSEPK		cytoplasm
Q88DW6	ACSA2_PSEPK		cytoplasm
Q88DW7	G6PI2_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88DW8	PANC_PSEPK		cytoplasm
Q88DW9	PANB_PSEPK		cytoplasm
Q88DX4	GLUQ_PSEPK		cytoplasm
Q88DX7	SFSA_PSEPK		cytoplasm
Q88DZ0	ILVC_PSEPK		cytoplasm
Q88DZ2	Y4676_PSEPK		cytoplasm
Q88DZ3	Y4675_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88E14	TRMA_PSEPK		cytoplasm
Q88E23	MSCL_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88E33	Y4635_PSEPK		cytoplasm
Q88E47	HGD_PSEPK		cytoplasm
Q88E69	MOAA_PSEPK		cytoplasm
Q88E79	Y4587_PSEPK		inner membrane
Q88E89	Y4577_PSEPK		inner membrane
Q88EA1	ACEK_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88EA5	Y4561_PSEPK		cytoplasm
Q88EA7	DEF2_PSEPK		cytoplasm
Q88EH2	PHS_PSEPK		cytoplasm
Q88EH6	ACSA1_PSEPK		cytoplasm
Q88EI7	ASTE_PSEPK		periplasm
Q88EM0	DADA1_PSEPK		cytoplasm
Q88ER9	FLGH_PSEPK	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q88ES0	FLGI_PSEPK	periplasm (by similarity).	periplasm
Q88ET3	FLIE_PSEPK		flagellar

Q88EV6	DDLA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88EX5	CCMA_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88EX9	CCME_PSEPK	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q88F17	AQPZ_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88F24	ZIPA_PSEPK	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q88F31	Y4268_PSEPK		inner membrane
Q88F32	RECR_PSEPK		cytoplasm
Q88F51	Y4248_PSEPK		inner membrane
Q88FB9	HTPG_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88FC4	FABA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88FC9	GPDA_PSEPK	cytoplasm (probable).	cytoplasm
Q88FD7	ATKA_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q88FD8	ATKC_PSEPK	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q88FF5	RNH_PSEPK	cytoplasm (potential).	extracellular
Q88FH2	NUOG_PSEPK		cytoplasm
Q88FJ7	UVRC_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88FN1	GLGB_PSEPK		extracellular
Q88FN9	GLGA_PSEPK		cytoplasm
Q88FQ4	RNZ_PSEPK		extracellular
Q88FR8	Y4015_PSEPK		inner membrane
Q88FR9	TRMU_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88FS4	CLPS_PSEPK		cytoplasm
Q88FS6	ATE_PSEPK	cytoplasm (potential).	cytoplasm
Q88FS7	LFTR_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88FS9	LOLA_PSEPK	periplasm (by similarity).	periplasm
Q88FT1	CRCB_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88G93	CSRA_PSEPK		cytoplasm
Q88G95	MODC_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88GQ7	Y3661_PSEPK	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q88GW8	KDGD_PSEPK		cytoplasm
Q88HA3	MOBA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88HL0	NIKE_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88HL1	NIKD_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q88HL5	NIKR_PSEPK		cytoplasm
Q88HM9	Y3324_PSEPK		inner membrane
Q88HV0	CAAL_PSEPK		cytoplasm
Q88I82	DNAE2_PSEPK	cytoplasm (probable).	cytoplasm
Q88IC7	ECOT_PSEPK	periplasm (by similarity).	periplasm
Q88IS4	MQO3_PSEPK		cytoplasm
Q88IS9	Y2920_PSEPK		inner membrane
Q88IU0	CSID_PSEPK		cytoplasm
Q88IU7	LPXH_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88IX6	Y2873_PSEPK		inner membrane
Q88IY7	UPPP_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88J03	UREE_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88J04	URE1_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88J05	URE2_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88J06	URE3_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88J85	MSUE_PSEPK		cytoplasm
Q88JG8	PQQD2_PSEPK		cytoplasm
Q88JJ0	PSTB1_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88JU7	Y2550_PSEPK		inner membrane
Q88JY1	GCH12_PSEPK		cytoplasm
Q88K24	RL20_PSEPK		cytoplasm
Q88K25	RL35_PSEPK		cytoplasm
Q88K39	ASPQ_PSEPK	periplasm (by similarity).	periplasm
Q88K63	SOTB_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88K84	AROQ2_PSEPK		cytoplasm
Q88K93	Y2398_PSEPK		inner membrane
Q88KB2	Y2378_PSEPK		inner membrane
Q88KH5	GREB_PSEPK		cytoplasm
Q88KI9	CLPX_PSEPK		cytoplasm
Q88KJ0	CLPP_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88KJ1	TIG_PSEPK		cytoplasm
Q88KX0	DUSA_PSEPK		cytoplasm
Q88KX1	TAL_PSEPK	cytoplasm (probable).	cytoplasm
Q88KX9	QUEF_PSEPK	cytoplasm (probable).	cytoplasm
Q88KY4	LOLD_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88KY8	STHA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88L01	FADA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88L02	FADB_PSEPK		cytoplasm
Q88L20	PDXB_PSEPK		cytoplasm
Q88L21	TUSA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88L23	MTFA_PSEPK		cytoplasm
Q88L40	PYRD_PSEPK	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q88L51	RRAA_PSEPK		cytoplasm
Q88L54	Y2081_PSEPK		cytoplasm
Q88L76	Y2059_PSEPK		cytoplasm
Q88LC3	PPNK_PSEPK	cytoplasm (by similarity).	cytoplasm

Q88LE0	TRPF_PSEPK		cytoplasm
Q88LE5	LEU3_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88LE7	LEUD_PSEPK		cytoplasm
Q88LE8	LEU2_PSEPK		cytoplasm
Q88LF2	DUSC_PSEPK		cytoplasm
Q88LF9	UVRB_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88LL5	ACP_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88LL8	PLSX_PSEPK		cytoplasm
Q88LL9	RL32_PSEPK		cytoplasm
Q88LM1	Y1909_PSEPK	cytoplasm (potential).	inner membrane
Q88LM5	MURB_PSEPK	cytoplasm (probable).	inner membrane
Q88LM7	KDSB_PSEPK	cytoplasm (potential).	cytoplasm
Q88LM9	LPXK_PSEPK		cytoplasm
Q88LQ6	MSRB_PSEPK		cytoplasm
Q88LQ8	HTPX_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q88LQ9	TPMT_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88LU7	AROC_PSEPK		cytoplasm
Q88LV4	GCH11_PSEPK		cytoplasm
Q88LW2	PYRF_PSEPK		cytoplasm
Q88LW9	G6PI1_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88LX0	KDSA2_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88M04	KCY_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88M07	SERC_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88M09	MTNA_PSEPK		cytoplasm
Q88M10	UBIG_PSEPK		cytoplasm
Q88M17	Y1758_PSEPK		inner membrane
Q88M20	FUMC_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88M24	Y1751_PSEPK		cytoplasm
Q88M28	Y1747_PSEPK		inner membrane
Q88M41	MINC_PSEPK		inner membrane
Q88M43	MINE_PSEPK		inner membrane
Q88M44	APEB_PSEPK		extracellular
Q88M72	RDGC_PSEPK	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q88M93	COBS_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88M95	COBT_PSEPK		cytoplasm
Q88M97	COBQ_PSEPK		cytoplasm
Q88MA9	PUR5_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88MB9	RUMA_PSEPK		cytoplasm
Q88MD4	SPRT_PSEPK	cytoplasm (potential).	extracellular
Q88ME3	RECX_PSEPK	cytoplasm (potential).	cytoplasm
Q88ME4	RECA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88ME7	MUTS_PSEPK		cytoplasm
Q88MF0	PIMT_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88MF1	SURE_PSEPK	cytoplasm (potential).	periplasm
Q88MF2	TRUD_PSEPK		cytoplasm
Q88MF3	ISPF_PSEPK		extracellular
Q88MF7	ISPD_PSEPK		cytoplasm
Q88MF9	ENO_PSEPK	cytoplasm (by similarity).	extracellular
Q88MG0	KDSA1_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88MG1	PYRG_PSEPK		cytoplasm

Q88MG3	TILS_PSEPK	cytoplasm (probable).	cytoplasm
Q88MG4	ACCA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88MG6	RNH2_PSEPK	cytoplasm (potential).	extracellular
Q88MG7	LPXB_PSEPK		cytoplasm
Q88MG8	LPXA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88MG9	FABZ_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88MH0	LPXD_PSEPK		cytoplasm
Q88MH4	DXR_PSEPK		cytoplasm
Q88MH6	UPPS_PSEPK		inner membrane
Q88MH7	RRF_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88MH8	PYRH_PSEPK		cytoplasm
Q88MI0	RS2_PSEPK		cytoplasm
Q88MI2	GLND_PSEPK		cytoplasm
Q88MQ0	PLSB_PSEPK	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q88MQ7	Y1513_PSEPK		inner membrane
Q88MR4	CAPP_PSEPK		cytoplasm
Q88MV0	XERD_PSEPK	cytoplasm (by similarity).	nucleoid
Q88MV3	RL19_PSEPK		cytoplasm
Q88MV4	TRMD_PSEPK	cytoplasm (potential).	cytoplasm
Q88MV5	RIMM_PSEPK	cytoplasm (potential).	cytoplasm
Q88MV6	RS16_PSEPK		cytoplasm
Q88MY2	PDXJ_PSEPK	cytoplasm (probable).	cytoplasm
Q88MY3	RECO_PSEPK		cytoplasm
Q88MY4	ERA_PSEPK		cytoplasm
Q88MY5	RNC_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88MY7	LEPA_PSEPK		cytoplasm
Q88MZ4	Y1424_PSEPK		nucleoid
Q88N05	UNG_PSEPK	cytoplasm (by similarity).	extracellular
Q88N29	TTGR_PSEPK		cytoplasm
Q88N30	TTGA_PSEPK	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
Q88N31	TTGB_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q88N32	TTGC_PSEPK	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
Q88N55	CH60_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88N56	CH10_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88N71	LPXC_PSEPK		cytoplasm
Q88N74	DDLB_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88N75	MURC_PSEPK	cytoplasm (probable).	cytoplasm
Q88N76	MURG_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88N78	MURD_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88N79	MRAY_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88N81	MURE_PSEPK	cytoplasm (probable).	cytoplasm
Q88N84	MRAW_PSEPK		cytoplasm
Q88N85	MRAZ_PSEPK		inner membrane
Q88N88	Y1324_PSEPK		inner membrane
Q88N96	RS9_PSEPK		cytoplasm
Q88NA9	CYSD_PSEPK		cytoplasm

Q88NB7	RHLB_PSEPK		cytoplasm
Q88NC0	MOAC_PSEPK		cytoplasm
Q88NC3	Y1289_PSEPK		inner membrane
Q88NC4	ALGD_PSEPK		cytoplasm
Q88NC5	ALG8_PSEPK	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q88NC6	ALG44_PSEPK		inner membrane
Q88NC7	ALGK_PSEPK	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q88NC8	ALGE_PSEPK	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
Q88NC9	ALGG_PSEPK	periplasm (probable).	extracellular
Q88ND0	ALGX_PSEPK	periplasm (potential).	periplasm
Q88ND1	ALGL_PSEPK	periplasm (potential).	periplasm
Q88ND2	ALGI_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88ND3	ALGJ_PSEPK	bacterial cell inner membrane; peripheral membrane protein; periplasmic side (by similarity).	periplasm
Q88ND4	ALGF_PSEPK	periplasm (by similarity).	periplasm
Q88ND5	ALGA_PSEPK		cytoplasm
Q88NF9	MQO2_PSEPK		cytoplasm
Q88NG9	PUR7_PSEPK		cytoplasm
Q88NJ0	RUVB_PSEPK		cytoplasm
Q88NJ1	RUVA_PSEPK		cytoplasm
Q88NJ2	RUVC_PSEPK		cytoplasm
Q88NJ3	Y1214_PSEPK		inner membrane
Q88NJ9	SLYX_PSEPK		inner membrane
Q88NK4	DPO4_PSEPK	cytoplasm (probable).	cytoplasm
Q88NK7	KUP_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88NL9	DCTA_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88NR0	RAPA_PSEPK		cytoplasm
Q88NS5	PDXH_PSEPK		cytoplasm
Q88NT5	Y1119_PSEPK		inner membrane
Q88NV4	DCD_PSEPK		extracellular
Q88NW7	PYRC_PSEPK		cytoplasm
Q88NW8	RNT_PSEPK		cytoplasm
Q88NX4	OTC_PSEPK	cytoplasm (probable).	cytoplasm
Q88NX8	GLPK_PSEPK		cytoplasm
Q88P21	GUAA_PSEPK		cytoplasm
Q88P26	EX7L_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88P28	LEU1_PSEPK		cytoplasm
Q88P52	ARCA_PSEPK	cytoplasm (potential).	extracellular
Q88P53	OTCC_PSEPK	cytoplasm (probable).	cytoplasm
Q88P61	Y992_PSEPK		inner membrane
Q88P64	GCSH1_PSEPK		cytoplasm
Q88P65	GCSP1_PSEPK		cytoplasm
Q88P73	AMPA_PSEPK	cytoplasm (by similarity).	cytoplasm

Q88P76	SYV_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88P80	NDPA_PSEPK	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q88P81	Y972_PSEPK		cytoplasm
Q88P86	HIS8_PSEPK		cytoplasm
Q88P87	HIS1_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88P88	MURA_PSEPK	cytoplasm (probable).	inner membrane
Q88PA1	Y949_PSEPK		cytoplasm
Q88PA9	Y941_PSEPK		inner membrane
Q88PB4	Y936_PSEPK	cytoplasm (potential).	inner membrane
Q88PB8	GATC_PSEPK		cytoplasm
Q88PB9	GATA_PSEPK		cytoplasm
Q88PC0	GATB_PSEPK		cytoplasm
Q88PD5	SODF_PSEPK		periplasm
Q88PI8	Y862_PSEPK		cytoplasm
Q88PJ3	ENGA_PSEPK		cytoplasm
Q88PJ7	ISPG_PSEPK		cytoplasm
Q88PK1	NDK_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88PK4	HSCA_PSEPK		cytoplasm
Q88PK5	HSCB_PSEPK		inner membrane
Q88PL7	TGT_PSEPK		cytoplasm
Q88PL8	QUEA_PSEPK	cytoplasm (potential).	cytoplasm
Q88PM5	PHNC_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88PM7	SELU_PSEPK		cytoplasm
Q88PP0	HMP_PSEPK		cytoplasm
Q88PU7	MQO1_PSEPK		cytoplasm
Q88PV2	UPP_PSEPK		cytoplasm
Q88PV4	HEMH_PSEPK	cytoplasm (by similarity).	extracellular
Q88PW2	MURI_PSEPK		cytoplasm
Q88PW6	HEM1_PSEPK		cytoplasm
Q88PX4	LOLB_PSEPK	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q88PX5	ISPE_PSEPK		cytoplasm
Q88PX6	KPRS_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88PX7	RL25_PSEPK		cytoplasm
Q88PX8	PTH_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88Q07	PROB_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88Q09	RL27_PSEPK		cytoplasm
Q88Q16	Y682_PSEPK	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q88Q27	GLYA2_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88Q44	MDH_PSEPK		cytoplasm
Q88Q65	COAE_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88Q66	Y630_PSEPK		cytoplasm
Q88Q71	CLPB_PSEPK	cytoplasm (probable).	cytoplasm
Q88Q89	ISPH_PSEPK		cytoplasm
Q88Q91	LSPA_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q88Q92	SYI_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88Q95	RS20_PSEPK		cytoplasm
Q88QA3	ADD_PSEPK		extracellular

Q88QC7	SPEA_PSEPK		cytoplasm
Q88QD4	AROQ1_PSEPK		cytoplasm
Q88QF2	EUTC_PSEPK		cytoplasm
Q88QF6	IPYR_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88QG4	RIBB_PSEPK		cytoplasm
Q88QG5	EX7S_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88QG7	DXS_PSEPK		cytoplasm
Q88QH1	GCH2_PSEPK		extracellular
Q88QH5	NUSB_PSEPK		cytoplasm
Q88QH6	RISB_PSEPK		cytoplasm
Q88QI0	NRDR_PSEPK		cytoplasm
Q88QJ8	SELA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88QJ9	FDHE_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88QK5	SSB_PSEPK		cytoplasm
Q88QK7	UVRA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88QL1	RPOA_PSEPK		cytoplasm
Q88QL2	RS4_PSEPK		cytoplasm
Q88QL3	RS13_PSEPK		cytoplasm
Q88QL6	RL15_PSEPK		cytoplasm
Q88QL8	RS5_PSEPK		cytoplasm
Q88QL9	RL18_PSEPK		cytoplasm
Q88QM1	RS8_PSEPK		cytoplasm
Q88QM3	RL5_PSEPK		cytoplasm
Q88QM4	RL24_PSEPK		cytoplasm
Q88QM6	RS17_PSEPK		cytoplasm
Q88QM7	RL29_PSEPK		cytoplasm
Q88QM8	RL16_PSEPK		cytoplasm
Q88QM9	RS3_PSEPK		cytoplasm
Q88QN0	RL22_PSEPK		cytoplasm
Q88QN1	RS19_PSEPK		cytoplasm
Q88QN2	RL2_PSEPK		cytoplasm
Q88QN4	RL4_PSEPK		cytoplasm
Q88QN5	RL3_PSEPK		cytoplasm
Q88QN6	RS10_PSEPK		cytoplasm
Q88QN9	RS7_PSEPK		cytoplasm
Q88QP0	RS12_PSEPK		cytoplasm
Q88QP1	RPOC_PSEPK		cytoplasm
Q88QP2	RPOB_PSEPK		cytoplasm
Q88QP3	RL10_PSEPK		cytoplasm
Q88QP4	RL1_PSEPK		cytoplasm
Q88QP5	RL11_PSEPK		cytoplasm
Q88QQ2	SYI_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88QQ6	ARGC1_PSEPK	cytoplasm (probable).	cytoplasm
Q88QR6	TRPC_PSEPK		cytoplasm
Q88QR7	TRPD_PSEPK		cytoplasm
Q88QS2	GPH_PSEPK		cytoplasm
Q88QT5	PDXA_PSEPK	cytoplasm (probable).	cytoplasm
Q88QT6	KSGA_PSEPK		cytoplasm
Q88QT7	APAG_PSEPK		inner membrane
Q88QT8	APAH_PSEPK		extracellular
Q88QT9	GLPE_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88QU2	CCA_PSEPK		cytoplasm
Q88QU5	Y391_PSEPK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q88QV3	PQQF_PSEPK		extracellular

Q88QV5	PQQB_PSEPK		cytoplasm
Q88QV6	PQQC_PSEPK		cytoplasm
Q88QV7	PQQD1_PSEPK		cytoplasm
Q88QV8	PQQE_PSEPK		cytoplasm
Q88QX8	MASZ_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88QZ4	GLNE_PSEPK		cytoplasm
Q88QZ8	MSRA_PSEPK		cytoplasm
Q88R12	GLYA1_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88R41	HIS6_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88R42	HIS4_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88R44	HIS5_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88R45	HIS7_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88R49	FETP_PSEPK		cytoplasm
Q88R68	AGUA_PSEPK		extracellular
Q88R81	HSLO_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88R85	RIMK_PSEPK		cytoplasm
Q88R90	GSH1_PSEPK		cytoplasm
Q88R95	SSUD_PSEPK		cytoplasm
Q88R97	SSUE_PSEPK		cytoplasm
Q88RA1	TAUB_PSEPK	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q88RE5	HEM3_PSEPK		cytoplasm
Q88RJ6	ALGB_PSEPK		cytoplasm
Q88RK5	ENGB_PSEPK		cytoplasm
Q88RP6	TRPB_PSEPK		cytoplasm
Q88RP7	TRPA_PSEPK		cytoplasm
Q88RQ6	HEM6_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88RR1	DEF1_PSEPK		cytoplasm
Q88RR2	FMT_PSEPK		cytoplasm
Q88RR3	RSMB_PSEPK		cytoplasm
Q88RW7	RECF_PSEPK	cytoplasm (by similarity).	cytoplasm
Q88RW8	GIDA_PSEPK	cytoplasm (by similarity).	cytoplasm
Q89A13	Y550_BUCBP		inner membrane
Q89A14	ENGA_BUCBP		cytoplasm
Q89A15	FER_BUCBP		cytoplasm
Q89A16	HSCA_BUCBP		cytoplasm
Q89A17	HSCB_BUCBP		inner membrane
Q89A18	NIFU_BUCBP		cytoplasm
Q89A19	ISCS_BUCBP		cytoplasm
Q89A20	ILVC_BUCBP		cytoplasm
Q89A21	REP_BUCBP		cytoplasm
Q89A22	RHO_BUCBP		inner membrane
Q89A23	Y532_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89A24	PIT_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89A25	USPA_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89A26	Y529_BUCBP		cytoplasm
Q89A27	SYK3_BUCBP		cytoplasm
Q89A28	FENR_BUCBP		cytoplasm
Q89A29	TRPE_BUCBP		cytoplasm
Q89A30	TRPG_BUCBP		cytoplasm
Q89A31	HSLU_BUCBP	cytoplasm (by similarity).	cytoplasm

Q89A32	RL31_BUCBP		cytoplasm
Q89A33	AMIB_BUCBP		extracellular
Q89A34	ORN_BUCBP	cytoplasm (potential).	cytoplasm
Q89A35	G6PI_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89A36	PTM3C_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89A37	MTLD_BUCBP		cytoplasm
Q89A38	MUTL_BUCBP		cytoplasm
Q89A39	HFLK_BUCBP		inner membrane
Q89A40	HFLC_BUCBP		extracellular
Q89A41	RS6_BUCBP		cytoplasm
Q89A42	RL9_BUCBP		cytoplasm
Q89A43	EX1_BUCBP		cytoplasm
Q89A44	FETP_BUCBP		cytoplasm
Q89A45	MUTY_BUCBP		cytoplasm
Q89A46	TRMB_BUCBP		cytoplasm
Q89A47	Y498_BUCBP		cytoplasm
Q89A48	Y497_BUCBP		cytoplasm
Q89A49	LEU1_BUCBP		cytoplasm
Q89A50	RUVX_BUCBP	cytoplasm (potential).	cytoplasm
Q89A51	Y491_BUCBP		inner membrane
Q89A52	DNAB_BUCBP		nucleoid
Q89A53	SSB_BUCBP		cytoplasm
Q89A54	BIOH_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89A55	Y486_BUCBP		inner membrane
Q89A57	DEOB_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89A58	DEOD_BUCBP		cytoplasm
Q89A59	RPE_BUCBP		cytoplasm
Q89A60	TSGA_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89A61	FKBA_BUCBP		cytoplasm
Q89A62	TUSD_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89A63	TUSC_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89A64	TUSB_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89A65	RS12_BUCBP		cytoplasm
Q89A66	RS7_BUCBP		cytoplasm
Q89A67	RS10_BUCBP		cytoplasm
Q89A68	RL3_BUCBP		cytoplasm
Q89A69	RL4_BUCBP		cytoplasm
Q89A70	RL23_BUCBP		cytoplasm
Q89A71	RL2_BUCBP		cytoplasm
Q89A72	RS19_BUCBP		cytoplasm
Q89A73	RL16_BUCBP		cytoplasm
Q89A74	RL29_BUCBP		cytoplasm
Q89A75	RS17_BUCBP		cytoplasm
Q89A76	RL14_BUCBP		cytoplasm
Q89A77	RL24_BUCBP		cytoplasm
Q89A78	RL5_BUCBP		cytoplasm
Q89A79	RS14_BUCBP		cytoplasm
Q89A80	RL6_BUCBP		cytoplasm
Q89A81	RL18_BUCBP		cytoplasm
Q89A82	RS5_BUCBP		cytoplasm
Q89A83	RL30_BUCBP		cytoplasm
Q89A84	RL15_BUCBP		cytoplasm

Q89A85	SECY_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89A87	RS13_BUCBP		cytoplasm
Q89A88	RL17_BUCBP		cytoplasm
Q89A89	Y438_BUCBP		inner membrane
Q89A90	CSPE_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89A91	Y432_BUCBP		inner membrane
Q89A92	FOLD_BUCBP		cytoplasm
Q89A93	HTPG_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89A94	Y426_BUCBP		inner membrane
Q89A95	DPO3X_BUCBP		cytoplasm
Q89A96	MDLB_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89A97	MDLA_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89A98	PPID_BUCBP	bacterial cell membrane; single-pass type ii membrane protein (potential).	outer membrane
Q89AA0	CLPX_BUCBP		cytoplasm
Q89AA1	CLPP_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AA2	Y418_BUCBP		cytoplasm
Q89AA3	CYOA_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AA4	CYOB_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AA5	CYOC_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AA6	CYOD_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AA7	CYOE_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AA8	TUSE_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AA9	Y411_BUCBP	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q89AB0	RIBD_BUCBP		cytoplasm
Q89AB1	RISB_BUCBP		cytoplasm
Q89AB2	EX5A_BUCBP		cytoplasm
Q89AB3	EX5B_BUCBP		cytoplasm
Q89AB4	EX5C_BUCBP		cytoplasm
Q89AB5	Y402_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AB6	ALF_BUCBP		cytoplasm
Q89AB7	Y399_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AB8	DHAS_BUCBP		cytoplasm

Q89AB9	TUSA_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AC0	HOLA_BUCBP		cytoplasm
Q89AC1	CORC_BUCBP		inner membrane
Q89AC2	Y392_BUCBP		cytoplasm
Q89AC3	Y391_BUCBP		cytoplasm
Q89AC6	DCDA_BUCBP		cytoplasm
Q89AC7	LGT_BUCBP	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q89AC8	KGUA_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AC9	TYPB_BUCBP		cytoplasm
Q89AD0	ENGB_BUCBP		cytoplasm
Q89AD1	EX53_BUCBP		cytoplasm
Q89AD2	DSBA_BUCBP		cytoplasm
Q89AD3	MUTS_BUCBP		cytoplasm
Q89AD5	RPIA_BUCBP		cytoplasm
Q89AD6	Y370_BUCBP		cytoplasm
Q89AD7	END1_BUCBP		extracellular
Q89AD8	GSH1_BUCBP		cytoplasm
Q89AD9	RLUD_BUCBP		cytoplasm
Q89AE0	AROQ_BUCBP		cytoplasm
Q89AE1	TLDD_BUCBP		cytoplasm
Q89AE2	RL19_BUCBP		cytoplasm
Q89AE3	RS16_BUCBP		cytoplasm
Q89AE4	SRP54_BUCBP		cytoplasm
Q89AE6	RL13_BUCBP		cytoplasm
Q89AE7	Y352_BUCBP		cytoplasm
Q89AE8	RL21_BUCBP		cytoplasm
Q89AE9	MURA_BUCBP	cytoplasm (probable).	inner membrane
Q89AF0	Y348_BUCBP		cytoplasm
Q89AF1	RRMJ_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AF2	FTSH_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AF3	GLMM_BUCBP		cytoplasm
Q89AF4	NUSA_BUCBP		cytoplasm
Q89AF6	TRUB_BUCBP		cytoplasm
Q89AF7	RS15_BUCBP		cytoplasm
Q89AF8	PNP_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AF9	DEAD_BUCBP	cytoplasm (probable).	cytoplasm
Q89AG0	Y334_BUCBP		cytoplasm
Q89AG1	OTC_BUCBP	cytoplasm (probable).	cytoplasm
Q89AG2	AMPA_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AG3	SYV_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AG4	Y330_BUCBP		inner membrane
Q89AG5	YHIT_BUCBP		cytoplasm
Q89AG6	PTGCB_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AG7	Y325_BUCBP		extracellular
Q89AG8	HOLB_BUCBP		cytoplasm
Q89AG9	FABG_BUCBP		cytoplasm
Q89AH0	FABD_BUCBP		cytoplasm
Q89AH1	RL32_BUCBP		cytoplasm
Q89AH2	RLUC_BUCBP		cytoplasm
Q89AH3	RNE_BUCBP	cytoplasm (by similarity).	cytoplasm

Q89AH4	FLGJ_BUCBP		inner membrane
Q89AH5	FLGI_BUCBP		periplasm
Q89AH6	FLGH_BUCBP	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q89AH7	FLGG_BUCBP		flagellar
Q89AH8	FLGF_BUCBP		inner membrane
Q89AH9	FLGC_BUCBP		flagellar
Q89AI0	FLGB_BUCBP		flagellar
Q89AI1	MVIN_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AI2	RSMC_BUCBP		cytoplasm
Q89AI3	MIND_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q89AI4	MINE_BUCBP		inner membrane
Q89AI5	Y301_BUCBP		cytoplasm
Q89AI6	Y300_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AI7	G6PD_BUCBP		cytoplasm
Q89AI8	KPYK_BUCBP		cytoplasm
Q89AI9	Y296_BUCBP		extracellular
Q89AJ0	ZNUC_BUCBP		inner membrane
Q89AJ1	ZNUB_BUCBP	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q89AJ3	RS1_BUCBP		cytoplasm
Q89AJ4	GPMA_BUCBP		cytoplasm
Q89AJ5	PAL_BUCBP	bacterial cell membrane; lipid-anchor (by similarity).	outer membrane
Q89AJ6	ODO2_BUCBP		inner membrane
Q89AJ7	ODO1_BUCBP		cytoplasm
Q89AJ8	Y279_BUCBP		cytoplasm
Q89AJ9	PHR_BUCBP		cytoplasm
Q89AK0	FLAV_BUCBP		cytoplasm
Q89AK2	MFD_BUCBP		cytoplasm
Q89AK3	6PGL_BUCBP		cytoplasm
Q89AK4	BIOA_BUCBP		cytoplasm
Q89AK5	BIOB_BUCBP		cytoplasm
Q89AK6	BIOF_BUCBP		cytoplasm
Q89AK7	BIOC_BUCBP		cytoplasm
Q89AK8	Y265_BUCBP		cytoplasm
Q89AK9	SUHB_BUCBP		extracellular
Q89AL0	SOHB_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	extracellular
Q89AL1	RLUB_BUCBP		cytoplasm
Q89AL2	Y256_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AL3	ISPZ_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AL4	Y254_BUCBP		cytoplasm

Q89AL5	CLS_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AL6	PYRF_BUCBP		cytoplasm
Q89AL7	LIPA_BUCBP	cytoplasm (potential).	cytoplasm
Q89AL8	LIPB_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AL9	Y248_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AM0	RNB_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AM1	FABI_BUCBP		cytoplasm
Q89AM2	MLTE_BUCBP		extracellular
Q89AM3	PUR8_BUCBP		cytoplasm
Q89AM4	Y243_BUCBP		inner membrane
Q89AM5	LEPA_BUCBP		cytoplasm
Q89AM6	LEP_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AM7	ERA_BUCBP		cytoplasm
Q89AM8	Y236_BUCBP		cytoplasm
Q89AM9	SSRP_BUCBP	cytoplasm (potential).	cytoplasm
Q89AN0	Y234_BUCBP		inner membrane
Q89AN1	GRPE1_BUCBP	cytoplasm (probable).	cytoplasm
Q89AN2	XGPT_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AN3	DPO3E_BUCBP		cytoplasm
Q89AN4	GLO2_BUCBP		cytoplasm
Q89AN5	FLHA_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AN6	FLHB_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AN8	DPO3A_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AN9	FABZ_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AP0	UPPS_BUCBP		inner membrane
Q89AP1	RRF_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AP2	RS2_BUCBP		cytoplasm
Q89AP3	AMPM_BUCBP		cytoplasm
Q89AP4	DAPD_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AP5	DEGP_BUCBP		cytoplasm
Q89AP6	APBE_BUCBP		cytoplasm
Q89AP7	ILVI_BUCBP		cytoplasm
Q89AP8	ILVH_BUCBP		cytoplasm
Q89AP9	FTSL_BUCBP		inner membrane
Q89AQ0	FTSI_BUCBP	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q89AQ1	MURF_BUCBP	cytoplasm (probable).	cytoplasm
Q89AQ2	MURD_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AQ3	FTSW_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AQ4	FTSA_BUCBP		inner membrane
Q89AQ6	Y193_BUCBP		inner membrane
Q89AQ7	MTNN_BUCBP		cytoplasm
Q89AQ8	DLDH_BUCBP	cytoplasm (by similarity).	cytoplasm

Q89AQ9	ODP2_BUCBP		inner membrane
Q89AR0	ODP1_BUCBP		cytoplasm
Q89AR1	SECA_BUCBP	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	inner membrane
Q89AR2	PBPB_BUCBP		inner membrane
Q89AR3	DKSA_BUCBP		cytoplasm
Q89AR4	AKH_BUCBP		cytoplasm
Q89AR5	THRC_BUCBP		cytoplasm
Q89AR6	ENGD_BUCBP		cytoplasm
Q89AR7	SODM_BUCBP		periplasm
Q89AR8	Y176_BUCBP		cytoplasm
Q89AR9	PPNK_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AS0	GRPE2_BUCBP	cytoplasm (probable).	cytoplasm
Q89AS1	TSAA_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AS2	Y170_BUCBP		extracellular
Q89AS3	GYRA_BUCBP		cytoplasm
Q89AS4	RIR1_BUCBP		cytoplasm
Q89AS5	RIR2_BUCBP		cytoplasm
Q89AS6	Y166_BUCBP		cytoplasm
Q89AS7	PTA_BUCBP	cytoplasm (potential).	inner membrane
Q89AS8	ACKA_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AS9	Y163_BUCBP		extracellular
Q89AT0	HEMK_BUCBP		cytoplasm
Q89AT1	CVPA_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AT2	FOLC_BUCBP		cytoplasm
Q89AT3	Y156_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AT5	NUOM_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AT6	NUOL_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AT7	NUOK_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AT8	NUOJ_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AT9	NUOI_BUCBP		cytoplasm
Q89AU1	NUOG_BUCBP		cytoplasm
Q89AU2	NUOF_BUCBP		periplasm
Q89AU3	NUOE_BUCBP		cytoplasm
Q89AU4	NUOCD_BUCBP		inner membrane
Q89AU5	NUOB_BUCBP		cytoplasm
Q89AU7	DNAJ_BUCBP	cytoplasm (probable).	cytoplasm
Q89AU8	RS20_BUCBP		cytoplasm
Q89AU9	SYI_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AV0	LSPA_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	extracellular

Q89AV1	ISPH_BUCBP		cytoplasm
Q89AV2	DYR_BUCBP		cytoplasm
Q89AV3	APAH_BUCBP		extracellular
Q89AV4	Y130_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AV5	RL25_BUCBP		cytoplasm
Q89AV7	Y125_BUCBP		inner membrane
Q89AV8	RL20_BUCBP		cytoplasm
Q89AV9	RL35_BUCBP		cytoplasm
Q89AW0	AROH_BUCBP		cytoplasm
Q89AW1	Y117_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AW2	Y116_BUCBP		inner membrane
Q89AW3	SYI_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AW4	END3_BUCBP		cytoplasm
Q89AW5	RNFE_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AW6	RNFG_BUCBP		inner membrane
Q89AW7	RNFD_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AW8	RNFC_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q89AX0	RNFA_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AX1	RISA_BUCBP		cytoplasm
Q89AX2	NORM_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89AX3	TILS_BUCBP	cytoplasm (probable).	cytoplasm
Q89AX5	6PGD_BUCBP		cytoplasm
Q89AX6	HIS2_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AX7	HIS8_BUCBP		cytoplasm
Q89AX8	Y092_BUCBP		inner membrane
Q89AX9	AROC_BUCBP		cytoplasm
Q89AY0	DAPA_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AY1	DAPE_BUCBP		cytoplasm
Q89AY2	TKT_BUCBP		extracellular
Q89AY3	TAL_BUCBP	cytoplasm (probable).	cytoplasm
Q89AY4	FABB_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89AY5	Y085_BUCBP		cytoplasm
Q89AY6	PMBA_BUCBP	cytoplasm (potential).	cytoplasm
Q89AY7	Y081_BUCBP		extracellular
Q89AY8	RL28_BUCBP		cytoplasm
Q89AY9	RL33_BUCBP		cytoplasm
Q89AZ0	FLIR_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AZ1	FLIQ_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q89AZ2	FLIP_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89AZ3	FLIN_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q89AZ4	FLIM_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q89AZ5	FLIK_BUCBP		inner membrane
Q89AZ6	FLIJ_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q89AZ7	FLII_BUCBP	cytoplasm (potential).	inner membrane
Q89AZ8	FLIH_BUCBP		inner membrane
Q89AZ9	FLIG_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q89B00	FLIF_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89B01	FLIE_BUCBP		flagellar
Q89B02	DNLJ_BUCBP		cytoplasm
Q89B03	PTHP_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89B05	PTGA_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89B06	CCA_BUCBP		cytoplasm
Q89B07	GCP_BUCBP		extracellular
Q89B08	RS21_BUCBP		cytoplasm
Q89B09	PRIM_BUCBP		cytoplasm
Q89B10	RPOD_BUCBP		cytoplasm
Q89B11	CYSE_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89B12	Y050_BUCBP		cytoplasm
Q89B13	METF_BUCBP		cytoplasm
Q89B14	SECE_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89B15	NUSG_BUCBP		cytoplasm
Q89B16	RL11_BUCBP		cytoplasm
Q89B17	RL1_BUCBP		cytoplasm
Q89B18	RL10_BUCBP		cytoplasm
Q89B19	RL7_BUCBP		cytoplasm
Q89B20	RPOB_BUCBP		cytoplasm
Q89B21	RPOC_BUCBP		cytoplasm
Q89B22	DBH_BUCBP		cytoplasm
Q89B23	PUR9_BUCBP		cytoplasm
Q89B24	METE_BUCBP		cytoplasm
Q89B25	Y030_BUCBP		cytoplasm
Q89B26	GLMU_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89B27	RP32_BUCBP		cytoplasm
Q89B28	FTSY_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q89B29	Y025_BUCBP		cytoplasm
Q89B30	DNAC_BUCBP		nucleoid
Q89B32	Y022_BUCBP		cytoplasm
Q89B33	MUTH_BUCBP	cytoplasm (by similarity).	extracellular

Q89B34	OXAA_BUCBP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89B36	DPO3B_BUCBP	cytoplasm (by similarity).	cytoplasm
Q89B37	GYRB_BUCBP		cytoplasm
Q89B38	ATPE_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q89B39	ATPB_BUCBP		inner membrane
Q89B40	ATPG_BUCBP		inner membrane
Q89B41	ATPA_BUCBP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q89B42	ATPD_BUCBP		inner membrane
Q89B43	ATPF_BUCBP		inner membrane
Q89B44	ATPL_BUCBP		inner membrane
Q89B45	ATP6_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89B46	REPA1_BUCBP		nucleoid
Q89B47	REPA2_BUCBP		nucleoid
Q89B48	YREP_BUCBP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89BJ8	LEPA_BRAJA		cytoplasm
Q89BK7	PPCK_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89BL4	PYRE_BRAJA		cytoplasm
Q89BN3	YUD3_BRAJA		inner membrane
Q89BN4	RECR_BRAJA		cytoplasm
Q89BN9	DEF_BRAJA		cytoplasm
Q89BP0	FMT_BRAJA		cytoplasm
Q89BP1	TRUA_BRAJA		cytoplasm
Q89BP4	DAPD_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89BP7	ARGB_BRAJA	cytoplasm (probable).	cytoplasm
Q89BQ0	OXAA_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89BU4	OPGG_BRAJA	periplasm (by similarity).	periplasm
Q89BU5	OPGH_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89C25	Y7973_BRAJA		cytoplasm
Q89C51	PHNC_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89CF5	Y7842_BRAJA		inner membrane
Q89CK8	HTPG_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89CN4	ADC1_BRAJA		cytoplasm
Q89D74	WRBA_BRAJA		cytoplasm
Q89DA6	CH607_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89DB4	HIS1_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89DC5	LPXK_BRAJA		cytoplasm
Q89DD2	EX7L_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89DF8	SYI_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89DI3	Y7456_BRAJA		cytoplasm
Q89DI7	RISB2_BRAJA		cytoplasm

Q89DJ1	KPRS_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89DJ5	LGT_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89DJ8	RL25_BRAJA		cytoplasm
Q89DJ9	PTH_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89DN1	GLMM_BRAJA		cytoplasm
Q89DR8	CARA_BRAJA		cytoplasm
Q89DV4	Y7333_BRAJA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89DX7	ARCA_BRAJA	cytoplasm (potential).	extracellular
Q89E33	Y7254_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89E48	UPP_BRAJA		cytoplasm
Q89EA6	NPD2_BRAJA	cytoplasm (probable).	cytoplasm
Q89ED5	TOLB_BRAJA	periplasm (potential).	periplasm
Q89EF0	Y7135_BRAJA		inner membrane
Q89EG9	AQPZ_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89EM1	PURA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89EM9	MSRB_BRAJA		cytoplasm
Q89EN5	NAPA_BRAJA	periplasm (by similarity).	periplasm
Q89EP4	ADC2_BRAJA		cytoplasm
Q89ER2	SSUD_BRAJA		cytoplasm
Q89ES2	FLIG_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89EW7	MODC2_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89F27	FLIE2_BRAJA		flagellar
Q89F30	FLG12_BRAJA	periplasm (by similarity).	periplasm
Q89F32	FLGH2_BRAJA	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q89F47	FLBT2_BRAJA		cytoplasm
Q89FC2	ATKA_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q89FC4	ATKC_BRAJA	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q89FD3	GLGB_BRAJA		extracellular
Q89FG1	PQQE_BRAJA		cytoplasm
Q89FG2	PQQD_BRAJA		cytoplasm
Q89FG3	PQQC_BRAJA		cytoplasm
Q89FG4	PQQB_BRAJA		cytoplasm
Q89FP0	THIC_BRAJA		cytoplasm
Q89FP2	THIG_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89FT9	MRAW_BRAJA		cytoplasm
Q89FU2	MURE_BRAJA	cytoplasm (probable).	cytoplasm
Q89FU4	MRAY_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q89FU5	MURD_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89FU7	MURG_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89FU8	MURC_BRAJA	cytoplasm (probable).	cytoplasm
Q89FU9	MURB_BRAJA	cytoplasm (probable).	inner membrane
Q89FW1	PRMA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89FY1	ASPD_BRAJA		cytoplasm
Q89G35	TYSY_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89G43	MIAA_BRAJA		cytoplasm
Q89G50	ILVC_BRAJA		cytoplasm
Q89G86	GLGA2_BRAJA		cytoplasm
Q89G87	GLGC_BRAJA		cytoplasm
Q89GA7	DDLA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89GB0	LEU1_BRAJA		cytoplasm
Q89GF9	AROQ2_BRAJA		cytoplasm
Q89GV3	HUTH_BRAJA	cytoplasm (potential).	cytoplasm
Q89GV4	HUTU_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89GX0	HIS81_BRAJA		cytoplasm
Q89GX9	GFA_BRAJA		extracellular
Q89H51	AMIF_BRAJA		cytoplasm
Q89H53	ADEC2_BRAJA		cytoplasm
Q89HA2	ILVD3_BRAJA		cytoplasm
Q89HP6	SAHH_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89HR3	HLDE_BRAJA		cytoplasm
Q89HW8	KDGD_BRAJA		cytoplasm
Q89HZ2	FLBT1_BRAJA		cytoplasm
Q89I01	FLG11_BRAJA	periplasm (by similarity).	periplasm
Q89I09	FLGH1_BRAJA	bacterial cell outer membrane (probable).	inner membrane
Q89I27	FLIE1_BRAJA		flagellar
Q89I84	RECA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89I86	GCSP_BRAJA		cytoplasm
Q89I87	GCSH_BRAJA		cytoplasm
Q89IB0	PUR71_BRAJA		cytoplasm
Q89IB6	PURQ_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89IC0	PURL_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89ID3	RS4_BRAJA		cytoplasm
Q89IK8	CH606_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89IM4	XGPT_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89IV9	Y5525_BRAJA		inner membrane
Q89IW2	GCH1_BRAJA		cytoplasm
Q89IW3	HIS3_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89J69	RL11_BRAJA		cytoplasm
Q89J70	RL1_BRAJA		cytoplasm
Q89J72	RL10_BRAJA		cytoplasm
Q89J74	RPOB_BRAJA		cytoplasm
Q89J75	RPOC_BRAJA		cytoplasm
Q89J79	RS12_BRAJA		cytoplasm
Q89J80	RS7_BRAJA		cytoplasm
Q89J83	RS10_BRAJA		cytoplasm
Q89J84	RL3_BRAJA		cytoplasm
Q89J85	RL4_BRAJA		cytoplasm
Q89J87	RL2_BRAJA		cytoplasm
Q89J88	RS19_BRAJA		cytoplasm
Q89J89	RL22_BRAJA		cytoplasm

Q89J90	RS3_BRAJA		cytoplasm
Q89J91	RL16_BRAJA		cytoplasm
Q89J93	RS17_BRAJA		cytoplasm
Q89J95	RL24_BRAJA		cytoplasm
Q89J96	RL5_BRAJA		cytoplasm
Q89J98	RS8_BRAJA		cytoplasm
Q89JA0	RL18_BRAJA		cytoplasm
Q89JA1	RS5_BRAJA		cytoplasm
Q89JA3	RL15_BRAJA		cytoplasm
Q89JA6	RS13_BRAJA		cytoplasm
Q89JA7	RPOA_BRAJA		cytoplasm
Q89JM6	LIPB_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89JW5	CLPS1_BRAJA		cytoplasm
Q89K13	ACPD_BRAJA		extracellular
Q89K17	Y5101_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89K19	PYRB_BRAJA		cytoplasm
Q89K24	RUVX_BRAJA	cytoplasm (potential).	cytoplasm
Q89K28	GATC_BRAJA		cytoplasm
Q89K29	GATA_BRAJA		cytoplasm
Q89K31	GATB_BRAJA		cytoplasm
Q89K46	MSCL_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89K67	MNTH_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q89K72	ATE_BRAJA	cytoplasm (potential).	cytoplasm
Q89K77	NRDR_BRAJA		cytoplasm
Q89K80	RISB1_BRAJA		cytoplasm
Q89K81	NUSB_BRAJA		cytoplasm
Q89K83	HPPA_BRAJA	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q89K88	PLSX_BRAJA		cytoplasm
Q89K89	FABH_BRAJA	cytoplasm (probable).	cytoplasm
Q89KE5	RS9_BRAJA		cytoplasm
Q89KG0	TIG_BRAJA		cytoplasm
Q89KG1	CLPP2_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89KG2	CLPX_BRAJA		cytoplasm
Q89KN0	LOLD_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89KP6	RRF_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89KP7	UPPS2_BRAJA		inner membrane
Q89KP9	DXR_BRAJA		cytoplasm
Q89KQ2	LPXD_BRAJA		cytoplasm
Q89KQ3	FABZ_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89KQ7	LPXB_BRAJA		cytoplasm
Q89KS7	LEXA_BRAJA		cytoplasm
Q89KU3	TPIS_BRAJA		cytoplasm
Q89KV0	KDSA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89KV2	GLSA2_BRAJA		extracellular
Q89KV4	QUEF_BRAJA	cytoplasm (probable).	cytoplasm
Q89KV6	ENO_BRAJA	cytoplasm (by similarity).	extracellular

Q89KY5	ILVD2_BRAJA		cytoplasm
Q89KZ0	DGTL1_BRAJA		cytoplasm
Q89L02	SURE_BRAJA	cytoplasm (potential).	periplasm
Q89L21	GID_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89L46	UVRA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89L50	SSB_BRAJA		cytoplasm
Q89L53	KUP3_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89L55	COAD_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89L59	QUEA_BRAJA	cytoplasm (potential).	cytoplasm
Q89LB1	CH605_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89LD7	GLMU_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89LG2	ARGD2_BRAJA	cytoplasm (probable).	cytoplasm
Q89LH6	Y4568_BRAJA		inner membrane
Q89LK8	ILVD1_BRAJA		cytoplasm
Q89LM5	KTHY_BRAJA		cytoplasm
Q89LQ8	ISPDF_BRAJA		cytoplasm
Q89LR6	LIPA1_BRAJA	cytoplasm (potential).	cytoplasm
Q89LR7	3MGH_BRAJA		cytoplasm
Q89LR8	SYV_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89LY4	NPD1_BRAJA	cytoplasm (probable).	cytoplasm
Q89M63	SYU_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89MA1	AROQ1_BRAJA		cytoplasm
Q89MA7	LFTR_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89MB9	NRDJ_BRAJA		cytoplasm
Q89MI2	MINC_BRAJA		inner membrane
Q89MQ0	Y4142_BRAJA		inner membrane
Q89MS3	NDK_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89MT4	AMPA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89MT9	PDXA2_BRAJA	cytoplasm (probable).	cytoplasm
Q89MU0	KSGA_BRAJA		cytoplasm
Q89MV4	KGUA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89MV8	ACP_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89MW3	RS6_BRAJA		cytoplasm
Q89MW4	RS18_BRAJA		cytoplasm
Q89MW6	RL9_BRAJA		cytoplasm
Q89MZ0	ENGA_BRAJA		cytoplasm
Q89MZ4	PANB2_BRAJA		cytoplasm
Q89N37	DEFL_BRAJA		cytoplasm
Q89N53	GUAU_BRAJA		cytoplasm
Q89N72	RRMJ_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89N76	Y3966_BRAJA		inner membrane
Q89NA7	GLSA1_BRAJA		extracellular
Q89ND7	Y3905_BRAJA		cytoplasm
Q89NF3	PDXA1_BRAJA	cytoplasm (probable).	cytoplasm
Q89NF8	RNZ_BRAJA		extracellular
Q89NN5	KUP2_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89NN6	KUP1_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89NP2	PSD_BRAJA		cytoplasm
Q89NS0	CAAL_BRAJA		cytoplasm
Q89NS9	RPIA_BRAJA		cytoplasm

Q89NW6	LIPA2_BRAJA	cytoplasm (potential).	cytoplasm
Q89P00	CH604_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89P49	Y3634_BRAJA		inner membrane
Q89PK4	Y3476_BRAJA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89PK5	Y3475_BRAJA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89PK9	Y3471_BRAJA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89PS5	DDLB_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89Q54	COBT_BRAJA		cytoplasm
Q89Q71	COBQ_BRAJA		cytoplasm
Q89Q75	COBS_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89QD9	Y3189_BRAJA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89QG3	ADEC1_BRAJA		cytoplasm
Q89QH2	FCTA_BRAJA		cytoplasm
Q89QJ5	GLNE_BRAJA		cytoplasm
Q89QK7	TYPH_BRAJA		cytoplasm
Q89QM1	Y3103_BRAJA		inner membrane
Q89QU8	DNAE2_BRAJA	cytoplasm (probable).	cytoplasm
Q89QW7	ISPH2_BRAJA		cytoplasm
Q89QX6	EUTC_BRAJA		cytoplasm
Q89R17	CAPP_BRAJA		cytoplasm
Q89R51	Y2921_BRAJA		inner membrane
Q89RB7	ARGD3_BRAJA	cytoplasm (probable).	cytoplasm
Q89RJ4	GLGA1_BRAJA		cytoplasm
Q89RW0	EX7S_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89RW1	DXS_BRAJA		cytoplasm
Q89RW4	Y2648_BRAJA		cytoplasm
Q89RX3	CRCB2_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89RX4	CRCB1_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89RX6	CLPS2_BRAJA		cytoplasm
Q89RY1	AROC_BRAJA		cytoplasm
Q89RY8	PDXH_BRAJA		cytoplasm
Q89S79	ISPE_BRAJA		cytoplasm
Q89S84	RNH2_BRAJA	cytoplasm (potential).	extracellular
Q89S87	PHK_BRAJA		cytoplasm
Q89SC2	HEM6_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89SJ4	Y2407_BRAJA		cytoplasm
Q89SJ7	Y2403_BRAJA		inner membrane
Q89SJ8	SLYX_BRAJA		inner membrane
Q89SK1	DCUP_BRAJA	cytoplasm (probable).	cytoplasm
Q89T28	DADA_BRAJA		cytoplasm
Q89T99	Y2140_BRAJA		cytoplasm
Q89TD1	ECTC_BRAJA		cytoplasm

Q89TQ9	MODC1_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89U80	RUVB_BRAJA		cytoplasm
Q89U81	RUVA_BRAJA		cytoplasm
Q89U82	RUVC_BRAJA		cytoplasm
Q89U83	Y1534_BRAJA		inner membrane
Q89U95	PGK_BRAJA	cytoplasm (potential).	cytoplasm
Q89UB0	RL31_BRAJA		cytoplasm
Q89UC1	Y1496_BRAJA		cytoplasm
Q89UC5	Y1492_BRAJA		inner membrane
Q89UD2	CYSA_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89UE3	MASZ_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89UE4	Y1473_BRAJA		cytoplasm
Q89UG0	URE1_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89UG2	URE2_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89UG3	URE3_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89UK6	GLPK_BRAJA		cytoplasm
Q89UL2	CLPB_BRAJA	cytoplasm (probable).	cytoplasm
Q89UL7	METX_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89UL9	HIS82_BRAJA		cytoplasm
Q89UU3	RNH_BRAJA	cytoplasm (potential).	extracellular
Q89UU4	KHSE_BRAJA		cytoplasm
Q89UU5	ISPH1_BRAJA		cytoplasm
Q89UZ9	Y1260_BRAJA		cytoplasm
Q89V00	Y1259_BRAJA	cytoplasm (potential).	inner membrane
Q89V02	Y1257_BRAJA		inner membrane
Q89VC7	XYLA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89VE6	APAG_BRAJA		inner membrane
Q89VE8	OTC_BRAJA	cytoplasm (probable).	cytoplasm
Q89VE9	ARGD1_BRAJA	cytoplasm (probable).	cytoplasm
Q89VF2	PSTB_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89VS2	AMIE_BRAJA		cytoplasm
Q89VU7	RL32_BRAJA		cytoplasm
Q89VU8	MTGA_BRAJA	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q89VV9	ISPG_BRAJA		cytoplasm
Q89VX1	MUTS_BRAJA		cytoplasm
Q89VX9	GLND_BRAJA		cytoplasm
Q89W41	UPPS1_BRAJA		inner membrane
Q89W61	RS20_BRAJA		cytoplasm
Q89W63	DNAA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89W71	MURA_BRAJA	cytoplasm (probable).	inner membrane
Q89W72	NORM_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89W81	PUR72_BRAJA		cytoplasm
Q89W99	Y793_BRAJA		inner membrane
Q89WA1	LNT_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q89WA4	TRMB_BRAJA		cytoplasm
Q89WA6	Y786_BRAJA		inner membrane
Q89WB0	RBFA_BRAJA	cytoplasm (potential).	cytoplasm
Q89WB1	TRUB_BRAJA		cytoplasm
Q89WB2	RS15_BRAJA		cytoplasm
Q89WC3	FABA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WC9	FPG_BRAJA		cytoplasm
Q89WD0	UBIE_BRAJA		cytoplasm
Q89WE4	TRPA_BRAJA		cytoplasm
Q89WE5	TRPB_BRAJA		cytoplasm
Q89WF1	KCY_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WF2	AROA_BRAJA	cytoplasm (probable).	cytoplasm
Q89WH1	UPPP_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q89WH9	RL35_BRAJA		cytoplasm
Q89WI0	RL20_BRAJA		cytoplasm
Q89WI8	NIKR_BRAJA		cytoplasm
Q89WK1	GPMA_BRAJA		cytoplasm
Q89WK2	DAPB_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WK4	RNPH_BRAJA		cytoplasm
Q89WK5	HAM1_BRAJA		extracellular
Q89WK9	Y669_BRAJA		inner membrane
Q89WL0	GSHB_BRAJA		cytoplasm
Q89WM2	COAA_BRAJA	cytoplasm (probable).	cytoplasm
Q89WM3	HIS21_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WM4	HIS6_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WM5	HIS4_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WM6	HIS5_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WM8	HIS7_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WM9	HSLV_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WN1	Y647_BRAJA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q89WN2	HSLU_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WN7	SECB_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WN9	COAE_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WP0	Y638_BRAJA	cytoplasm (potential).	inner membrane
Q89WP1	Y637_BRAJA		cytoplasm
Q89WP5	GIDA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WP6	GIDB_BRAJA		cytoplasm
Q89WR7	CLPP1_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WU7	PUR9_BRAJA		cytoplasm
Q89WV5	ACSA_BRAJA		cytoplasm
Q89WW0	GPDA_BRAJA	cytoplasm (probable).	cytoplasm
Q89WY0	IPYR_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89WZ6	PHS_BRAJA		cytoplasm
Q89X19	LEU3_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89X27	LEUD_BRAJA		cytoplasm
Q89X34	LEU22_BRAJA		cytoplasm
Q89X35	RL19_BRAJA		cytoplasm
Q89X36	TRMD_BRAJA	cytoplasm (potential).	cytoplasm
Q89X39	RIMM_BRAJA	cytoplasm (potential).	cytoplasm
Q89X40	RS16_BRAJA		cytoplasm
Q89X45	DAPF_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89X59	MDH_BRAJA		cytoplasm

Q89X68	XERC_BRAJA	cytoplasm (by similarity).	nucleoid
Q89X72	ATPA_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89X73	ATPG_BRAJA		inner membrane
Q89X75	ATPE_BRAJA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q89X78	NUDH_BRAJA		cytoplasm
Q89X82	Y432_BRAJA		inner membrane
Q89X84	NADD_BRAJA		cytoplasm
Q89X85	PROA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89X86	PROB_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89X93	RL27_BRAJA		cytoplasm
Q89X98	LEU21_BRAJA		cytoplasm
Q89XH1	HGD_BRAJA		cytoplasm
Q89XJ6	NOSZ_BRAJA	periplasm (by similarity).	periplasm
Q89XM2	FUMC2_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89XM4	MQO_BRAJA		cytoplasm
Q89XN9	MDCC_BRAJA	cytoplasm (by similarity).	cytoplasm
Q89XP2	MDCG_BRAJA		cytoplasm
Q89XR6	1A1D_BRAJA		cytoplasm
Q89XU2	UBIG_BRAJA		cytoplasm
Q89XW5	XERD_BRAJA	cytoplasm (by similarity).	nucleoid
Q89XW7	AROK_BRAJA	cytoplasm (probable).	cytoplasm
Q89XW8	AROB_BRAJA	cytoplasm (probable).	cytoplasm
Q89XZ3	RL28_BRAJA		cytoplasm
Q8CVG4	NANC_ECOL6	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8CVG7	C562_ECOL6	periplasm (by similarity).	periplasm
Q8CVH8	MDTP_ECOL6	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
Q8CVI4	LAMB_ECOL6	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8CVI9	ZRAP_ECOL6	periplasm (by similarity).	periplasm
Q8CVJ0	BTUB_ECOL6	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8CVL1	MDTE_ECOL6	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
Q8CVS2	PTRA_ECOL6	periplasm (by similarity).	extracellular
Q8CVU7	FADL_ECOL6	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8CVV4	MEPA_ECOL6	periplasm (by similarity).	extracellular
Q8CVW3	ECOT_ECOL6	periplasm (by similarity).	periplasm
Q8CVW4	NAPA_ECOL6	periplasm (by similarity).	periplasm
Q8CVX8	MDTA_ECOL6	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8CVZ3	TORZ_ECOL6	periplasm (by similarity).	periplasm
Q8CVZ5	YEBF_ECOL6	secreted protein (by similarity).	inner membrane
Q8CW10	YDHL_ECOL6		cytoplasm

Q8CW15	ASR_ECOL6		inner membrane
Q8CW16	DMSD_ECOL6	under anaerobic conditions: bacterial cell inner membrane; peripheral membrane protein (by similarity). under aerobic conditions; cytoplasm. mainly (by similarity).	cytoplasm
Q8CW29	YDDW_ECOL6	bacterial cell membrane; lipid-anchor (potential).	cytoplasm
Q8CW34	OPGD_ECOL6	periplasm (by similarity).	periplasm
Q8CW45	LOLB_ECOL6	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q8CW46	TREA_ECOL6	periplasm (by similarity).	periplasm
Q8CW54	FLGI_ECOL6	periplasm (by similarity).	periplasm
Q8CW55	FLGH_ECOL6	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8CW58	YCEI_ECOL6	periplasm (potential).	periplasm
Q8CW65	YCDX_ECOL6		cytoplasm
Q8CW73	TORA_ECOL6	periplasm (by similarity).	periplasm
Q8CWA2	CUSB_ECOL6		inner membrane
Q8CWA3	CUSF_ECOL6	periplasm (by similarity).	periplasm
Q8CWA4	CUSC_ECOL6	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
Q8CWD2	BTUF_ECOL6	periplasm (by similarity).	periplasm
Q8CWE6	OSTA_ECOL6	bacterial cell outer membrane (by similarity).	outer membrane
Q8CWI5	DCUP_WIGBR	cytoplasm (probable).	cytoplasm
Q8CWI9	CH102_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8CWJ0	CH602_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8CWJ5	LUXO_VIBVU		cytoplasm
Q8CWJ6	TRMU_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8CWK1	METE_VIBVU		cytoplasm
Q8CWK3	TRUA_VIBVU		cytoplasm
Q8CWK5	TRMD_VIBVU	cytoplasm (potential).	cytoplasm
Q8CWK6	CYSH_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8CWK7	DUSA_VIBVU		cytoplasm
Q8CWK9	MIAA_VIBVU		cytoplasm
Q8CWL1	CH101_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8CWL2	DUSB_VIBVU		cytoplasm
Q8CWL6	CCA_VIBVU		cytoplasm
Q8CWL9	DGTL1_VIBVU		cytoplasm
Q8CX33	GLMS_SHEON	cytoplasm (by similarity).	cytoplasm
Q8CX35	MURG_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8CX40	TRMU_SHEON	cytoplasm (by similarity).	cytoplasm
Q8CX42	HIS6_SHEON	cytoplasm (by similarity).	cytoplasm
Q8CX44	TRMD_SHEON	cytoplasm (potential).	cytoplasm
Q8CX46	NADD_SHEON		cytoplasm
Q8CX48	CH60_SHEON	cytoplasm (by similarity).	cytoplasm
Q8CX49	CH10_SHEON	cytoplasm (by similarity).	cytoplasm
Q8CX50	MIAA_SHEON		cytoplasm
Q8CX51	TRMA_SHEON		cytoplasm
Q8CX52	TRME_SHEON		cytoplasm
Q8CXV3	MIAA_ECOL6		cytoplasm
Q8CXW3	TRMA_ECOL6		cytoplasm

Q8CXW5	RHAS_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8CXX5	MALT_ECOL6		cytoplasm
Q8CXX6	CCA_ECOL6		cytoplasm
Q8CXZ8	TRMU_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8CXZ9	PHOP_ECOL6	cytoplasm (probable).	cytoplasm
Q8CY19	GLND_ECOL6		cytoplasm
Q8CY30	GLMS_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8CY35	TRMD_BRUSU	cytoplasm (potential).	cytoplasm
Q8CY36	NADD_BRUSU		cytoplasm
Q8CY38	TRMU_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8CY39	MURG_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8CY40	MIAA_BRUSU		cytoplasm
Q8CZS4	Y674_YERPE		cytoplasm
Q8D074	Y1514_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D0D7	ARGM_YERPE		cytoplasm
Q8D0M6	SUFS_YERPE	cytoplasm (by similarity).	cytoplasm
Q8D0P1	CHEY_YERPE	cytoplasm (by similarity).	cytoplasm
Q8D0U3	PDXB_YERPE		cytoplasm
Q8D0W8	CYSA_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D124	LNT_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D170	Y3224_YERPE		cytoplasm
Q8D1E4	SUGE_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D1G8	ACSA_YERPE		cytoplasm
Q8D1H3	RPOC_YERPE		cytoplasm
Q8D1I3	UBIE_YERPE		cytoplasm
Q8D1K0	XERC_YERPE	cytoplasm (by similarity).	nucleoid
Q8D1Q8	KDSD_YERPE		cytoplasm
Q8D1U5	NRDI_WIGBR		inner membrane
Q8D1U8	KUP_WIGBR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D1V0	GUAA_WIGBR		cytoplasm
Q8D1V2	MQO_WIGBR		cytoplasm
Q8D1V8	G6PI_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D1W6	PYRI_WIGBR		cytoplasm
Q8D1X1	BIOH_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D1X2	HEM6_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D1X3	TAL_WIGBR	cytoplasm (probable).	cytoplasm
Q8D1X8	AROK_WIGBR	cytoplasm (probable).	cytoplasm
Q8D1X9	PPCK_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D1Y3	ISPG_WIGBR		cytoplasm
Q8D1Y6	NDK_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D1Y8	RPOA_WIGBR		cytoplasm
Q8D1Y9	RS4_WIGBR		cytoplasm
Q8D1Z0	RS13_WIGBR		cytoplasm
Q8D1Z5	RS5_WIGBR		cytoplasm

Q8D1Z6	RL18_WIGBR		cytoplasm
Q8D1Z8	RS8_WIGBR		cytoplasm
Q8D200	RL5_WIGBR		cytoplasm
Q8D201	RL24_WIGBR		cytoplasm
Q8D203	RS17_WIGBR		cytoplasm
Q8D204	RL29_WIGBR		cytoplasm
Q8D205	RL16_WIGBR		cytoplasm
Q8D206	RS3_WIGBR		cytoplasm
Q8D207	RL22_WIGBR		cytoplasm
Q8D208	RS19_WIGBR		cytoplasm
Q8D209	RL2_WIGBR		cytoplasm
Q8D211	RL4_WIGBR		cytoplasm
Q8D212	RL3_WIGBR		cytoplasm
Q8D213	RS10_WIGBR		cytoplasm
Q8D216	GPDA_WIGBR	cytoplasm (probable).	cytoplasm
Q8D218	DCD_WIGBR		extracellular
Q8D223	ISPD_WIGBR		cytoplasm
Q8D224	ISPF_WIGBR		extracellular
Q8D226	HEMH_WIGBR	cytoplasm (by similarity).	extracellular
Q8D227	KAD_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D228	HTPG_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D229	YGG0_WIGBR		inner membrane
Q8D232	RPOC_WIGBR		cytoplasm
Q8D233	RPOB_WIGBR		cytoplasm
Q8D234	RL7_WIGBR		cytoplasm
Q8D235	RL10_WIGBR		cytoplasm
Q8D236	RL1_WIGBR		cytoplasm
Q8D237	RL11_WIGBR		cytoplasm
Q8D243	MURB_WIGBR	cytoplasm (probable).	inner membrane
Q8D244	PUR9_WIGBR		cytoplasm
Q8D245	PUR2_WIGBR		cytoplasm
Q8D247	THIC_WIGBR		cytoplasm
Q8D249	THIG_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D252	RPIA_WIGBR		cytoplasm
Q8D253	GLYA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D254	SSB_WIGBR		cytoplasm
Q8D258	DEF_WIGBR		cytoplasm
Q8D259	FMT_WIGBR		cytoplasm
Q8D264	LOLA_WIGBR	periplasm (by similarity).	periplasm
Q8D268	SERC_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D274	IPYR_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D278	RL27_WIGBR		cytoplasm
Q8D281	UPPP_WIGBR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D284	RS21_WIGBR		cytoplasm
Q8D291	RISB_WIGBR		cytoplasm
Q8D293	PYRB_WIGBR		cytoplasm
Q8D294	SYV_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D295	AMPA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D298	BIOD_WIGBR		cytoplasm
Q8D2A5	PANB_WIGBR		cytoplasm
Q8D2A7	Y447_WIGBR		inner membrane
Q8D2A8	LNT_WIGBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q8D2B6	Y438_WIGBR		cytoplasm
Q8D2C5	ORN_WIGBR	cytoplasm (potential).	cytoplasm
Q8D2C6	PSD_WIGBR		cytoplasm
Q8D2E2	TOLB_WIGBR	periplasm (potential).	periplasm
Q8D2E8	KGUA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2E9	RPOZ_WIGBR		cytoplasm
Q8D2F0	RL33_WIGBR		cytoplasm
Q8D2F1	RL28_WIGBR		cytoplasm
Q8D2F3	DUT_WIGBR		extracellular
Q8D2F5	TILS_WIGBR	cytoplasm (probable).	cytoplasm
Q8D2G0	DAPD_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2G2	RS2_WIGBR		cytoplasm
Q8D2G5	RRF_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2G6	DXR_WIGBR		cytoplasm
Q8D2G7	UPPS_WIGBR		inner membrane
Q8D2H2	LPXD_WIGBR		cytoplasm
Q8D2H3	LPXA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2H4	LPXB_WIGBR		cytoplasm
Q8D2H6	ACCA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2H8	ALR_WIGBR		cytoplasm
Q8D2H9	MINC_WIGBR		inner membrane
Q8D2I1	MINE_WIGBR		inner membrane
Q8D2I8	CLS_WIGBR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D2I9	Y365_WIGBR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D2J0	GCH2_WIGBR		extracellular
Q8D2J1	PYRF_WIGBR		cytoplasm
Q8D2J7	SUFS_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2J8	SUFE_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2K0	PYRG_WIGBR		cytoplasm
Q8D2K1	ENO_WIGBR	cytoplasm (by similarity).	extracellular
Q8D2K4	PTH_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2K6	ISPE_WIGBR		cytoplasm
Q8D2K8	HEM1_WIGBR		cytoplasm
Q8D2L3	FUMC_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2L6	SY Y_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2L7	PDXH_WIGBR		cytoplasm
Q8D2L8	SLYA_WIGBR		cytoplasm
Q8D2L9	RNT_WIGBR		cytoplasm
Q8D2M3	DAPA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2M4	PUR7_WIGBR		cytoplasm
Q8D2M5	MURA_WIGBR	cytoplasm (probable).	inner membrane
Q8D2N4	TYSY_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2N5	LGT_WIGBR	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8D2N7	GCH1_WIGBR		cytoplasm
Q8D2N9	PGSA_WIGBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D2P5	TRUA_WIGBR		cytoplasm
Q8D2P6	PDXB_WIGBR		cytoplasm
Q8D2P7	SURE_WIGBR	cytoplasm (potential).	periplasm

Q8D2P9	PGK_WIGBR	cytoplasm (potential).	cytoplasm
Q8D2Q3	Y301_WIGBR		cytoplasm
Q8D2Q4	FABA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2Q5	DNAK_WIGBR		cytoplasm
Q8D2Q6	DNAJ_WIGBR	cytoplasm (probable).	cytoplasm
Q8D2Q8	RS20_WIGBR		cytoplasm
Q8D2R0	SYI_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2R1	LSPA_WIGBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8D2R2	ISPH_WIGBR		cytoplasm
Q8D2R5	COAD_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2S5	RL31_WIGBR		cytoplasm
Q8D2S6	HSLV_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2S7	HSLU_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2T0	TPIS_WIGBR		cytoplasm
Q8D2T1	CORA_WIGBR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D2T4	DAPF_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2U2	RS6_WIGBR		cytoplasm
Q8D2U3	RS18_WIGBR		cytoplasm
Q8D2U4	RL9_WIGBR		cytoplasm
Q8D2U8	MSBA_WIGBR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D2U9	LPXK_WIGBR		cytoplasm
Q8D2V2	RL19_WIGBR		cytoplasm
Q8D2V3	RS16_WIGBR		cytoplasm
Q8D2V5	GSH1_WIGBR		periplasm
Q8D2V8	LPXH_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2W4	CCA_WIGBR		cytoplasm
Q8D2W5	RIBB_WIGBR		cytoplasm
Q8D2W7	RECA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2W9	GREA_WIGBR		cytoplasm
Q8D2X0	RRMJ_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2X3	GLMM_WIGBR		cytoplasm
Q8D2X7	RBFA_WIGBR	cytoplasm (potential).	cytoplasm
Q8D2X8	TRUB_WIGBR		cytoplasm
Q8D2X9	RS15_WIGBR		cytoplasm
Q8D2Y4	GPMA_WIGBR		cytoplasm
Q8D2Y5	DDL_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2Y8	MRAW_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2Z1	MURE_WIGBR	cytoplasm (probable).	cytoplasm
Q8D2Z3	MRAY_WIGBR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D2Z4	MURD_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D2Z6	MURG_WIGBR	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D2Z7	MURC_WIGBR	cytoplasm (probable).	cytoplasm
Q8D303	ACPS_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D304	PDXJ_WIGBR	cytoplasm (probable).	cytoplasm
Q8D305	RNC_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D307	LEPA_WIGBR		cytoplasm

Q8D308	COAE_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D317	HFQ_WIGBR		cytoplasm
Q8D318	MIAA_WIGBR		cytoplasm
Q8D320	ACKA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D322	PURA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D325	LIPA_WIGBR	cytoplasm (potential).	cytoplasm
Q8D326	LIPB_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D330	NADD_WIGBR		cytoplasm
Q8D335	GSHB_WIGBR		cytoplasm
Q8D336	Y165_WIGBR		inner membrane
Q8D337	RUVX_WIGBR	cytoplasm (potential).	cytoplasm
Q8D338	ENGB_WIGBR		cytoplasm
Q8D339	ARNT_WIGBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D341	ARNA_WIGBR		cytoplasm
Q8D342	ARNC_WIGBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D343	NADA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D346	CLPP_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D347	CLPX_WIGBR		cytoplasm
Q8D357	DXS_WIGBR		cytoplasm
Q8D362	RS9_WIGBR		cytoplasm
Q8D382	UBIE_WIGBR		cytoplasm
Q8D387	RUVC_WIGBR		cytoplasm
Q8D391	PPNK_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D392	GRPE_WIGBR	cytoplasm (probable).	cytoplasm
Q8D396	HTPX_WIGBR	bacterial cell membrane; multi-pass membrane protein (by similarity).	extracellular
Q8D397	TRMU_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D3A0	LOLD_WIGBR	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D3A7	KTHY_WIGBR		cytoplasm
Q8D3A9	ACP_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D3B2	RL32_WIGBR		cytoplasm
Q8D3B7	RL20_WIGBR		cytoplasm
Q8D3B8	RL35_WIGBR		cytoplasm
Q8D3C3	HAM1_WIGBR		extracellular
Q8D3C5	FETP_WIGBR		cytoplasm
Q8D3D3	RNH_WIGBR	cytoplasm (potential).	extracellular
Q8D3D5	OPGH_WIGBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D3D6	Y065_WIGBR		inner membrane
Q8D3F6	FLGI_WIGBR		periplasm
Q8D3F7	FLGH_WIGBR	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8D3H1	FLHD_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D3H3	RS7_WIGBR		cytoplasm
Q8D3H4	RS12_WIGBR		cytoplasm
Q8D3H6	DAPB_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D3H7	CARA_WIGBR		cytoplasm
Q8D3I0	APAH_WIGBR		extracellular

Q8D3I1	KSGA_WIGBR		cytoplasm
Q8D3I8	OXAA_WIGBR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D3I9	TRME_WIGBR		cytoplasm
Q8D3J0	GLMS_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D3J1	GLMU_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D3J2	ATPE_WIGBR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D3J4	ATPG_WIGBR		inner membrane
Q8D3J5	ATPA_WIGBR	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D3K0	GIDA_WIGBR	cytoplasm (by similarity).	cytoplasm
Q8D3K2	BETA_VIBVU		cytoplasm
Q8D3K3	BETB_VIBVU		cytoplasm
Q8D3K4	BETI_VIBVU		cytoplasm
Q8D3Q3	SPEE_VIBVU		cytoplasm
Q8D3T7	NADE_VIBVU		cytoplasm
Q8D3T9	PYRC_VIBVU		cytoplasm
Q8D3U7	MALG_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D3U8	MALF_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D3V0	MALK_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D3X4	PSTB2_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D3Z2	DEOD2_VIBVU		cytoplasm
Q8D3Z9	CBIO_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D442	TDH_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D452	CATA_VIBVU	periplasm (probable).	periplasm
Q8D456	Y5469_VIBVU		cytoplasm
Q8D461	YBG4_VIBVU		inner membrane
Q8D477	Y5434_VIBVU		inner membrane
Q8D485	RIBB_VIBVU		cytoplasm
Q8D4B4	RUMB_VIBVU		cytoplasm
Q8D4F7	NORW_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D4F8	NORV_VIBVU	cytoplasm (probable).	cytoplasm
Q8D4F9	NORR_VIBVU		cytoplasm
Q8D4N5	PCP_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D4P0	GLGB_VIBVU		extracellular
Q8D4P3	MALT_VIBVU		cytoplasm
Q8D4Q0	ENGC2_VIBVU		cytoplasm
Q8D4S0	Y5219_VIBVU		inner membrane
Q8D4T9	NAGB_VIBVU		extracellular
Q8D4U6	Y8J3_VIBVU		cytoplasm
Q8D4V5	Y5184_VIBVU		inner membrane

Q8D4Y9	COLA_VIBVU	secreted protein (by similarity).	extracellular
Q8D4Z4	Y8E1_VIBVU		inner membrane
Q8D510	PDXH_VIBVU		cytoplasm
Q8D556	UXAC_VIBVU		cytoplasm
Q8D562	UXUA_VIBVU		cytoplasm
Q8D5G0	Y4960_VIBVU		inner membrane
Q8D5H3	Y4946_VIBVU		inner membrane
Q8D5N2	SY Y1_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D5P5	DEF2_VIBVU		cytoplasm
Q8D5Q4	TNAA_VIBVU		cytoplasm
Q8D5R2	RIMK_VIBVU		cytoplasm
Q8D5Z0	Y4759_VIBVU		inner membrane
Q8D5Z5	LUXP_VIBVU	periplasm (probable).	periplasm
Q8D5Z6	LUXQ_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8D612	NANK_VIBVU		cytoplasm
Q8D613	NANE_VIBVU		cytoplasm
Q8D623	NAPA_VIBVU	periplasm (by similarity).	periplasm
Q8D632	GUAC_VIBVU		cytoplasm
Q8D653	CYSA_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D6H8	TKT2_VIBVU		extracellular
Q8D6H9	TAL_VIBVU	cytoplasm (probable).	cytoplasm
Q8D6I7	GCH1_VIBVU		cytoplasm
Q8D6J0	CLCA_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8D6Q0	ALR2_VIBVU		cytoplasm
Q8D6Q8	ADD_VIBVU		extracellular
Q8D6S1	PHS_VIBVU		cytoplasm
Q8D719	FABH2_VIBVU	cytoplasm (probable).	cytoplasm
Q8D737	COBQ_VIBVU		cytoplasm
Q8D740	MODC_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D757	Y4310_VIBVU		inner membrane
Q8D781	DDL_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D788	Y4273_VIBVU		inner membrane
Q8D795	Y4265_VIBVU		cytoplasm
Q8D7C6	Y4231_VIBVU	periplasm (potential).	periplasm
Q8D7D8	FTHS_VIBVU		cytoplasm
Q8D7E0	GLGC2_VIBVU		cytoplasm
Q8D7G5	GLYA2_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D7G6	GCSH_VIBVU		cytoplasm
Q8D7G7	GCSP_VIBVU		cytoplasm
Q8D7K4	ACKA2_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D7K6	RNB_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D7M9	MGSA_VIBVU		cytoplasm
Q8D7V1	Y4047_VIBVU		inner membrane
Q8D7V4	GBPA_VIBVU	secreted protein (by similarity).	cytoplasm
Q8D7Y4	GLPB_VIBVU		cytoplasm
Q8D7Y9	Y4006_VIBVU		cytoplasm

Q8D825	Y3166_VIBVU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8D849	MSRB_VIBVU		cytoplasm
Q8D866	Y3121_VIBVU		inner membrane
Q8D867	NDPA_VIBVU	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q8D882	RNT_VIBVU		cytoplasm
Q8D885	RNFE_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D887	RNFD_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8D890	RNFA_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D891	UVRB_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D892	LUXU_VIBVU		inner membrane
Q8D894	MOAA_VIBVU		cytoplasm
Q8D896	MOAC_VIBVU		cytoplasm
Q8D8B1	TRPA_VIBVU		cytoplasm
Q8D8B2	TRPB_VIBVU		cytoplasm
Q8D8B4	TRPD_VIBVU		cytoplasm
Q8D8C0	RLUB_VIBVU		cytoplasm
Q8D8C1	Y3059_VIBVU		inner membrane
Q8D8C2	EFPL_VIBVU		cytoplasm
Q8D8C7	UVRC_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8E0	UBIG_VIBVU		cytoplasm
Q8D8F4	URK_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8G1	RLUC_VIBVU		cytoplasm
Q8D8G2	Y3015_VIBVU	cytoplasm (potential).	inner membrane
Q8D8G4	RL32_VIBVU		cytoplasm
Q8D8G5	PLSX_VIBVU		cytoplasm
Q8D8G6	FABH1_VIBVU	cytoplasm (probable).	cytoplasm
Q8D8G9	ACP_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8H5	KTHY_VIBVU		cytoplasm
Q8D8I6	DNAE2_VIBVU	cytoplasm (probable).	cytoplasm
Q8D8J1	KCY_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8J3	IHFB_VIBVU		cytoplasm
Q8D8J6	PYRF_VIBVU		cytoplasm
Q8D8M3	LOLA_VIBVU	periplasm (by similarity).	periplasm
Q8D8N2	BIOD_VIBVU		cytoplasm
Q8D8P4	Y2927_VIBVU		inner membrane
Q8D8P9	HIS1_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8Q0	HISX_VIBVU		cytoplasm
Q8D8Q1	HIS8_VIBVU		cytoplasm
Q8D8Q2	HIS7_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8Q3	HIS5_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8Q4	HIS4_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8Q5	HIS6_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8Q6	HIS2_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8Q7	Y2912_VIBVU		inner membrane
Q8D8Q9	LPXH_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8R2	KITH_VIBVU	cytoplasm (potential).	cytoplasm

Q8D8S3	TORA_VIBVU	periplasm (by similarity).	periplasm
Q8D8V3	HCP_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D8W6	RL25_VIBVU		cytoplasm
Q8D8X2	RSUA_VIBVU		cytoplasm
Q8D8Y6	Y2829_VIBVU		cytoplasm
Q8D8Y7	Y2828_VIBVU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8D900	SERC_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D911	MAO1_VIBVU		cytoplasm
Q8D915	PUR7_VIBVU		cytoplasm
Q8D921	COBT_VIBVU		cytoplasm
Q8D922	COBS_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D926	ASTE_VIBVU		periplasm
Q8D927	BTUC_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8D928	BTUD_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D937	META_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D9E8	Y2655_VIBVU		inner membrane
Q8D9F8	PROQ_VIBVU	cytoplasm (potential).	inner membrane
Q8D9G5	PYRD_VIBVU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8D9H3	FABA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D9H5	Y2627_VIBVU		inner membrane
Q8D9J9	NPD_VIBVU	cytoplasm (probable).	cytoplasm
Q8D9M4	MOBA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8D9N5	Y2562_VIBVU		cytoplasm
Q8D9N8	NORM_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DA11	RL35_VIBVU		cytoplasm
Q8DA18	HUTI_VIBVU	cytoplasm (potential).	cytoplasm
Q8DA19	HUTG_VIBVU		cytoplasm
Q8DA20	HUTU_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DA21	HUTH_VIBVU	cytoplasm (potential).	cytoplasm
Q8DA31	CDD_VIBVU		cytoplasm
Q8DA33	TPMT_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DA35	IHFA_VIBVU		cytoplasm
Q8DA38	PNCB_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DA54	SPEA_VIBVU		cytoplasm
Q8DA60	LIFO_VIBVU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
Q8DA68	ACPD_VIBVU		extracellular
Q8DAD7	Y2269_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DAG7	GCH2_VIBVU		extracellular
Q8DAG8	FADR_VIBVU	cytoplasm (potential).	cytoplasm
Q8DAH1	DUSC_VIBVU		cytoplasm

Q8DAH7	Y2222_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DAH8	ACKA1_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DAI5	Y2212_VIBVU		cytoplasm
Q8DAM0	NADA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DAM3	TOLB_VIBVU	periplasm (potential).	periplasm
Q8DAN4	RUVA_VIBVU		cytoplasm
Q8DAN6	RUVC_VIBVU		cytoplasm
Q8DAQ0	MUKF_VIBVU	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q8DAQ4	TORD_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DAR0	GLGA_VIBVU		cytoplasm
Q8DAR1	GLGC1_VIBVU		cytoplasm
Q8DAR7	LFRR_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DAS0	CLPS_VIBVU		cytoplasm
Q8DAS3	RLUE_VIBVU		cytoplasm
Q8DAS4	Y2115_VIBVU		inner membrane
Q8DAU5	Y2093_VIBVU		inner membrane
Q8DAU9	KDSB_VIBVU	cytoplasm (potential).	cytoplasm
Q8DAV1	LPXK_VIBVU		cytoplasm
Q8DAV2	MSBA_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DAV6	LOLD_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8DB14	Y2014_VIBVU		inner membrane
Q8DB17	AQPZ_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DB22	RECR_VIBVU		cytoplasm
Q8DB23	Y2004_VIBVU		inner membrane
Q8DB36	PDXB_VIBVU		cytoplasm
Q8DB38	Y1985_VIBVU		cytoplasm
Q8DB42	AROC_VIBVU		cytoplasm
Q8DB44	Y1979_VIBVU		inner membrane
Q8DB47	FADJ_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DB48	FADI_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DB58	CCME_VIBVU	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8DB62	CCMA_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8DBB0	DAPA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBC2	PUR5_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBC5	GMHA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBD5	RNH_VIBVU	cytoplasm (potential).	extracellular
Q8DBE1	Y1880_VIBVU	cytoplasm (potential).	inner membrane
Q8DBE4	TILS_VIBVU	cytoplasm (probable).	cytoplasm
Q8DBE5	ACCA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBE7	RNH2_VIBVU	cytoplasm (potential).	extracellular
Q8DBE8	LPXB_VIBVU		cytoplasm

Q8DBE9	LPXA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBF0	FABZ_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBF1	LPXD_VIBVU		cytoplasm
Q8DBF5	DXR_VIBVU		cytoplasm
Q8DBF7	UPPS_VIBVU		inner membrane
Q8DBF9	PYRH_VIBVU		cytoplasm
Q8DBG1	RS2_VIBVU		cytoplasm
Q8DBG3	GLND_VIBVU		cytoplasm
Q8DBG4	Y1856_VIBVU		inner membrane
Q8DBG5	TRUC_VIBVU		cytoplasm
Q8DBH0	Y1847_VIBVU		inner membrane
Q8DBH7	RL31B_VIBVU		cytoplasm
Q8DBI1	Y1836_VIBVU		inner membrane
Q8DBI7	DPO4_VIBVU	cytoplasm (probable).	cytoplasm
Q8DBJ1	NQRF_VIBVU	bacterial cell inner membrane (potential).	inner membrane
Q8DBJ2	NQRE_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8DBJ3	NQRD_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8DBJ4	NQRC_VIBVU	bacterial cell inner membrane (potential).	inner membrane
Q8DBJ5	NQRB_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8DBJ6	NQRA_VIBVU		inner membrane
Q8DBM6	GLPK_VIBVU		cytoplasm
Q8DBN9	GAL1_VIBVU	cytoplasm (potential).	cytoplasm
Q8DBS9	DEOD1_VIBVU		cytoplasm
Q8DBT0	DEOB_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBT1	TYPH_VIBVU		cytoplasm
Q8DBT2	DEOC_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBV5	RS15_VIBVU		cytoplasm
Q8DBV8	TRUB_VIBVU		cytoplasm
Q8DBV9	RBFA_VIBVU	cytoplasm (potential).	cytoplasm
Q8DBW2	Y1694_VIBVU		inner membrane
Q8DBW4	GLMM_VIBVU		cytoplasm
Q8DBW7	RRMJ_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBW9	GREA_VIBVU		cytoplasm
Q8DBX0	OMPU_VIBVU	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8DBX3	SY Y2_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DBX8	GSA_VIBVU	cytoplasm (potential).	cytoplasm
Q8DBZ8	Y1657_VIBVU		inner membrane
Q8DC06	SFSA_VIBVU		cytoplasm
Q8DC08	GLUQ_VIBVU		cytoplasm
Q8DC11	PANB_VIBVU		cytoplasm
Q8DC12	PANC_VIBVU		cytoplasm
Q8DC30	Y1619_VIBVU		cytoplasm
Q8DC31	RL19_VIBVU		cytoplasm
Q8DC32	RIMM_VIBVU	cytoplasm (potential).	cytoplasm
Q8DC33	RS16_VIBVU		cytoplasm
Q8DC38	GSH1_VIBVU		cytoplasm

Q8DC44	OADG_VIBVU	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q8DC47	CSRA_VIBVU		cytoplasm
Q8DC50	RECX_VIBVU	cytoplasm (potential).	cytoplasm
Q8DC51	RECA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DC53	MUTS_VIBVU		cytoplasm
Q8DC56	PIMT_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DC57	SURE_VIBVU	cytoplasm (potential).	periplasm
Q8DC58	TRUD_VIBVU		cytoplasm
Q8DC59	ISPF_VIBVU		extracellular
Q8DC60	ISPD_VIBVU		cytoplasm
Q8DC61	FTSB_VIBVU	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q8DC62	ENO_VIBVU	cytoplasm (by similarity).	extracellular
Q8DC63	PYRG_VIBVU		cytoplasm
Q8DC67	RUMA_VIBVU		cytoplasm
Q8DC72	ACPS_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DC73	PDXJ_VIBVU	cytoplasm (probable).	cytoplasm
Q8DC74	RECO_VIBVU		cytoplasm
Q8DC75	ERA_VIBVU		cytoplasm
Q8DC76	RNC_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DC78	LEPA_VIBVU		cytoplasm
Q8DC84	Y1557_VIBVU		cytoplasm
Q8DC90	Y1551_VIBVU		inner membrane
Q8DC93	RPIA_VIBVU		cytoplasm
Q8DC96	ICIA_VIBVU		cytoplasm
Q8DCA0	PGK_VIBVU	cytoplasm (potential).	cytoplasm
Q8DCA2	TKT1_VIBVU		extracellular
Q8DCA5	SPRT_VIBVU	cytoplasm (potential).	extracellular
Q8DCA9	GSHB_VIBVU		cytoplasm
Q8DCB0	Y1529_VIBVU		inner membrane
Q8DCB1	RUVX_VIBVU	cytoplasm (potential).	cytoplasm
Q8DCB7	Y1522_VIBVU		inner membrane
Q8DCB9	HAM1_VIBVU		extracellular
Q8DCC2	GLSA_VIBVU		extracellular
Q8DCC3	TRMB_VIBVU		cytoplasm
Q8DCC5	FETP_VIBVU		cytoplasm
Q8DCD5	Y1486_VIBVU		inner membrane
Q8DCE5	AMPA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCE7	SYV_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCF4	ARCA_VIBVU	cytoplasm (potential).	extracellular
Q8DCF5	OTC_VIBVU	cytoplasm (probable).	cytoplasm
Q8DCF6	PYRB_VIBVU		cytoplasm
Q8DCF7	PYRI_VIBVU		cytoplasm
Q8DCG0	RLUA_VIBVU		cytoplasm
Q8DCG1	RAPA_VIBVU		cytoplasm
Q8DCG8	Y1452_VIBVU	cytoplasm (potential).	inner membrane
Q8DCJ0	SSB_VIBVU		cytoplasm
Q8DCJ3	UVRA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCJ7	METH_VIBVU		cytoplasm
Q8DCK1	CYSI_VIBVU		cytoplasm
Q8DCK2	CYSJ_VIBVU		cytoplasm

Q8DCK7	G6PI_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCL0	ALR1_VIBVU		cytoplasm
Q8DCL3	RL9_VIBVU		cytoplasm
Q8DCL5	PRIB_VIBVU		cytoplasm
Q8DCL6	RS6_VIBVU		cytoplasm
Q8DCM0	AROB_VIBVU	cytoplasm (probable).	cytoplasm
Q8DCM1	AROK_VIBVU	cytoplasm (probable).	cytoplasm
Q8DCM9	ARLY_VIBVU	cytoplasm (probable).	cytoplasm
Q8DCN0	ASSY_VIBVU	cytoplasm (probable).	cytoplasm
Q8DCN1	ARGE_VIBVU	cytoplasm (probable).	extracellular
Q8DCN2	CAPP_VIBVU		cytoplasm
Q8DCN7	METJ_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCN9	RL31_VIBVU		cytoplasm
Q8DCP3	HSLV_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCP4	HSLU_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCP6	RRAA2_VIBVU		cytoplasm
Q8DCQ3	TPIS_VIBVU		cytoplasm
Q8DCQ9	RS7_VIBVU		cytoplasm
Q8DCR0	RS12_VIBVU		cytoplasm
Q8DCR2	TUSC_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCR3	TUSD_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCR7	SLYX_VIBVU		inner membrane
Q8DCS6	Y1320_VIBVU		inner membrane
Q8DCT7	GPH_VIBVU		cytoplasm
Q8DCT8	RLMB_VIBVU		cytoplasm
Q8DCU2	HMP_VIBVU		cytoplasm
Q8DCU4	PURA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCU9	HFQ_VIBVU		cytoplasm
Q8DCV0	MUTL_VIBVU		cytoplasm
Q8DCV6	ORN_VIBVU	cytoplasm (potential).	cytoplasm
Q8DCV7	ENGC1_VIBVU		cytoplasm
Q8DCV8	PSD_VIBVU		cytoplasm
Q8DCW1	GPMI_VIBVU		cytoplasm
Q8DCW3	SECB_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCW4	GPDA_VIBVU	cytoplasm (probable).	cytoplasm
Q8DCX0	SYK3_VIBVU		cytoplasm
Q8DCX3	FRDC_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DCX4	FRDD_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DCY1	K6PF_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DCZ0	DSBD_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DCZ9	ACSA_VIBVU		cytoplasm
Q8DD00	AROQ_VIBVU		cytoplasm
Q8DD03	PRMA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DD06	PUR9_VIBVU		cytoplasm
Q8DD07	PUR2_VIBVU		cytoplasm
Q8DD14	DCUP_VIBVU	cytoplasm (probable).	cytoplasm
Q8DD17	NUDC_VIBVU		cytoplasm
Q8DD19	RPOC_VIBVU		cytoplasm
Q8DD20	RPOB_VIBVU		cytoplasm
Q8DD21	RL7_VIBVU		cytoplasm

Q8DD22	RL10_VIBVU		cytoplasm
Q8DD23	RL1_VIBVU		cytoplasm
Q8DD24	RL11_VIBVU		cytoplasm
Q8DD25	NUSG_VIBVU		cytoplasm
Q8DD30	COAA_VIBVU	cytoplasm (probable).	cytoplasm
Q8DD33	MURB_VIBVU	cytoplasm (probable).	inner membrane
Q8DD39	MURI_VIBVU		cytoplasm
Q8DD41	BTUB_VIBVU	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8DD43	TRMA_VIBVU		cytoplasm
Q8DD46	STHA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DD47	LEXA_VIBVU		cytoplasm
Q8DD48	PLSB_VIBVU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8DD53	GLPE_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DD54	RP32_VIBVU		cytoplasm
Q8DD81	DAPF_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DD83	CYAY_VIBVU		inner membrane
Q8DD85	HEM3_VIBVU		cytoplasm
Q8DD90	USPB_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DD95	Y1112_VIBVU		inner membrane
Q8DDC8	ILVC_VIBVU		cytoplasm
Q8DDD4	AROE_VIBVU		cytoplasm
Q8DDD5	HEM6_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDD7	PUR6_VIBVU		cytoplasm
Q8DDD8	PURK_VIBVU		cytoplasm
Q8DDE0	SMG_VIBVU		cytoplasm
Q8DDE3	DEF1_VIBVU		cytoplasm
Q8DDE4	FMT_VIBVU		cytoplasm
Q8DDE5	RSMB_VIBVU		cytoplasm
Q8DDG1	ILVD_VIBVU		cytoplasm
Q8DDG6	GLMU_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDG7	ATPE_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8DDG9	ATPG_VIBVU		inner membrane
Q8DDH0	ATPA_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8DDH2	ATPF_VIBVU		inner membrane
Q8DDH8	GIDB_VIBVU		cytoplasm
Q8DDH9	GIDA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDI1	TRME_VIBVU		cytoplasm
Q8DDI2	OXAA_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DDI3	RNPA_VIBVU		extracellular
Q8DDI9	DNAA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDJ1	RECF_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDK1	TUSA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDK5	FADA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDK6	FADB_VIBVU		cytoplasm

Q8DDL3	CRCB_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DDL4	THIC_VIBVU		cytoplasm
Q8DDL8	THIG_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDM6	Y952_VIBVU		inner membrane
Q8DDN5	GPPA_VIBVU		extracellular
Q8DDN6	RHLB_VIBVU		cytoplasm
Q8DDP9	UBIE_VIBVU		cytoplasm
Q8DDQ1	UBIB_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DDQ2	TATA_VIBVU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8DDQ3	TATB_VIBVU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8DDQ8	ENGB_VIBVU		cytoplasm
Q8DDR1	Y896_VIBVU		inner membrane
Q8DDS2	RBN_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DDS3	DTD_VIBVU	cytoplasm (probable).	cytoplasm
Q8DDS6	PPCK_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDS7	HSLO_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDU2	Y864_VIBVU		inner membrane
Q8DDU4	BIOH_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDU7	GREB_VIBVU		cytoplasm
Q8DDV5	RPOZ_VIBVU		cytoplasm
Q8DDV6	KGUA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDX4	RNPH_VIBVU		cytoplasm
Q8DDX5	PYRE_VIBVU		cytoplasm
Q8DDX7	SLMA_VIBVU	nucleoid-associated (by similarity).	cytoplasm
Q8DDX8	COABC_VIBVU		cytoplasm
Q8DDY0	RADC_VIBVU		cytoplasm
Q8DDY1	RL28_VIBVU		cytoplasm
Q8DDY2	RL33_VIBVU		cytoplasm
Q8DDY4	FPG_VIBVU		cytoplasm
Q8DDY6	COAD_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DDZ1	KDKA_VIBVU	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q8DE09	HLDD_VIBVU		cytoplasm
Q8DE39	RL3_VIBVU		cytoplasm
Q8DE40	RL4_VIBVU		cytoplasm
Q8DE42	RL2_VIBVU		cytoplasm
Q8DE43	RS19_VIBVU		cytoplasm
Q8DE44	RL22_VIBVU		cytoplasm
Q8DE45	RS3_VIBVU		cytoplasm
Q8DE46	RL16_VIBVU		cytoplasm
Q8DE47	RL29_VIBVU		cytoplasm
Q8DE48	RS17_VIBVU		cytoplasm
Q8DE50	RL24_VIBVU		cytoplasm
Q8DE51	RL5_VIBVU		cytoplasm

Q8DE54	RS8_VIBVU		cytoplasm
Q8DE56	RL18_VIBVU		cytoplasm
Q8DE57	RS5_VIBVU		cytoplasm
Q8DE59	RL15_VIBVU		cytoplasm
Q8DE61	RS13_VIBVU		cytoplasm
Q8DE62	RS4_VIBVU		cytoplasm
Q8DE63	RPOA_VIBVU		cytoplasm
Q8DE72	CYSD_VIBVU		cytoplasm
Q8DE73	CYSN_VIBVU		cytoplasm
Q8DE75	CYSC_VIBVU		cytoplasm
Q8DE84	MSRA_VIBVU		cytoplasm
Q8DE89	IPYR_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DE96	Y700_VIBVU		cytoplasm
Q8DEA0	Y695_VIBVU		cytoplasm
Q8DEB6	MURA_VIBVU	cytoplasm (probable).	inner membrane
Q8DEC1	ARGR_VIBVU	cytoplasm (probable).	cytoplasm
Q8DEC2	MDH_VIBVU		cytoplasm
Q8DEC5	RL27_VIBVU		cytoplasm
Q8DED0	APAH_VIBVU		extracellular
Q8DED1	APAG_VIBVU		inner membrane
Q8DED2	KSGA_VIBVU		cytoplasm
Q8DED3	PDXA_VIBVU	cytoplasm (probable).	cytoplasm
Q8DED5	OSTA_VIBVU	bacterial cell outer membrane (by similarity).	outer membrane
Q8DED6	DJLA_VIBVU	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q8DED8	LEUD_VIBVU		cytoplasm
Q8DED9	LEU2_VIBVU		cytoplasm
Q8DEE0	LEU3_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DEE1	LEU1_VIBVU		cytoplasm
Q8DEF3	GLMS_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DEF5	MTLD_VIBVU		cytoplasm
Q8DEG8	UPPP_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DEH6	GLNE_VIBVU		cytoplasm
Q8DEH7	HLDE_VIBVU		cytoplasm
Q8DEJ0	RS9_VIBVU		cytoplasm
Q8DEJ8	Y590_VIBVU		inner membrane
Q8DEK2	MRAW_VIBVU		cytoplasm
Q8DEK5	MURE_VIBVU	cytoplasm (probable).	cytoplasm
Q8DEK7	MRAY_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DEK8	MURD_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DEL0	MURG_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8DEL1	MURC_VIBVU	cytoplasm (probable).	cytoplasm
Q8DEL6	LPXC_VIBVU		cytoplasm
Q8DEM0	DAPB_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DEM1	CARA_VIBVU		cytoplasm
Q8DEM2	CARB_VIBVU		cytoplasm
Q8DEM7	BTUF_VIBVU	periplasm (by similarity).	periplasm
Q8DEP0	RRAA1_VIBVU		cytoplasm

Q8DEP3	KHSE_VIBVU	cytoplasm (potential).	cytoplasm
Q8DEP5	GRCA_VIBVU		cytoplasm
Q8DEP6	END4_VIBVU		extracellular
Q8DEP7	UNG_VIBVU	cytoplasm (by similarity).	extracellular
Q8DEQ1	Y535_VIBVU		inner membrane
Q8DER4	MUTH_VIBVU	cytoplasm (by similarity).	extracellular
Q8DER5	NUDH_VIBVU		cytoplasm
Q8DER8	LGT_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DER9	TYSY_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DES4	RS20_VIBVU		cytoplasm
Q8DES7	SYI_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DES8	LSPA_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8DET0	ISPH_VIBVU		cytoplasm
Q8DET1	Y503_VIBVU		cytoplasm
Q8DEU5	TRPR_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DEU6	Y488_VIBVU		inner membrane
Q8DEV0	RLUD_VIBVU		cytoplasm
Q8DEV2	CLPB_VIBVU	cytoplasm (probable).	cytoplasm
Q8DEV7	RDGC_VIBVU	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q8DEW2	PPK_VIBVU		cytoplasm
Q8DEW5	PSTB1_VIBVU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8DEX2	CUTC_VIBVU	cytoplasm (probable).	cytoplasm
Q8DEX9	QUEA_VIBVU	cytoplasm (potential).	cytoplasm
Q8DEY0	TGT_VIBVU		cytoplasm
Q8DEY7	ISCS_VIBVU		cytoplasm
Q8DEZ0	HSCB_VIBVU		inner membrane
Q8DEZ1	HSCA_VIBVU		cytoplasm
Q8DEZ4	PEPB_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DEZ5	NDK_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DEZ8	ISPG_VIBVU		cytoplasm
Q8DF02	ENGA_VIBVU		cytoplasm
Q8DF05	EX7L_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DF07	GUAA_VIBVU		cytoplasm
Q8DF11	DADA_VIBVU		cytoplasm
Q8DF53	SSRP_VIBVU	cytoplasm (potential).	cytoplasm
Q8DF55	Y369_VIBVU		inner membrane
Q8DF58	PPNK_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DF59	GRPE_VIBVU	cytoplasm (probable).	cytoplasm
Q8DF66	DNAK_VIBVU		cytoplasm
Q8DF67	DNAJ_VIBVU	cytoplasm (probable).	cytoplasm
Q8DF90	XGPT_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DF91	Y328_VIBVU		cytoplasm
Q8DF93	PROB_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DF94	PROA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DF95	NRDR_VIBVU		cytoplasm
Q8DF99	RISB_VIBVU		cytoplasm
Q8DFA0	NUSB_VIBVU		cytoplasm
Q8DFA3	DXS_VIBVU		cytoplasm

Q8DFA5	EX7S_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DFA8	THII_VIBVU	cytoplasm (potential).	cytoplasm
Q8DFB2	MTFA_VIBVU		cytoplasm
Q8DFB8	QUEF_VIBVU	cytoplasm (probable).	cytoplasm
Q8DFC9	GLYA1_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DFD1	LIPA_VIBVU	cytoplasm (potential).	cytoplasm
Q8DFD2	LIPB_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DFD3	Y282_VIBVU		inner membrane
Q8DFD8	Y277_VIBVU		inner membrane
Q8DFE4	LNT_VIBVU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8DFE6	Y268_VIBVU		inner membrane
Q8DFF4	PTH_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DFF5	KPRS_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DFF6	ISPE_VIBVU		cytoplasm
Q8DFF7	LOLB_VIBVU	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q8DFF8	HEM1_VIBVU		cytoplasm
Q8DFG3	KDSA_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DFG4	5NTD_VIBVU	bacterial cell outer membrane; lipid-anchor (probable).	periplasm
Q8DFH4	Y237_VIBVU		cytoplasm
Q8DFJ1	FLGH_VIBVU	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8DFJ2	FLGI_VIBVU	periplasm (by similarity).	periplasm
Q8DFK4	ZIPA_VIBVU	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q8DFM0	HTPG_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DFM1	KAD_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DFM2	HEMH_VIBVU	cytoplasm (by similarity).	extracellular
Q8DFQ1	SUCC_VIBVU		cytoplasm
Q8DFS2	MINE_VIBVU		inner membrane
Q8DFS4	MINC_VIBVU		inner membrane
Q8DG12	Y039_VIBVU		inner membrane
Q8DG25	TIG_VIBVU		cytoplasm
Q8DG26	CLPP_VIBVU	cytoplasm (by similarity).	cytoplasm
Q8DG27	CLPX_VIBVU		cytoplasm
Q8DG35	Y013_VIBVU		inner membrane
Q8E8A9	GIDA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E8B0	GIDB_SHEON		cytoplasm
Q8E8B8	ATPA_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8E8B9	ATPG_SHEON		inner membrane
Q8E8C1	ATPE_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8E8C2	GLMU_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E8D4	ADD_SHEON		extracellular
Q8E8D7	Y4728_SHEON		inner membrane
Q8E8D9	FPG_SHEON		cytoplasm
Q8E8E3	MOBA_SHEON	cytoplasm (by similarity).	cytoplasm

Q8E8F7	Y4708_SHEON	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8E8I0	COAD_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E8J1	TDH_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E8J2	GLPE_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E8K6	Y4658_SHEON		inner membrane
Q8E8K8	CYSA2_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8E8N6	BIOH_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E8P2	Y4619_SHEON		inner membrane
Q8E8Q8	LEXA_SHEON		cytoplasm
Q8E8Q9	PLSB_SHEON	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8E927	Y4467_SHEON		inner membrane
Q8E942	MOAC_SHEON		cytoplasm
Q8E985	RBN_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8E988	DTD_SHEON	cytoplasm (probable).	cytoplasm
Q8E9D5	ILVC_SHEON		cytoplasm
Q8E9D9	ILVD_SHEON		cytoplasm
Q8E9H0	HEM3_SHEON		cytoplasm
Q8E9H2	CYAY_SHEON		inner membrane
Q8E9H5	DAPF_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E9I8	PSTB2_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8E9K1	UPPP2_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8E9L4	RNPH_SHEON		cytoplasm
Q8E9L5	PYRE_SHEON		cytoplasm
Q8E9L6	GCH1_SHEON		cytoplasm
Q8E9L9	SLMA_SHEON	nucleoid-associated (by similarity).	cytoplasm
Q8E9M0	DUT_SHEON		extracellular
Q8E9M2	RADC_SHEON		cytoplasm
Q8E9M3	RL28_SHEON		cytoplasm
Q8E9N2	LEU1_SHEON		cytoplasm
Q8E9N3	LEU3_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E9N4	LEU2_SHEON		cytoplasm
Q8E9N5	LEUD_SHEON		cytoplasm
Q8E9N7	GLPK_SHEON		cytoplasm
Q8E9N9	MRAZ_SHEON		inner membrane
Q8E9P0	MRAW_SHEON		cytoplasm
Q8E9P3	MURE_SHEON	cytoplasm (probable).	cytoplasm
Q8E9P5	MRAY_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8E9P6	MURD_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E9P8	MURC_SHEON	cytoplasm (probable).	cytoplasm
Q8E9Q2	LPXC_SHEON		cytoplasm

Q8E9R3	TATB_SHEON	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8E9R7	UBIE_SHEON		cytoplasm
Q8E9R9	RRAA2_SHEON		cytoplasm
Q8E9U9	HSLU_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E9V0	HSLV_SHEON	cytoplasm (by similarity).	cytoplasm
Q8E9Z0	RL31_SHEON		cytoplasm
Q8EA14	Y4095_SHEON	cytoplasm (potential).	inner membrane
Q8EA30	Y4079_SHEON		inner membrane
Q8EA43	PUR7_SHEON		cytoplasm
Q8EA52	METJ_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EA81	SSB_SHEON		cytoplasm
Q8EA87	MDTL_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EAB0	Y3997_SHEON	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8EAC4	Y3983_SHEON		inner membrane
Q8EAE2	Y3964_SHEON		cytoplasm
Q8EAF7	MURA_SHEON	cytoplasm (probable).	inner membrane
Q8EAG3	RS9_SHEON		cytoplasm
Q8EAG5	PURA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EAG8	RLMB_SHEON		cytoplasm
Q8EAH2	RS6_SHEON		cytoplasm
Q8EAH3	PRIB_SHEON		cytoplasm
Q8EAH4	RS18_SHEON		cytoplasm
Q8EAH5	RL9_SHEON		cytoplasm
Q8EAI9	Y3913_SHEON		cytoplasm
Q8EAJ0	DUSA_SHEON		cytoplasm
Q8EAL4	Y3888_SHEON		extracellular
Q8EAM8	DSBI_SHEON	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8EAN3	MODC_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8EAP2	MAO1_SHEON		cytoplasm
Q8EAQ9	KPRS_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EAR0	ISPE_SHEON		cytoplasm
Q8EAR1	LOLB_SHEON	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q8EAR2	HEM1_SHEON		cytoplasm
Q8EAR9	KDSA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EAS7	Y3815_SHEON		inner membrane
Q8EAX8	SPEE_SHEON		cytoplasm
Q8EAY1	GLNE_SHEON		cytoplasm
Q8EAZ0	CVRA_SHEON	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8EAZ2	HLDE_SHEON		cytoplasm
Q8EAZ9	CYSJ_SHEON		cytoplasm
Q8EB00	CYSI_SHEON		cytoplasm
Q8EB01	CYSH_SHEON	cytoplasm (by similarity).	cytoplasm

Q8EB02	MTGA_SHEON	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8EB10	CYSN_SHEON		cytoplasm
Q8EB13	CYSC_SHEON		cytoplasm
Q8EB78	UNG_SHEON	cytoplasm (by similarity).	extracellular
Q8EB81	RL27_SHEON		cytoplasm
Q8EB91	APAH_SHEON		extracellular
Q8EB92	APAG_SHEON		inner membrane
Q8EB93	KSGA_SHEON		cytoplasm
Q8EB94	PDXA_SHEON	cytoplasm (probable).	cytoplasm
Q8EB99	DJLA_SHEON	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q8EBC3	CYSA1_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8EBE6	CLPB_SHEON	cytoplasm (probable).	cytoplasm
Q8EBF9	GSH1_SHEON		cytoplasm
Q8EBH1	G6PI_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EBH2	TAL_SHEON	cytoplasm (probable).	cytoplasm
Q8EBH6	Y3540_SHEON		inner membrane
Q8EBH9	RS20_SHEON		cytoplasm
Q8EBI4	SYI_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EBI7	ISPH_SHEON		cytoplasm
Q8EBN8	GLYA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EBN9	NRDR_SHEON		cytoplasm
Q8EBP3	RISB_SHEON		cytoplasm
Q8EBP4	NUSB_SHEON		cytoplasm
Q8EBQ3	RUMA_SHEON		cytoplasm
Q8EBR0	ENO_SHEON	cytoplasm (by similarity).	extracellular
Q8EBR1	FTSB_SHEON	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q8EBR2	ISPD_SHEON		cytoplasm
Q8EBR3	ISPF_SHEON		extracellular
Q8EBR4	TRUD_SHEON		cytoplasm
Q8EBR5	SURE_SHEON	cytoplasm (potential).	periplasm
Q8EBR9	MUTS_SHEON		cytoplasm
Q8EBS0	RECA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EBS3	CSRA_SHEON		cytoplasm
Q8EBS5	SYV_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EBU8	Y3401_SHEON		inner membrane
Q8EBX5	Y3370_SHEON	periplasm (potential).	periplasm
Q8EBX6	FETP_SHEON		cytoplasm
Q8EBX8	TRMB_SHEON		cytoplasm
Q8EBY0	GLSA_SHEON		extracellular
Q8EBY7	HAM1_SHEON		extracellular
Q8EBY9	Y3356_SHEON		inner membrane
Q8EBZ7	HEMH2_SHEON	cytoplasm (by similarity).	extracellular
Q8EBZ8	RUVX_SHEON	cytoplasm (potential).	cytoplasm
Q8EBZ9	Y3346_SHEON		inner membrane
Q8EC32	ISPG_SHEON		cytoplasm
Q8EC36	ENGA_SHEON		cytoplasm

Q8EC50	EX7L_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EC52	GUAA_SHEON		cytoplasm
Q8ECA1	FLGH_SHEON	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8ECA2	FLGI_SHEON	periplasm (by similarity).	periplasm
Q8ECK0	KITH_SHEON	cytoplasm (by similarity).	cytoplasm
Q8ECM2	QUEA_SHEON	cytoplasm (potential).	cytoplasm
Q8ECM3	TGT_SHEON		cytoplasm
Q8ECP6	FADI_SHEON	cytoplasm (by similarity).	cytoplasm
Q8ECP7	FADJ_SHEON	cytoplasm (by similarity).	cytoplasm
Q8ECR2	PDXB_SHEON		cytoplasm
Q8ECU9	TRPA_SHEON		cytoplasm
Q8ECV0	TRPB_SHEON		cytoplasm
Q8ECV2	TRPD_SHEON		cytoplasm
Q8ED55	ACKA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8ED56	Y2914_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ED69	ZIPA_SHEON	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q8ED71	PDXH_SHEON		cytoplasm
Q8ED72	Y2894_SHEON		inner membrane
Q8ED80	FADR_SHEON	cytoplasm (potential).	cytoplasm
Q8EDD1	GCH2_SHEON		extracellular
Q8EDD7	Y2823_SHEON		cytoplasm
Q8EDF0	MSBA_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EDF1	LPXK_SHEON		cytoplasm
Q8EDG1	CDD_SHEON		cytoplasm
Q8EDG7	Y2782_SHEON	cytoplasm (potential).	inner membrane
Q8EDG9	RL32_SHEON		cytoplasm
Q8EDH0	PLSX_SHEON		cytoplasm
Q8EDH1	FABH_SHEON	cytoplasm (probable).	cytoplasm
Q8EDH4	ACP_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EDI8	PUR5_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EDI9	UPP_SHEON		cytoplasm
Q8EDJ2	RNT_SHEON		cytoplasm
Q8EDJ8	TOLB_SHEON	periplasm (potential).	periplasm
Q8EDK3	ACSA_SHEON		cytoplasm
Q8EDL2	OPGG2_SHEON	periplasm (by similarity).	periplasm
Q8EDL3	PEPE_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EDL5	HTPX_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8EDM4	DEOD3_SHEON		cytoplasm
Q8EDU8	Y2645_SHEON		cytoplasm
Q8EDV9	Y2634_SHEON		inner membrane
Q8EDW4	CLPS_SHEON		cytoplasm
Q8EDW7	ATE_SHEON	cytoplasm (potential).	cytoplasm
Q8EDW8	LFTR_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EDX4	URK_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EDX5	DCD_SHEON		extracellular
Q8EDX8	KTHY_SHEON		cytoplasm
Q8EDY8	PROQ_SHEON	cytoplasm (potential).	inner membrane

Q8EDZ8	PYRD_SHEON	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8EE14	MINC_SHEON		inner membrane
Q8EE17	Y2573_SHEON		inner membrane
Q8EE23	RRAA1_SHEON		cytoplasm
Q8EE30	RNH_SHEON	cytoplasm (potential).	extracellular
Q8EE60	DEF3_SHEON		cytoplasm
Q8EEA2	DGTL1_SHEON		cytoplasm
Q8EEA3	Y2484_SHEON		inner membrane
Q8EED7	THIC_SHEON		cytoplasm
Q8EEE1	THIG_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EEF0	Y2432_SHEON		inner membrane
Q8EEF1	RUVC_SHEON		cytoplasm
Q8EEF2	RUVA_SHEON		cytoplasm
Q8EEF3	RUVB_SHEON		cytoplasm
Q8EEG9	UBIG_SHEON		cytoplasm
Q8EEH2	SERC_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EEH8	AROA_SHEON	cytoplasm (probable).	cytoplasm
Q8EEH9	KCY_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EEI4	PYRF_SHEON		cytoplasm
Q8EEN5	NADA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EEN9	ASTE_SHEON		periplasm
Q8EEQ7	ECOT_SHEON	periplasm (by similarity).	periplasm
Q8EER0	CRCB_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EER2	LOLA_SHEON	periplasm (by similarity).	periplasm
Q8EER6	RL20_SHEON		cytoplasm
Q8EER7	RL35_SHEON		cytoplasm
Q8EES3	NORM_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EES8	Y2290_SHEON		inner membrane
Q8EEU0	NDK_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EEU3	RIMK2_SHEON		cytoplasm
Q8EEU5	HSCA_SHEON		cytoplasm
Q8EEU6	HSCB_SHEON		inner membrane
Q8EEV5	LOLD_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8EEW2	NAGZ_SHEON	cytoplasm (by similarity).	extracellular
Q8EEZ2	DDL_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EF25	NDPA_SHEON	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q8EF26	Y2176_SHEON		inner membrane
Q8EF50	Y2143_SHEON	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8EF74	RL25_SHEON		cytoplasm
Q8EF78	OPGH_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EF79	OPGG1_SHEON	periplasm (by similarity).	periplasm
Q8EF96	HYP_A_SHEON		cytoplasm

Q8EFB0	HIS1_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFB1	HISX_SHEON		cytoplasm
Q8EFB2	HIS8_SHEON		cytoplasm
Q8EFB3	HIS7_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFB4	HIS5_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFB5	HIS4_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFB6	HIS2_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFF2	NADE_SHEON		cytoplasm
Q8EFF4	HEMH1_SHEON	cytoplasm (by similarity).	extracellular
Q8EFF5	KAD_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFF7	HTPG_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFF8	RECR_SHEON		cytoplasm
Q8EFF9	Y2014_SHEON		inner membrane
Q8EFG5	Y2008_SHEON		inner membrane
Q8EFG7	DUSC_SHEON		cytoplasm
Q8EFN2	NPD_SHEON	cytoplasm (probable).	cytoplasm
Q8EFN8	SUCC_SHEON		cytoplasm
Q8EFP8	Y1922_SHEON		inner membrane
Q8EFT7	DAPA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFU5	SPEA_SHEON		cytoplasm
Q8EFV4	UVRC_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFV9	FABA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EFZ8	Y1816_SHEON		inner membrane
Q8EG03	Y1811_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EG14	Y1800_SHEON		cytoplasm
Q8EG18	CLPX_SHEON		cytoplasm
Q8EG19	CLPP_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EG20	TIG_SHEON		cytoplasm
Q8EG24	LPXH_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EG82	PSTB1_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8EGC8	META_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EGD7	PHS_SHEON		cytoplasm
Q8EGF2	Y1652_SHEON		cytoplasm
Q8EGF9	TILS_SHEON	cytoplasm (probable).	cytoplasm
Q8EGG1	RNH2_SHEON	cytoplasm (potential).	extracellular
Q8EGG2	LPXB_SHEON		cytoplasm
Q8EGG4	FABZ_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EGG5	LPXD_SHEON		cytoplasm
Q8EGG9	DXR_SHEON		cytoplasm
Q8EGH1	UPPS_SHEON		inner membrane
Q8EGH2	RRF_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EGH5	RS2_SHEON		cytoplasm
Q8EGH8	GLND_SHEON		cytoplasm
Q8EGH9	DAPD_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EGJ4	QUEF_SHEON	cytoplasm (probable).	cytoplasm
Q8EGP3	RDGC_SHEON	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q8EGQ9	MTFA_SHEON		cytoplasm
Q8EGR4	THII_SHEON	cytoplasm (potential).	cytoplasm
Q8EGR9	DXS_SHEON		cytoplasm
Q8EGS0	GRPE_SHEON	cytoplasm (probable).	cytoplasm

Q8EGS1	PPNK_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EGU3	GLGC_SHEON		cytoplasm
Q8EGU7	GLGB_SHEON		extracellular
Q8EGW6	Y1475_SHEON		inner membrane
Q8EGW8	SSRP_SHEON	cytoplasm (potential).	cytoplasm
Q8EH62	AMPA2_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EH67	HCP_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EH70	RL19_SHEON		cytoplasm
Q8EH71	RIMM_SHEON	cytoplasm (potential).	cytoplasm
Q8EH72	RS16_SHEON		cytoplasm
Q8EH77	ACPS_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EH78	PDXJ_SHEON	cytoplasm (probable).	cytoplasm
Q8EH81	RNC_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EH83	LEPA_SHEON		cytoplasm
Q8EH90	Y1339_SHEON		cytoplasm
Q8EH94	TYSY_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EH95	LGT_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EH98	NUDH_SHEON		cytoplasm
Q8EH99	MUTH_SHEON	cytoplasm (by similarity).	extracellular
Q8EHB3	SYI_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EHC1	AQPZ_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EHC8	GSA_SHEON	cytoplasm (potential).	cytoplasm
Q8EHD3	UPPP1_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EHI9	TORA_SHEON	periplasm (by similarity).	periplasm
Q8EHJ0	TORD_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EHK0	DEOD2_SHEON		cytoplasm
Q8EHK2	DEOB_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EHK3	TYPH_SHEON		cytoplasm
Q8EHK4	DEOC_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EHL3	RS15_SHEON		cytoplasm
Q8EHL4	RBFA_SHEON	cytoplasm (potential).	cytoplasm
Q8EHL7	Y1202_SHEON		inner membrane
Q8EHL9	TPIS_SHEON		cytoplasm
Q8EHM0	GLMM_SHEON		cytoplasm
Q8EHM3	RRMJ_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EHN5	PTH_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EHN9	Y1179_SHEON		inner membrane
Q8EHP1	LNT_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EHP8	Y1169_SHEON		inner membrane
Q8EHQ4	Y1163_SHEON		inner membrane
Q8EHQ5	LIPB_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EHQ6	LIPA_SHEON	cytoplasm (potential).	cytoplasm
Q8EHR7	RPIA_SHEON		cytoplasm
Q8EHS6	CARA_SHEON		cytoplasm
Q8EHS7	DAPB_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EHT6	DNAJ_SHEON	cytoplasm (probable).	cytoplasm
Q8EHT7	DNAK_SHEON		cytoplasm
Q8EHU1	PROA_SHEON	cytoplasm (by similarity).	cytoplasm

Q8EHU2	PROB_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EHW1	LUXS_SHEON		extracellular
Q8EHY2	GBPA_SHEON	secreted protein (by similarity).	cytoplasm
Q8EHZ1	SLYX_SHEON		inner membrane
Q8EHZ2	DEF2_SHEON		cytoplasm
Q8EI02	OPGD_SHEON	periplasm (by similarity).	periplasm
Q8EI15	COBQ_SHEON		cytoplasm
Q8EI17	COBS_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EI18	COBT_SHEON		cytoplasm
Q8EI34	NUOG_SHEON		cytoplasm
Q8EI67	RUMB_SHEON		cytoplasm
Q8EI85	AMPA1_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EIB1	PGK_SHEON	cytoplasm (potential).	cytoplasm
Q8EIC6	Y917_SHEON		inner membrane
Q8EIF2	AGUA_SHEON		extracellular
Q8EIG4	SFSA_SHEON		cytoplasm
Q8EIG6	GLUQ_SHEON		cytoplasm
Q8EIG9	PANB_SHEON		cytoplasm
Q8EIH0	PANC_SHEON		cytoplasm
Q8EIJ1	NAPA_SHEON	periplasm (by similarity).	periplasm
Q8EIK5	SPRT_SHEON	cytoplasm (potential).	extracellular
Q8EIK8	GSHB_SHEON		cytoplasm
Q8EIM0	METE_SHEON		cytoplasm
Q8EIM7	RIHA_SHEON		extracellular
Q8EIQ6	GCSP_SHEON		cytoplasm
Q8EIQ7	GCSH_SHEON		cytoplasm
Q8EIQ8	GCST_SHEON		cytoplasm
Q8EIR6	ARGR_SHEON	cytoplasm (probable).	cytoplasm
Q8EIV3	Y727_SHEON		inner membrane
Q8EJ69	HFQ_SHEON		cytoplasm
Q8EJ78	ORN_SHEON	cytoplasm (potential).	cytoplasm
Q8EJ79	ENGC_SHEON		cytoplasm
Q8EJ86	TPMT_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EJ93	RAPA_SHEON		cytoplasm
Q8EJA7	FTHS_SHEON		cytoplasm
Q8EJC1	RIMK1_SHEON		cytoplasm
Q8EJE5	MSCL_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8EJF5	AROQ_SHEON		cytoplasm
Q8EJM1	PUR9_SHEON		cytoplasm
Q8EJM8	DCUP_SHEON	cytoplasm (probable).	cytoplasm
Q8EJP9	COAE_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EJQ0	Y412_SHEON		inner membrane
Q8EJQ1	Y411_SHEON		cytoplasm
Q8EJQ5	RHLB_SHEON		cytoplasm
Q8EJR7	PRMA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EJR8	DUSB_SHEON		cytoplasm
Q8EJR9	FIS_SHEON		cytoplasm
Q8EJU6	KGUA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EJU7	RPOZ_SHEON		cytoplasm
Q8EJX6	Y330_SHEON		inner membrane
Q8EJZ5	Y311_SHEON		cytoplasm

Q8EK07	Y299_SHEON		inner membrane
Q8EK19	AROB_SHEON	cytoplasm (probable).	cytoplasm
Q8EK20	AROK_SHEON	cytoplasm (probable).	cytoplasm
Q8EK27	ARLY_SHEON	cytoplasm (probable).	cytoplasm
Q8EK28	ASSY_SHEON	cytoplasm (probable).	cytoplasm
Q8EK29	OTC_SHEON	cytoplasm (probable).	cytoplasm
Q8EK30	CAPP_SHEON		cytoplasm
Q8EK40	CCMA_SHEON	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8EK44	CCME_SHEON	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8EK47	RPOA_SHEON		cytoplasm
Q8EK48	RS13_SHEON		cytoplasm
Q8EK51	RL15_SHEON		cytoplasm
Q8EK53	RL18_SHEON		cytoplasm
Q8EK55	RS8_SHEON		cytoplasm
Q8EK57	RL5_SHEON		cytoplasm
Q8EK58	RL24_SHEON		cytoplasm
Q8EK60	RS17_SHEON		cytoplasm
Q8EK61	RL29_SHEON		cytoplasm
Q8EK62	RL16_SHEON		cytoplasm
Q8EK63	RL22_SHEON		cytoplasm
Q8EK64	RS19_SHEON		cytoplasm
Q8EK65	RL2_SHEON		cytoplasm
Q8EK67	RL4_SHEON		cytoplasm
Q8EK68	RL3_SHEON		cytoplasm
Q8EK69	RS10_SHEON		cytoplasm
Q8EK72	RS7_SHEON		cytoplasm
Q8EK73	RPOC_SHEON		cytoplasm
Q8EK74	RPOB_SHEON		cytoplasm
Q8EK76	RL10_SHEON		cytoplasm
Q8EK77	RL1_SHEON		cytoplasm
Q8EK78	RL11_SHEON		cytoplasm
Q8EK83	COAA_SHEON	cytoplasm (probable).	cytoplasm
Q8EK85	MURB_SHEON	cytoplasm (probable).	inner membrane
Q8EK90	MURI_SHEON		cytoplasm
Q8EK99	SELD_SHEON		cytoplasm
Q8EKA0	SELU_SHEON		cytoplasm
Q8EKD2	HSLO_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKD3	PPCK_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKF2	RIBB_SHEON		cytoplasm
Q8EKI6	FDHD_SHEON	cytoplasm (potential).	cytoplasm
Q8EKI8	SELA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKJ4	HUTH_SHEON	cytoplasm (potential).	cytoplasm
Q8EKJ5	HUTU_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKK0	DEOD1_SHEON		cytoplasm
Q8EKN9	GPDA_SHEON	cytoplasm (probable).	cytoplasm
Q8EKP0	SECB_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKQ2	HEM6_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKQ5	SMG_SHEON		cytoplasm
Q8EKQ8	DEF1_SHEON		cytoplasm
Q8EKQ9	FMT_SHEON		cytoplasm
Q8EKR0	RSMB_SHEON		cytoplasm

Q8EKR9	FADB_SHEON		cytoplasm
Q8EKS0	FADA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKS1	TUSA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKT0	RECF_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKT2	DNAA_SHEON	cytoplasm (by similarity).	cytoplasm
Q8EKT4	RNPA_SHEON		extracellular
Q8EKT5	Y005_SHEON		inner membrane
Q8EKT6	OXAA_SHEON	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FA40	GPMB_ECOL6		cytoplasm
Q8FA41	YJXJ_ECOL6		inner membrane
Q8FA42	TRPR_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FA49	LPLA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FA51	DEOD_ECOL6		cytoplasm
Q8FA52	TYPH_ECOL6		cytoplasm
Q8FA72	DNAT_ECOL6		nucleoid
Q8FA73	OPGB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FAD1	SYV_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FAD6	OTC_ECOL6	cytoplasm (probable).	cytoplasm
Q8FAD9	ARCA2_ECOL6	cytoplasm (by similarity).	extracellular
Q8FAE1	OTCC_ECOL6	cytoplasm (probable).	cytoplasm
Q8FAF3	YJGA_ECOL6		cytoplasm
Q8FAG0	IPYR_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FAG4	MSRA_ECOL6		cytoplasm
Q8FAG5	CYSQ_ECOL6	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	cytoplasm
Q8FAH0	RCMNS_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FAI6	ULAF_ECOL6		cytoplasm
Q8FAI7	ULAE_ECOL6		cytoplasm
Q8FAI9	ULAD_ECOL6		cytoplasm
Q8FAJ0	ULAC_ECOL6	cytoplasm (probable).	cytoplasm
Q8FAJ1	ULAB_ECOL6	cytoplasm (potential).	cytoplasm
Q8FAJ3	ULAA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FAJ4	ULAG_ECOL6	cytoplasm (potential).	extracellular
Q8FAK9	MUTL_ECOL6		cytoplasm
Q8FAL3	ENG_C_ECOL6		cytoplasm
Q8FAL7	FRDD_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FAL8	SUGE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FAM7	CUTA_ECOL6	cytoplasm (by similarity).	periplasm
Q8FAV1	PHNC_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FAW3	YJDP_ECOL6		nucleoid

Q8FAX1	MDTN_ECOL6	bacterial cell inner membrane; single-pass type ii membrane protein (potential).	inner membrane
Q8FAX2	MDTO_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FAY8	ACSA_ECOL6		cytoplasm
Q8FAZ0	ACTP_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FB02	UVRA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FB20	ALR1_ECOL6		cytoplasm
Q8FB22	DNAB_ECOL6		cytoplasm
Q8FB30	DUSA_ECOL6		cytoplasm
Q8FB32	YJBJ_ECOL6		cytoplasm
Q8FB37	MALK_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FB38	MALF_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FB44	G6PI_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FB47	RLUF_ECOL6		cytoplasm
Q8FB55	PEPE_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FB61	ACEK_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FB64	META_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FB68	PUR9_ECOL6		cytoplasm
Q8FB69	PUR2_ECOL6		cytoplasm
Q8FB74	DCUP_ECOL6	cytoplasm (probable).	cytoplasm
Q8FB75	NUDC_ECOL6		cytoplasm
Q8FB77	THIC_ECOL6		cytoplasm
Q8FB78	THIE_ECOL6		cytoplasm
Q8FB81	THIG_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FB83	RPOC_ECOL6		cytoplasm
Q8FB88	MURB_ECOL6	cytoplasm (probable).	inner membrane
Q8FB89	MURI_ECOL6		cytoplasm
Q8FB93	STHA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FB96	ARLY_ECOL6	cytoplasm (probable).	cytoplasm
Q8FB97	ARGE_ECOL6	cytoplasm (probable).	extracellular
Q8FB98	CAPP_ECOL6		cytoplasm
Q8FBA7	FSAB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBC0	HSLU_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBC3	GLPK_ECOL6		cytoplasm
Q8FBC8	CDH_ECOL6	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8FBD0	K6PF1_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBD1	FIEF_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FBD6	RHAT_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FBD7	RHAR_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBD9	RHAB_ECOL6		cytoplasm
Q8FBE0	RHAA_ECOL6	cytoplasm (by similarity).	cytoplasm

Q8FBE1	RHAD_ECOL6	cytoplasm (by similarity).	extracellular
Q8FBE5	FDHD_ECOL6	cytoplasm (potential).	cytoplasm
Q8FBE8	FDHE_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBG9	YIHI_ECOL6		inner membrane
Q8FBH7	MOBA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBI2	FADB_ECOL6		cytoplasm
Q8FBI3	FADA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBI6	TATC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FBI7	TATB_ECOL6	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8FBI8	UBIB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FBJ0	UBIE_ECOL6		cytoplasm
Q8FBM1	METE_ECOL6		cytoplasm
Q8FBN6	DAPF_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBN9	CYAY_ECOL6		inner membrane
Q8FBP0	CYAA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBP1	HEM3_ECOL6		cytoplasm
Q8FBP7	WECG_ECOL6		cytoplasm
Q8FBP8	WZYE_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8FBP9	WECF_ECOL6	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8FBR0	GPPA_ECOL6		extracellular
Q8FBR2	ILVC_ECOL6		cytoplasm
Q8FBR5	ILVD_ECOL6		cytoplasm
Q8FBS5	RAVA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBS6	VIAA_ECOL6		nucleoid
Q8FBT3	GLMU_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBT4	GLMS_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FBU9	YIDZ_ECOL6		cytoplasm
Q8FBV0	MDTL_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FBV2	TNAA_ECOL6		cytoplasm
Q8FBV5	RNPA_ECOL6		extracellular
Q8FBW6	YIDE_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8FBX7	ADEC_ECOL6		cytoplasm
Q8FC86	SLMA_ECOL6	nucleoid-associated (by similarity).	cytoplasm
Q8FC87	FPG_ECOL6		cytoplasm
Q8FC88	COAD_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FCA0	HLDD_ECOL6		cytoplasm
Q8FCA2	TDH_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FCA6	GPMI_ECOL6		cytoplasm
Q8FCB1	LLDD_ECOL6		cytoplasm
Q8FCB7	MTLD_ECOL6		cytoplasm
Q8FCC1	SELA_ECOL6	cytoplasm (by similarity).	cytoplasm

Q8FCD6	DLGD_ECOL6	cytoplasm (potential).	cytoplasm
Q8FCE3	XYLA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FCI4	TREF_ECOL6	cytoplasm (by similarity).	extracellular
Q8FCI6	GADX_ECOL6		cytoplasm
Q8FCI7	GADW_ECOL6		cytoplasm
Q8FCI8	MDTF_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FCK2	DCTR_ECOL6		cytoplasm
Q8FCL1	YHIQ_ECOL6		inner membrane
Q8FCM9	NIKE_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FCN0	NIKD_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FCN5	YHHS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FCR6	DHAS_ECOL6		cytoplasm
Q8FCR7	GLGB_ECOL6		extracellular
Q8FCR8	GLGX_ECOL6		extracellular
Q8FCS8	RTCA_ECOL6	cytoplasm (potential).	cytoplasm
Q8FCT3	GNTX_ECOL6		cytoplasm
Q8FCT4	BIOH_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FCT7	FEOB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FCU4	PPCK_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FCV7	AROB_ECOL6	cytoplasm (probable).	cytoplasm
Q8FCW0	GPH_ECOL6		cytoplasm
Q8FCY0	KEFB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FCY1	TUSD_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FCY2	TUSC_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FCY3	TUSB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FD03	RL24_ECOL6		cytoplasm
Q8FD11	MSCL_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FD12	RSMB_ECOL6		cytoplasm
Q8FD13	FMT_ECOL6		cytoplasm
Q8FD15	SMG_ECOL6		cytoplasm
Q8FD18	AROE_ECOL6		cytoplasm
Q8FD47	YHDE_ECOL6	cytoplasm (potential).	inner membrane
Q8FD49	AAEX_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8FD50	AAEA_ECOL6	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q8FD51	AAEB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FD52	ARGR_ECOL6	cytoplasm (probable).	cytoplasm

Q8FD57	NANR_ECOL6		cytoplasm
Q8FD58	NANA1_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FD59	NANT1_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FD60	NANK1_ECOL6		cytoplasm
Q8FD68	MTGA_ECOL6	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8FD73	KDSD_ECOL6		cytoplasm
Q8FD84	GLMM_ECOL6		cytoplasm
Q8FD94	YHBQ_ECOL6		cytoplasm
Q8FD95	YHBP_ECOL6		inner membrane
Q8FD98	DIAA_ECOL6		cytoplasm
Q8FDA0	YRAN_ECOL6		inner membrane
Q8FDE1	ALX_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FDG5	PRIM_ECOL6		cytoplasm
Q8FDH4	GLNE_ECOL6		cytoplasm
Q8FDH5	HLDE_ECOL6		cytoplasm
Q8FDH9	RIBB_ECOL6		cytoplasm
Q8FDI3	DSBI_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8FDK9	YGIQ_ECOL6		cytoplasm
Q8FDL6	YQHA_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8FDM1	HYBF_ECOL6		cytoplasm
Q8FDN6	MASZ_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FDU7	NANA2_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FDU8	NANK2_ECOL6		cytoplasm
Q8FDU9	NANT2_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FDZ8	HLYB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8FE19	FETP_ECOL6		cytoplasm
Q8FE22	TRMB_ECOL6		cytoplasm
Q8FE27	HAM1_ECOL6		extracellular
Q8FE28	YGGU_ECOL6		inner membrane
Q8FE30	GSHB_ECOL6		cytoplasm
Q8FE31	SPRT_ECOL6	cytoplasm (potential).	extracellular
Q8FE34	SPEA_ECOL6	periplasm (by similarity).	cytoplasm
Q8FE36	SPEB_ECOL6		cytoplasm
Q8FE55	ARGO_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FE65	GCST_ECOL6		cytoplasm
Q8FE66	GCSH_ECOL6		cytoplasm
Q8FE67	GCSP_ECOL6		cytoplasm
Q8FE69	YQFB_ECOL6		inner membrane
Q8FE75	IDI_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FE91	YGEW_ECOL6		cytoplasm

Q8FE97	KDUI_ECOL6		cytoplasm
Q8FEA6	AAS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FEA9	MUTH_ECOL6	cytoplasm (by similarity).	extracellular
Q8FEE5	MTFA_ECOL6		cytoplasm
Q8FEE7	FUCI_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FEF0	FUCA_ECOL6		extracellular
Q8FEF7	QUEF_ECOL6	cytoplasm (probable).	cytoplasm
Q8FEF9	TRUC_ECOL6		cytoplasm
Q8FEG6	RUMA_ECOL6		cytoplasm
Q8FEI7	CYSJ_ECOL6		cytoplasm
Q8FEI8	CYSI_ECOL6		cytoplasm
Q8FEI9	CYSH_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FEJ0	CYSD_ECOL6		cytoplasm
Q8FEJ1	CYSN_ECOL6		cytoplasm
Q8FEJ2	CYSC_ECOL6		cytoplasm
Q8FEJ4	FTSB_ECOL6	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q8FEJ5	ISPD_ECOL6		cytoplasm
Q8FEJ6	ISPF_ECOL6		extracellular
Q8FEJ7	TRUD_ECOL6		cytoplasm
Q8FEJ8	PIMT_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FEL3	MUTS_ECOL6		cytoplasm
Q8FEL9	HYP_A_ECOL6		cytoplasm
Q8FEN4	NORW_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FEN6	NORR_ECOL6		cytoplasm
Q8FEP8	LUXS_ECOL6		extracellular
Q8FES6	CSID_ECOL6		cytoplasm
Q8FEY5	YFJF_ECOL6		inner membrane
Q8FEY7	PPNK_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FEY9	GRPE_ECOL6	cytoplasm (probable).	cytoplasm
Q8FF08	UNG_ECOL6	cytoplasm (by similarity).	extracellular
Q8FF10	GRCA_ECOL6		cytoplasm
Q8FF17	ERA_ECOL6		cytoplasm
Q8FF18	PDXJ_ECOL6	cytoplasm (probable).	cytoplasm
Q8FF19	ACPS_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FF24	TADA_ECOL6		cytoplasm
Q8FF30	HMP_ECOL6		cytoplasm
Q8FF44	HSCA_ECOL6		cytoplasm
Q8FF46	ISCX_ECOL6		cytoplasm
Q8FF47	PEPB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FF53	NDK_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FF59	ENGA_ECOL6		cytoplasm
Q8FF64	EX7L_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FF70	PPK_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FF72	PUR5_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FFA1	EUTC_ECOL6		cytoplasm
Q8FFA3	HEM6_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FFB0	YFEU_ECOL6		cytoplasm

Q8FFB3	CYSA_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FFB9	CYSZ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FFC0	ZIPA_ECOL6	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q8FFD3	YFEO_ECOL6	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8FFD4	GLK_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FFD5	PTFC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FFD8	PTFAX_ECOL6	cytoplasm (probable).	cytoplasm
Q8FFE0	YPDB_ECOL6		cytoplasm
Q8FFE8	FCTA_ECOL6		cytoplasm
Q8FFF4	SDHD_ECOL6		cytoplasm
Q8FFG3	FADI_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FFG4	FADJ_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FFH0	YFCK_ECOL6		cytoplasm
Q8FFH2	PDXB_ECOL6		cytoplasm
Q8FFJ2	YFBV_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FFJ5	YFBR_ECOL6	cytoplasm (potential).	periplasm
Q8FFJ9	NUOG_ECOL6		cytoplasm
Q8FFK8	RNZ_ECOL6		extracellular
Q8FFL7	YFBJ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FFL9	ARNT_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FFM1	ARNA_ECOL6		cytoplasm
Q8FFM2	ARNC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FFM3	ARNB_ECOL6		cytoplasm
Q8FFP0	UBIG_ECOL6		cytoplasm
Q8FFQ5	MQO_ECOL6		cytoplasm
Q8FFR2	CCMA_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FFR9	NDPA_ECOL6	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q8FFS1	RL25_ECOL6		cytoplasm
Q8FFT8	RIHB_ECOL6		extracellular
Q8FFU0	END4_ECOL6		extracellular
Q8FFU1	YEIH_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FFV3	CDD_ECOL6		cytoplasm

Q8FFV5	DUSC_ECOL6		cytoplasm
Q8FFV6	MDTQ_ECOL6	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
Q8FFX9	RCNA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FFY0	THIM_ECOL6		cytoplasm
Q8FFY2	YEGX_ECOL6		extracellular
Q8FG00	YEGP_ECOL6		extracellular
Q8FG03	MDTC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FG04	MDTB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8FG45	UDG_ECOL6		cytoplasm
Q8FG47	HIS2_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FG48	HIS6_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FG49	HIS4_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FG50	HIS7_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FG51	HIS8_ECOL6		cytoplasm
Q8FG52	HISX_ECOL6		cytoplasm
Q8FG53	YOEB_ECOL6		extracellular
Q8FGA4	COBS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FGA5	COBT_ECOL6		cytoplasm
Q8FGI6	YEDZ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FGI7	YEDY_ECOL6		cytoplasm
Q8FGL1	FLIE_ECOL6		flagellar
Q8FGN7	USPC_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FGP6	CHEY_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FGQ2	CUTC_ECOL6	cytoplasm (probable).	periplasm
Q8FGR3	RUVB_ECOL6		cytoplasm
Q8FGT3	PROQ_ECOL6	cytoplasm (potential).	inner membrane
Q8FGU0	YEBN_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FGZ9	ARGM_ECOL6		cytoplasm
Q8FH03	ASTE_ECOL6		periplasm
Q8FH06	NADE_ECOL6		cytoplasm
Q8FH10	CHBG_ECOL6		inner membrane
Q8FH26	BTUC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FH28	BTUD_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FH30	YDIU_ECOL6		inner membrane
Q8FH32	AROH_ECOL6		cytoplasm
Q8FH42	AROD_ECOL6		cytoplasm
Q8FH54	SUFS_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FH55	SUFE_ECOL6	cytoplasm (by similarity).	cytoplasm

Q8FH68	MDTK_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FH87	PDXH_ECOL6		cytoplasm
Q8FH88	SY Y_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FH91	TPPB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FH92	RNFE_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FH99	ADD_ECOL6		extracellular
Q8FHA7	FUMC_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FHA8	TUS_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FHB4	MDTJ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FHB5	MDTI_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FHC1	CLCB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FHC4	YNFC_ECOL6	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8FHC6	YNFA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FHD1	YDFH_ECOL6		cytoplasm
Q8FHD2	YDFG_ECOL6		cytoplasm
Q8FHE6	SOTB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FHF8	YDEP_ECOL6		periplasm
Q8FHF9	Y1929_ECOL6	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8FHG0	YDEO_ECOL6		cytoplasm
Q8FHG5	DCEB_ECOL6		cytoplasm
Q8FHG6	GADC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FHH1	MAO1_ECOL6		cytoplasm
Q8FHN0	ACPD_ECOL6		extracellular
Q8FHT6	RNB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FHU2	PYRF_ECOL6		cytoplasm
Q8FHU5	GCH2_ECOL6		cytoplasm
Q8FHV4	RLUB_ECOL6		cytoplasm
Q8FHV9	TRPB_ECOL6		cytoplasm
Q8FHW0	TRPA_ECOL6		cytoplasm
Q8FHW3	YCIC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FHX4	KITH_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FHX8	YCHJ_ECOL6		inner membrane
Q8FHZ8	KDSA_ECOL6	cytoplasm (by similarity).	cytoplasm

Q8FI03	HEM1_ECOL6		cytoplasm
Q8FI04	ISPE_ECOL6		cytoplasm
Q8FI27	HLYEL_ECOL6		cytoplasm
Q8FI28	YCGN_ECOL6		inner membrane
Q8FI31	MINC_ECOL6		inner membrane
Q8FIB6	RLUE_ECOL6		cytoplasm
Q8FIB7	YCFC_ECOL6		inner membrane
Q8FIB8	PHOQ_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FIM4	NPD_ECOL6	cytoplasm (probable).	cytoplasm
Q8FIM7	LOLD_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FIN2	NAGZ_ECOL6	cytoplasm (by similarity).	extracellular
Q8FIN3	THIK_ECOL6		cytoplasm
Q8FIN9	KTHY_ECOL6		cytoplasm
Q8FIP6	YCEF_ECOL6	cytoplasm (potential).	inner membrane
Q8FIP7	RLUC_ECOL6		cytoplasm
Q8FIP8	YCEQ_ECOL6		cytoplasm
Q8FIR2	PYRC_ECOL6		cytoplasm
Q8FIR3	MTOX_ECOL6		cytoplasm
Q8FIR7	YCEA_ECOL6		inner membrane
Q8FIR9	MDTG_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8FIS3	OPGH_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FJ50	CBPA_ECOL6	nucleoid (by similarity).	inner membrane
Q8FJ51	CBPM_ECOL6		cytoplasm
Q8FJ52	TORD_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJ69	TUSE_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJ74	MGSA_ECOL6		cytoplasm
Q8FJ81	YCBG_ECOL6		inner membrane
Q8FJ83	FABA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJ91	PYRD_ECOL6	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FJ93	SSUD_ECOL6		cytoplasm
Q8FJ98	PNCB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJA4	MUKF_ECOL6	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q8FJA9	KDSB_ECOL6	cytoplasm (potential).	cytoplasm
Q8FJB1	MSBA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FJB6	AROA_ECOL6	cytoplasm (probable).	cytoplasm
Q8FJB7	SERC_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJC1	YCAD_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FJD1	LFTR_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJE0	HCP_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJE7	RUMB_ECOL6		cytoplasm

Q8FJF0	RIMK_ECOL6		cytoplasm
Q8FJL3	FSAA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJM6	YBIX_ECOL6		cytoplasm
Q8FJN1	DING_ECOL6		cytoplasm
Q8FJN6	YBHG_ECOL6	periplasm (potential).	inner membrane
Q8FJN9	YBHO_ECOL6	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FJP8	UVRB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJR2	6PGL_ECOL6		cytoplasm
Q8FJR4	MODC_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FJS1	GAL1_ECOL6	cytoplasm (potential).	cytoplasm
Q8FJS7	NADA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FJU5	END8_ECOL6		cytoplasm
Q8FJU6	YBGL_ECOL6		inner membrane
Q8FJV3	ATKA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8FJV4	ATKB_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8FJV5	ATKC_ECOL6	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8FJX7	NAGB_ECOL6		extracellular
Q8FJY3	YBEY_ECOL6		inner membrane
Q8FJY4	LNT_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FJY8	RIHA_ECOL6		extracellular
Q8FJZ1	NADD_ECOL6		cytoplasm
Q8FJZ8	DCUC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FK03	CITX_ECOL6		cytoplasm
Q8FK04	CITG_ECOL6		cytoplasm
Q8FK07	USPG_ECOL6		cytoplasm
Q8FK20	ENTS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FK30	CAAL_ECOL6		cytoplasm
Q8FK36	CUSA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8FK37	CUSS_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FK47	LPXH_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FK51	ARCC_ECOL6	cytoplasm (potential).	cytoplasm
Q8FK65	ALLA_ECOL6		extracellular
Q8FK67	SELU_ECOL6		cytoplasm
Q8FK74	CUER_ECOL6	cytoplasm (probable).	cytoplasm
Q8FK76	GLSA1_ECOL6		extracellular
Q8FK83	HEMH_ECOL6	cytoplasm (by similarity).	extracellular

Q8FK84	KAD_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FKA7	TIG_ECOL6		cytoplasm
Q8FKB7	THII_ECOL6	cytoplasm (potential).	cytoplasm
Q8FKB9	DXS_ECOL6		cytoplasm
Q8FKC5	QUEA_ECOL6	cytoplasm (potential).	cytoplasm
Q8FKC6	ACPH_ECOL6		cytoplasm
Q8FKD5	RDGC_ECOL6	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q8FKE3	DDLA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FKF5	TAUB_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FKI7	BETI_ECOL6		cytoplasm
Q8FKI8	BETB_ECOL6		cytoplasm
Q8FKI9	BETA_ECOL6	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FKM3	PROA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FKM5	YAFA_ECOL6		cytoplasm
Q8FKM7	XGPT_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FKZ1	GMHB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FKZ8	YAEP_ECOL6		inner membrane
Q8FL00	TILS_ECOL6	cytoplasm (probable).	cytoplasm
Q8FL03	ACCA_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FL06	RNH2_ECOL6	cytoplasm (potential).	extracellular
Q8FL07	LPXB_ECOL6		cytoplasm
Q8FL11	DAPD_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FL13	DGTP_ECOL6		cytoplasm
Q8FL15	CLCA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8FL16	GSA_ECOL6	cytoplasm (potential).	cytoplasm
Q8FL24	SFSA_ECOL6		cytoplasm
Q8FL25	GLUQ_ECOL6		cytoplasm
Q8FL30	PANB_ECOL6		cytoplasm
Q8FL31	PANC_ECOL6		cytoplasm
Q8FL39	SPED_ECOL6		cytoplasm
Q8FL40	YAQL_ECOL6		inner membrane
Q8FL49	AROP_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FL55	COAE_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FL56	YACF_ECOL6		inner membrane
Q8FL57	YACG_ECOL6		cytoplasm
Q8FL63	DDLB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FL64	MURG_ECOL6	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FL65	MURD_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FL67	MURE_ECOL6	cytoplasm (probable).	cytoplasm
Q8FL68	MRAW_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FL75	LEU1_ECOL6		cytoplasm
Q8FL76	LEU3_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FL79	LEUD_ECOL6		cytoplasm
Q8FL88	ARAB_ECOL6		cytoplasm

Q8FL89	ARAA_ECOL6		cytoplasm
Q8FL92	RAPA_ECOL6		cytoplasm
Q8FL93	RLUA_ECOL6		cytoplasm
Q8FL94	DJLA_ECOL6	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q8FL95	PDXA_ECOL6	cytoplasm (probable).	cytoplasm
Q8FL96	KSGA_ECOL6		cytoplasm
Q8FL98	APAH_ECOL6		extracellular
Q8FLA1	KEFC_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FLA3	FIXA_ECOL6		cytoplasm
Q8FLA4	CAIB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FLA5	CAIC_ECOL6		cytoplasm
Q8FLA6	CAID_ECOL6		cytoplasm
Q8FLA7	CAIE_ECOL6		cytoplasm
Q8FLB0	CARB_ECOL6		cytoplasm
Q8FLB1	CARA_ECOL6		cytoplasm
Q8FLB4	DAPB_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FLB6	LSPA_ECOL6	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8FLB7	SYI_ECOL6	cytoplasm (by similarity).	cytoplasm
Q8FLC5	DNAJ_ECOL6	cytoplasm (probable).	cytoplasm
Q8FLD1	TALB_ECOL6	cytoplasm (probable).	cytoplasm
Q8FLD3	YAAA_ECOL6		inner membrane
Q8FLD8	KHSE_ECOL6	cytoplasm (potential).	cytoplasm
Q8FUQ2	Y4168_BRUSU		inner membrane
Q8FUS3	FLIF_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FUS9	FLGE_BRUSU		flagellar
Q8FUT3	FLBT_BRUSU		cytoplasm
Q8FUZ0	MSRA_BRUSU		cytoplasm
Q8FUZ1	SLYX_BRUSU		inner membrane
Q8FUZ2	UPP_BRUSU		cytoplasm
Q8FUZ3	UBIE_BRUSU		cytoplasm
Q8FV07	DAPB_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FV09	GLK_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FV17	LEPA_BRUSU		cytoplasm
Q8FV25	DAPD_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FV29	OXAA_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FV30	RNPA_BRUSU		extracellular
Q8FV59	Y3991_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FVB2	HUTU_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FVB4	HUTH_BRUSU	cytoplasm (potential).	cytoplasm
Q8FVB5	HUTI_BRUSU	cytoplasm (potential).	cytoplasm
Q8FVC0	DADA_BRUSU		cytoplasm
Q8FVC1	ALR_BRUSU		cytoplasm
Q8FVD4	Y3910_BRUSU		cytoplasm
Q8FVE4	OCD_BRUSU		cytoplasm

Q8FVF3	LEU3_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FVH2	TPIS2_BRUSU		cytoplasm
Q8FVL9	CRCB4_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FVM0	CRCB3_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FVM2	UXUA_BRUSU		cytoplasm
Q8FVM4	UXAC_BRUSU		cytoplasm
Q8FVM9	NIKD_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FVN0	NIKE_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FVR1	EX7L_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FVU9	GCSP_BRUSU		cytoplasm
Q8FVV5	FBPC_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FVX5	SP39_BRUSU	periplasm (probable).	periplasm
Q8FVY1	RRMJ_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FW12	ADEC_BRUSU		cytoplasm
Q8FW50	DPO4_BRUSU	cytoplasm (probable).	cytoplasm
Q8FW65	RUVX_BRUSU	cytoplasm (potential).	cytoplasm
Q8FW67	GATA_BRUSU		cytoplasm
Q8FW72	LIPB_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FW78	GLMU_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FW95	BFR_BRUSU		cytoplasm
Q8FWG5	BIOD_BRUSU		cytoplasm
Q8FWG9	META_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FWK8	GLPK_BRUSU		cytoplasm
Q8FWN5	NANEK_BRUSU		cytoplasm
Q8FWR0	PHK_BRUSU		cytoplasm
Q8FWT4	GUAA_BRUSU		cytoplasm
Q8FWU0	CATA_BRUSU	periplasm (by similarity).	periplasm
Q8FWV5	GLSA_BRUSU		extracellular
Q8FWX2	MINC_BRUSU		inner membrane
Q8FX16	NOSZ_BRUSU	periplasm (by similarity).	periplasm
Q8FX81	SYI_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FX87	CH60_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FX90	FUMC_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FX92	HIS1_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FXC2	FLGH_BRUSU	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8FXI7	MODC_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FXJ1	Y3086_BRUSU		inner membrane
Q8FXR6	FPG_BRUSU		cytoplasm
Q8FXT0	RS15_BRUSU		cytoplasm
Q8FXT1	RBFA_BRUSU	cytoplasm (potential).	cytoplasm
Q8FXT6	TRMB_BRUSU		cytoplasm

Q8FXT9	LNT_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FXW6	GSHB_BRUSU		cytoplasm
Q8FXW9	PYRF_BRUSU		cytoplasm
Q8FXX1	DNAJ_BRUSU	cytoplasm (probable).	cytoplasm
Q8FXX2	DNAK_BRUSU		cytoplasm
Q8FXY4	TRPB_BRUSU		cytoplasm
Q8FXY6	TRPA_BRUSU		cytoplasm
Q8FXZ7	SAHH_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY05	PPCK_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY07	HIS6_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY08	HIS4_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY09	HIS5_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY11	HSLV_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY12	HSLU_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY19	SECB_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY21	AROE_BRUSU		cytoplasm
Q8FY22	Y2068_BRUSU	cytoplasm (potential).	inner membrane
Q8FY24	DCUP_BRUSU	cytoplasm (probable).	cytoplasm
Q8FY28	GIDA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY29	GIDB_BRUSU		cytoplasm
Q8FY57	ACCA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FY59	AROK_BRUSU	cytoplasm (probable).	cytoplasm
Q8FY85	AQPZ_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FY98	HIS8_BRUSU		cytoplasm
Q8FYA4	ARLY_BRUSU	cytoplasm (probable).	cytoplasm
Q8FYA6	Y1979_BRUSU		inner membrane
Q8FYE2	ARGJ_BRUSU	cytoplasm (probable).	cytoplasm
Q8FYF0	DAPF_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FYF4	MDH_BRUSU		cytoplasm
Q8FYF7	ODO1_BRUSU		cytoplasm
Q8FYG9	LEU2_BRUSU		cytoplasm
Q8FYI6	HEM3_BRUSU		cytoplasm
Q8FYK0	UBIG_BRUSU		cytoplasm
Q8FYL8	RL27_BRUSU		cytoplasm
Q8FYM3	PROA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FYM9	NUDH_BRUSU		cytoplasm
Q8FYQ1	HTPX_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8FYQ3	ACSA_BRUSU		cytoplasm
Q8FYQ9	TAL_BRUSU	cytoplasm (probable).	cytoplasm
Q8FYR3	ATPA_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FYR4	ATPG_BRUSU		inner membrane
Q8FYT2	ISPG_BRUSU		cytoplasm
Q8FYT4	MTGA_BRUSU	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8FYV6	Y1752_BRUSU		inner membrane
Q8FYW3	PUR6_BRUSU		cytoplasm
Q8FYX8	PGK_BRUSU	cytoplasm (potential).	cytoplasm

Q8FZ01	RUVC_BRUSU		cytoplasm
Q8FZ02	RUVB_BRUSU		cytoplasm
Q8FZ07	TOLB_BRUSU	periplasm (potential).	periplasm
Q8FZ11	TILS_BRUSU	cytoplasm (probable).	cytoplasm
Q8FZ13	GLMM_BRUSU		cytoplasm
Q8FZ45	FUR_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FZ50	MASZ_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FZC4	LEU1_BRUSU		cytoplasm
Q8FZE8	RL25_BRUSU		cytoplasm
Q8FZF0	KPRS_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FZF5	LGT_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FZJ3	CARB_BRUSU		cytoplasm
Q8FZJ8	CARA_BRUSU		cytoplasm
Q8FZK4	Y1477_BRUSU		inner membrane
Q8FZM5	TAM_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FZP0	MURE_BRUSU	cytoplasm (probable).	cytoplasm
Q8FZP2	MURD_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FZP4	MURB_BRUSU	cytoplasm (probable).	inner membrane
Q8FZP5	DDLB_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FZP9	LPXC_BRUSU		cytoplasm
Q8FZQ6	PRMA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FZR2	Y1411_BRUSU		inner membrane
Q8FZT6	PDXJ_BRUSU	cytoplasm (probable).	cytoplasm
Q8FZT8	KUP_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FZU1	ILVC_BRUSU		cytoplasm
Q8FZV0	CRCB2_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FZV1	CRCB1_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8FZV2	CBIO_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8FZW2	URE12_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FZW3	URE22_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8FZW4	URE32_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G005	COBQ_BRUSU		cytoplasm
Q8G018	CBID_BRUSU		cytoplasm
Q8G020	COBB_BRUSU		cytoplasm
Q8G022	COBD_BRUSU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8G034	SFSA_BRUSU		cytoplasm
Q8G044	DDLA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G065	RL11_BRUSU		cytoplasm
Q8G066	RL1_BRUSU		cytoplasm
Q8G067	RL10_BRUSU		cytoplasm
Q8G069	RPOB_BRUSU		cytoplasm
Q8G070	RPOC_BRUSU		cytoplasm
Q8G074	RS7_BRUSU		cytoplasm
Q8G076	RL3_BRUSU		cytoplasm

Q8G077	RL4_BRUSU		cytoplasm
Q8G079	RL2_BRUSU		cytoplasm
Q8G080	RL22_BRUSU		cytoplasm
Q8G081	RL16_BRUSU		cytoplasm
Q8G082	RS17_BRUSU		cytoplasm
Q8G084	RL24_BRUSU		cytoplasm
Q8G085	RL5_BRUSU		cytoplasm
Q8G086	RS8_BRUSU		cytoplasm
Q8G088	RL18_BRUSU		cytoplasm
Q8G090	RL15_BRUSU		cytoplasm
Q8G092	KAD_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G094	RPOA_BRUSU		cytoplasm
Q8G0B6	QUEF_BRUSU	cytoplasm (probable).	cytoplasm
Q8G0C9	CLPS_BRUSU		cytoplasm
Q8G0D6	RS2_BRUSU		cytoplasm
Q8G0D7	PYRH_BRUSU		cytoplasm
Q8G0D8	RRF_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0D9	UPPS_BRUSU		inner membrane
Q8G0E0	CDSA_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8G0E1	Y1156_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8G0E4	FABZ_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0F1	LEXA_BRUSU		cytoplasm
Q8G0F3	MOAC_BRUSU		cytoplasm
Q8G0F5	TRPD_BRUSU		cytoplasm
Q8G0F7	TPIS1_BRUSU		cytoplasm
Q8G0G1	PYRG_BRUSU		cytoplasm
Q8G0G2	KDSA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0G3	ENO_BRUSU	cytoplasm (by similarity).	extracellular
Q8G0H4	ISPDF_BRUSU		cytoplasm
Q8G0I4	CLPP_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0I5	CLPX_BRUSU		cytoplasm
Q8G0I7	LON_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0I9	UVRA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0J1	SSB_BRUSU		cytoplasm
Q8G0K1	TGT_BRUSU		cytoplasm
Q8G0L3	HIS3_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0L4	GCH1_BRUSU		cytoplasm
Q8G0L6	Y1073_BRUSU		inner membrane
Q8G0P0	WRBA_BRUSU		cytoplasm
Q8G0P3	XGPT_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0Q5	Y1033_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8G0S7	RPIA_BRUSU		cytoplasm
Q8G0U3	KTHY_BRUSU		cytoplasm
Q8G0X4	MOAA_BRUSU		cytoplasm
Q8G0X5	MOBA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0Y3	SYV_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G0Z4	PPNK_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G105	SYV_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G120	AROQ_BRUSU		cytoplasm
Q8G129	TIG_BRUSU		cytoplasm

Q8G155	COBT_BRUSU		cytoplasm
Q8G156	COBS_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8G178	PUR7_BRUSU		cytoplasm
Q8G180	PURQ_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G183	PURL_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G195	LOLD_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8G1C9	RS9_BRUSU		cytoplasm
Q8G1E0	FABH_BRUSU	cytoplasm (probable).	cytoplasm
Q8G1E1	PLSX_BRUSU		cytoplasm
Q8G1E6	HPPA_BRUSU	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8G1E7	NUSB_BRUSU		cytoplasm
Q8G1E8	RISB1_BRUSU		cytoplasm
Q8G1F1	GLYA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G1K5	PUR5_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G1L5	UVRC_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G1L9	NDK_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G1M4	AMPA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G1N0	KSGA_BRUSU		cytoplasm
Q8G1P8	RECO_BRUSU		cytoplasm
Q8G1P9	ERA_BRUSU		cytoplasm
Q8G1Q6	RPOZ_BRUSU		cytoplasm
Q8G1R0	DAPA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G1U0	GLNE_BRUSU		cytoplasm
Q8G1U5	CCME_BRUSU	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8G1Z7	BETI_BRUSU		cytoplasm
Q8G1Z8	BETA_BRUSU		cytoplasm
Q8G1Z9	BETB_BRUSU		cytoplasm
Q8G204	XYLA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G228	Y508_BRUSU		cytoplasm
Q8G253	Y480_BRUSU		inner membrane
Q8G256	KHSE_BRUSU		cytoplasm
Q8G257	ISPH_BRUSU		cytoplasm
Q8G261	COXZ_BRUSU		inner membrane
Q8G274	RS6_BRUSU		cytoplasm
Q8G276	RL9_BRUSU		cytoplasm
Q8G285	PSD_BRUSU		cytoplasm
Q8G290	Y438_BRUSU		cytoplasm
Q8G292	DXS_BRUSU		cytoplasm
Q8G2A9	PDXH_BRUSU		cytoplasm
Q8G2B1	PUR2_BRUSU		cytoplasm
Q8G2D0	ISPE_BRUSU		cytoplasm
Q8G2E8	ENGA_BRUSU		cytoplasm
Q8G2I1	NORM_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8G2J0	PANC_BRUSU		cytoplasm

Q8G2K1	MSCL_BRUSU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8G2L5	APAG_BRUSU		inner membrane
Q8G2L7	OTC_BRUSU	cytoplasm (probable).	cytoplasm
Q8G2N3	G6PI_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G2P7	UREE_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G2P8	URE11_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G2P9	URE21_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G2Q0	URE31_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G2R2	HISX_BRUSU		cytoplasm
Q8G2R6	Y248_BRUSU	cytoplasm (potential).	inner membrane
Q8G2U9	THIG_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G2Y5	RNPH_BRUSU		cytoplasm
Q8G2Y6	GRPE_BRUSU	cytoplasm (probable).	cytoplasm
Q8G304	IHFB_BRUSU		cytoplasm
Q8G310	MUTS_BRUSU		cytoplasm
Q8G312	GLND_BRUSU		cytoplasm
Q8G340	PNCB_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G353	ILVD_BRUSU		cytoplasm
Q8G358	CCMA_BRUSU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8G376	ASSY_BRUSU	cytoplasm (probable).	cytoplasm
Q8G381	DNAE2_BRUSU	cytoplasm (probable).	cytoplasm
Q8G3B7	RECR_BRUSU		cytoplasm
Q8G3C1	Y028_BRUSU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8G3C3	KCY_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G3C4	ARO_A_BRUSU	cytoplasm (probable).	cytoplasm
Q8G3E5	RECF_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G3E7	DNAA_BRUSU	cytoplasm (by similarity).	cytoplasm
Q8G969	HCP_PHOPO	cytoplasm (by similarity).	cytoplasm
Q8G9C6	TRMB_AZOSE		cytoplasm
Q8G9G3	NUSB_VIBFI		cytoplasm
Q8G9G4	RISB_VIBFI		cytoplasm
Q8G9P0	MERB_PSEPU		inner membrane
Q8GAV9	CPNA_COMS9		cytoplasm
Q8GAW0	CPNB_COMS9		cytoplasm
Q8GAZ4	HMP_BURST		cytoplasm
Q8GB16	CAIE_PROSL		cytoplasm
Q8GB17	CAID_PROSL		cytoplasm
Q8GB18	CAIC_PROSL		cytoplasm
Q8GB19	CAIB_PROSL	cytoplasm (by similarity).	cytoplasm
Q8GB20	CAIA_PROSL		cytoplasm
Q8GBB4	CH60_AZOVI	cytoplasm (by similarity).	cytoplasm
Q8GBD2	CH60_ACEAC	cytoplasm (by similarity).	cytoplasm
Q8GBD3	CH10_ACEAC	cytoplasm (by similarity).	cytoplasm
Q8GBD4	GBPA_YEREN	secreted protein (by similarity).	extracellular
Q8GCY1	ODO2_BARVB		inner membrane
Q8GDE1	CBID_RHORT		cytoplasm
Q8GE63	ENO_AERHY	cytoplasm (by similarity).	extracellular
Q8GEI4	THIG_ERWPY	cytoplasm (by similarity).	cytoplasm
Q8GEK8	Y264_METEX		inner membrane

Q8GF77	RNH_PHOLU	cytoplasm (potential).	extracellular
Q8GF87	PGK_PHOLU	cytoplasm (potential).	cytoplasm
Q8GH23	RS12_BRUME		cytoplasm
Q8GH95	UREE_HELBI	cytoplasm (by similarity).	cytoplasm
Q8GH98	URE23_HELBI	cytoplasm (by similarity).	cytoplasm
Q8GHA0	GIDB_HELBI		cytoplasm
Q8GHM3	BDBC_PSERE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8GHV6	NDPA_PSERE	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q8GJK1	TRME_SHEPU		cytoplasm
Q8GLE4	HSCA_XENNE		cytoplasm
Q8GLE5	HSCB_XENNE		inner membrane
Q8GLE6	ISCA_XENNE		cytoplasm
Q8GLH3	Y2295_PHOTE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8GLH5	Y2291_PHOTE		cytoplasm
Q8GLU7	HLDE_ACTPL		cytoplasm
Q8GNC1	K6PF_HAEDU	cytoplasm (by similarity).	cytoplasm
Q8GNH6	NODI2_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8GNM0	MDH_FRANO		cytoplasm
Q8GNS3	FEOB_LEGPN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8GNT2	DKE1_ACIJO		cytoplasm
Q8GP19	RSSA_SERMA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8GP20	RSSB_SERMA	cytoplasm (by similarity).	cytoplasm
Q8GPG2	DDHD_RHOSU	cytoplasm (probable).	cytoplasm
Q8GQ92	CCMA2_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8GQ96	CCME2_PSEAE	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8GQC5	GLGB_ERWCH		extracellular
Q8K900	CYAY_BUCAP		inner membrane
Q8K901	DAPF_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K902	Y567_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K903	PIT_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K904	Y565_BUCAP		inner membrane
Q8K905	Y564_BUCAP		cytoplasm
Q8K906	HSLV_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K907	RL31_BUCAP		cytoplasm
Q8K908	AMIB_BUCAP		extracellular

Q8K909	ORN_BUCAP	cytoplasm (potential).	cytoplasm
Q8K910	G6PI_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K911	PTM3C_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K912	MTLD_BUCAP		cytoplasm
Q8K913	MUTL_BUCAP		cytoplasm
Q8K914	HFLK_BUCAP		inner membrane
Q8K915	HFLC_BUCAP		extracellular
Q8K916	PURA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K917	RNR_BUCAP		cytoplasm
Q8K918	RS6_BUCAP		cytoplasm
Q8K919	RS18_BUCAP		cytoplasm
Q8K920	RL9_BUCAP		cytoplasm
Q8K921	DUT_BUCAP		extracellular
Q8K922	Y538_BUCAP		inner membrane
Q8K923	EX1_BUCAP		cytoplasm
Q8K924	MURI_BUCAP		cytoplasm
Q8K925	FETP_BUCAP		cytoplasm
Q8K926	MUTY_BUCAP		cytoplasm
Q8K927	TRMB_BUCAP		cytoplasm
Q8K928	Y532_BUCAP		cytoplasm
Q8K929	Y531_BUCAP		cytoplasm
Q8K930	RUVX_BUCAP	cytoplasm (potential).	cytoplasm
Q8K931	GSHB_BUCAP		cytoplasm
Q8K932	DNAB_BUCAP		cytoplasm
Q8K933	SSB_BUCAP		cytoplasm
Q8K934	Y525_BUCAP		inner membrane
Q8K936	DEOB_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K937	DEOD_BUCAP		cytoplasm
Q8K938	AROK_BUCAP	cytoplasm (probable).	cytoplasm
Q8K939	AROB_BUCAP	cytoplasm (probable).	cytoplasm
Q8K940	RPE_BUCAP		cytoplasm
Q8K942	TSGA_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K943	FKBA_BUCAP		cytoplasm
Q8K944	TUSD_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K945	TUSC_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K946	TUSB_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K947	RS12_BUCAP		cytoplasm
Q8K949	RS10_BUCAP		cytoplasm
Q8K950	RL3_BUCAP		cytoplasm
Q8K951	RL4_BUCAP		cytoplasm
Q8K952	RL23_BUCAP		cytoplasm
Q8K953	RL2_BUCAP		cytoplasm
Q8K954	RS19_BUCAP		cytoplasm
Q8K955	RL22_BUCAP		cytoplasm
Q8K956	RS3_BUCAP		cytoplasm
Q8K957	RL16_BUCAP		cytoplasm
Q8K958	RL29_BUCAP		cytoplasm
Q8K959	RS17_BUCAP		cytoplasm
Q8K960	RL14_BUCAP		cytoplasm
Q8K961	RL24_BUCAP		cytoplasm
Q8K962	RL5_BUCAP		cytoplasm
Q8K963	RS14_BUCAP		cytoplasm

Q8K964	RL6_BUCAP		cytoplasm
Q8K965	RL18_BUCAP		cytoplasm
Q8K966	RS5_BUCAP		cytoplasm
Q8K967	RL30_BUCAP		cytoplasm
Q8K968	RL15_BUCAP		cytoplasm
Q8K971	RS13_BUCAP		cytoplasm
Q8K972	RS11_BUCAP		cytoplasm
Q8K973	RL17_BUCAP		cytoplasm
Q8K974	FMT_BUCAP		cytoplasm
Q8K975	DEF_BUCAP		cytoplasm
Q8K976	SMG_BUCAP		cytoplasm
Q8K977	Y475_BUCAP		inner membrane
Q8K978	Y472_BUCAP		inner membrane
Q8K979	FOLD_BUCAP		cytoplasm
Q8K980	KAD_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K981	HTPG_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K982	Y467_BUCAP		inner membrane
Q8K983	DPO3X_BUCAP		cytoplasm
Q8K984	MDLB_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K985	MDLA_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K986	Y463_BUCAP		inner membrane
Q8K987	PPID_BUCAP	bacterial cell membrane; single-pass type ii membrane protein (potential).	outer membrane
Q8K989	CLPX_BUCAP		cytoplasm
Q8K990	CLPP_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K991	TIG_BUCAP		cytoplasm
Q8K992	Y457_BUCAP		cytoplasm
Q8K993	CYOA_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K994	CYOB_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K995	CYOC_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K996	CYOD_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K997	CYOE_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K998	TUSE_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K999	Y450_BUCAP	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8K9A0	ISPA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9A1	DXS_BUCAP		cytoplasm
Q8K9A2	NUSB_BUCAP		cytoplasm
Q8K9A3	RIBD2_BUCAP		cytoplasm
Q8K9A4	RIBD1_BUCAP		cytoplasm

Q8K9A5	THIL_BUCAP		cytoplasm
Q8K9A6	RISB_BUCAP		cytoplasm
Q8K9A8	EX5A_BUCAP		cytoplasm
Q8K9A9	EX5B_BUCAP		cytoplasm
Q8K9B0	EX5C_BUCAP		cytoplasm
Q8K9B1	Y437_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9B2	ALF_BUCAP		cytoplasm
Q8K9B3	PGK_BUCAP	cytoplasm (potential).	cytoplasm
Q8K9B4	Y434_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9B5	DHAS_BUCAP		cytoplasm
Q8K9B6	TUSA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9B7	NADD_BUCAP		cytoplasm
Q8K9B8	HOLA_BUCAP		cytoplasm
Q8K9C0	CORC_BUCAP		inner membrane
Q8K9C1	Y427_BUCAP		inner membrane
Q8K9C2	Y426_BUCAP		cytoplasm
Q8K9C3	TYSY_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9C4	DCDA_BUCAP		cytoplasm
Q8K9C6	Y420_BUCAP		cytoplasm
Q8K9C7	KGUA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9C8	TYPB_BUCAP		cytoplasm
Q8K9C9	ENGB_BUCAP		cytoplasm
Q8K9D0	EX53_BUCAP		cytoplasm
Q8K9D1	DSBA_BUCAP		periplasm
Q8K9D2	MUTS_BUCAP		cytoplasm
Q8K9D3	CYSJ_BUCAP		cytoplasm
Q8K9D4	CYSC_BUCAP		cytoplasm
Q8K9D6	ISPD_BUCAP		cytoplasm
Q8K9D7	ISPF_BUCAP		extracellular
Q8K9D8	PTPS_BUCAP		extracellular
Q8K9D9	Y401_BUCAP		cytoplasm
Q8K9E0	ENO_BUCAP	cytoplasm (by similarity).	extracellular
Q8K9E2	RPIA_BUCAP		cytoplasm
Q8K9E3	Y396_BUCAP		cytoplasm
Q8K9E4	Y395_BUCAP		cytoplasm
Q8K9E6	CSRA_BUCAP		cytoplasm
Q8K9E8	Y389_BUCAP		cytoplasm
Q8K9E9	RLUD_BUCAP		cytoplasm
Q8K9F0	FIS_BUCAP		cytoplasm
Q8K9F1	AROQ_BUCAP		cytoplasm
Q8K9F2	TLDD_BUCAP		cytoplasm
Q8K9F3	RL19_BUCAP		cytoplasm
Q8K9F4	TRMD_BUCAP	cytoplasm (potential).	cytoplasm
Q8K9F5	RIMM_BUCAP	cytoplasm (potential).	cytoplasm
Q8K9F6	RS16_BUCAP		cytoplasm
Q8K9F7	SRP54_BUCAP		cytoplasm
Q8K9F9	RL13_BUCAP		cytoplasm
Q8K9G0	RS9_BUCAP		cytoplasm
Q8K9G1	Y376_BUCAP		cytoplasm
Q8K9G2	RL27_BUCAP		cytoplasm
Q8K9G3	RL21_BUCAP		cytoplasm
Q8K9G4	MURA_BUCAP	cytoplasm (probable).	inner membrane

Q8K9G5	Y372_BUCAP		cytoplasm
Q8K9G6	GREB_BUCAP		cytoplasm
Q8K9G7	RRMJ_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9G8	FTSH_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9G9	SECG_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K9H0	NUSA_BUCAP		cytoplasm
Q8K9H2	RBFA_BUCAP	cytoplasm (potential).	cytoplasm
Q8K9H3	TRUB_BUCAP		cytoplasm
Q8K9H4	RS15_BUCAP		cytoplasm
Q8K9H5	PNP_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9H6	DEAD_BUCAP	cytoplasm (probable).	cytoplasm
Q8K9H7	Y359_BUCAP		cytoplasm
Q8K9H8	PYRI_BUCAP		cytoplasm
Q8K9H9	PYRB_BUCAP		cytoplasm
Q8K9I0	AMPA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9I1	SYV_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9I2	Y353_BUCAP		inner membrane
Q8K9I3	UUP_BUCAP		inner membrane
Q8K9I4	Y351_BUCAP		cytoplasm
Q8K9I5	PYRD_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8K9I6	PNCB_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9I8	PORL_BUCAP		outer membrane
Q8K9I9	YHIT_BUCAP		cytoplasm
Q8K9J0	PTGCB_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K9J1	Y343_BUCAP		extracellular
Q8K9J2	HOLB_BUCAP		cytoplasm
Q8K9J3	KTHY_BUCAP		cytoplasm
Q8K9J4	ACP_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9J5	FABG_BUCAP		cytoplasm
Q8K9J6	FABD_BUCAP		cytoplasm
Q8K9J7	RL32_BUCAP		cytoplasm
Q8K9J8	RLUC_BUCAP		cytoplasm
Q8K9J9	RNE_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9K0	FLGK_BUCAP		flagellar
Q8K9K1	FLGJ_BUCAP		inner membrane
Q8K9K2	FLGI_BUCAP		periplasm
Q8K9K3	FLGH_BUCAP	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8K9K4	FLGG_BUCAP		flagellar
Q8K9K5	FLGF_BUCAP		flagellar
Q8K9K6	FLGE_BUCAP		flagellar
Q8K9K7	FLGD_BUCAP		cytoplasm
Q8K9K8	FLGC_BUCAP		flagellar
Q8K9K9	FLGB_BUCAP		flagellar
Q8K9L0	FLGA_BUCAP	periplasm (probable).	flagellar
Q8K9L1	FLGN_BUCAP		flagellar
Q8K9L2	PYRC_BUCAP		cytoplasm

Q8K9L3	MVIN_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K9L5	RSMC_BUCAP		cytoplasm
Q8K9L6	MINC_BUCAP		inner membrane
Q8K9L7	MIND_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8K9L8	MINE_BUCAP		inner membrane
Q8K9L9	Y315_BUCAP		cytoplasm
Q8K9M0	Y314_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9M1	HTPX_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	extracellular
Q8K9M2	G6PD_BUCAP		cytoplasm
Q8K9M3	KPYK_BUCAP		cytoplasm
Q8K9M4	Y310_BUCAP		extracellular
Q8K9M5	ZNUA_BUCAP		periplasm
Q8K9M6	ZNUC_BUCAP		inner membrane
Q8K9M7	ZNUB_BUCAP	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8K9M9	GLPF_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K9N0	K6PF_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9N1	GPMA_BUCAP		cytoplasm
Q8K9N2	ODO2_BUCAP		inner membrane
Q8K9N3	ODO1_BUCAP		cytoplasm
Q8K9N4	Y291_BUCAP		cytoplasm
Q8K9N5	FLAV_BUCAP		cytoplasm
Q8K9N6	QUEF_BUCAP	cytoplasm (probable).	cytoplasm
Q8K9N7	Y286_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9N8	LOLC_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K9N9	6PGL_BUCAP		cytoplasm
Q8K9P0	BIOA_BUCAP		cytoplasm
Q8K9P1	BIOB_BUCAP		cytoplasm
Q8K9P2	GLYA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9P4	ISPG_BUCAP		cytoplasm
Q8K9P5	Y275_BUCAP		cytoplasm
Q8K9P6	SUHB_BUCAP		extracellular
Q8K9P7	TOP1_BUCAP		cytoplasm
Q8K9P8	SOHB_BUCAP		extracellular
Q8K9P9	CLS_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K9Q0	GCH2_BUCAP		extracellular
Q8K9Q1	PYRF_BUCAP		cytoplasm
Q8K9Q2	LIPA_BUCAP	cytoplasm (potential).	cytoplasm
Q8K9Q3	LIPB_BUCAP	cytoplasm (by similarity).	cytoplasm

Q8K9Q4	Y257_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9Q5	RNB_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9Q6	FABI_BUCAP		cytoplasm
Q8K9Q7	PUR8_BUCAP		cytoplasm
Q8K9Q8	TRMU_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9Q9	LEPA_BUCAP		cytoplasm
Q8K9R0	LEP_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K9R1	RNC_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9R2	ERA_BUCAP		cytoplasm
Q8K9R3	ACPS_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9R4	Y246_BUCAP		cytoplasm
Q8K9R5	SSRP_BUCAP	cytoplasm (potential).	cytoplasm
Q8K9R6	Y244_BUCAP		inner membrane
Q8K9R7	GRPE1_BUCAP	cytoplasm (probable).	cytoplasm
Q8K9R8	XGPT_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9R9	GMHA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9S0	FLHA_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K9S1	FLHB_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9S3	DPO3A_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9S4	FABZ_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9S5	Y231_BUCAP		outer membrane
Q8K9S6	UPPS_BUCAP		inner membrane
Q8K9S7	DXR_BUCAP		cytoplasm
Q8K9S8	RRF_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9T0	RS2_BUCAP		cytoplasm
Q8K9T1	AMPM_BUCAP		cytoplasm
Q8K9T2	MURD_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9T3	FTSW_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8K9T4	MURG_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8K9T5	SPEE_BUCAP		cytoplasm
Q8K9T6	SPED_BUCAP		cytoplasm
Q8K9T7	DLDH_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9T8	ODP2_BUCAP		inner membrane
Q8K9T9	ODP1_BUCAP		cytoplasm
Q8K9U0	GUAC_BUCAP		cytoplasm
Q8K9U1	COAE_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9U2	MUTT_BUCAP		cytoplasm
Q8K9U3	SECA_BUCAP	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	inner membrane
Q8K9U4	TRUA_BUCAP		cytoplasm
Q8K9U5	DKSA_BUCAP		cytoplasm
Q8K9U6	PANB_BUCAP		cytoplasm

Q8K9U7	PANC_BUCAP		cytoplasm
Q8K9U8	HPRT_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9U9	AKH_BUCAP		cytoplasm
Q8K9V0	KHSE_BUCAP	cytoplasm (potential).	cytoplasm
Q8K9V1	THRC_BUCAP		cytoplasm
Q8K9V2	ENGD_BUCAP		cytoplasm
Q8K9V3	PTH_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9V4	SODM_BUCAP		periplasm
Q8K9V5	RNT_BUCAP		cytoplasm
Q8K9V6	Y181_BUCAP		cytoplasm
Q8K9V7	SMPA_BUCAP	bacterial cell membrane; lipid- anchor (by similarity).	outer membrane
Q8K9V8	PPNK_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9V9	GRPE2_BUCAP	cytoplasm (probable).	cytoplasm
Q8K9W0	TSAA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9W1	Y175_BUCAP		extracellular
Q8K9W2	GYRA_BUCAP		cytoplasm
Q8K9W3	RIR1_BUCAP		cytoplasm
Q8K9W4	RIR2_BUCAP		cytoplasm
Q8K9W5	PTA_BUCAP	cytoplasm (potential).	inner membrane
Q8K9W6	ACKA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9W7	NADE_BUCAP		cytoplasm
Q8K9W8	Y167_BUCAP		extracellular
Q8K9W9	HEMK_BUCAP		cytoplasm
Q8K9X1	ISPE_BUCAP		cytoplasm
Q8K9X2	KPRS_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9X3	FOLC_BUCAP		cytoplasm
Q8K9X4	Y160_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9X6	NUOM_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9X7	NUOL_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9X8	NUOK_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9X9	NUOJ_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8K9Y0	NUOI_BUCAP		cytoplasm
Q8K9Y2	NUOG_BUCAP		cytoplasm
Q8K9Y3	NUOF_BUCAP		periplasm
Q8K9Y4	NUOE_BUCAP		cytoplasm
Q8K9Y5	NUOCD_BUCAP		inner membrane
Q8K9Y6	NUOB_BUCAP		cytoplasm
Q8K9Y8	DNAK_BUCAP		cytoplasm
Q8K9Y9	DNAJ_BUCAP	cytoplasm (probable).	cytoplasm
Q8K9Z0	RS20_BUCAP		cytoplasm
Q8K9Z1	RIBF_BUCAP		cytoplasm
Q8K9Z2	SYI_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9Z3	LSPA_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	extracellular

Q8K9Z4	ISPH_BUCAP		cytoplasm
Q8K9Z5	DAPB_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8K9Z6	CARA_BUCAP		cytoplasm
Q8K9Z7	CARB_BUCAP		cytoplasm
Q8K9Z8	DYR_BUCAP		cytoplasm
Q8K9Z9	APAH_BUCAP		extracellular
Q8KA00	KSGA_BUCAP		cytoplasm
Q8KA01	SURA_BUCAP		periplasm
Q8KA02	Y132_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8KA03	RL25_BUCAP		cytoplasm
Q8KA05	GCH1_BUCAP		cytoplasm
Q8KA08	Y126_BUCAP		inner membrane
Q8KA09	TGT_BUCAP		cytoplasm
Q8KA10	QUEA_BUCAP	cytoplasm (potential).	cytoplasm
Q8KA11	IHFA_BUCAP		cytoplasm
Q8KA12	Y115_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8KA13	Y114_BUCAP		inner membrane
Q8KA14	SYI_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8KA15	PRIA_BUCAP		cytoplasm
Q8KA16	END3_BUCAP		cytoplasm
Q8KA17	RNFE_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8KA18	RNFG_BUCAP		inner membrane
Q8KA19	RNFD_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8KA21	RNFA_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8KA22	RISA_BUCAP		cytoplasm
Q8KA23	TILS_BUCAP	cytoplasm (probable).	cytoplasm
Q8KA24	DAPA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8KA25	DAPE_BUCAP		cytoplasm
Q8KA26	TKT_BUCAP		extracellular
Q8KA27	TAL_BUCAP	cytoplasm (probable).	cytoplasm
Q8KA28	FABB_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8KA29	Y083_BUCAP		cytoplasm
Q8KA30	PMBA_BUCAP	cytoplasm (potential).	cytoplasm
Q8KA31	IPYR_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8KA32	Y080_BUCAP		inner membrane
Q8KA33	RL28_BUCAP		cytoplasm
Q8KA34	RL33_BUCAP		cytoplasm
Q8KA35	FLIR_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8KA36	FLIQ_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8KA37	FLIP_BUCAP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q8KA38	FLIN_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8KA39	FLIM_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8KA40	FLIK_BUCAP		flagellar
Q8KA41	FLIJ_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8KA42	FLII_BUCAP	cytoplasm (potential).	inner membrane
Q8KA43	FLIH_BUCAP		inner membrane
Q8KA44	FLIG_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8KA45	FLIF_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8KA46	FLIE_BUCAP		flagellar
Q8KA48	CYSK_BUCAP		cytoplasm
Q8KA49	PTHP_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8KA51	PTGA_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8KA52	UPPP_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8KA53	CCA_BUCAP		cytoplasm
Q8KA54	Y057_BUCAP		cytoplasm
Q8KA55	RIBB_BUCAP		cytoplasm
Q8KA56	GCP_BUCAP		extracellular
Q8KA57	RS21_BUCAP		cytoplasm
Q8KA58	Y049_BUCAP		cytoplasm
Q8KA59	ARLY_BUCAP	cytoplasm (probable).	cytoplasm
Q8KA60	ASSY_BUCAP	cytoplasm (probable).	cytoplasm
Q8KA61	ARGC_BUCAP	cytoplasm (probable).	cytoplasm
Q8KA62	METF_BUCAP		cytoplasm
Q8KA63	MURB_BUCAP	cytoplasm (probable).	inner membrane
Q8KA64	SECE_BUCAP	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8KA65	NUSG_BUCAP		cytoplasm
Q8KA66	RL11_BUCAP		cytoplasm
Q8KA67	RL1_BUCAP		cytoplasm
Q8KA68	RL10_BUCAP		cytoplasm
Q8KA69	DBH_BUCAP		cytoplasm
Q8KA70	PUR9_BUCAP		cytoplasm
Q8KA71	METE_BUCAP		cytoplasm
Q8KA72	METR_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8KA73	Y029_BUCAP		cytoplasm
Q8KA74	GLMU_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8KA75	GLMS_BUCAP	cytoplasm (by similarity).	cytoplasm
Q8KA76	RP32_BUCAP		cytoplasm
Q8KA77	FTSY_BUCAP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8KA78	DNAT_BUCAP		nucleoid
Q8KA79	DNAC_BUCAP		nucleoid

Q8KIL1	SLDA2_GLUOX	secreted protein (probable).	cytoplasm
Q8KIU8	WBJE_PSEAE		cytoplasm
Q8KIX0	CH60_BUCCM	cytoplasm (by similarity).	cytoplasm
Q8KIX1	CH60_BUCTT	cytoplasm (by similarity).	cytoplasm
Q8KIX2	CH60_BUCTS	cytoplasm (by similarity).	cytoplasm
Q8KIX3	CH60_BUCTC	cytoplasm (by similarity).	cytoplasm
Q8KIX4	CH60_BUCPP	cytoplasm (by similarity).	cytoplasm
Q8KKH0	LAMB_AERHY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8KL44	GPMA_RHIEC		cytoplasm
Q8KLS9	HADD_RHISH		extracellular
Q8KLU8	ACEK_RALEU	cytoplasm (by similarity).	cytoplasm
Q8KLV8	YPTO_PSEST		cytoplasm
Q8KM01	LUXS_PHOLL		extracellular
Q8KM02	SLYA_PHOLL		cytoplasm
Q8KM30	CH60_OLEAN	cytoplasm (by similarity).	cytoplasm
Q8KM31	CH10_OLEAN	cytoplasm (by similarity).	cytoplasm
Q8KMJ5	MCH_METDI	cytoplasm (by similarity).	cytoplasm
Q8KN28	TFDB_COMAC		cytoplasm
Q8KNX8	RS12_BARHE		cytoplasm
Q8KR41	YEEX_ESCFE		inner membrane
Q8KR69	GLGX_ERWCH	cytoplasm (potential).	extracellular
Q8KRC9	DNAJ_MYXXA	cytoplasm (probable).	cytoplasm
Q8KRD1	RS12_MYXXA		cytoplasm
Q8KRG6	METE_VIBHA		cytoplasm
Q8KRM1	GPDA_SERMA	cytoplasm (probable).	cytoplasm
Q8KRM2	SECB_SERMA	cytoplasm (by similarity).	cytoplasm
Q8KRR5	4OT_PSEFL		cytoplasm
Q8KRT5	DEOD_XENNE		cytoplasm
Q8KRZ2	XERC_PSESX	cytoplasm (by similarity).	nucleoid
Q8KSB4	CPS1_ACTPL		cytoplasm
Q8KSV0	LGT_AZOBR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8KT78	LEXA_PSECL		cytoplasm
Q8KTE1	FUMC_METEX	cytoplasm (by similarity).	cytoplasm
Q8KTH8	RPOC_XANOR		cytoplasm
Q8KTI4	SSRP_AZOVI	cytoplasm (potential).	cytoplasm
Q8KTP6	RS12_TREPR		cytoplasm
Q8KTP9	RL7_TREPR		cytoplasm
Q8KTQ9	ARLY_TREPR	cytoplasm (probable).	cytoplasm
Q8KTR0	RRMJ_TREPR	cytoplasm (by similarity).	cytoplasm
Q8KTR3	MRAW_TREPR		cytoplasm
Q8KTR6	ILVC_TREPR		cytoplasm
Q8KTR9	CH10_TREPR	cytoplasm (by similarity).	cytoplasm
Q8KTS0	CH60_TREPR	cytoplasm (by similarity).	cytoplasm
Q8KTS4	RS16_TREPR		cytoplasm
Q8KTS9	ILVD_TREPR		cytoplasm
Q8KU07	AZOR_XENAZ		cytoplasm
Q8KUM1	HRPA_ERWCH	extracellular and secreted via type iii secretion system (by similarity).	extracellular
Q8KWX2	RPOB_EHRCR		cytoplasm
Q8KWX4	RPOB_ANAMA		cytoplasm

Q8KY01	HPPA2_RHOPA	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8KY27	HIS7_AMIAM	cytoplasm (by similarity).	cytoplasm
Q8KZA0	CARA_DESVM		cytoplasm
Q8KZR4	TAUB_PSEPU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8L0V4	CHS_ECOLI		cytoplasm
Q8L1B1	SYI2_PSEFL	cytoplasm (by similarity).	cytoplasm
Q8L1D5	SLDB2_GLUOX	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8L1H5	RRF_PARZE	cytoplasm (by similarity).	cytoplasm
Q8L1H9	EX7S_PARZE	cytoplasm (by similarity).	cytoplasm
Q8L1I4	IDI2_PARZE	cytoplasm (by similarity).	cytoplasm
Q8L1Z5	TPIS_BARHE		cytoplasm
Q8L1Z7	GPMA_BARHE		cytoplasm
Q8L1Z8	PGK_BARHE	cytoplasm (by similarity).	cytoplasm
Q8L1Z9	G6PI_BARHE	cytoplasm (by similarity).	cytoplasm
Q8L202	ENO_BARHE	cytoplasm (by similarity).	extracellular
Q8L2A6	SSB_PROVU		cytoplasm
Q8L2F8	RPOA_PSEPU		cytoplasm
Q8L3B9	CHBG_SERMA		inner membrane
Q8L3D3	DNAJ_COLMA	cytoplasm (probable).	cytoplasm
Q8NKX3	RS13_XANAC		cytoplasm
Q8NKY0	RL22_XANAC		cytoplasm
Q8P336	CAPP_XANCP		cytoplasm
Q8P337	RNPA_XANCP		extracellular
Q8P338	OXAA_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8P340	TRME_XANCP		cytoplasm
Q8P3B0	GCH1_XANCP		cytoplasm
Q8P3B7	Y4159_XANCP		inner membrane
Q8P3C4	FPG_XANCP		cytoplasm
Q8P3C6	OPGD_XANCP	periplasm (by similarity).	periplasm
Q8P3C8	KITH_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P3D7	PYRF_XANCP		cytoplasm
Q8P3E3	PLSB_XANCP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8P3F4	UXAC_XANCP		cytoplasm
Q8P3H1	XYLA2_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P3H3	RDGC_XANCP	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q8P3H6	HEMH_XANCP	cytoplasm (by similarity).	extracellular
Q8P3H8	TATA_XANCP	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8P3H9	TATB_XANCP	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane

Q8P3J6	Y4075_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8P3L1	ACSA_XANCP		cytoplasm
Q8P3N0	GIDB_XANCP		cytoplasm
Q8P3Q0	HEM6_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P3W6	AROE_XANCP		cytoplasm
Q8P3X4	DUSA_XANCP		cytoplasm
Q8P3Y6	KTHY_XANCP		cytoplasm
Q8P3Z3	Y3924_XANCP		cytoplasm
Q8P447	SPEE_XANCP		cytoplasm
Q8P448	SPEA_XANCP		cytoplasm
Q8P456	RADC_XANCP		cytoplasm
Q8P458	DUT_XANCP		extracellular
Q8P469	PYRE_XANCP		cytoplasm
Q8P475	SYX_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P4C5	QUEF_XANCP	cytoplasm (probable).	cytoplasm
Q8P4D0	ACEK_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P4D4	RHLB_XANCP		cytoplasm
Q8P4D7	UNG_XANCP	cytoplasm (by similarity).	extracellular
Q8P4E6	NORM_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8P4F6	SMG_XANCP		cytoplasm
Q8P4F9	DEF2_XANCP		cytoplasm
Q8P4G0	FMT_XANCP		cytoplasm
Q8P4G1	RSMB_XANCP		cytoplasm
Q8P4H1	DTD_XANCP	cytoplasm (probable).	cytoplasm
Q8P4H2	RPOD_XANCP		cytoplasm
Q8P4Q9	DADA_XANCP		cytoplasm
Q8P4R0	ALR_XANCP		cytoplasm
Q8P4S5	Y3632_XANCP		inner membrane
Q8P4U3	MDCG_XANCP		cytoplasm
Q8P4U6	MDCC_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P4V1	Y3605_XANCP		cytoplasm
Q8P4V7	GLNE_XANCP		cytoplasm
Q8P519	TREA_XANCP	periplasm (by similarity).	periplasm
Q8P532	OPGH_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8P536	HEM3_XANCP		cytoplasm
Q8P548	DAPF_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P550	XERC_XANCP	cytoplasm (by similarity).	nucleoid
Q8P551	HSLV_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P552	HSLU_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P558	UBIE_XANCP		cytoplasm
Q8P588	Y3453_XANCP		inner membrane
Q8P589	LIPB_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P590	LIPA_XANCP	cytoplasm (potential).	cytoplasm
Q8P5D4	CCA_XANCP		cytoplasm
Q8P5D7	BETA_XANCP		cytoplasm
Q8P5D8	BETB_XANCP		cytoplasm
Q8P5E4	ENGB_XANCP		cytoplasm
Q8P5F3	GFA_XANCP		extracellular
Q8P5F9	Y3382_XANCP		cytoplasm
Q8P5H9	ENGC_XANCP		cytoplasm

Q8P5J5	DCTA_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8P5K9	LEUD_XANCP		cytoplasm
Q8P5L1	LEU3_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P5L5	ILVC_XANCP		cytoplasm
Q8P5L9	THIC_XANCP		cytoplasm
Q8P5M4	IPYR_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P5M6	HPPA_XANCP	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8P5P5	KAD_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P5Q4	ARGD_XANCP	cytoplasm (probable).	cytoplasm
Q8P5R4	GSA_XANCP	cytoplasm (potential).	cytoplasm
Q8P5R7	THIE_XANCP		cytoplasm
Q8P5S1	RPIA_XANCP		cytoplasm
Q8P5S6	Y3260_XANCP		inner membrane
Q8P5T3	HAM1_XANCP		extracellular
Q8P5T5	RNPH_XANCP		cytoplasm
Q8P5T7	KGUA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P5U0	KDGD_XANCP		cytoplasm
Q8P5U9	RL31B_XANCP		cytoplasm
Q8P5Z4	PGK_XANCP	cytoplasm (potential).	cytoplasm
Q8P5Z7	ALF1_XANCP		cytoplasm
Q8P5Z8	Y3184_XANCP		inner membrane
Q8P607	CYSH_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P626	MSCL_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8P631	TRMB_XANCP		cytoplasm
Q8P632	THIG_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P665	RIMK_XANCP		cytoplasm
Q8P676	SUCC_XANCP		cytoplasm
Q8P682	RLUD_XANCP		cytoplasm
Q8P698	Y3072_XANCP		cytoplasm
Q8P6A0	CLPB_XANCP	cytoplasm (probable).	cytoplasm
Q8P6B1	COBT_XANCP		cytoplasm
Q8P6E3	Y3027_XANCP		inner membrane
Q8P6E4	RUVC_XANCP		cytoplasm
Q8P6E5	RUVA_XANCP		cytoplasm
Q8P6E6	KUP_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8P6E7	RUVB_XANCP		cytoplasm
Q8P6F2	TOLB_XANCP	periplasm (potential).	periplasm
Q8P6M7	PQQE_XANCP		cytoplasm
Q8P6M8	PQQD_XANCP		cytoplasm
Q8P6M9	PQQC_XANCP		cytoplasm
Q8P6N0	PQQB_XANCP		cytoplasm
Q8P6P1	GSHB_XANCP		cytoplasm
Q8P6Q4	CUTC_XANCP	cytoplasm (probable).	periplasm
Q8P6T5	Y2880_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8P6T8	GCST_XANCP		cytoplasm
Q8P6T9	GCSH_XANCP		cytoplasm

Q8P6V1	MTGA_XANCP	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8P6X1	DCUP_XANCP	cytoplasm (probable).	cytoplasm
Q8P6X3	AROB_XANCP	cytoplasm (probable).	cytoplasm
Q8P6X4	AROK_XANCP	cytoplasm (probable).	cytoplasm
Q8P6X5	PDXH_XANCP		cytoplasm
Q8P708	Y2806_XANCP		cytoplasm
Q8P709	HPRK_XANCP		cytoplasm
Q8P719	MURA_XANCP	cytoplasm (probable).	inner membrane
Q8P725	PUR5_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P741	Y2773_XANCP		cytoplasm
Q8P765	Y2748_XANCP		inner membrane
Q8P766	RUVX_XANCP	cytoplasm (potential).	cytoplasm
Q8P767	PYRB_XANCP		cytoplasm
Q8P775	MURD_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P778	SSB_XANCP		cytoplasm
Q8P7A0	NFI_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P7A1	GPMA_XANCP		cytoplasm
Q8P7H4	RRAA_XANCP		cytoplasm
Q8P7J8	Y2613_XANCP		inner membrane
Q8P7K1	Y2610_XANCP	cytoplasm (potential).	inner membrane
Q8P7K6	Y2605_XANCP		inner membrane
Q8P7L2	EX7S_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P7L3	TILS_XANCP	cytoplasm (probable).	cytoplasm
Q8P7P1	DCD_XANCP		extracellular
Q8P7P8	GREB_XANCP		cytoplasm
Q8P7Q6	PSD_XANCP		cytoplasm
Q8P7R0	AROC_XANCP		cytoplasm
Q8P7R5	TRUA_XANCP		cytoplasm
Q8P7R6	TRPF_XANCP		cytoplasm
Q8P7R8	TRPB_XANCP		cytoplasm
Q8P7S0	TRPA_XANCP		cytoplasm
Q8P7S2	GLMM_XANCP		cytoplasm
Q8P7T0	TPIS_XANCP		cytoplasm
Q8P7U5	Y2513_XANCP		inner membrane
Q8P7U8	RBFA_XANCP	cytoplasm (potential).	cytoplasm
Q8P7U9	TRUB_XANCP		cytoplasm
Q8P7V0	RS15_XANCP		cytoplasm
Q8P7X1	UVRB_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P7Z4	RL20_XANCP		cytoplasm
Q8P815	DXS_XANCP		cytoplasm
Q8P829	FETP_XANCP		cytoplasm
Q8P855	HTPG_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P857	COAD_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P867	QUEA_XANCP	cytoplasm (potential).	cytoplasm
Q8P868	TGT_XANCP		cytoplasm
Q8P888	FDHD_XANCP	cytoplasm (potential).	cytoplasm
Q8P8B2	Y2331_XANCP		inner membrane
Q8P8C7	3MGH_XANCP		cytoplasm
Q8P8D9	EX7L_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P8E4	MUTL_XANCP		cytoplasm
Q8P8E9	GLUQ_XANCP		cytoplasm
Q8P8F0	HTPX_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular

Q8P8F4	Y2288_XANCP		inner membrane
Q8P8F8	Y2284_XANCP		inner membrane
Q8P8H2	UBIG_XANCP		cytoplasm
Q8P8H3	GPH_XANCP		cytoplasm
Q8P8I0	EUTC_XANCP		cytoplasm
Q8P8I5	Y2256_XANCP		inner membrane
Q8P8J2	AOTC_XANCP	cytoplasm (probable).	cytoplasm
Q8P8J4	ASSY_XANCP	cytoplasm (probable).	cytoplasm
Q8P8J6	ARGB_XANCP	cytoplasm (probable).	cytoplasm
Q8P8J8	ARGC_XANCP	cytoplasm (probable).	cytoplasm
Q8P8J9	ARLY_XANCP	cytoplasm (probable).	cytoplasm
Q8P8K2	PROB_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P8K3	PROA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P8L2	METX_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P8L7	CCME2_XANCP	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8P8M1	CCMA_XANCP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8P8M6	RLUB_XANCP		cytoplasm
Q8P8P4	KCY_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P8P6	IHFB_XANCP		cytoplasm
Q8P8Q6	GUAA_XANCP		cytoplasm
Q8P8R8	MNTH_XANCP	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8P8S1	ORN_XANCP	cytoplasm (potential).	cytoplasm
Q8P8S4	Y2165_XANCP		cytoplasm
Q8P8U7	GLK_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P8V6	Y2128_XANCP		cytoplasm
Q8P8V9	LOLD_XANCP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8P8W4	MSBA_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8P8W5	LPXK_XANCP		cytoplasm
Q8P8Y5	Y2094_XANCP		cytoplasm
Q8P8Z2	Y2086_XANCP		inner membrane
Q8P8Z6	ZUPT_XANCP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8P8Z8	PHS_XANCP		cytoplasm
Q8P8Z9	Y2079_XANCP		inner membrane
Q8P979	ENGA_XANCP		cytoplasm
Q8P991	LOLA_XANCP	periplasm (by similarity).	periplasm
Q8P996	LFTR_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P999	CLPS_XANCP		cytoplasm
Q8P9A1	TRMU_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9A2	Y1963_XANCP		inner membrane
Q8P9B9	FLGH_XANCP	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8P9C0	FLGI_XANCP	periplasm (by similarity).	periplasm
Q8P9L3	DAPB_XANCP	cytoplasm (by similarity).	cytoplasm

Q8P9M3	EFPL_XANCP		cytoplasm
Q8P9N8	HIS2_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9N9	HIS6_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9P0	HIS4_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9P1	HIS5_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9P2	HISX_XANCP		cytoplasm
Q8P9P3	HIS1_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9Q0	KHSE_XANCP	cytoplasm (potential).	cytoplasm
Q8P9R0	PYRD_XANCP	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8P9R1	MURB_XANCP	cytoplasm (probable).	inner membrane
Q8P9R7	ISPG_XANCP		cytoplasm
Q8P9S7	G6PI_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9S8	PAND_XANCP		cytoplasm
Q8P9S9	PANC_XANCP		cytoplasm
Q8P9T0	PANB_XANCP		cytoplasm
Q8P9T3	Y1764_XANCP		inner membrane
Q8P9T9	XYLA1_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9V6	DAPA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9W9	CSRA_XANCP		cytoplasm
Q8P9X1	RECX_XANCP	cytoplasm (potential).	cytoplasm
Q8P9X2	LEXA2_XANCP		cytoplasm
Q8P9X7	HFQ_XANCP		cytoplasm
Q8P9X8	MIAA_XANCP		cytoplasm
Q8P9Y1	RRMJ_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9Y7	SURE_XANCP	cytoplasm (potential).	periplasm
Q8P9Y9	TRUD_XANCP		cytoplasm
Q8P9Z0	ISPF_XANCP		extracellular
Q8P9Z1	ISPD_XANCP		cytoplasm
Q8P9Z2	FTSB_XANCP	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q8P9Z3	ENO_XANCP	cytoplasm (by similarity).	extracellular
Q8P9Z5	KDSA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8P9Z6	PYRG_XANCP		cytoplasm
Q8PA33	CCME1_XANCP	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8PA95	AROAXANCP	cytoplasm (probable).	cytoplasm
Q8PA97	SERC_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PA98	Y1588_XANCP		cytoplasm
Q8PA99	Y1587_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PAA6	HUTI_XANCP	cytoplasm (potential).	cytoplasm
Q8PAA7	HUTH_XANCP	cytoplasm (potential).	cytoplasm
Q8PAB7	ZIPA_XANCP	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q8PAC0	RL9_XANCP		cytoplasm
Q8PAC1	RS18_XANCP		cytoplasm
Q8PAC2	RS6_XANCP		cytoplasm

Q8PAD9	PPNK_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PAG0	PSTB_XANCP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PAG3	RNT_XANCP		cytoplasm
Q8PAG5	RLMB_XANCP		cytoplasm
Q8PAH9	SLYX_XANCP		inner membrane
Q8PAK8	DNAJ_XANCP	cytoplasm (probable).	cytoplasm
Q8PAK9	DNAK_XANCP		cytoplasm
Q8PAL0	GRPE_XANCP	cytoplasm (probable).	cytoplasm
Q8PAL1	HRCA_XANCP		cytoplasm
Q8PAL7	SSRP_XANCP	cytoplasm (potential).	cytoplasm
Q8PAT3	TPMT_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PAU4	GLND_XANCP		cytoplasm
Q8PAV2	RS2_XANCP		cytoplasm
Q8PAV6	RRF_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PAV7	UPPS_XANCP		inner membrane
Q8PAV9	DXR_XANCP		cytoplasm
Q8PAW3	LPXD_XANCP		cytoplasm
Q8PAW4	FABZ_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PAW5	LPXA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PAW6	LPXB_XANCP		cytoplasm
Q8PAW7	RNH2_XANCP	cytoplasm (potential).	extracellular
Q8PAW9	ACCA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PB42	NAGZ_XANCP	cytoplasm (by similarity).	extracellular
Q8PB48	RUMA_XANCP		cytoplasm
Q8PB50	RECO_XANCP		cytoplasm
Q8PB51	ERA_XANCP		cytoplasm
Q8PB52	RNC_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PB55	LEPA_XANCP		cytoplasm
Q8PBB5	MUTS_XANCP		cytoplasm
Q8PBC0	RL19_XANCP		cytoplasm
Q8PBC1	TRMD_XANCP	cytoplasm (potential).	cytoplasm
Q8PBC2	RIMM_XANCP	cytoplasm (potential).	cytoplasm
Q8PBC3	RS16_XANCP		cytoplasm
Q8PBG4	ISPH_XANCP		cytoplasm
Q8PBG6	SYI_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PBG9	RS20_XANCP		cytoplasm
Q8PBH1	RL27_XANCP		cytoplasm
Q8PBH3	UVRA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PBH8	ATE_XANCP	cytoplasm (potential).	cytoplasm
Q8PBJ4	MINC_XANCP		inner membrane
Q8PBK7	GCSP_XANCP		cytoplasm
Q8PBL8	DNAE2_XANCP	cytoplasm (probable).	cytoplasm
Q8PBM1	LEXA1_XANCP		cytoplasm
Q8PBR6	PURA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PBV1	FABH_XANCP	cytoplasm (probable).	cytoplasm
Q8PBV2	RL32_XANCP		cytoplasm
Q8PBV4	Y1013_XANCP	cytoplasm (potential).	inner membrane
Q8PBW4	RECR_XANCP		cytoplasm
Q8PBW5	Y1002_XANCP		inner membrane
Q8PBW9	MOAC_XANCP		cytoplasm
Q8PBX1	MOAA_XANCP		cytoplasm
Q8PBX3	KDKA_XANCP	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm

Q8PBX8	RNH_XANCP	cytoplasm (potential).	extracellular
Q8PBY5	CLPX_XANCP		cytoplasm
Q8PBY6	CLPP_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PBY7	TIG_XANCP		cytoplasm
Q8PC11	CYSA_XANCP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PC25	MDH_XANCP		cytoplasm
Q8PC32	DSBB_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PC34	RL15_XANCP		cytoplasm
Q8PC36	RL18_XANCP		cytoplasm
Q8PC38	RS8_XANCP		cytoplasm
Q8PC39	RL5_XANCP		cytoplasm
Q8PC40	RL24_XANCP		cytoplasm
Q8PC41	RS17_XANCP		cytoplasm
Q8PC42	RL29_XANCP		cytoplasm
Q8PC43	RL16_XANCP		cytoplasm
Q8PC44	RS3_XANCP		cytoplasm
Q8PC45	RS19_XANCP		cytoplasm
Q8PC46	RL2_XANCP		cytoplasm
Q8PC48	RL4_XANCP		cytoplasm
Q8PC49	RL3_XANCP		cytoplasm
Q8PC50	RS10_XANCP		cytoplasm
Q8PC53	RS7_XANCP		cytoplasm
Q8PC54	RS12_XANCP		cytoplasm
Q8PC55	RPOC_XANCP		cytoplasm
Q8PC56	RPOB_XANCP		cytoplasm
Q8PC57	RL7_XANCP		cytoplasm
Q8PC61	PTH_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PC62	RL25_XANCP		cytoplasm
Q8PC63	KPRS_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PC64	ISPE_XANCP		cytoplasm
Q8PC65	LOLB_XANCP	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q8PC67	HEM1_XANCP		cytoplasm
Q8PC75	RBN_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PCA4	TAL_XANCP	cytoplasm (probable).	cytoplasm
Q8PCA6	MSRA_XANCP		cytoplasm
Q8PCB8	MTFA_XANCP		cytoplasm
Q8PCE0	OSTA_XANCP	bacterial cell outer membrane (by similarity).	outer membrane
Q8PCE2	PDXA_XANCP	cytoplasm (probable).	cytoplasm
Q8PCE3	KSGA_XANCP		cytoplasm
Q8PCE4	APAG_XANCP		inner membrane
Q8PCE5	APAH_XANCP		extracellular
Q8PCE7	TYSY_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCE8	LGT_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PCH1	DUSB_XANCP		cytoplasm
Q8PCH5	SAHH_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCJ5	LPXC_XANCP		cytoplasm

Q8PCJ8	DDLB_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCJ9	MURC_XANCP	cytoplasm (probable).	cytoplasm
Q8PCK0	MURG_XANCP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PCK2	MRAY_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PCK4	MURE_XANCP	cytoplasm (probable).	cytoplasm
Q8PCK7	MRAW_XANCP		cytoplasm
Q8PCK8	MRAZ_XANCP		inner membrane
Q8PCL4	Y710_XANCP		inner membrane
Q8PCM0	ATKC_XANCP	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8PCM1	ATKB_XANCP	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8PCM6	NUSB_XANCP		cytoplasm
Q8PCM7	RISB_XANCP		cytoplasm
Q8PCN3	NRDR_XANCP		cytoplasm
Q8PCN4	GLYA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCN7	DEF1_XANCP		cytoplasm
Q8PCP3	PNCB_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCQ9	XERD_XANCP	cytoplasm (by similarity).	nucleoid
Q8PCR4	AMPA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCR7	SYV_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCW4	BIOD_XANCP		cytoplasm
Q8PCW8	DPO4_XANCP	cytoplasm (probable).	cytoplasm
Q8PCW9	FABA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCY1	GLMS_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCZ1	GLMU_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PCZ4	ATPE_XANCP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PCZ6	ATPG_XANCP		inner membrane
Q8PCZ7	ATPA_XANCP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PD23	CH60_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PD27	AROQ_XANCP		cytoplasm
Q8PD33	PRMA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PD37	FISL_XANCP		cytoplasm
Q8PD47	PUR9_XANCP		cytoplasm
Q8PD48	PUR2_XANCP		cytoplasm
Q8PD65	NUDH_XANCP		cytoplasm
Q8PD66	RS9_XANCP		cytoplasm
Q8PD70	TRPC_XANCP		cytoplasm
Q8PD71	TRPD_XANCP		cytoplasm
Q8PD87	PUR7_XANCP		cytoplasm
Q8PDA2	HGD_XANCP		cytoplasm
Q8PDD1	GLGB2_XANCP		extracellular
Q8PDD2	GLGA_XANCP		cytoplasm
Q8PDD7	OPGB_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q8PDF3	BIOH_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PDG1	GIDA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PDI0	GLPK_XANCP		cytoplasm
Q8PDJ3	ILVD_XANCP		cytoplasm
Q8PDM9	NPD_XANCP	cytoplasm (probable).	cytoplasm
Q8PDR2	RLUE_XANCP		cytoplasm
Q8PDW1	UBIB_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PDW3	DDLA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PDY0	GPDA_XANCP	cytoplasm (probable).	cytoplasm
Q8PDY1	SECB_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PDZ9	UPPP_XANCP	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PE32	KDUI_XANCP		cytoplasm
Q8PE48	GLGB1_XANCP		extracellular
Q8PE66	Y115_XANCP		cytoplasm
Q8PE75	LLDD_XANCP		cytoplasm
Q8PEB1	Y070_XANCP		cytoplasm
Q8PEG8	PDXJ_XANCP	cytoplasm (probable).	cytoplasm
Q8PEH3	RECF_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PEH5	DNAA_XANCP	cytoplasm (by similarity).	cytoplasm
Q8PEH6	RNPA_XANAC		extracellular
Q8PEH7	OXAA_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PEH9	TRME_XANAC		cytoplasm
Q8PEP3	GCH1_XANAC		cytoplasm
Q8PEP8	Y4297_XANAC		inner membrane
Q8PEQ4	FPG_XANAC		cytoplasm
Q8PEQ6	OPGD_XANAC	periplasm (by similarity).	periplasm
Q8PEQ8	KITH_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PER4	PYRF_XANAC		cytoplasm
Q8PES0	PLSB_XANAC	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PET9	UXAC_XANAC		cytoplasm
Q8PEW5	XYLA2_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PEW7	RDGC_XANAC	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q8PEX0	HEMH_XANAC	cytoplasm (by similarity).	extracellular
Q8PEX2	TATA_XANAC	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8PEX3	TATB_XANAC	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8PEZ6	Y4192_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PF09	ACSA_XANAC		cytoplasm
Q8PF23	GIDB_XANAC		cytoplasm
Q8PF76	HEM6_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PFE9	AROE_XANAC		cytoplasm

Q8PFF8	DUSA_XANAC		cytoplasm
Q8PFG7	KTHY_XANAC		cytoplasm
Q8PFH4	Y4007_XANAC		cytoplasm
Q8PFQ4	SPEE_XANAC		cytoplasm
Q8PFQ5	SPEA_XANAC		cytoplasm
Q8PFR3	RADC_XANAC		cytoplasm
Q8PFR5	DUT_XANAC		extracellular
Q8PFS5	PYRE_XANAC		cytoplasm
Q8PFT1	SYU_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PFX6	QUEF_XANAC	cytoplasm (probable).	cytoplasm
Q8PFZ0	ACEK_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PFZ3	RHLB_XANAC		cytoplasm
Q8PFZ6	UNG_XANAC	cytoplasm (by similarity).	extracellular
Q8PG07	NORM_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PG17	SMG_XANAC		cytoplasm
Q8PG20	DEF2_XANAC		cytoplasm
Q8PG21	FMT_XANAC		cytoplasm
Q8PG22	RSMB_XANAC		cytoplasm
Q8PG32	DTD_XANAC	cytoplasm (probable).	cytoplasm
Q8PG33	RPOD_XANAC		cytoplasm
Q8PGC9	DADA_XANAC		cytoplasm
Q8PGD0	ALR_XANAC		cytoplasm
Q8PGE6	Y3671_XANAC		inner membrane
Q8PGG5	ATPA_XANAC	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PGG6	ATPG_XANAC		inner membrane
Q8PGG8	ATPE_XANAC	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PGH2	GLMU_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PGH9	GLMS_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PGJ2	FABA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PGJ4	DPO4_XANAC	cytoplasm (probable).	cytoplasm
Q8PGK0	BIOD_XANAC		cytoplasm
Q8PGQ7	SYV_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PGR0	AMPA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PGR5	XERD_XANAC	cytoplasm (by similarity).	nucleoid
Q8PGU5	PNCB_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PGV2	DEF1_XANAC		cytoplasm
Q8PGV3	Y3508_XANAC		cytoplasm
Q8PGW9	ENGC_XANAC		cytoplasm
Q8PGZ1	DCTA_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PH04	LEUD_XANAC		cytoplasm
Q8PH05	LEU3_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PH09	ILVC_XANAC		cytoplasm
Q8PH13	THIC_XANAC		cytoplasm
Q8PH18	IPYR_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PH20	HPPA_XANAC	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8PH23	KAD_XANAC	cytoplasm (by similarity).	cytoplasm

Q8PH31	ARGD_XANAC	cytoplasm (probable).	cytoplasm
Q8PH40	GSA_XANAC	cytoplasm (potential).	cytoplasm
Q8PH45	THIE_XANAC		cytoplasm
Q8PH49	RPIA_XANAC		cytoplasm
Q8PH54	Y3406_XANAC		inner membrane
Q8PH61	HAM1_XANAC		extracellular
Q8PH63	RNPH_XANAC		cytoplasm
Q8PH65	KGUA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PH70	RL31B_XANAC		cytoplasm
Q8PHB2	PGK_XANAC	cytoplasm (potential).	cytoplasm
Q8PHB5	ALF1_XANAC		cytoplasm
Q8PHB6	Y3343_XANAC		inner membrane
Q8PHC7	CYSH_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PHE9	MSCL_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PHF4	TRMB_XANAC		cytoplasm
Q8PHF5	THIG_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PHK1	RIMK_XANAC		cytoplasm
Q8PHK7	COAE_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PHL5	SUCC_XANAC		cytoplasm
Q8PHN2	RLUD_XANAC		cytoplasm
Q8PHP7	Y3202_XANAC		cytoplasm
Q8PHQ4	CLPB_XANAC	cytoplasm (probable).	cytoplasm
Q8PHR1	COBQ_XANAC		cytoplasm
Q8PHR3	COBT_XANAC		cytoplasm
Q8PHU8	Y3151_XANAC		inner membrane
Q8PHU9	RUVC_XANAC		cytoplasm
Q8PHV0	RUVA_XANAC		cytoplasm
Q8PHV1	KUP_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PHV2	RUVB_XANAC		cytoplasm
Q8PHV7	TOLB_XANAC	periplasm (potential).	periplasm
Q8PHY1	PQQE_XANAC		cytoplasm
Q8PHY2	PQQD_XANAC		cytoplasm
Q8PHY3	PQQC_XANAC		cytoplasm
Q8PHY4	PQQB_XANAC		cytoplasm
Q8PHZ5	GSHB_XANAC		cytoplasm
Q8PI07	CUTC_XANAC	cytoplasm (probable).	periplasm
Q8PI34	Y3064_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PI37	GCST_XANAC		cytoplasm
Q8PI38	GCSH_XANAC		cytoplasm
Q8PI51	MTGA_XANAC	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8PI85	DCUP_XANAC	cytoplasm (probable).	cytoplasm
Q8PI87	AROB_XANAC	cytoplasm (probable).	cytoplasm
Q8PI88	AROK_XANAC	cytoplasm (probable).	cytoplasm
Q8PI89	PDXH_XANAC		cytoplasm
Q8PIC2	Y2976_XANAC		cytoplasm
Q8PIC3	HPRK_XANAC		cytoplasm
Q8PID3	MURA_XANAC	cytoplasm (probable).	inner membrane
Q8PID9	PUR5_XANAC	cytoplasm (by similarity).	cytoplasm

Q8PIF6	Y2942_XANAC		cytoplasm
Q8PIH9	Y2918_XANAC		inner membrane
Q8PII0	RUVX_XANAC	cytoplasm (potential).	cytoplasm
Q8PII1	PYRB_XANAC		cytoplasm
Q8PII9	MURD_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PIJ2	SSB_XANAC		cytoplasm
Q8PIM0	NFI_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PIM1	GPMA_XANAC		cytoplasm
Q8PIT8	RRAA_XANAC		cytoplasm
Q8PIX0	Y2775_XANAC		inner membrane
Q8PIX4	Y2771_XANAC	cytoplasm (potential).	inner membrane
Q8PIX9	Y2766_XANAC		inner membrane
Q8PIY4	EX7S_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PIY5	TILS_XANAC	cytoplasm (probable).	cytoplasm
Q8PJ05	DCD_XANAC		extracellular
Q8PJ11	GREB_XANAC		cytoplasm
Q8PJ17	PSD_XANAC		cytoplasm
Q8PJ20	AROC_XANAC		cytoplasm
Q8PJ25	TRUA_XANAC		cytoplasm
Q8PJ26	TRPF_XANAC		cytoplasm
Q8PJ28	TRPB_XANAC		cytoplasm
Q8PJ29	TRPA_XANAC		cytoplasm
Q8PJ31	GLMM_XANAC		cytoplasm
Q8PJ38	TPIS_XANAC		cytoplasm
Q8PJ53	Y2689_XANAC		inner membrane
Q8PJ56	RBFA_XANAC	cytoplasm (potential).	cytoplasm
Q8PJ57	TRUB_XANAC		cytoplasm
Q8PJ58	RS15_XANAC		cytoplasm
Q8PJB1	UVRB_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PJE3	RL20_XANAC		cytoplasm
Q8PJG7	DXS_XANAC		cytoplasm
Q8PJH7	FETP_XANAC		cytoplasm
Q8PJK3	HTPG_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PJK5	COAD_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PJL6	QUEA_XANAC	cytoplasm (potential).	cytoplasm
Q8PJL7	TGT_XANAC		cytoplasm
Q8PJP3	FDHD_XANAC	cytoplasm (potential).	cytoplasm
Q8PJR6	Y2464_XANAC		inner membrane
Q8PJT1	3MGH_XANAC		cytoplasm
Q8PJW7	EX7L_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PJX2	MUTL_XANAC		cytoplasm
Q8PJX7	GLUQ_XANAC		cytoplasm
Q8PJX8	HTPX_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8PJY5	Y2392_XANAC		inner membrane
Q8PK00	UBIG_XANAC		cytoplasm
Q8PK11	EUTC_XANAC		cytoplasm
Q8PK18	Y2359_XANAC		inner membrane
Q8PK25	AOTC_XANAC	cytoplasm (probable).	cytoplasm
Q8PK26	ASSY_XANAC	cytoplasm (probable).	cytoplasm
Q8PK29	ARGB_XANAC	cytoplasm (probable).	cytoplasm
Q8PK31	ARGC_XANAC	cytoplasm (probable).	cytoplasm
Q8PK32	ARLY_XANAC	cytoplasm (probable).	cytoplasm
Q8PK34	PROB_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PK35	PROA_XANAC	cytoplasm (by similarity).	cytoplasm

Q8PK44	METX_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PK49	CCME2_XANAC	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8PK53	CCMA_XANAC	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PK58	RLUB_XANAC		cytoplasm
Q8PK76	KCY_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PK78	IHFB_XANAC		cytoplasm
Q8PK88	GUA_A_XANAC		cytoplasm
Q8PKQ2	Y2117_XANAC		inner membrane
Q8PKQ3	PHS_XANAC		cytoplasm
Q8PKQ5	ZUPT_XANAC	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8PKR0	Y2109_XANAC		inner membrane
Q8PKS4	LPXK_XANAC		cytoplasm
Q8PKS5	MSBA_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PKT0	LOLD_XANAC	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PKT3	Y2079_XANAC		cytoplasm
Q8PKU2	GLK_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PKW5	Y2042_XANAC		cytoplasm
Q8PKW8	ORN_XANAC	cytoplasm (potential).	cytoplasm
Q8PKX1	MNTH_XANAC	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8PKY6	ENGA_XANAC		cytoplasm
Q8PKZ5	CRCB_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PKZ8	LOLA_XANAC	periplasm (by similarity).	periplasm
Q8PL03	LFTR_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PL06	CLPS_XANAC		cytoplasm
Q8PL08	TRMU_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PL09	Y1997_XANAC		inner membrane
Q8PL26	FLGH_XANAC	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8PL27	FLGI_XANAC	periplasm (by similarity).	periplasm
Q8PLD9	GRE_A_XANAC		cytoplasm
Q8PLE0	DAPB_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLF1	EFPL_XANAC		cytoplasm
Q8PLG5	HIS2_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLG6	HIS6_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLG7	HIS4_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLG8	HIS5_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLG9	HISX_XANAC		cytoplasm
Q8PLH0	HIS1_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLJ2	PYRD_XANAC	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane

Q8PLJ3	MURB_XANAC	cytoplasm (probable).	inner membrane
Q8PLJ8	ISPG_XANAC		cytoplasm
Q8PLK9	PAND_XANAC		cytoplasm
Q8PLL0	PANC_XANAC		cytoplasm
Q8PLL1	PANB_XANAC		cytoplasm
Q8PLL4	Y1781_XANAC		inner membrane
Q8PLL9	XYLA1_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLN5	DAPA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLP9	CSRA_XANAC		cytoplasm
Q8PLQ4	HFQ_XANAC		cytoplasm
Q8PLQ5	MIAA_XANAC		cytoplasm
Q8PLQ8	RRMJ_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLR4	SURE_XANAC	cytoplasm (potential).	periplasm
Q8PLR6	TRUD_XANAC		cytoplasm
Q8PLR7	ISPF_XANAC		extracellular
Q8PLR8	ISPD_XANAC		cytoplasm
Q8PLR9	FTSB_XANAC	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q8PLS0	ENO_XANAC	cytoplasm (by similarity).	extracellular
Q8PLS2	KDSA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLS3	PYRG_XANAC		cytoplasm
Q8PLW2	CCME1_XANAC	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8PLY5	AROA_XANAC	cytoplasm (probable).	cytoplasm
Q8PLY7	SERC_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PLY8	Y1647_XANAC		cytoplasm
Q8PLY9	Y1646_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PLZ7	HUTI_XANAC	cytoplasm (potential).	cytoplasm
Q8PLZ8	HUTH_XANAC	cytoplasm (potential).	cytoplasm
Q8PM10	ZIPA_XANAC	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q8PM12	RL9_XANAC		cytoplasm
Q8PM13	RS18_XANAC		cytoplasm
Q8PM14	RS6_XANAC		cytoplasm
Q8PM39	PPNK_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PM59	PSTB_XANAC	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PM62	RNT_XANAC		cytoplasm
Q8PM64	RLMB_XANAC		cytoplasm
Q8PM80	SLYX_XANAC		inner membrane
Q8PMA9	DNAJ_XANAC	cytoplasm (probable).	cytoplasm
Q8PMB0	DNAK_XANAC		cytoplasm
Q8PMB1	GRPE_XANAC	cytoplasm (probable).	cytoplasm
Q8PMB2	HRCA_XANAC		cytoplasm
Q8PMB9	SSRP_XANAC	cytoplasm (potential).	cytoplasm
Q8PMI8	TPMT_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PMJ8	GLND_XANAC		cytoplasm

Q8PMK5	RS2_XANAC		cytoplasm
Q8PMK8	RRF_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PMK9	UPPS_XANAC		inner membrane
Q8PML1	DXR_XANAC		cytoplasm
Q8PML5	LPXD_XANAC		cytoplasm
Q8PML6	FABZ_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PML7	LPXA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PML8	LPXB_XANAC		cytoplasm
Q8PML9	RNH2_XANAC	cytoplasm (potential).	extracellular
Q8PMM1	ACCA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PMU1	NAGZ_XANAC	cytoplasm (by similarity).	extracellular
Q8PMU6	RUMA_XANAC		cytoplasm
Q8PMU8	RECO_XANAC		cytoplasm
Q8PMU9	ERA_XANAC		cytoplasm
Q8PMV0	RNC_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PMV3	LEPA_XANAC		cytoplasm
Q8PMX2	MUTS_XANAC		cytoplasm
Q8PMY0	RL19_XANAC		cytoplasm
Q8PMY1	TRMD_XANAC	cytoplasm (potential).	cytoplasm
Q8PMY2	RIMM_XANAC	cytoplasm (potential).	cytoplasm
Q8PMY3	RS16_XANAC		cytoplasm
Q8PN17	ISPH_XANAC		cytoplasm
Q8PN19	SYI_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PN22	RS20_XANAC		cytoplasm
Q8PN24	RL27_XANAC		cytoplasm
Q8PN26	UVRA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PN34	ATE_XANAC	cytoplasm (potential).	cytoplasm
Q8PN47	MINC_XANAC		cytoplasm
Q8PN59	GCSP_XANAC		cytoplasm
Q8PN74	DNAE2_XANAC	cytoplasm (probable).	cytoplasm
Q8PN77	LEXA1_XANAC		cytoplasm
Q8PNB5	PURA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PNE8	FABH_XANAC	cytoplasm (probable).	cytoplasm
Q8PNE9	RL32_XANAC		cytoplasm
Q8PNF1	Y1120_XANAC	cytoplasm (potential).	inner membrane
Q8PNG0	RECR_XANAC		cytoplasm
Q8PNG1	Y1110_XANAC		inner membrane
Q8PNH0	MOAC_XANAC		cytoplasm
Q8PNH1	MOAA_XANAC		cytoplasm
Q8PNH3	KDKA_XANAC	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q8PNH8	RNH_XANAC	cytoplasm (potential).	extracellular
Q8PNI4	CLPX_XANAC		cytoplasm
Q8PNI5	CLPP_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PNI6	TIG_XANAC		cytoplasm
Q8PNN2	TDH_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PNN4	CYSA_XANAC	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PNP8	MDH_XANAC		cytoplasm
Q8PNQ6	DSBB_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PNR1	RL18_XANAC		cytoplasm
Q8PNR3	RS8_XANAC		cytoplasm

Q8PNR4	RL5_XANAC		cytoplasm
Q8PNR5	RL24_XANAC		cytoplasm
Q8PNR6	RS17_XANAC		cytoplasm
Q8PNR7	RL29_XANAC		cytoplasm
Q8PNR8	RL16_XANAC		cytoplasm
Q8PNR9	RS3_XANAC		cytoplasm
Q8PNS0	RS19_XANAC		cytoplasm
Q8PNS1	RL2_XANAC		cytoplasm
Q8PNS3	RL4_XANAC		cytoplasm
Q8PNS4	RL3_XANAC		cytoplasm
Q8PNS5	RS10_XANAC		cytoplasm
Q8PNS7	RS7_XANAC		cytoplasm
Q8PNS8	RS12_XANAC		cytoplasm
Q8PNS9	RPOC_XANAC		cytoplasm
Q8PNT0	RPOB_XANAC		cytoplasm
Q8PNT1	RL7_XANAC		cytoplasm
Q8PNT2	RL10_XANAC		cytoplasm
Q8PNT3	RL1_XANAC		cytoplasm
Q8PNT4	RL11_XANAC		cytoplasm
Q8PNT8	PTH_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PNT9	RL25_XANAC		cytoplasm
Q8PNU0	KPRS_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PNU1	ISPE_XANAC		cytoplasm
Q8PNU2	LOLB_XANAC	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q8PNU4	HEM1_XANAC		cytoplasm
Q8PNV2	RBN_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PNY6	TAL_XANAC	cytoplasm (probable).	cytoplasm
Q8PNY8	MSRA_XANAC		cytoplasm
Q8PNZ9	MTFA_XANAC		cytoplasm
Q8PP22	OSTA_XANAC	bacterial cell outer membrane (by similarity).	outer membrane
Q8PP24	PDXA_XANAC	cytoplasm (probable).	cytoplasm
Q8PP25	KSGA_XANAC		cytoplasm
Q8PP26	APAG_XANAC		inner membrane
Q8PP27	APAH_XANAC		extracellular
Q8PP38	SSUD_XANAC		cytoplasm
Q8PP46	TYSY_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PP47	LGT_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PP72	DUSB_XANAC		cytoplasm
Q8PP82	CAPP_XANAC		cytoplasm
Q8PP84	SAHH_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PPA3	LPXC_XANAC		cytoplasm
Q8PPA6	DDL_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PPA7	MURC_XANAC	cytoplasm (probable).	cytoplasm
Q8PPA8	MURG_XANAC	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8PPB0	MRAY_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PPB2	MURE_XANAC	cytoplasm (probable).	cytoplasm

Q8PPB5	MRAW_XANAC		cytoplasm
Q8PPB6	MRAZ_XANAC		inner membrane
Q8PPC2	Y764_XANAC		inner membrane
Q8PPC8	ATKC_XANAC	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8PPC9	ATKB_XANAC	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8PPD5	NUSB_XANAC		cytoplasm
Q8PPD6	RISB_XANAC		cytoplasm
Q8PPE2	NRDR_XANAC		cytoplasm
Q8PPE3	GLYA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PPF1	GFA_XANAC		extracellular
Q8PPG1	ENGB_XANAC		cytoplasm
Q8PPG7	BETB_XANAC		cytoplasm
Q8PPG8	BETA_XANAC		cytoplasm
Q8PPG9	CCA_XANAC		cytoplasm
Q8PPL8	LIPA_XANAC	cytoplasm (potential).	cytoplasm
Q8PPL9	LIPB_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PPM0	Y666_XANAC		inner membrane
Q8PPP2	UBIE_XANAC		cytoplasm
Q8PPP7	HSLU_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PPP8	HSLV_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PPP9	XERC_XANAC	cytoplasm (by similarity).	nucleoid
Q8PPQ1	DAPF_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PPR3	HEM3_XANAC		cytoplasm
Q8PPR7	OPGH_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PPT1	TREA_XANAC	periplasm (by similarity).	periplasm
Q8PPW9	MDCG_XANAC		cytoplasm
Q8PPX2	MDCC_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PPX9	Y554_XANAC		cytoplasm
Q8PPY3	GLNE_XANAC		cytoplasm
Q8PPZ1	CH60_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PPZ9	AROQ_XANAC		cytoplasm
Q8PQ06	PRMA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PQ10	FISL_XANAC		cytoplasm
Q8PQ19	PUR9_XANAC		cytoplasm
Q8PQ21	PUR2_XANAC		cytoplasm
Q8PQ40	NUDH_XANAC		cytoplasm
Q8PQ41	RS9_XANAC		cytoplasm
Q8PQ44	SPED_XANAC		cytoplasm
Q8PQ47	TRPC_XANAC		cytoplasm
Q8PQ48	TRPD_XANAC		cytoplasm
Q8PQ58	PUR7_XANAC		cytoplasm
Q8PQ74	HGD_XANAC		cytoplasm
Q8PQA2	GLGB2_XANAC		extracellular
Q8PQA3	GLGA_XANAC		cytoplasm
Q8PQA7	OPGB_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PQE0	BIOH_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PQE8	GIDA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PQG7	GLPK_XANAC		cytoplasm

Q8PQI0	ILVD_XANAC		cytoplasm
Q8PQI6	KDGT_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PQK3	NPD_XANAC	cytoplasm (probable).	cytoplasm
Q8PQN3	RLUE_XANAC		cytoplasm
Q8PQT0	UBIB_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PQU9	GPDA_XANAC	cytoplasm (probable).	cytoplasm
Q8PQV0	SECB_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PQW6	UPPP_XANAC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8PR01	KDUI_XANAC		cytoplasm
Q8PR13	GLGB1_XANAC		extracellular
Q8PR25	Y141_XANAC		cytoplasm
Q8PR33	LLDD_XANAC		cytoplasm
Q8PR66	Y100_XANAC		cytoplasm
Q8PRF1	PDXJ_XANAC	cytoplasm (probable).	cytoplasm
Q8PRG0	RECF_XANAC	cytoplasm (by similarity).	cytoplasm
Q8PRG2	DNAA_XANAC	cytoplasm (by similarity).	cytoplasm
Q8RIT7	CH60_XANCH	cytoplasm (by similarity).	cytoplasm
Q8RKT5	LIFO_PSEME	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
Q8RLV9	AROAXENNE	cytoplasm (probable).	cytoplasm
Q8RLW0	SERC_XENNE	cytoplasm (by similarity).	cytoplasm
Q8RLY6	RPIA_ENTCL		cytoplasm
Q8RNM2	PURA_LEGPN	cytoplasm (by similarity).	cytoplasm
Q8RNP4	TYPH_LEGPN		cytoplasm
Q8RP98	LEUD_METEX		cytoplasm
Q8RPN9	CCME_LEGPN	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8RPP4	CCMA_LEGPN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8RPX9	UREE_RHILV	cytoplasm (by similarity).	cytoplasm
Q8RPY7	URE3_RHILV	cytoplasm (by similarity).	cytoplasm
Q8RPZ9	RL20_AZOVI		cytoplasm
Q8RQ00	RL35_AZOVI		cytoplasm
Q8RQD1	GLND_AZOBR		cytoplasm
Q8RQD2	4OT2_PSEPU		cytoplasm
Q8RQS4	AAEX_SERMA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8RR17	FIEF_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8RSS3	CARB_HALER		cytoplasm
Q8RSS4	CARA_HALER		cytoplasm
Q8RSX4	COAD_PROMI	cytoplasm (by similarity).	cytoplasm
Q8RSY9	GUN_PSEFL	secreted protein (by similarity).	extracellular

Q8RT64	RRF_BARBA	cytoplasm (by similarity).	cytoplasm
Q8RT65	PYRH_BARBA		cytoplasm
Q8RT67	COAD_BARBA	cytoplasm (by similarity).	cytoplasm
Q8RTI2	RBL2_MAGMG		extracellular
Q8RTJ3	RL10_XANCP		cytoplasm
Q8RTJ4	RL1_XANCP		cytoplasm
Q8RTJ5	RL11_XANCP		cytoplasm
Q8U527	Y2691_AGRT5	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8U530	YR5A_AGRT5	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8U642	DNE23_AGRT5	cytoplasm (probable).	cytoplasm
Q8U671	DHAA_AGRT5		cytoplasm
Q8U6M1	FBPC_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8U6T2	Y4724_AGRT5		cytoplasm
Q8U705	ARLY2_AGRT5	cytoplasm (probable).	cytoplasm
Q8U7B8	AROQ2_AGRT5		cytoplasm
Q8U7F8	Y4491_AGRT5		inner membrane
Q8U7G6	XYLA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8U7I5	TAL_AGRT5	cytoplasm (probable).	cytoplasm
Q8U7P1	NAPA_AGRT5	periplasm (by similarity).	periplasm
Q8U7Y1	XYLD_AGRT5		cytoplasm
Q8U7Y5	GLYA2_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8U822	Y4277_AGRT5		inner membrane
Q8U8I2	SDH_AGRT5		cytoplasm
Q8U8I9	KCY_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8U8J3	NADA_AGRT5	cytoplasm (by similarity).	inner membrane
Q8U8J4	NADB_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8U8L4	GLGB_AGRT5		extracellular
Q8U8L5	GLGC_AGRT5		cytoplasm
Q8U8M6	RS21A_AGRT5		cytoplasm
Q8U8Q0	Y4040_AGRT5		inner membrane
Q8U8R2	RECX_AGRT5	cytoplasm (potential).	cytoplasm
Q8U8T9	BIOD_AGRT5		cytoplasm
Q8U8Z6	HUTI_AGRT5	cytoplasm (potential).	cytoplasm
Q8U8Z7	HUTH_AGRT5	cytoplasm (potential).	cytoplasm
Q8U8Z9	HUTU_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8U919	Y3911_AGRT5		inner membrane
Q8U920	Y3910_AGRT5		inner membrane
Q8U940	GLPK_AGRT5		cytoplasm
Q8U9D8	ATKA_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8U9D9	ATKB_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8U9E0	ATKC_AGRT5	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8U9F7	HEMH_AGRT5	cytoplasm (by similarity).	extracellular
Q8U9F9	OMP10_AGRT5	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane

Q8U9H0	Y3758_AGRT5		inner membrane
Q8U9I5	RL31_AGRT5		cytoplasm
Q8U9I9	PGK_AGRT5	cytoplasm (potential).	cytoplasm
Q8U9K1	Y3727_AGRT5		inner membrane
Q8U9K4	RUVC_AGRT5		cytoplasm
Q8U9K5	RUVA_AGRT5		cytoplasm
Q8U9K6	RUVB_AGRT5		cytoplasm
Q8U9L4	TOLB_AGRT5	periplasm (potential).	periplasm
Q8U9L5	OMP16_AGRT5	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q8U9L7	TILS_AGRT5	cytoplasm (probable).	cytoplasm
Q8U9L9	GLMM_AGRT5		cytoplasm
Q8U9M5	TRMU_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8U9Q9	PAND_AGRT5		cytoplasm
Q8U9T8	RS21C_AGRT5		cytoplasm
Q8U9U5	ACCA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8U9U6	XERD_AGRT5	cytoplasm (by similarity).	nucleoid
Q8U9V0	AROB_AGRT5	cytoplasm (probable).	cytoplasm
Q8U9V8	RL28_AGRT5		cytoplasm
Q8U9W3	HIS8_AGRT5		cytoplasm
Q8U9X6	ARLY1_AGRT5	cytoplasm (probable).	cytoplasm
Q8U9Z2	KPTA_AGRT5		cytoplasm
Q8UA46	UXUA2_AGRT5		cytoplasm
Q8UA56	ARGJ_AGRT5	cytoplasm (probable).	cytoplasm
Q8UA66	UBIG_AGRT5		cytoplasm
Q8UA73	CYSA2_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UA91	PANB_AGRT5		cytoplasm
Q8UA92	PANC_AGRT5		cytoplasm
Q8UAA5	GPMI_AGRT5		cytoplasm
Q8UAE8	SSUD_AGRT5		cytoplasm
Q8UAN8	3MGH_AGRT5		cytoplasm
Q8UAS8	THIE_AGRT5		cytoplasm
Q8UAS9	THIM_AGRT5		cytoplasm
Q8UAW0	UXUA1_AGRT5		cytoplasm
Q8UAW8	MINC_AGRT5		inner membrane
Q8UAX0	MINE_AGRT5		inner membrane
Q8UAY9	DNE21_AGRT5	cytoplasm (probable).	cytoplasm
Q8UB75	KDUI_AGRT5		cytoplasm
Q8UB77	KDGD_AGRT5		cytoplasm
Q8UB93	RS21B_AGRT5		cytoplasm
Q8UBL6	DCUP_AGRT5	cytoplasm (probable).	cytoplasm
Q8UBM0	GIDA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UBM1	GIDB_AGRT5		cytoplasm
Q8UBM5	HTPX_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8UBM8	PUR9_AGRT5		cytoplasm
Q8UBP3	COBQ_AGRT5		cytoplasm
Q8UBQ6	CBID_AGRT5		cytoplasm
Q8UBQ8	COBB_AGRT5		cytoplasm
Q8UBR0	LEUD_AGRT5		cytoplasm
Q8UBR6	RL27_AGRT5		cytoplasm
Q8UBS0	PROB_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UBS1	PROA_AGRT5	cytoplasm (by similarity).	cytoplasm

Q8UBS2	NADD_AGRT5		cytoplasm
Q8UBS4	Y2776_AGRT5		inner membrane
Q8UBS8	NUDH_AGRT5		cytoplasm
Q8UBV5	ACSA_AGRT5		cytoplasm
Q8UBX8	MTGA_AGRT5	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8UBY9	LEU2_AGRT5		cytoplasm
Q8UBZ5	RL19_AGRT5		cytoplasm
Q8UBZ7	TRMD_AGRT5	cytoplasm (potential).	cytoplasm
Q8UBZ8	RIMM_AGRT5	cytoplasm (potential).	cytoplasm
Q8UBZ9	RS16_AGRT5		cytoplasm
Q8UC03	DAPF_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UC06	ISPZ_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UC12	CCMA_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UC22	PHS_AGRT5		cytoplasm
Q8UC31	ASSY_AGRT5	cytoplasm (probable).	cytoplasm
Q8UC37	IPYR_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UC38	Y2660_AGRT5		inner membrane
Q8UC46	HEM3_AGRT5		cytoplasm
Q8UC48	GPDA_AGRT5	cytoplasm (probable).	cytoplasm
Q8UC57	OMP19_AGRT5	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q8UC60	SUCC_AGRT5		cytoplasm
Q8UC70	XERC_AGRT5	cytoplasm (by similarity).	nucleoid
Q8UC74	ATPA_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UC75	ATPG_AGRT5		inner membrane
Q8UC77	ATPE_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UC79	FDHD_AGRT5	cytoplasm (potential).	cytoplasm
Q8UC86	DXR_AGRT5		cytoplasm
Q8UCA5	CUTC_AGRT5	cytoplasm (probable).	periplasm
Q8UCC9	THIC_AGRT5		cytoplasm
Q8UCD2	THIG_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UCD5	MODC_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UCE7	PUR72_AGRT5		cytoplasm
Q8UCM8	UNG_AGRT5	cytoplasm (by similarity).	extracellular
Q8UCN6	PURA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UCS6	URE3_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UCS8	URE2_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UCT2	URE1_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UCT5	UREE_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UD04	ALLA_AGRT5		extracellular
Q8UD28	ENGA_AGRT5		cytoplasm
Q8UD54	QUEF_AGRT5	cytoplasm (probable).	cytoplasm
Q8UD57	CLPP3_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UD63	LEU1_AGRT5		cytoplasm

Q8UD80	HEM6_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UD95	CLPS2_AGRT5		cytoplasm
Q8UD97	PTH_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UDA0	RL25_AGRT5		cytoplasm
Q8UDA9	KPRS_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UDB4	LGT_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UDE5	GREA_AGRT5		cytoplasm
Q8UDE9	CARB_AGRT5		cytoplasm
Q8UDF5	NORM_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UDF7	CARA_AGRT5		cytoplasm
Q8UDM3	MURE_AGRT5	cytoplasm (probable).	cytoplasm
Q8UDM5	MRAY_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UDM6	MURD_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UDM8	MURG_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UDM9	MURC_AGRT5	cytoplasm (probable).	cytoplasm
Q8UDN0	MURB_AGRT5	cytoplasm (probable).	inner membrane
Q8UDN3	DDL_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UDP9	PRMA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UDS3	TYSY_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UDU5	PDXJ_AGRT5	cytoplasm (probable).	cytoplasm
Q8UDV0	ILVC_AGRT5		cytoplasm
Q8UE04	RL11_AGRT5		cytoplasm
Q8UE05	RL1_AGRT5		cytoplasm
Q8UE06	RL10_AGRT5		cytoplasm
Q8UE07	RL7_AGRT5		cytoplasm
Q8UE08	RPOB_AGRT5		cytoplasm
Q8UE09	RPOC_AGRT5		cytoplasm
Q8UE13	RS12_AGRT5		cytoplasm
Q8UE14	RS7_AGRT5		cytoplasm
Q8UE17	RS10_AGRT5		cytoplasm
Q8UE18	RL3_AGRT5		cytoplasm
Q8UE19	RL4_AGRT5		cytoplasm
Q8UE21	RL2_AGRT5		cytoplasm
Q8UE22	RS19_AGRT5		cytoplasm
Q8UE23	RL22_AGRT5		cytoplasm
Q8UE24	RS3_AGRT5		cytoplasm
Q8UE25	RL16_AGRT5		cytoplasm
Q8UE26	RL29_AGRT5		cytoplasm
Q8UE27	RS17_AGRT5		cytoplasm
Q8UE29	RL24_AGRT5		cytoplasm
Q8UE30	RL5_AGRT5		cytoplasm
Q8UE32	RS8_AGRT5		cytoplasm
Q8UE34	RL18_AGRT5		cytoplasm
Q8UE35	RS5_AGRT5		cytoplasm
Q8UE36	RL15_AGRT5		cytoplasm
Q8UE38	KAD_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UE39	RS13_AGRT5		cytoplasm
Q8UE40	RS11_AGRT5		cytoplasm

Q8UE43	ILVD_AGRT5		cytoplasm
Q8UE66	RIBB_AGRT5		cytoplasm
Q8UE71	COBT_AGRT5		cytoplasm
Q8UE93	MURI_AGRT5		cytoplasm
Q8UEA0	RS4_AGRT5		cytoplasm
Q8UEA1	GLSA_AGRT5		extracellular
Q8UEB0	PURL_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEB7	PUR71_AGRT5		cytoplasm
Q8UED2	SYU_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEH0	GLMU_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEH1	GLMS_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEK7	HIS3_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEK8	GCH1_AGRT5		cytoplasm
Q8UEL0	Y1747_AGRT5		inner membrane
Q8UEM1	MNTH_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8UEM4	NRDJ_AGRT5		cytoplasm
Q8UEM6	XGPT_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEN6	SYV_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEP3	DGTL1_AGRT5		cytoplasm
Q8UEP9	TATA_AGRT5	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8UEQ0	TATB_AGRT5	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8UEQ3	SURE_AGRT5	cytoplasm (potential).	periplasm
Q8UEQ7	MOBA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UER0	MOAA_AGRT5		cytoplasm
Q8UER6	MOAC_AGRT5		cytoplasm
Q8UER8	TRPD_AGRT5		cytoplasm
Q8UES4	COAD_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UES7	QUEA_AGRT5	cytoplasm (potential).	cytoplasm
Q8UES8	TGT_AGRT5		cytoplasm
Q8UET6	GID_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEU0	TIG_AGRT5		cytoplasm
Q8UEU8	Y1656_AGRT5		outer membrane
Q8UEX6	CLPP1_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEY3	TPIS_AGRT5		cytoplasm
Q8UEY5	PYRG_AGRT5		cytoplasm
Q8UEY7	FUMC_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UEY9	GPH_AGRT5		cytoplasm
Q8UEZ0	RPIA_AGRT5		cytoplasm
Q8UEZ6	RADC_AGRT5		cytoplasm
Q8UF02	ACPXL_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UF44	LIPB_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UF49	DEFL_AGRT5		cytoplasm
Q8UF86	UVRA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UF87	SSB_AGRT5		cytoplasm
Q8UFA0	KTHY_AGRT5		cytoplasm
Q8UFC0	Y1478_AGRT5		inner membrane
Q8UFC8	CRCB_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UFD5	GCSH_AGRT5		cytoplasm

Q8UFD6	GCSP_AGRT5		cytoplasm
Q8UFE8	HFQ_AGRT5		cytoplasm
Q8UFF4	ISPDF_AGRT5		cytoplasm
Q8UFG1	LIPA_AGRT5	cytoplasm (potential).	cytoplasm
Q8UFH1	ENO_AGRT5	cytoplasm (by similarity).	extracellular
Q8UFH3	KDSA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UFI8	ALLH_AGRT5		extracellular
Q8UFK2	LEXA_AGRT5		cytoplasm
Q8UFL3	LPXA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UFL4	FABZ_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UFL5	LPXD_AGRT5		cytoplasm
Q8UFL7	Y1380_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8UFL9	UPPS_AGRT5		inner membrane
Q8UFM0	RRF_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UFM1	PYRH_AGRT5		cytoplasm
Q8UFM3	RS2_AGRT5		cytoplasm
Q8UFN4	CLPS1_AGRT5		cytoplasm
Q8UFP1	Y1356_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UFR5	AROQ1_AGRT5		cytoplasm
Q8UFR8	LFTR_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UFS3	GATB_AGRT5		cytoplasm
Q8UFS8	GATA_AGRT5		cytoplasm
Q8UFS9	GATC_AGRT5		cytoplasm
Q8UFT1	RUVX_AGRT5	cytoplasm (potential).	cytoplasm
Q8UFT9	PYRB_AGRT5		cytoplasm
Q8UFU1	Y1306_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UFU8	RL33_AGRT5		cytoplasm
Q8UFV3	DPO41_AGRT5	cytoplasm (probable).	cytoplasm
Q8UFV7	LOLD_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UFY5	CLPX_AGRT5		cytoplasm
Q8UFY6	CLPP2_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UFZ8	RS9_AGRT5		cytoplasm
Q8UG00	ARGC_AGRT5	cytoplasm (probable).	cytoplasm
Q8UG34	ACPD_AGRT5		extracellular
Q8UG61	IHFA_AGRT5		cytoplasm
Q8UG62	FABH_AGRT5	cytoplasm (probable).	cytoplasm
Q8UG63	PLSX_AGRT5		cytoplasm
Q8UG69	NUSB_AGRT5		cytoplasm
Q8UG70	RISB_AGRT5		cytoplasm
Q8UG74	NRDR_AGRT5		cytoplasm
Q8UG75	GLYA1_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UG81	ATE_AGRT5	cytoplasm (potential).	cytoplasm
Q8UG98	PUR5_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UGB6	NDK_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UGC8	AMPA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UGD4	PDXA_AGRT5	cytoplasm (probable).	cytoplasm
Q8UGD5	KSGA_AGRT5		cytoplasm
Q8UGD7	KGUA_AGRT5	cytoplasm (by similarity).	cytoplasm

Q8UGE2	ACP_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UGE7	RS6_AGRT5		cytoplasm
Q8UGE8	RS18_AGRT5		cytoplasm
Q8UGF0	RL9_AGRT5		cytoplasm
Q8UGH4	PSD_AGRT5		cytoplasm
Q8UGH9	Y1058_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UGJ8	RECO_AGRT5		cytoplasm
Q8UGK1	ERA_AGRT5		cytoplasm
Q8UGK2	RNC_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UGK4	ACPS_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UGK8	RPOZ_AGRT5		cytoplasm
Q8UGL2	SSRP_AGRT5	cytoplasm (potential).	cytoplasm
Q8UGL3	DAPA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UGQ4	GLNE_AGRT5		cytoplasm
Q8UGR1	CCME_AGRT5	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8UGU0	Y945_AGRT5		inner membrane
Q8UGV3	Y932_AGRT5		inner membrane
Q8UGX7	MSRB_AGRT5		cytoplasm
Q8UH04	C556_AGRT5		periplasm
Q8UH15	TAM_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UH55	BETA_AGRT5		cytoplasm
Q8UH56	BETB_AGRT5		cytoplasm
Q8UH57	BETI_AGRT5		cytoplasm
Q8UH62	CYSA1_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UH67	CYSH_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UH69	CYSN_AGRT5		cytoplasm
Q8UH73	MQO_AGRT5		cytoplasm
Q8UHA1	Y782_AGRT5		cytoplasm
Q8UHA2	Y781_AGRT5		inner membrane
Q8UHA7	RNH_AGRT5	cytoplasm (potential).	extracellular
Q8UHA8	KHSE_AGRT5		cytoplasm
Q8UHB2	COXZ_AGRT5		inner membrane
Q8UHC2	PDXH_AGRT5		cytoplasm
Q8UHD5	EX7S_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UHD7	DXS_AGRT5		cytoplasm
Q8UHE7	Y735_AGRT5		cytoplasm
Q8UHG2	RNH2_AGRT5	cytoplasm (potential).	extracellular
Q8UHH1	KUP1_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UHI3	MUTL_AGRT5		cytoplasm
Q8UHI5	LPXK_AGRT5		cytoplasm
Q8UHI6	SYI_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UHK1	HIS1_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UHK2	HISZ_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UHN3	PUR2_AGRT5		cytoplasm
Q8UHP8	ISPE_AGRT5		cytoplasm
Q8UHR0	RRMJ_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UHV1	FLBT_AGRT5		cytoplasm

Q8UHW9	MURA_AGRT5	cytoplasm (probable).	inner membrane
Q8UHX1	HISX_AGRT5		cytoplasm
Q8UHX2	Y536_AGRT5		inner membrane
Q8UHX5	Y533_AGRT5	cytoplasm (potential).	inner membrane
Q8UHY0	MSCL_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UI24	AQPZ1_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UI68	APAG_AGRT5		inner membrane
Q8UI70	OTC_AGRT5	cytoplasm (probable).	cytoplasm
Q8UI71	ARGD_AGRT5	cytoplasm (probable).	cytoplasm
Q8UI76	PSTB_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UI94	G6PI_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UI98	PYRE_AGRT5		cytoplasm
Q8UI99	PYRC_AGRT5		cytoplasm
Q8UIB2	RNPA_AGRT5		extracellular
Q8UIB3	OXAA_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UIB4	ENGB_AGRT5		cytoplasm
Q8UIB7	ARGB_AGRT5	cytoplasm (probable).	cytoplasm
Q8UIC6	DAPD_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UIC9	TRUA_AGRT5		cytoplasm
Q8UID0	FMT_AGRT5		cytoplasm
Q8UID1	DEF_AGRT5		cytoplasm
Q8UID4	TRMB_AGRT5		cytoplasm
Q8UID7	LNT_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UID9	Y358_AGRT5		inner membrane
Q8UIF1	GLND_AGRT5		cytoplasm
Q8UIF2	MUTS_AGRT5		cytoplasm
Q8UIF9	IHFB_AGRT5		cytoplasm
Q8UIG8	RNPH_AGRT5		cytoplasm
Q8UIG9	HAM1_AGRT5		extracellular
Q8UIH1	DNAA_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UIH2	RS20_AGRT5		cytoplasm
Q8UIH4	FPG_AGRT5		cytoplasm
Q8UIH5	UBIE_AGRT5		cytoplasm
Q8UII1	DUT_AGRT5		extracellular
Q8UII5	GSHB_AGRT5		cytoplasm
Q8UIJ2	Y303_AGRT5		inner membrane
Q8UIL2	GUAA_AGRT5		cytoplasm
Q8UIM4	EX7L_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UIN7	RL20_AGRT5		cytoplasm
Q8UIN8	RL35_AGRT5		cytoplasm
Q8UIQ2	LEPA_AGRT5		cytoplasm
Q8UIR1	Y232_AGRT5		cytoplasm
Q8UIS9	PNCB_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UIV6	MGSA_AGRT5		cytoplasm
Q8UIV7	GLK_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UIV8	DAPB_AGRT5	cytoplasm (by similarity).	cytoplasm

Q8UIW7	PHNC_AGRT5	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8UJ04	DEOB_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJ05	ADD_AGRT5		extracellular
Q8UJ06	UPP_AGRT5		cytoplasm
Q8UJ08	TYPH_AGRT5		cytoplasm
Q8UJ09	DEOC_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJ19	SLYX_AGRT5		inner membrane
Q8UJ26	RL32_AGRT5		cytoplasm
Q8UJ37	CYCM_AGRT5	bacterial cell membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q8UJ43	Y095_AGRT5		inner membrane
Q8UJ45	RECR_AGRT5		cytoplasm
Q8UJ48	Y090_AGRT5		inner membrane
Q8UJ52	RBFA_AGRT5	cytoplasm (potential).	cytoplasm
Q8UJ53	TRUB_AGRT5		cytoplasm
Q8UJ54	RS15_AGRT5		cytoplasm
Q8UJ65	RECF_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJ69	NRDI_AGRT5		inner membrane
Q8UJ85	MASZ_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJ87	HSLU_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJ88	HSLV_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJ89	HIS7_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJ91	HIS2_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJ92	COAA_AGRT5	cytoplasm (probable).	cytoplasm
Q8UJ94	PPCK_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJ99	SAHH_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJA9	TRPA_AGRT5		cytoplasm
Q8UJB0	TRPB_AGRT5		cytoplasm
Q8UJB1	TRPF_AGRT5		cytoplasm
Q8UJC2	SECB_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJC4	COAE_AGRT5	cytoplasm (by similarity).	cytoplasm
Q8UJC5	AROE_AGRT5		cytoplasm
Q8UJC6	Y002_AGRT5	cytoplasm (potential).	inner membrane
Q8UJC7	Y001_AGRT5		cytoplasm
Q8UJK7	DPO42_AGRT5	cytoplasm (probable).	cytoplasm
Q8UJL0	KUP2_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UJW4	AQPZ2_AGRT5	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8UJW6	Y5359_AGRT5		cytoplasm
Q8UK38	GLGA2_AGRT5		cytoplasm
Q8UK62	Y5261_AGRT5		periplasm
Q8UKK2	DNE22_AGRT5	cytoplasm (probable).	cytoplasm
Q8VL02	NAPA_ACTAC	periplasm (by similarity).	periplasm
Q8VLP6	INH_YERRU	periplasm (by similarity).	periplasm
Q8VM97	CRCB_RALEU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8VMM4	SSBW_PSEPU		cytoplasm
Q8VMM5	REXP_PSEPU	cytoplasm (potential).	cytoplasm
Q8VMR3	HUTH_PSEFL	cytoplasm (potential).	cytoplasm

Q8VNZ2	MVIN_BDEBA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8VP65	AROA_PASPI	cytoplasm (probable).	cytoplasm
Q8VPB5	ACSF_RHOGE		inner membrane
Q8VQ22	FABZ_BARHE	cytoplasm (by similarity).	cytoplasm
Q8VQ23	LPXD_BARHE		cytoplasm
Q8VQ25	Y627_BARHE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8VQ84	RBL_NITST		extracellular
Q8VQP8	FLIE_BURCE		flagellar
Q8VQX6	RECX_MYXXA	cytoplasm (potential).	cytoplasm
Q8VQZ3	YDFE_MYXXA		inner membrane
Q8VR31	PYRE_METCA		cytoplasm
Q8VRZ3	HPPA_RHIME	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8VS06	XERC_PSEFL	cytoplasm (by similarity).	nucleoid
Q8VS50	MSRA_XANCH		cytoplasm
Q8VSR0	CYSI_ACTPL		cytoplasm
Q8VTT5	HYDA_BURPI		extracellular
Q8VV76	ATPE_COLMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8VV78	ATPG_COLMA		inner membrane
Q8VW58	RL27_BRUAB		cytoplasm
Q8VW75	COAD_PASPI	cytoplasm (by similarity).	cytoplasm
Q8VW76	FPG_PASPI		cytoplasm
Q8VW94	RBL_NITS1		extracellular
Q8VWE6	LRHA_ECO57		cytoplasm
Q8X3J5	MDTF_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X3J8	YJDP_ECO57		nucleoid
Q8X3T0	RHAT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X3U5	RCNA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X400	ZITB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X444	UNG_ECO57	cytoplasm (by similarity).	extracellular
Q8X492	ZIPA_ECO57	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q8X4B4	WRBA_ECO57		cytoplasm
Q8X4I9	ALR2_ECO57		cytoplasm
Q8X4L0	MDTE_ECO57	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
Q8X4L6	NIKE_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q8X4L7	BTUC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X4L9	RLUF_ECO57		cytoplasm
Q8X4P8	FUMA_ECO57		cytoplasm
Q8X4Q7	KDGT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X4Q8	RLUB_ECO57		cytoplasm
Q8X4S4	TSGA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X4S6	ARGD_ECO57	cytoplasm (probable).	cytoplasm
Q8X4T6	XERC_ECO57	cytoplasm (by similarity).	nucleoid
Q8X4V5	GSA_ECO57	cytoplasm (potential).	cytoplasm
Q8X4V6	YGAY_ECO57	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8X4V7	MODC_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8X4W4	NANR_ECO57		cytoplasm
Q8X4Z7	RUTR_ECO57		cytoplasm
Q8X4Z9	YDEO_ECO57		cytoplasm
Q8X505	YDFG_ECO57		cytoplasm
Q8X524	QSEC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X529	YGHZ_ECO57		inner membrane
Q8X534	ENTS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X560	USPC_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X564	YFCJ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X574	XERD_ECO57	cytoplasm (by similarity).	nucleoid
Q8X580	SLMA_ECO57	nucleoid-associated (by similarity).	cytoplasm
Q8X583	YCGV_ECO57		outer membrane
Q8X598	ARGM_ECO57		cytoplasm
Q8X5C4	YFCN_ECO57		inner membrane
Q8X5I5	LFTR_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X5I6	TAUB_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8X5L7	BCSA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8X5L8	BCSB_ECO57	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q8X5L9	GUN_ECO57	secreted protein (by similarity).	extracellular
Q8X5M0	BCSC_ECO57		outer membrane

Q8X5M2	DCTA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X5P3	DCTR_ECO57		cytoplasm
Q8X5R2	MDTN_ECO57	bacterial cell inner membrane; single-pass type ii membrane protein (potential).	inner membrane
Q8X5R8	MDTO_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X5R9	MDTP_ECO57	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
Q8X5S3	NRFG_ECO57		extracellular
Q8X5S5	NRFF_ECO57	periplasm (probable).	periplasm
Q8X5T5	ACSA_ECO57		cytoplasm
Q8X5T7	ACTP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X5U1	NIKD_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8X5U4	ACPT_ECO57		cytoplasm
Q8X5U9	UVRA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X5V2	ALR1_ECO57		cytoplasm
Q8X5V3	DNAB_ECO57		cytoplasm
Q8X5V6	DUSA_ECO57		cytoplasm
Q8X5W0	BTUD_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8X5W3	YDIU_ECO57		inner membrane
Q8X5W4	AROH_ECO57		cytoplasm
Q8X5W7	LAMB_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8X5X7	AROD_ECO57		cytoplasm
Q8X5X8	PSIE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X5Y4	YDIB_ECO57		cytoplasm
Q8X5Z7	YJBB_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8X607	ACEK_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X610	META_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X611	PUR9_ECO57		cytoplasm
Q8X612	PUR2_ECO57		cytoplasm
Q8X613	ZRAR_ECO57	cytoplasm (probable).	cytoplasm
Q8X614	ZRAS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X616	YDHX_ECO57		periplasm
Q8X621	YD HQ_ECO57		extracellular
Q8X625	YDHP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q8X638	YDHK_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8X645	YEIT_ECO57		inner membrane
Q8X647	PDXH_ECO57		cytoplasm
Q8X648	CDD_ECO57		cytoplasm
Q8X651	TPPB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X653	YOHJ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X659	MDTQ_ECO57	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
Q8X661	ADD_ECO57		extracellular
Q8X693	PRPD_ECO57		cytoplasm
Q8X6C0	YGEW_ECO57		cytoplasm
Q8X6C3	BETI_ECO57		cytoplasm
Q8X6C4	XDHC_ECO57		cytoplasm
Q8X6C5	XDHB_ECO57		cytoplasm
Q8X6C6	BETA_ECO57	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8X6C7	XDHA_ECO57		periplasm
Q8X6E3	Y387_ECO57		inner membrane
Q8X6H3	PBL_ECO57	periplasm (potential).	periplasm
Q8X6I8	KDUI_ECO57		cytoplasm
Q8X6I9	YAGT_ECO57		cytoplasm
Q8X6J0	YAGS_ECO57		cytoplasm
Q8X6J4	YAGR_ECO57		periplasm
Q8X6J8	AAS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X6K2	MUTH_ECO57	cytoplasm (by similarity).	extracellular
Q8X6K6	AQPZ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X6L0	HCP_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X6M8	PTRA_ECO57	periplasm (by similarity).	extracellular
Q8X6P4	YHHS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X6Q5	RUMB_ECO57		cytoplasm
Q8X6R3	FUCK_ECO57		cytoplasm
Q8X6R5	FUCI_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X6R8	FUCA_ECO57		extracellular
Q8X6S9	QUEF_ECO57	cytoplasm (probable).	cytoplasm
Q8X6T6	TRUC_ECO57		cytoplasm
Q8X6V3	YLIF_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8X6X5	DCUP_ECO57	cytoplasm (probable).	cytoplasm
Q8X6X6	GLGB_ECO57		extracellular
Q8X6X7	NUDC_ECO57		cytoplasm
Q8X6X8	GLGX_ECO57		extracellular
Q8X6X9	THIC_ECO57		cytoplasm

Q8X6Y0	THIE_ECO57		cytoplasm
Q8X6Z5	GLPE_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X701	MALT_ECO57		cytoplasm
Q8X711	MURB_ECO57	cytoplasm (probable).	inner membrane
Q8X714	BTUB_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8X715	GNTX_ECO57		cytoplasm
Q8X716	BIOH_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X723	TRMA_ECO57		cytoplasm
Q8X724	RLUE_ECO57		cytoplasm
Q8X727	STHA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X730	ARLY_ECO57	cytoplasm (probable).	cytoplasm
Q8X732	ARGC_ECO57	cytoplasm (probable).	cytoplasm
Q8X733	PPCK_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X735	TRMU_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X736	YCFC_ECO57		inner membrane
Q8X738	PHOP_ECO57	cytoplasm (probable).	cytoplasm
Q8X739	PHOQ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X742	ARGE_ECO57	cytoplasm (probable).	extracellular
Q8X743	CAPP_ECO57		cytoplasm
Q8X769	FUMC_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X783	ASR_ECO57		periplasm
Q8X794	CLCB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X799	YNFC_ECO57	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8X7A5	CDH_ECO57	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8X7A6	YNFA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X7B3	RHAR_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X7B5	TRPA_ECO57		cytoplasm
Q8X7B6	TRPB_ECO57		cytoplasm
Q8X7B7	TRPC_ECO57		cytoplasm
Q8X7D6	RNB_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X7G3	THIM_ECO57		cytoplasm
Q8X7H0	YEGX_ECO57		extracellular
Q8X7I0	YEGP_ECO57		extracellular
Q8X7J4	MDTB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8X7J5	MDTA_ECO57	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8X7J8	YEGI_ECO57		cytoplasm
Q8X7L9	WZC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X7N4	PROA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X7N7	YAFA_ECO57		cytoplasm

Q8X7P0	YKFJ_ECO57		inner membrane
Q8X7P7	GNE_ECO57		cytoplasm
Q8X7Q1	DPO4_ECO57	cytoplasm (probable).	cytoplasm
Q8X7R2	FADE_ECO57		cytoplasm
Q8X7S0	YGCU_ECO57		cytoplasm
Q8X7U1	CYSJ_ECO57		cytoplasm
Q8X7U2	CYSI_ECO57		cytoplasm
Q8X7U3	CYSH_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X7U4	MNTR_ECO57	cytoplasm (probable).	cytoplasm
Q8X7U9	HOKG_ECO57		inner membrane
Q8X7W8	YBIX_ECO57		cytoplasm
Q8X7X4	CYSD_ECO57		cytoplasm
Q8X7X7	CYSN_ECO57		cytoplasm
Q8X7X9	DING_ECO57		cytoplasm
Q8X7Y4	ISPD_ECO57		cytoplasm
Q8X7Y6	TRUD_ECO57		cytoplasm
Q8X7Y9	YBHG_ECO57	periplasm (potential).	inner membrane
Q8X7Z7	DKGB_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X800	METI_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X821	BIOD_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X824	AROB_ECO57	cytoplasm (probable).	cytoplasm
Q8X826	DAMX_ECO57		inner membrane
Q8X835	FRLR_ECO57		cytoplasm
Q8X839	FRLD_ECO57		cytoplasm
Q8X840	FRLC_ECO57		cytoplasm
Q8X844	FRLB_ECO57		cytoplasm
Q8X845	FRLA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8X850	NORW_ECO57	cytoplasm (probable).	cytoplasm
Q8X852	NORV_ECO57	cytoplasm (probable).	cytoplasm
Q8X854	NORR_ECO57		cytoplasm
Q8X878	KEFB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X883	TUSD_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X884	TUSC_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X885	TUSB_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X897	RHAS_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X899	RHAB_ECO57		cytoplasm
Q8X8A1	RHAD_ECO57	cytoplasm (by similarity).	extracellular
Q8X8B5	FDHD_ECO57	cytoplasm (potential).	cytoplasm
Q8X8B9	FDHE_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X8D5	YIHT_ECO57		cytoplasm
Q8X8E0	NPD_ECO57	cytoplasm (probable).	cytoplasm
Q8X8E3	LOLD_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8X8F1	FMT_ECO57		cytoplasm
Q8X8F2	OMPL_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8X8F6	SMG_ECO57		cytoplasm
Q8X8F9	AROE_ECO57		cytoplasm

Q8X8G6	THIK_ECO57		cytoplasm
Q8X8G9	YIHI_ECO57		inner membrane
Q8X8H0	ENGB_ECO57		cytoplasm
Q8X8H6	KTHY_ECO57		cytoplasm
Q8X8H9	RDOA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X8I2	FADB_ECO57		cytoplasm
Q8X8I7	FABD_ECO57		cytoplasm
Q8X8J3	RLUC_ECO57		cytoplasm
Q8X8J4	FADA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X8K9	YCJF_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X8L4	DLHH_ECO57		extracellular
Q8X8L5	METE_ECO57		cytoplasm
Q8X8L6	YIGL_ECO57		cytoplasm
Q8X8N8	PYRC_ECO57		cytoplasm
Q8X8P2	YCEA_ECO57		inner membrane
Q8X8Q3	YDAM_ECO57		inner membrane
Q8X8T3	HISX_ECO57		cytoplasm
Q8X8T4	HIS1_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X8U9	COBS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X8V0	COBT_ECO57		cytoplasm
Q8X8V5	YEEN_ECO57		inner membrane
Q8X8V7	YEEJ_ECO57		extracellular
Q8X8V9	METQ_ECO57	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8X8W8	TILS_ECO57	cytoplasm (probable).	cytoplasm
Q8X8X6	RNH2_ECO57	cytoplasm (potential).	extracellular
Q8X8X7	LPXB_ECO57		cytoplasm
Q8X8X8	LPXA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X8Y1	DXR_ECO57		cytoplasm
Q8X8Y6	GLND_ECO57		cytoplasm
Q8X8Y7	DAPD_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X8Y9	DGTP_ECO57		cytoplasm
Q8X8Z0	BTUF_ECO57	periplasm (by similarity).	periplasm
Q8X900	GSH1_ECO57		cytoplasm
Q8X902	LUXS_ECO57		extracellular
Q8X909	GLUQ_ECO57		cytoplasm
Q8X926	6PGL_ECO57		cytoplasm
Q8X929	PANB_ECO57		cytoplasm
Q8X930	PANC_ECO57		cytoplasm
Q8X954	CSID_ECO57		cytoplasm
Q8X963	NADA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X968	AROP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X984	GUAC_ECO57		cytoplasm
Q8X988	YACF_ECO57		inner membrane
Q8X9A9	DHSD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X9B6	ACCC_ECO57		cytoplasm
Q8X9C6	END8_ECO57		cytoplasm
Q8X9C8	YBGL_ECO57		inner membrane

Q8X9E0	AAEX_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8X9E5	AAEA_ECO57	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q8X9E6	AAEB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X9F0	RLUD_ECO57		cytoplasm
Q8X9F8	ATKA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8X9F9	ATKB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8X9G0	ATKC_ECO57	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8X9G8	NANT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X9G9	NANE_ECO57		cytoplasm
Q8X9H0	NANK_ECO57		cytoplasm
Q8X9I1	MTGA_ECO57	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8X9I3	MDTG_ECO57	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8X9I6	OPGC_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8X9J0	KDSD_ECO57		cytoplasm
Q8X9J6	YCDX_ECO57		cytoplasm
Q8X9J9	MURA_ECO57	cytoplasm (probable).	inner membrane
Q8X9L0	FTSH_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8X9L2	GLMM_ECO57		cytoplasm
Q8X9M0	ASSY_ECO57	cytoplasm (probable).	cytoplasm
Q8X9M2	RS15_ECO57		cytoplasm
Q8X9S9	ACPD_ECO57		extracellular
Q8X9T8	R31B2_ECO57		cytoplasm
Q8X9U6	UREE_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X9V1	OPGD_ECO57	periplasm (by similarity).	periplasm
Q8X9X2	YNCE_ECO57		cytoplasm
Q8X9X8	YDCE_ECO57		cytoplasm
Q8X9Y6	DDLB_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X9Y8	MURG_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8X9Y9	MURD_ECO57	cytoplasm (by similarity).	cytoplasm
Q8X9Z2	MURE_ECO57	cytoplasm (probable).	cytoplasm
Q8X9Z8	LEU1_ECO57		cytoplasm
Q8X9Z9	LEU3_ECO57	cytoplasm (by similarity).	cytoplasm

Q8XA00	LEU2_ECO57		cytoplasm
Q8XA01	LEUD_ECO57		cytoplasm
Q8XA10	RLUA_ECO57		cytoplasm
Q8XA13	OSTA_ECO57	bacterial cell outer membrane (by similarity).	outer membrane
Q8XA14	KSGA_ECO57		cytoplasm
Q8XA15	APAH_ECO57		extracellular
Q8XA20	KEFC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XA23	NADB_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XA24	KEFF_ECO57		cytoplasm
Q8XA27	FIXB_ECO57		cytoplasm
Q8XA30	CAIT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XA32	CAIB_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XA34	CAIC_ECO57		cytoplasm
Q8XA35	CAID_ECO57		cytoplasm
Q8XA36	CAIE_ECO57		cytoplasm
Q8XA39	ACPS_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XA41	RIHC_ECO57		extracellular
Q8XA44	TADA_ECO57		cytoplasm
Q8XA48	LSPA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8XA49	SYI_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XA55	GLYA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XA65	DNAJ_ECO57	cytoplasm (probable).	cytoplasm
Q8XA76	YAAA_ECO57		inner membrane
Q8XA81	YHFR_ECO57		inner membrane
Q8XA82	KHSE_ECO57	cytoplasm (potential).	cytoplasm
Q8XA83	YFHQ_ECO57		cytoplasm
Q8XA87	DEAD_ECO57	cytoplasm (probable).	cytoplasm
Q8XA92	YHBQ_ECO57		cytoplasm
Q8XA94	YHBP_ECO57		inner membrane
Q8XAA3	YRAN_ECO57		inner membrane
Q8XAB0	EX7L_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAC5	PUR5_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAD2	YFGC_ECO57		inner membrane
Q8XAF1	TDCD_ECO57		cytoplasm
Q8XAG0	URE1_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAG1	URE2_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAG2	URE3_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAI7	UXAC_ECO57		cytoplasm
Q8XAJ0	ALX_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XAN4	MUG_ECO57	cytoplasm (potential).	extracellular
Q8XAN7	ACPS2_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAP0	CYAY_ECO57		inner membrane
Q8XAP1	CYAA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAP3	HEM3_ECO57		cytoplasm
Q8XAQ3	WECG_ECO57		cytoplasm

Q8XAQ5	WZYE_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8XAQ8	WECF_ECO57	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8XAR3	PGAB_ECO57	bacterial cell membrane; lipid-anchor (probable).	cytoplasm
Q8XAR5	PGAC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XAR8	WECB_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAS7	WECA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XAS9	MAO1_ECO57		cytoplasm
Q8XAT4	RHLB_ECO57		cytoplasm
Q8XAT5	GPPA_ECO57		extracellular
Q8XAU1	RUTA_ECO57		cytoplasm
Q8XAU3	RUTB_ECO57		cytoplasm
Q8XAU7	RUTD_ECO57		cytoplasm
Q8XAU9	RUTE_ECO57		cytoplasm
Q8XAV0	RUTF_ECO57		cytoplasm
Q8XAV1	ILVD_ECO57		cytoplasm
Q8XAW1	HDFR_ECO57		cytoplasm
Q8XAW8	RBSD_ECO57	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8XAW9	KUP_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XAX0	Y2208_ECO57	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8XAX1	YDEP_ECO57		periplasm
Q8XAX4	RAVA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAY0	GIDA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAZ2	TAM_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XAZ3	YNEE_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8XAZ4	YNEF_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8XB14	YJJX_ECO57		inner membrane
Q8XB23	YIDZ_ECO57		cytoplasm
Q8XB24	MDTL_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XB33	TNAB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XB34	TNAA_ECO57		cytoplasm
Q8XB35	TYPH_ECO57		cytoplasm
Q8XB36	DEOC_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XB43	RNPA_ECO57		extracellular

Q8XB55	DNAT_ECO57		nucleoid
Q8XB59	OPGB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XB73	YEDZ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XB74	YEDY_ECO57		cytoplasm
Q8XB75	YEDX_ECO57	periplasm (potential).	periplasm
Q8XB78	HCHA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XB84	MDTM_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XB92	YEDQ_ECO57	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8XB99	MPGP_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XBC7	DCYD_ECO57		cytoplasm
Q8XBI4	HEM6_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XBJ0	YFEW_ECO57		extracellular
Q8XBJ2	YFEU_ECO57		cytoplasm
Q8XBJ4	UCPA_ECO57		cytoplasm
Q8XBJ8	CYSA_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XBK1	YBEY_ECO57		inner membrane
Q8XBK2	LNT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XBK5	TTDB_ECO57		cytoplasm
Q8XBL4	CCA_ECO57		cytoplasm
Q8XBL8	RIHA_ECO57		extracellular
Q8XBM3	GLNE_ECO57		cytoplasm
Q8XBP0	NADD_ECO57		cytoplasm
Q8XBP8	PTFC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XBQ1	RLPA_ECO57	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8XBQ8	PTFAX_ECO57	cytoplasm (probable).	cytoplasm
Q8XBR7	FCTA_ECO57		cytoplasm
Q8XBS3	QSEB_ECO57	cytoplasm (potential).	cytoplasm
Q8XBS4	YGIV_ECO57		cytoplasm
Q8XBS7	YGIQ_ECO57		cytoplasm
Q8XBT3	USPG_ECO57		cytoplasm
Q8XBT6	DKGA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XBV3	ENTE_ECO57		cytoplasm
Q8XBV9	ENTF_ECO57		cytoplasm
Q8XBW0	HYBE_ECO57		cytoplasm
Q8XBW1	HYBF_ECO57		cytoplasm
Q8XBW9	HOKE_ECO57		inner membrane
Q8XBX0	HOKF_ECO57		inner membrane
Q8XBX1	CAAL_ECO57		cytoplasm
Q8XBY1	CUSA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane

Q8XBY2	CUSF_ECO57	periplasm (by similarity).	periplasm
Q8XBY3	CUSC_ECO57	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
Q8XBY4	CUSS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XBZ3	DNAA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XBZ8	YIDS_ECO57		cytoplasm
Q8XC04	IBPB_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XC17	SFA_ECO57		inner membrane
Q8XC28	ETK_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XC50	NLPA_ECO57	bacterial cell inner membrane; lipid-anchor (by similarity).	inner membrane
Q8XCB3	SYV_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XCB7	YCIC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XCB8	OTC_ECO57	cytoplasm (probable).	cytoplasm
Q8XCC6	CLS_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XCE7	OTSA_ECO57		cytoplasm
Q8XCE8	FLHD_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XCG3	MSRA_ECO57		cytoplasm
Q8XCG6	CYSQ_ECO57	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	cytoplasm
Q8XCH4	CUTC_ECO57	cytoplasm (probable).	periplasm
Q8XCH9	RCMNS_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XCI6	YECE_ECO57		cytoplasm
Q8XCJ6	G6PD_ECO57		cytoplasm
Q8XCK0	YEBF_ECO57	secreted protein (by similarity).	inner membrane
Q8XCK5	SDHD_ECO57		cytoplasm
Q8XCM4	PROQ_ECO57	cytoplasm (potential).	inner membrane
Q8XCN6	FADL_ECO57	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XCN8	YEBO_ECO57	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8XCN9	FADI_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XCP1	YEBN_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XCP2	FADJ_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XCP5	YOBD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XCQ5	MEPA_ECO57	periplasm (by similarity).	extracellular
Q8XCQ7	YFCK_ECO57		cytoplasm
Q8XCR0	PDXB_ECO57		cytoplasm

Q8XCS6	MLTC_ECO57	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q8XCU1	LPXH_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XCU5	HAM1_ECO57		extracellular
Q8XCU6	YGGU_ECO57		inner membrane
Q8XCV0	RUVX_ECO57	cytoplasm (potential).	cytoplasm
Q8XCV1	YFBU_ECO57		inner membrane
Q8XCV3	YFBR_ECO57	cytoplasm (potential).	periplasm
Q8XCV5	ARCC_ECO57	cytoplasm (potential).	cytoplasm
Q8XCW1	YQGE_ECO57		inner membrane
Q8XCW6	SPRT_ECO57	cytoplasm (potential).	extracellular
Q8XCX2	NUOG_ECO57		cytoplasm
Q8XCX9	SPEA_ECO57	periplasm (by similarity).	cytoplasm
Q8XCZ0	RNZ_ECO57		extracellular
Q8XCZ2	SELU_ECO57		cytoplasm
Q8XD03	PGK_ECO57	cytoplasm (potential).	cytoplasm
Q8XD09	CUER_ECO57	cytoplasm (probable).	cytoplasm
Q8XD10	ARGO_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XD23	GLSA1_ECO57		extracellular
Q8XD24	ATCU_ECO57	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8XD32	GCST_ECO57		cytoplasm
Q8XD33	GCSP_ECO57		cytoplasm
Q8XD39	HEMH_ECO57	cytoplasm (by similarity).	extracellular
Q8XD58	IDI_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XD64	XDHD_ECO57		cytoplasm
Q8XD75	YGFK_ECO57		cytoplasm
Q8XD82	TUSE_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XD86	RECG_ECO57		cytoplasm
Q8XD88	KGUA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XD99	PYRE_ECO57		cytoplasm
Q8XDB3	YCHJ_ECO57		inner membrane
Q8XDD9	SSUD_ECO57		cytoplasm
Q8XDE7	KDSA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XDE8	PNCB_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XDE9	GPMI_ECO57		cytoplasm
Q8XDF7	LLDD_ECO57		cytoplasm
Q8XDG2	MUKF_ECO57	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q8XDG6	KDSB_ECO57	cytoplasm (potential).	cytoplasm
Q8XDG9	MTLD_ECO57		cytoplasm
Q8XDH0	LPXK_ECO57		cytoplasm
Q8XDH7	TREA_ECO57	periplasm (by similarity).	periplasm
Q8XDI4	ULAF_ECO57		cytoplasm
Q8XDI5	ULAE_ECO57		cytoplasm
Q8XDI7	ULAD_ECO57		cytoplasm
Q8XDJ5	ULAA_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XDJ6	ULAG_ECO57	cytoplasm (potential).	extracellular
Q8XDJ8	LDCA_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XDL7	YCGN_ECO57		inner membrane

Q8XDN0	MINC_ECO57		inner membrane
Q8XDN3	MIAA_ECO57		cytoplasm
Q8XDN4	MUTL_ECO57		cytoplasm
Q8XDP0	ORN_ECO57	cytoplasm (potential).	cytoplasm
Q8XDP1	ENGC_ECO57		cytoplasm
Q8XDQ5	SUGE_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XDR6	LCFA_ECO57	membrane; peripheral membrane protein (potential). partially membrane-associated (potential).	cytoplasm
Q8XDS0	ASPA_ECO57		cytoplasm
Q8XDT8	YEAH_ECO57		inner membrane
Q8XDV7	PHNC_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XDY7	YFBJ_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XDZ0	ASTE_ECO57		periplasm
Q8XDZ1	ARNT_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XDZ3	ARNA_ECO57		cytoplasm
Q8XDZ5	ARNC_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XDZ7	CHO_ECO57		cytoplasm
Q8XDZ9	NADE_ECO57		cytoplasm
Q8XE10	CHBG_ECO57		inner membrane
Q8XE29	UBIG_ECO57		cytoplasm
Q8XE35	YFAS_ECO57		cytoplasm
Q8XE45	MQO_ECO57		cytoplasm
Q8XE46	ECOT_ECO57	periplasm (by similarity).	periplasm
Q8XE47	NAPA_ECO57	periplasm (by similarity).	periplasm
Q8XE58	CCMA_ECO57	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XE68	NDPA_ECO57	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q8XE69	RL25_ECO57		cytoplasm
Q8XE72	PANE_ECO57	cytoplasm (potential).	cytoplasm
Q8XE74	THII_ECO57	cytoplasm (potential).	cytoplasm
Q8XE76	DXS_ECO57		cytoplasm
Q8XE87	THIL_ECO57		cytoplasm
Q8XE94	RIHB_ECO57		extracellular
Q8XE96	ACPH_ECO57		cytoplasm
Q8XEA1	RDGC_ECO57	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q8XEA7	SERC_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XEB7	YCAD_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q8XEC6	DUSC_ECO57		cytoplasm
Q8XED4	FEOB_ECO57	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XED8	RUMA_ECO57		cytoplasm
Q8XEE5	RSMB_ECO57		cytoplasm
Q8XEG2	GLMS_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XEG9	UXUA_ECO57		cytoplasm
Q8XEH3	EUTC_ECO57		cytoplasm
Q8XEH9	CUSB_ECO57		inner membrane
Q8XEJ1	TDH_ECO57	cytoplasm (by similarity).	cytoplasm
Q8XEK3	AAEX_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8XEL4	YBJJ_SALTI		cytoplasm
Q8XEL6	YEJL_SALTI		inner membrane
Q8XEM4	DJLA_SALTI	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q8XEN3	CCME_SALTI	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8XEN6	YGFY_SALTI		cytoplasm
Q8XEQ0	HDA_SALTI		nucleoid
Q8XET6	RL16_SALTI		cytoplasm
Q8XEW9	HLDE_SALTI		cytoplasm
Q8XEY8	GRPE_SALTI	cytoplasm (probable).	cytoplasm
Q8XEZ7	TUSE_SALTI	cytoplasm (by similarity).	cytoplasm
Q8XEZ8	ACPH_SALTI		cytoplasm
Q8XF46	TYPH_SALTI		cytoplasm
Q8XF77	SUFS_SALTI	cytoplasm (by similarity).	cytoplasm
Q8XF81	GLMM_SALTI		cytoplasm
Q8XF83	AAEA_SALTI	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q8XFD0	PGSA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XFE0	GRCA_SALTI		cytoplasm
Q8XFE5	DSBE_SALTI	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q8XFE7	RL25_SALTI		cytoplasm
Q8XFF4	CITD2_SALTI	cytoplasm (by similarity).	cytoplasm
Q8XFG0	MDTM_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XFG2	MUG_SALTI	cytoplasm (potential).	extracellular
Q8XFI3	MTFA_SALTI		cytoplasm
Q8XFK6	DSBE_SALTY	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q8XFQ8	KITH_SALTI	cytoplasm (by similarity).	cytoplasm

Q8XFZ8	ISCA_SALTI		cytoplasm
Q8XG35	CCME_SALTY	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8XG70	SYI_SALTI	cytoplasm (by similarity).	cytoplasm
Q8XG87	ULAG_SALTI	cytoplasm (potential).	extracellular
Q8XG95	ATPA_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XG96	ULAR_SALTI	cytoplasm (potential).	cytoplasm
Q8XGE0	CUTA_SALTI	cytoplasm (by similarity).	periplasm
Q8XGG5	SECB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8XGJ8	HNSB2_SALTI		cytoplasm
Q8XGN1	CITD1_SALTI	cytoplasm (by similarity).	cytoplasm
Q8XGP4	RL18_SALTI		cytoplasm
Q8XGQ8	CBPM_SALTI		cytoplasm
Q8XGT8	SUGE_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XGW7	IBPA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8XGW8	YCDX_SALTI		cytoplasm
Q8XGX8	DCOA_SALTI		inner membrane
Q8XGY4	TADA_SALTI		cytoplasm
Q8XPI1	ACP3_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XPV7	EFE_RALSO		cytoplasm
Q8XQ73	FLHD_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XQ89	AROQ2_RALSO		cytoplasm
Q8XQB8	NOSZ_RALSO	periplasm (by similarity).	periplasm
Q8XQC5	SPEE1_RALSO		cytoplasm
Q8XQC8	SPEE2_RALSO		cytoplasm
Q8XQE8	FUMC_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XQF2	Y4775_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XQG4	NADB2_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XQG9	ACPD_RALSO		extracellular
Q8XQN1	RIBB_RALSO		cytoplasm
Q8XQR9	Y4651_RALSO		cytoplasm
Q8XQS6	GLSA_RALSO		extracellular
Q8XR10	ADC_RALSO		cytoplasm
Q8XR16	KUP1_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XR33	Y4535_RALSO		inner membrane
Q8XR66	DCTA3_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XRC2	Y4436_RALSO		inner membrane
Q8XRG1	Y4393_RALSO		cytoplasm
Q8XRK5	KDGD_RALSO		cytoplasm
Q8XRL7	MQO_RALSO		cytoplasm
Q8XRM1	ATPE2_RALSO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XRN1	DNAE2_RALSO	cytoplasm (probable).	cytoplasm

Q8XRV9	ASPD_RALSO		cytoplasm
Q8XRZ0	HGD_RALSO		cytoplasm
Q8XS00	TRPD2_RALSO		cytoplasm
Q8XS01	TRPC2_RALSO		cytoplasm
Q8XS03	PDXH2_RALSO		cytoplasm
Q8XS05	METE_RALSO		cytoplasm
Q8XS35	1A1D_RALSO		cytoplasm
Q8XS59	CBID_RALSO		cytoplasm
Q8XSF6	MNTH_RALSO	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8XSS8	FLIE_RALSO		flagellar
Q8XSU8	ACP2_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XSW8	FLGI_RALSO	periplasm (by similarity).	periplasm
Q8XSW9	FLGH_RALSO	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8XSZ5	LEU12_RALSO		cytoplasm
Q8XT38	TREA_RALSO	periplasm (by similarity).	periplasm
Q8XT53	OPGD2_RALSO	periplasm (by similarity).	periplasm
Q8XT56	PCP2_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XT73	GLGA_RALSO		cytoplasm
Q8XT76	GLGB_RALSO		extracellular
Q8XTL6	XERC2_RALSO	cytoplasm (by similarity).	nucleoid
Q8XTQ1	GLYA2_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XTT4	CLCL_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XTV4	DNAA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XTY1	Y3414_RALSO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8XU10	ATKC_RALSO	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8XU11	ATKB_RALSO	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8XU12	ATKA_RALSO	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8XU30	MINC_RALSO		inner membrane
Q8XU38	PHS_RALSO		cytoplasm
Q8XU39	PH4H_RALSO		cytoplasm
Q8XU65	GIDA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XU66	GIDB_RALSO		cytoplasm
Q8XU74	ATPA_RALSO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XU75	ATPG_RALSO		inner membrane
Q8XU77	ATPE1_RALSO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XU90	DCUP_RALSO	cytoplasm (probable).	cytoplasm
Q8XU95	CAAL_RALSO		cytoplasm
Q8XU98	GCSP_RALSO		cytoplasm
Q8XU99	GCSH_RALSO		cytoplasm

Q8XUA0	GCST_RALSO		cytoplasm
Q8XUB6	DCTA2_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XUB7	ALLC_RALSO		extracellular
Q8XUC6	Y3265_RALSO		inner membrane
Q8XUR0	EUTC_RALSO		cytoplasm
Q8XUU7	HIP_RALSO	periplasm (potential).	periplasm
Q8XUX6	OPGD1_RALSO	periplasm (by similarity).	periplasm
Q8XUY1	GLPK_RALSO		cytoplasm
Q8XUZ4	RL11_RALSO		cytoplasm
Q8XUZ5	RL1_RALSO		cytoplasm
Q8XUZ6	RL10_RALSO		cytoplasm
Q8XUZ7	RL7_RALSO		cytoplasm
Q8XUZ8	RPOB_RALSO		cytoplasm
Q8XUZ9	RPOC_RALSO		cytoplasm
Q8XV02	Y3030_RALSO		cytoplasm
Q8XV08	RS12_RALSO		cytoplasm
Q8XV09	RS7_RALSO		cytoplasm
Q8XV11	RS10_RALSO		cytoplasm
Q8XV12	RL3_RALSO		cytoplasm
Q8XV13	RL4_RALSO		cytoplasm
Q8XV15	RL2_RALSO		cytoplasm
Q8XV16	RS19_RALSO		cytoplasm
Q8XV17	RL22_RALSO		cytoplasm
Q8XV18	RS3_RALSO		cytoplasm
Q8XV19	RL16_RALSO		cytoplasm
Q8XV20	RL29_RALSO		cytoplasm
Q8XV21	RS17_RALSO		cytoplasm
Q8XV23	RL24_RALSO		cytoplasm
Q8XV24	RL5_RALSO		cytoplasm
Q8XV25	RS14_RALSO		cytoplasm
Q8XV26	RS8_RALSO		cytoplasm
Q8XV28	RL18_RALSO		cytoplasm
Q8XV29	RS5_RALSO		cytoplasm
Q8XV31	RL15_RALSO		cytoplasm
Q8XV35	RS13_RALSO		cytoplasm
Q8XV36	RS11_RALSO		cytoplasm
Q8XV37	RS4_RALSO		cytoplasm
Q8XV38	RPOA_RALSO		cytoplasm
Q8XV41	DSBD_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XV43	ENGB_RALSO		cytoplasm
Q8XV50	Y2981_RALSO		cytoplasm
Q8XV51	Y2980_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XV54	CYAY_RALSO		inner membrane
Q8XV61	AROK_RALSO	cytoplasm (probable).	cytoplasm
Q8XV62	AROB_RALSO	cytoplasm (probable).	cytoplasm
Q8XV63	DGTL1_RALSO		cytoplasm
Q8XV78	MURA_RALSO	cytoplasm (probable).	inner membrane
Q8XV79	HISX_RALSO		cytoplasm
Q8XV80	HIS81_RALSO		cytoplasm
Q8XV81	HIS7_RALSO	cytoplasm (by similarity).	cytoplasm

Q8XV83	HIS5_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XV84	HIS4_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XV85	HIS6_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XV86	HIS3_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XV87	HIS2_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XV89	TATA_RALSO	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8XV90	TATB_RALSO	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8XVA0	Y2931_RALSO		cytoplasm
Q8XVA1	MSCL_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XVC2	OPGH_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XVC3	OPGG_RALSO	periplasm (by similarity).	periplasm
Q8XVD1	HEM1_RALSO		cytoplasm
Q8XVE4	UNG_RALSO	cytoplasm (by similarity).	extracellular
Q8XVE6	TRPC1_RALSO		cytoplasm
Q8XVE7	TRPD1_RALSO		cytoplasm
Q8XVF3	APAG_RALSO		inner membrane
Q8XVH8	MRAZ_RALSO		inner membrane
Q8XVH9	MRAW_RALSO		cytoplasm
Q8XVI2	MURE_RALSO	cytoplasm (probable).	cytoplasm
Q8XVI4	MRAY_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XVI5	MURD_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XVI7	MURG_RALSO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XVI8	MURC_RALSO	cytoplasm (probable).	cytoplasm
Q8XVI9	DDL_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XVJ7	ARGJ_RALSO	cytoplasm (probable).	cytoplasm
Q8XVK0	Y2830_RALSO		cytoplasm
Q8XVK1	Y2829_RALSO		inner membrane
Q8XVK2	COAE_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XVK9	RL27_RALSO		cytoplasm
Q8XVL1	PROB_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XVL3	NUDH_RALSO		cytoplasm
Q8XVP0	TPX_RALSO		cytoplasm
Q8XVP2	PRMA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XVP5	AROQ1_RALSO		cytoplasm
Q8XVQ4	MTGA_RALSO	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8XVT2	DAPB_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XVT6	PROA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XVU6	PAND_RALSO		cytoplasm
Q8XVV4	TRMU_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XVW4	TGT_RALSO		cytoplasm
Q8XVW5	QUEA_RALSO	cytoplasm (potential).	cytoplasm
Q8XW19	GLNE_RALSO		cytoplasm

Q8XW25	PPNK_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XW26	HRCR_RALSO		cytoplasm
Q8XW28	HUTU_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XW29	HUTH_RALSO	cytoplasm (potential).	cytoplasm
Q8XW30	HUTG_RALSO		cytoplasm
Q8XW31	HUTI_RALSO	cytoplasm (potential).	cytoplasm
Q8XW32	HEMH_RALSO	cytoplasm (by similarity).	extracellular
Q8XW36	GRPE_RALSO	cytoplasm (probable).	cytoplasm
Q8XW40	DNAK_RALSO		cytoplasm
Q8XW41	DNAJ_RALSO	cytoplasm (probable).	cytoplasm
Q8XW45	PANB_RALSO		cytoplasm
Q8XW52	PUR5_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XWB0	MIAA_RALSO		cytoplasm
Q8XWB1	MUTL_RALSO		cytoplasm
Q8XWB8	RS20_RALSO		cytoplasm
Q8XWC0	OTC_RALSO	cytoplasm (probable).	cytoplasm
Q8XWC1	ASSY_RALSO	cytoplasm (probable).	cytoplasm
Q8XWC2	Y2552_RALSO		inner membrane
Q8XWC4	MURB_RALSO	cytoplasm (probable).	inner membrane
Q8XWC5	Y2549_RALSO		inner membrane
Q8XWC8	Y2546_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XWD0	XERD_RALSO	cytoplasm (by similarity).	nucleoid
Q8XWE1	KAD_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XWE4	LPXK_RALSO		cytoplasm
Q8XWE7	EX7L_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XWK9	CLPS_RALSO		cytoplasm
Q8XWL1	DUT_RALSO		extracellular
Q8XWM5	NADA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XWM7	NADB1_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XWM8	RL33_RALSO		cytoplasm
Q8XWM9	RL28_RALSO		cytoplasm
Q8XWN0	RADC_RALSO		cytoplasm
Q8XWN8	ARGD_RALSO	cytoplasm (probable).	cytoplasm
Q8XWQ8	AMPA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XWR1	ILVD_RALSO		cytoplasm
Q8XWR3	LGT_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XWS3	COBT_RALSO		cytoplasm
Q8XWS4	COBS_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XWT0	COBQ_RALSO		cytoplasm
Q8XWT3	PANC_RALSO		cytoplasm
Q8XWV4	DCD_RALSO		extracellular
Q8XWV7	ARLY_RALSO	cytoplasm (probable).	cytoplasm
Q8XWW2	CAPP_RALSO		cytoplasm
Q8XWW3	HEM3_RALSO		cytoplasm
Q8XWX1	IPYR_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XWY0	LOLA_RALSO	periplasm (by similarity).	periplasm
Q8XX54	DADA2_RALSO		cytoplasm
Q8XX80	SYV_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XX83	KHSE_RALSO		cytoplasm
Q8XX93	EX7S_RALSO	cytoplasm (by similarity).	cytoplasm

Q8XX95	DXS_RALSO		cytoplasm
Q8XX96	Y2220_RALSO		inner membrane
Q8XXA6	GLUQ_RALSO		cytoplasm
Q8XXB0	Y2206_RALSO		inner membrane
Q8XXB6	MSBA_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XXC0	Y2195_RALSO		inner membrane
Q8XXC3	HEM6_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XXC4	PUR2_RALSO		cytoplasm
Q8XXC5	Y2190_RALSO		inner membrane
Q8XXC7	UPP_RALSO		cytoplasm
Q8XXD0	MGSA_RALSO		cytoplasm
Q8XXE5	Y2169_RALSO		inner membrane
Q8XXE9	PCP1_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XXF4	HAM1_RALSO		extracellular
Q8XXF5	RNPH_RALSO		cytoplasm
Q8XXF9	KGUA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XXG0	RPOZ_RALSO		cytoplasm
Q8XXJ7	Y2116_RALSO		cytoplasm
Q8XXL5	ADD_RALSO		extracellular
Q8XXN8	ILVC_RALSO		cytoplasm
Q8XXN9	PSD_RALSO		cytoplasm
Q8XXP1	LEU11_RALSO		cytoplasm
Q8XXP4	RS15_RALSO		cytoplasm
Q8XXP9	TPIS_RALSO		cytoplasm
Q8XXS8	URE3_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XXT0	URE2_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XXT1	URE1_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XXT2	UREE_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XXV4	Y2009_RALSO		inner membrane
Q8XXW5	MDH_RALSO		cytoplasm
Q8XXX1	Y1992_RALSO		nucleoid
Q8XXX3	LEU2_RALSO		cytoplasm
Q8XXX4	LEUD_RALSO		cytoplasm
Q8XXX5	LEU3_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XXX8	TRUA_RALSO		cytoplasm
Q8XXX9	TRPF_RALSO		cytoplasm
Q8XXY0	TRPB_RALSO		cytoplasm
Q8XXY2	TRPA_RALSO		cytoplasm
Q8XY07	MURI_RALSO		cytoplasm
Q8XY11	ACSA_RALSO		cytoplasm
Q8XYH4	KTHY_RALSO		cytoplasm
Q8XYL0	Y1748_RALSO		inner membrane
Q8XYL1	MSRB_RALSO		cytoplasm
Q8XYL2	ISPZ_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XYN9	G6PI_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XYP6	CLPX_RALSO		cytoplasm
Q8XYP7	CLPP_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XYP8	TIG_RALSO		cytoplasm
Q8XYP9	Y1709_RALSO		cytoplasm
Q8XYW2	ISPF_RALSO		extracellular
Q8XYW3	ISPD_RALSO		cytoplasm
Q8XYX1	DUSC_RALSO		cytoplasm

Q8XYX8	RLUD_RALSO		cytoplasm
Q8XYY5	KUP2_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XYY7	PYRD_RALSO	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XYY9	LFTR_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XYZ0	ATE_RALSO	cytoplasm (potential).	cytoplasm
Q8XZ19	DPO4_RALSO	cytoplasm (probable).	cytoplasm
Q8XZ23	IHFA_RALSO		cytoplasm
Q8XZ26	RL20_RALSO		cytoplasm
Q8XZ27	RL35_RALSO		cytoplasm
Q8XZ40	AROC_RALSO		cytoplasm
Q8XZ72	PSTB_RALSO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XZ76	GLMM_RALSO		cytoplasm
Q8XZ79	RRMJ_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XZ82	GREA_RALSO		cytoplasm
Q8XZ83	CARB_RALSO		cytoplasm
Q8XZ85	CARA_RALSO		cytoplasm
Q8XZ91	RNH_RALSO	cytoplasm (potential).	extracellular
Q8XZC2	BIOD_RALSO		cytoplasm
Q8XZG4	GUAA_RALSO		cytoplasm
Q8XZG9	Y1426_RALSO		inner membrane
Q8XZH1	SSRP_RALSO	cytoplasm (potential).	cytoplasm
Q8XZH5	Y1420_RALSO		cytoplasm
Q8XZH7	RNH2_RALSO	cytoplasm (potential).	extracellular
Q8XZH8	LPXB_RALSO		cytoplasm
Q8XZH9	LPXA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XZI0	FABZ_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XZI1	LPXD_RALSO		cytoplasm
Q8XZI4	Y1411_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8XZI5	DXR_RALSO		cytoplasm
Q8XZI7	UPPS_RALSO		inner membrane
Q8XZI8	RRF_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XZI9	PYRH_RALSO		cytoplasm
Q8XZJ1	RS2_RALSO		cytoplasm
Q8XZJ3	GLND_RALSO		cytoplasm
Q8XZJ6	DEF2_RALSO		cytoplasm
Q8XZK2	DAPD_RALSO	cytoplasm (by similarity).	cytoplasm
Q8XZK8	GCH1_RALSO		cytoplasm
Q8XZM2	DSBB_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XZM4	ALR_RALSO		cytoplasm
Q8XZP1	RRAA_RALSO		cytoplasm
Q8XZP8	CYSA_RALSO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8XZQ6	SSUD_RALSO		cytoplasm
Q8XZR0	CLPB_RALSO	cytoplasm (probable).	cytoplasm

Q8XZR2	CRCB_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8XZR3	MOAE_RALSO		cytoplasm
Q8XZT5	RL9_RALSO		cytoplasm
Q8XZT6	RS18_RALSO		cytoplasm
Q8XZT8	RS6_RALSO		cytoplasm
Q8XZU1	LEXA_RALSO		cytoplasm
Q8XZV4	TRUB_RALSO		cytoplasm
Q8XZV5	RBFA_RALSO	cytoplasm (potential).	cytoplasm
Q8XZV8	Y1287_RALSO		inner membrane
Q8XZZ5	ARGA_RALSO	cytoplasm (probable).	cytoplasm
Q8Y010	FETP_RALSO		cytoplasm
Q8Y013	RPIA_RALSO		cytoplasm
Q8Y014	TAL_RALSO	cytoplasm (probable).	cytoplasm
Q8Y015	NPD_RALSO	cytoplasm (probable).	cytoplasm
Q8Y016	RLMB_RALSO		cytoplasm
Q8Y019	PURA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y020	HISZ_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y025	HFQ_RALSO		cytoplasm
Q8Y026	ENGA_RALSO		cytoplasm
Q8Y033	NDK_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y035	RUMA_RALSO		cytoplasm
Q8Y040	SURE_RALSO	cytoplasm (potential).	periplasm
Q8Y050	RECR_RALSO		cytoplasm
Q8Y052	Y1192_RALSO		inner membrane
Q8Y062	RL31B_RALSO		cytoplasm
Q8Y074	TILS_RALSO	cytoplasm (probable).	cytoplasm
Q8Y075	ACCA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y081	LPXH_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y093	MUTS_RALSO		cytoplasm
Q8Y099	DAPA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0B4	FTSB_RALSO	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q8Y0B5	ENO_RALSO	cytoplasm (by similarity).	extracellular
Q8Y0B7	KDSA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0B8	PYRG_RALSO		cytoplasm
Q8Y0C6	LOLD_RALSO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Y0D3	Y1111_RALSO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Y0H3	UVRC_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0H7	ACPS_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0H8	PDXJ_RALSO	cytoplasm (probable).	cytoplasm
Q8Y0H9	RECO_RALSO		cytoplasm
Q8Y0I0	ERA_RALSO		cytoplasm
Q8Y0I1	RNC_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0I4	LEPA_RALSO		cytoplasm
Q8Y0J1	ACP1_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0J4	FABH_RALSO	cytoplasm (probable).	cytoplasm
Q8Y0J5	PLSX_RALSO		cytoplasm

Q8Y0J6	RL32_RALSO		cytoplasm
Q8Y0K4	MOAA_RALSO		cytoplasm
Q8Y0K5	MOBA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0L2	PNCB_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0L9	HSCA_RALSO		cytoplasm
Q8Y0M0	HSCB_RALSO		inner membrane
Q8Y0N2	UVRB_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0N9	Y1004_RALSO		cytoplasm
Q8Y0Q3	HTPG_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0U6	TYSY_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0U9	Y944_RALSO		inner membrane
Q8Y0V1	ORN_RALSO	cytoplasm (potential).	cytoplasm
Q8Y0V3	ENGC_RALSO		cytoplasm
Q8Y0V7	RL19_RALSO		cytoplasm
Q8Y0V8	TRMD_RALSO	cytoplasm (potential).	cytoplasm
Q8Y0V9	RIMM_RALSO	cytoplasm (potential).	cytoplasm
Q8Y0W0	RS16_RALSO		cytoplasm
Q8Y0W7	DADA1_RALSO		cytoplasm
Q8Y0X8	HLDD_RALSO		cytoplasm
Q8Y0Y3	IHFB_RALSO		cytoplasm
Q8Y0Y5	KCY_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0Y6	AROA_RALSO	cytoplasm (probable).	cytoplasm
Q8Y0Y8	HIS82_RALSO		cytoplasm
Q8Y0Z0	SERC_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y0Z5	UBIG_RALSO		cytoplasm
Q8Y183	Y807_RALSO		cytoplasm
Q8Y1B3	HEMTL_RALSO		cytoplasm
Q8Y1C3	PDXH1_RALSO		cytoplasm
Q8Y1C6	MSRA_RALSO		cytoplasm
Q8Y1F5	TOLB_RALSO	periplasm (potential).	periplasm
Q8Y1G1	GLYA1_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y1G2	NRDR_RALSO		cytoplasm
Q8Y1H8	RISB_RALSO		cytoplasm
Q8Y1H9	NUSB_RALSO		cytoplasm
Q8Y1I7	TRMB_RALSO		cytoplasm
Q8Y1I9	UPPP_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Y1K9	APAH_RALSO		extracellular
Q8Y1L2	PYRB_RALSO		cytoplasm
Q8Y1L3	PYRR_RALSO		cytoplasm
Q8Y1L4	RUVX_RALSO	cytoplasm (potential).	cytoplasm
Q8Y1L5	Y675_RALSO		inner membrane
Q8Y1P8	CH60_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y1P9	CH10_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y1S1	RDGC_RALSO	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q8Y1W3	PUR7_RALSO		cytoplasm
Q8Y1W6	PGK_RALSO	cytoplasm (potential).	cytoplasm
Q8Y1X7	MOAC_RALSO		cytoplasm
Q8Y1Y3	SUCC_RALSO		cytoplasm
Q8Y1Y5	RECX_RALSO	cytoplasm (potential).	cytoplasm
Q8Y1Y6	RECA_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y208	Y529_RALSO		inner membrane

Q8Y210	LNT_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Y219	KSGA_RALSO		cytoplasm
Q8Y231	FISL_RALSO		cytoplasm
Q8Y232	PUR9_RALSO		cytoplasm
Q8Y233	RUVC_RALSO		cytoplasm
Q8Y235	RUVA_RALSO		cytoplasm
Q8Y236	RUVB_RALSO		cytoplasm
Q8Y239	DTD_RALSO	cytoplasm (probable).	cytoplasm
Q8Y240	SYT_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y245	RS9_RALSO		cytoplasm
Q8Y249	PYRC_RALSO		cytoplasm
Q8Y275	UBIB_RALSO	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8Y278	UBIE_RALSO		cytoplasm
Q8Y288	QUEF_RALSO	cytoplasm (probable).	cytoplasm
Q8Y2B4	SSB_RALSO		cytoplasm
Q8Y2D1	HPRK_RALSO		cytoplasm
Q8Y2D3	Y403_RALSO		cytoplasm
Q8Y2D7	FPG_RALSO		cytoplasm
Q8Y2D9	LOLB_RALSO	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q8Y2E0	ISPE_RALSO		cytoplasm
Q8Y2E1	KPRS_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y2E2	RL25_RALSO		cytoplasm
Q8Y2E3	PTH_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y2E6	COAD_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y2H9	GPDA_RALSO	cytoplasm (probable).	cytoplasm
Q8Y2I0	SECB_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y2I3	GPMA_RALSO		cytoplasm
Q8Y2K1	Y334_RALSO		inner membrane
Q8Y2K5	DCTA1_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Y2K9	Y326_RALSO		inner membrane
Q8Y2L2	LIPB_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y2L3	LIPA_RALSO	cytoplasm (potential).	cytoplasm
Q8Y2Q7	ACEK_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y2R5	Y270_RALSO		inner membrane
Q8Y2V6	RTCA_RALSO	cytoplasm (potential).	cytoplasm
Q8Y2X7	TYPH_RALSO		cytoplasm
Q8Y303	GLMS_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y304	GLMU_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y339	ARGC_RALSO	cytoplasm (probable).	cytoplasm
Q8Y342	PYRE_RALSO		cytoplasm
Q8Y344	DAPF_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y368	THIC_RALSO		cytoplasm
Q8Y372	THIG_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y387	SAHH_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y395	CCA_RALSO		cytoplasm
Q8Y3A6	HTPX_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8Y3A8	FMT_RALSO		cytoplasm

Q8Y3B0	DEF1_RALSO		cytoplasm
Q8Y3C2	GATC_RALSO		cytoplasm
Q8Y3C3	GATA_RALSO		cytoplasm
Q8Y3C6	GATB_RALSO		cytoplasm
Q8Y3C8	XERC1_RALSO	cytoplasm (by similarity).	nucleoid
Q8Y3D7	HSLV_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y3D8	HSLU_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y3E7	ARGB_RALSO	cytoplasm (probable).	cytoplasm
Q8Y3F3	METX_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y3G3	PPCK_RALSO	cytoplasm (by similarity).	cytoplasm
Q8Y3H2	RS21_RALSO		cytoplasm
Q8Y3H5	TRME_RALSO		cytoplasm
Q8Y3H6	OXAA_RALSO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Y3H7	Y003_RALSO		inner membrane
Q8YB19	FLGH_BRUME	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8YB47	HIS1_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YB48	GLUP_BRUME	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8YB50	FUMC_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YB53	CH60_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YB57	SYI_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YBC6	NOSZ_BRUME	periplasm (by similarity).	periplasm
Q8YBH3	MINC_BRUME		inner membrane
Q8YBL2	GUAA_BRUME		cytoplasm
Q8YBL8	PHK_BRUME		cytoplasm
Q8YBP2	NANEK_BRUME		cytoplasm
Q8YBR2	GLPK_BRUME		cytoplasm
Q8YBV5	META_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YBV9	BIOD_BRUME		cytoplasm
Q8YC47	GLMS_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YC48	GLMU_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YC54	LIPB_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YC57	GATA_BRUME		cytoplasm
Q8YC60	RUVX_BRUME	cytoplasm (potential).	cytoplasm
Q8YC64	Y3668_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YC76	DPO4_BRUME	cytoplasm (probable).	cytoplasm
Q8YCA5	ADEC_BRUME		cytoplasm
Q8YCD5	RRMJ_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YCE2	SP39_BRUME	periplasm (probable).	periplasm
Q8YCE3	FDHD_BRUME	cytoplasm (potential).	cytoplasm
Q8YCG3	FBPC_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8YCK1	EX7L_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YCN7	NIKE_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8YCN8	NIKD_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q8YCP2	NIKR_BRUME		cytoplasm
Q8YCQ2	UXAC_BRUME		cytoplasm
Q8YCQ4	UXUA_BRUME		cytoplasm
Q8YCQ8	CRCB4_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YCQ9	CRCB3_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YCV3	TPIS2_BRUME		cytoplasm
Q8YCX4	LEU3_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YCY1	OCD_BRUME		cytoplasm
Q8Y CZ1	Y3386_BRUME		cytoplasm
Q8YD03	ALR_BRUME		cytoplasm
Q8YD04	DADA_BRUME		cytoplasm
Q8YD09	HUTIH_BRUME	cytoplasm (potential).	cytoplasm
Q8YD12	HUTU_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YD73	Y3304_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YDA2	RNPA_BRUME		extracellular
Q8YDA3	OXAA_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YDA5	ARGB_BRUME	cytoplasm (probable).	cytoplasm
Q8YDA8	DAPD_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YDB2	TRUA_BRUME		cytoplasm
Q8YDB8	LEPA_BRUME		cytoplasm
Q8YDC6	GLK_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YDC8	DAPB_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YDC9	GPMA_BRUME		cytoplasm
Q8YDE4	UBIE_BRUME		cytoplasm
Q8YDE5	UPP_BRUME		cytoplasm
Q8YDE6	SLYX_BRUME		inner membrane
Q8YDE7	MSRA_BRUME		cytoplasm
Q8YDL2	FLBT_BRUME		cytoplasm
Q8YDL6	FLGE_BRUME		flagellar
Q8YDM4	FLIF_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YDP7	Y3128_BRUME		cytoplasm
Q8YE11	Y007_BRUME		inner membrane
Q8YE15	MODC_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8YE19	Y2059_BRUME	cytoplasm (potential).	inner membrane
Q8YE20	AROE_BRUME		cytoplasm
Q8YE23	SECB_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YE30	HSLU_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YE31	HSLV_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YE35	HIS5_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YE36	HIS4_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YE37	HIS6_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YE41	PPCK_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YE49	SAHH_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YE59	TRPA_BRUME		cytoplasm

Q8YE60	TRPB_BRUME		cytoplasm
Q8YE76	DNAK_BRUME		cytoplasm
Q8YE77	DNAJ_BRUME	cytoplasm (probable).	cytoplasm
Q8YE79	PYRF_BRUME		cytoplasm
Q8YE82	GSHB_BRUME		cytoplasm
Q8YEA6	LNT_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YEA9	TRMB_BRUME		cytoplasm
Q8YEB4	RBFA_BRUME	cytoplasm (potential).	cytoplasm
Q8YEB5	TRUB_BRUME		cytoplasm
Q8YEB6	RS15_BRUME		cytoplasm
Q8YED2	FPG_BRUME		cytoplasm
Q8YED5	DNAA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YED7	RECF_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YEG1	AROA_BRUME	cytoplasm (probable).	cytoplasm
Q8YEG2	KCY_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YEG4	Y1914_BRUME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8YEG8	RECR_BRUME		cytoplasm
Q8YEK2	DNAE2_BRUME	cytoplasm (probable).	cytoplasm
Q8YEK8	ASSY_BRUME	cytoplasm (probable).	cytoplasm
Q8YEM5	CCMA_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8YEN0	ILVD_BRUME		cytoplasm
Q8YEP2	PNCB_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YES3	GLND_BRUME		cytoplasm
Q8YES6	MUTS_BRUME		cytoplasm
Q8YET3	IHFB_BRUME		cytoplasm
Q8YEV0	GRPE_BRUME	cytoplasm (probable).	cytoplasm
Q8YEV2	RNPH_BRUME		cytoplasm
Q8YEZ2	THIG_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YF55	Y1672_BRUME	cytoplasm (potential).	inner membrane
Q8YF59	HISX_BRUME		cytoplasm
Q8YF71	URE31_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YF72	URE11_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YF86	G6PI_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YFA2	OTC_BRUME	cytoplasm (probable).	cytoplasm
Q8YFA4	APAG_BRUME		inner membrane
Q8YFB7	MSCL_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YFC9	PANC_BRUME		cytoplasm
Q8YFD7	NORM_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YFH2	ENGA_BRUME		cytoplasm
Q8YFI3	ISPE_BRUME		cytoplasm
Q8YFK1	PUR2_BRUME		cytoplasm
Q8YFK3	PDXH_BRUME		cytoplasm
Q8YFM2	DXS_BRUME		cytoplasm
Q8YFM4	Y1496_BRUME		cytoplasm
Q8YFM9	PSD_BRUME		cytoplasm
Q8YFN7	RL9_BRUME		cytoplasm

Q8YFP0	RS6_BRUME		cytoplasm
Q8YFQ7	COXZ_BRUME		inner membrane
Q8YFR1	ISPH_BRUME		cytoplasm
Q8YFR2	KHSE_BRUME		cytoplasm
Q8YFR6	Y1454_BRUME		inner membrane
Q8YFU0	ALLA_BRUME		extracellular
Q8YFU1	Y1429_BRUME		cytoplasm
Q8YFX5	XYLA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YFY0	BETB_BRUME		cytoplasm
Q8YFY2	BETA_BRUME		cytoplasm
Q8YFY3	BETI_BRUME		cytoplasm
Q8YG29	CCME_BRUME	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q8YG32	DEGP_BRUME	periplasm (potential).	periplasm
Q8YG35	GLNE_BRUME		cytoplasm
Q8YG60	DAPA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YG64	RPOZ_BRUME		cytoplasm
Q8YG75	ERA_BRUME		cytoplasm
Q8YG76	RECO_BRUME		cytoplasm
Q8YG94	KSGA_BRUME		cytoplasm
Q8YG99	AMPA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGA4	NDK_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGA9	UVRC_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGB9	PUR5_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGG7	GLYA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGH2	RISB1_BRUME		cytoplasm
Q8YGH3	NUSB_BRUME		cytoplasm
Q8YGH4	HPPA_BRUME	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8YGH8	PLSX_BRUME		cytoplasm
Q8YGH9	FABH_BRUME	cytoplasm (probable).	cytoplasm
Q8YGI8	ARGC_BRUME	cytoplasm (probable).	cytoplasm
Q8YGJ0	RS9_BRUME		cytoplasm
Q8YGM0	LOLD_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8YGM5	RS4_BRUME		cytoplasm
Q8YGN1	PURL_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGN4	PURQ_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGN6	PUR7_BRUME		cytoplasm
Q8YQG8	COBS_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YQG9	COBT_BRUME		cytoplasm
Q8YGT8	TIG_BRUME		cytoplasm
Q8YGU6	AROQ_BRUME		cytoplasm
Q8YGV8	SYV_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGW9	PPNK_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGX8	SYV_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGY5	MOBA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YGY6	MOAA_BRUME		cytoplasm
Q8YH15	KTHY_BRUME		cytoplasm
Q8YH30	RPIA_BRUME		cytoplasm

Q8YH52	Y952_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YH64	XGPT_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YH68	WRBA_BRUME		cytoplasm
Q8YH92	Y912_BRUME		inner membrane
Q8YH94	GCH1_BRUME		cytoplasm
Q8YH95	HIS3_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHB2	TGT_BRUME		cytoplasm
Q8YHC2	SSB_BRUME		cytoplasm
Q8YHC4	UVRA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHC6	LON_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHC7	CLPX_BRUME		cytoplasm
Q8YHC8	CLPP_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHD8	ISPDF_BRUME		cytoplasm
Q8YHF0	ENO_BRUME	cytoplasm (by similarity).	extracellular
Q8YHF1	KDSA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHF2	PYRG_BRUME		cytoplasm
Q8YHF5	TPIS1_BRUME		cytoplasm
Q8YHF7	TRPD_BRUME		cytoplasm
Q8YHF9	MOAC_BRUME		cytoplasm
Q8YHG1	LEXA_BRUME		cytoplasm
Q8YHG9	FABZ_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHH1	Y829_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8YHH2	CDSA_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YHH3	UPPS_BRUME		inner membrane
Q8YHH4	PYRH_BRUME		cytoplasm
Q8YHH6	RS2_BRUME		cytoplasm
Q8YHI3	CLPS_BRUME		cytoplasm
Q8YHJ4	QUEF_BRUME	cytoplasm (probable).	cytoplasm
Q8YHL6	RPOA_BRUME		cytoplasm
Q8YHL9	KAD_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHM1	RL15_BRUME		cytoplasm
Q8YHM4	RL18_BRUME		cytoplasm
Q8YHM6	RS8_BRUME		cytoplasm
Q8YHM8	RL5_BRUME		cytoplasm
Q8YHM9	RL24_BRUME		cytoplasm
Q8YHN1	RS17_BRUME		cytoplasm
Q8YHN3	RL16_BRUME		cytoplasm
Q8YHN4	RS3_BRUME		cytoplasm
Q8YHN5	RL22_BRUME		cytoplasm
Q8YHN7	RL2_BRUME		cytoplasm
Q8YHN9	RL4_BRUME		cytoplasm
Q8YHP0	RL3_BRUME		cytoplasm
Q8YHP4	RS7_BRUME		cytoplasm
Q8YHP7	RPOC_BRUME		cytoplasm
Q8YHP8	RPOB_BRUME		cytoplasm
Q8YHP9	RL10_BRUME		cytoplasm
Q8YHQ0	RL1_BRUME		cytoplasm
Q8YHQ1	RL11_BRUME		cytoplasm
Q8YHR9	DDLA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHS6	SFSA_BRUME		cytoplasm

Q8YHT9	COBD_BRUME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8YHU1	COBB_BRUME		cytoplasm
Q8YHU3	CBID_BRUME		cytoplasm
Q8YHV6	COBQ_BRUME		cytoplasm
Q8YHZ6	URE32_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHZ7	URE22_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YHZ8	URE12_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YI11	CRCB2_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YI12	CRCB1_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YI21	ILVC_BRUME		cytoplasm
Q8YI23	KUP_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YI24	PDXJ_BRUME		cytoplasm
Q8YI29	MIAA_BRUME		cytoplasm
Q8YI47	Y598_BRUME		inner membrane
Q8YI53	PRMA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YI59	LPXC_BRUME		cytoplasm
Q8YI63	DDLB_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YI64	MURB_BRUME	cytoplasm (probable).	inner membrane
Q8YI66	MURG_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8YI68	MURD_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YI71	MURE_BRUME	cytoplasm (probable).	cytoplasm
Q8YI91	TAM_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YIB0	Y534_BRUME		inner membrane
Q8YIB8	CARA_BRUME		cytoplasm
Q8YIC2	CARB_BRUME		cytoplasm
Q8YIF6	LGT_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YIG1	KPRS_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YIG3	RL25_BRUME		cytoplasm
Q8YIJ3	LEU1_BRUME		cytoplasm
Q8YIL6	TRMU_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YIR3	MASZ_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YIR8	FUR_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YIU8	GLMM_BRUME		cytoplasm
Q8YIV0	TILS_BRUME	cytoplasm (probable).	cytoplasm
Q8YIV5	RUVB_BRUME		cytoplasm
Q8YIV7	RUVC_BRUME		cytoplasm
Q8YIY0	PGK_BRUME	cytoplasm (potential).	cytoplasm
Q8YIZ8	Y289_BRUME		inner membrane
Q8YJ15	MTGA_BRUME	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8YJ17	ISPG_BRUME		cytoplasm
Q8YJ36	ATPG_BRUME		inner membrane

Q8YJ37	ATPA_BRUME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8YJ42	TAL_BRUME	cytoplasm (probable).	cytoplasm
Q8YJ48	ACSA_BRUME		cytoplasm
Q8YJ50	HTPX_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8YJ71	NUDH_BRUME		cytoplasm
Q8YJ77	NADD_BRUME		cytoplasm
Q8YJ78	PROA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YJ84	RL27_BRUME		cytoplasm
Q8YJ91	CLPB_BRUME	cytoplasm (probable).	cytoplasm
Q8YJ98	UBIG_BRUME		cytoplasm
Q8YJB0	HEM3_BRUME		cytoplasm
Q8YJC9	LEU2_BRUME		cytoplasm
Q8YJD7	TRMD_BRUME	cytoplasm (potential).	cytoplasm
Q8YJD9	XERC_BRUME	cytoplasm (by similarity).	nucleoid
Q8YJE4	ODO1_BRUME		cytoplasm
Q8YJE7	MDH_BRUME		cytoplasm
Q8YJF0	DAPF_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YJF3	ISPZ_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8YJF9	ARGJ_BRUME	cytoplasm (probable).	cytoplasm
Q8YJJ5	Y088_BRUME		inner membrane
Q8YJJ7	ARLY_BRUME	cytoplasm (probable).	cytoplasm
Q8YJK3	HIS8_BRUME		cytoplasm
Q8YJP2	XERD_BRUME	cytoplasm (by similarity).	nucleoid
Q8YJP3	ACCA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YJS4	GIDB_BRUME		cytoplasm
Q8YJS5	GIDA_BRUME	cytoplasm (by similarity).	cytoplasm
Q8YJT1	DCUP_BRUME	cytoplasm (probable).	cytoplasm
Q8Z0T4	GPMB_SALTI		cytoplasm
Q8Z0T9	SMP_SALTI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z0U0	LPLA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z0U2	DEOD_SALTI		cytoplasm
Q8Z0U3	DEOC_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z0W2	OPGB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z0Y7	YEEN_SALTI		inner membrane
Q8Z116	AMPA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z118	SYV_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z123	OTC_SALTI	cytoplasm (probable).	cytoplasm
Q8Z125	ARCA_SALTI	cytoplasm (by similarity).	extracellular
Q8Z127	OTCC_SALTI	cytoplasm (probable).	cytoplasm
Q8Z130	PYRI_SALTI		cytoplasm
Q8Z138	NRDG_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z150	MSRA_SALTI		cytoplasm
Q8Z153	CYSQ_SALTI	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	cytoplasm

Q8Z157	RCMNS_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z163	RL9_SALTI		cytoplasm
Q8Z164	PRIB_SALTI		cytoplasm
Q8Z167	ULAF_SALTI		cytoplasm
Q8Z168	ULAE_SALTI		cytoplasm
Q8Z169	ULAD_SALTI		cytoplasm
Q8Z170	ULAC_SALTI	cytoplasm (probable).	cytoplasm
Q8Z171	ULAB_SALTI	cytoplasm (potential).	cytoplasm
Q8Z172	ULAA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z182	RLMB_SALTI		cytoplasm
Q8Z186	MIAA_SALTI		cytoplasm
Q8Z187	MUTL_SALTI		cytoplasm
Q8Z193	ENGC_SALTI		cytoplasm
Q8Z194	PSD_SALTI		cytoplasm
Q8Z197	SYK3_SALTI		cytoplasm
Q8Z1A8	DSBD_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z1Q9	NRFA_SALTI	periplasm (by similarity).	periplasm
Q8Z1R0	ACSA_SALTI		cytoplasm
Q8Z1R2	ACTP_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z1T1	DUSA_SALTI		cytoplasm
Q8Z1T6	PLSB_SALTI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z1T9	LAMB_SALTI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8Z1U0	MALK_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z1U2	MALF_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z1U3	MALG_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z1U7	G6PI_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z1V3	RLUF_SALTI		cytoplasm
Q8Z1V8	ACEK_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z1W1	META_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z1W5	AROE_SALTI		cytoplasm
Q8Z1W9	DEF_SALTI		cytoplasm
Q8Z1X0	FMT_SALTI		cytoplasm
Q8Z1X1	RSMB_SALTI		cytoplasm
Q8Z1X4	RS4_SALTI		cytoplasm
Q8Z1X5	RS11_SALTI		cytoplasm
Q8Z1X6	RS13_SALTI		cytoplasm
Q8Z1X8	RL24_SALTI		cytoplasm
Q8Z1Y1	TUSB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z1Y2	TUSC_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z1Y3	TUSD_SALTI	cytoplasm (by similarity).	cytoplasm

Q8Z1Y5	SLYX_SALTI		inner membrane
Q8Z1Y7	KEFB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z1Z3	ARGD_SALTI	cytoplasm (probable).	cytoplasm
Q8Z1Z8	TSGA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z202	GPH_SALTI		cytoplasm
Q8Z205	AROB_SALTI	cytoplasm (probable).	cytoplasm
Q8Z214	HSLO_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z216	PPCK_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z217	GREB_SALTI		cytoplasm
Q8Z221	BIOH_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z222	GNTX_SALTI		cytoplasm
Q8Z223	GNTY_SALTI		inner membrane
Q8Z227	MALT_SALTI		cytoplasm
Q8Z232	GLGA_SALTI		cytoplasm
Q8Z233	GLGC_SALTI		cytoplasm
Q8Z234	GLGX_SALTI		extracellular
Q8Z235	GLGB_SALTI		extracellular
Q8Z257	YHHS_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z259	ACPT_SALTI		cytoplasm
Q8Z268	USPA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z270	YHIQ_SALTI		inner membrane
Q8Z277	TREF_SALTI	cytoplasm (by similarity).	extracellular
Q8Z286	YHJJ_SALTI	periplasm (potential).	extracellular
Q8Z287	DCTA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z290	BCSB_SALTI	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q8Z291	BCSA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z292	YHJQ_SALTI		inner membrane
Q8Z2C5	DLGD_SALTI	cytoplasm (potential).	cytoplasm
Q8Z2D8	SELA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2E0	MTLD_SALTI		cytoplasm
Q8Z2E3	LLDP_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z2E5	LLDD_SALTI		cytoplasm
Q8Z2F0	GPMI_SALTI		cytoplasm
Q8Z2F4	TDH_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2H1	COAD_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2H2	FPG_SALTI		cytoplasm
Q8Z2H4	SLMA_SALTI	nucleoid-associated (by similarity).	cytoplasm
Q8Z2H5	PYRE_SALTI		cytoplasm
Q8Z2H9	KGUA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2L4	SDHD_SALTI		cytoplasm

Q8Z2L7	YIDE_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z2L8	IBPB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2M3	TORD_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2M4	TORA_SALTI	periplasm (by similarity).	periplasm
Q8Z2N4	RECF_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2N6	DNAA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2N7	OXAA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z2N8	TRME_SALTI		cytoplasm
Q8Z2N9	MDTL_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z2P0	YIDZ_SALTI		cytoplasm
Q8Z2Q2	GLMS_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2Q3	GLMU_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2Q5	ATPG_SALTI		inner membrane
Q8Z2Q7	GIDA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2Q9	ASNA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2R0	VIAA_SALTI		cytoplasm
Q8Z2R1	RAVA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2R2	KUP_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z2R9	MOBA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2S0	RDOA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2T0	OMPL_SALTI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8Z2T5	YIHT_SALTI		cytoplasm
Q8Z2T8	RBN_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z2U4	FDHE_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2U7	FDHD_SALTI	cytoplasm (potential).	cytoplasm
Q8Z2V3	RHAA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2V4	RHAB_SALTI		cytoplasm
Q8Z2V5	RHAR_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2V6	RHAT_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z2V9	SODM_SALTI		periplasm
Q8Z2W4	FIEF_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z2W6	CDH_SALTI	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8Z2Y2	TPIS_SALTI		cytoplasm
Q8Z2Y6	GLPK_SALTI		cytoplasm
Q8Z2Z0	HSLU_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2Z4	METJ_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z2Z9	GLYA2_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z300	ALR3_SALTI		cytoplasm

Q8Z303	CATA_SALTI		periplasm
Q8Z305	FSA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z307	CAPP_SALTI		cytoplasm
Q8Z308	ARGE_SALTI	cytoplasm (probable).	extracellular
Q8Z309	ARGC_SALTI	cytoplasm (probable).	cytoplasm
Q8Z311	ARLY_SALTI	cytoplasm (probable).	cytoplasm
Q8Z313	TRMA_SALTI		cytoplasm
Q8Z314	BTUB_SALTI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8Z315	MURI_SALTI		cytoplasm
Q8Z316	MURB_SALTI	cytoplasm (probable).	inner membrane
Q8Z318	COAA_SALTI	cytoplasm (probable).	cytoplasm
Q8Z320	RPOB_SALTI		cytoplasm
Q8Z322	THIG_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z325	THIE_SALTI		cytoplasm
Q8Z326	THIC_SALTI		cytoplasm
Q8Z328	NUDC_SALTI		cytoplasm
Q8Z329	DCUP_SALTI	cytoplasm (probable).	cytoplasm
Q8Z331	ZRAP_SALTI	periplasm (by similarity).	periplasm
Q8Z332	ZRAS_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z333	ZRAR_SALTI	cytoplasm (probable).	cytoplasm
Q8Z334	PUR2_SALTI		cytoplasm
Q8Z335	PUR9_SALTI		cytoplasm
Q8Z377	ILVD_SALTI		cytoplasm
Q8Z381	ILVC_SALTI		cytoplasm
Q8Z386	WECA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z388	WECB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z389	WECC_SALTI		cytoplasm
Q8Z395	WECF_SALTI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8Z396	WZYE_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z397	WECG_SALTI		cytoplasm
Q8Z3A4	CYAY_SALTI		inner membrane
Q8Z3A8	XERC_SALTI	cytoplasm (by similarity).	nucleoid
Q8Z3B1	RARD_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z3B3	RHTC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z3B4	RHTB_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z3B6	METE_SALTI		cytoplasm
Q8Z3B8	DLHH_SALTI		extracellular
Q8Z3C0	TATB_SALTI	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane

Q8Z3C6	FADB_SALTI		cytoplasm
Q8Z3D4	YEDZ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z3D8	AAEB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z3E0	MDH_SALTI		cytoplasm
Q8Z3E4	OADG2_SALTI	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q8Z3F0	NANA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z3F1	NANK_SALTI		cytoplasm
Q8Z3G0	MTGA_SALTI	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8Z3G1	YHBJ_SALTI		cytoplasm
Q8Z3G5	KDSC_SALTI		cytoplasm
Q8Z3G6	KDSD_SALTI		cytoplasm
Q8Z3H3	RRMJ_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z3H5	ASSY_SALTI	cytoplasm (probable).	cytoplasm
Q8Z3H8	RBFA_SALTI	cytoplasm (potential).	cytoplasm
Q8Z3H9	TRUB_SALTI		cytoplasm
Q8Z3I7	DIAA_SALTI		cytoplasm
Q8Z3I8	YRAN_SALTI		inner membrane
Q8Z3K5	TDCD_SALTI		cytoplasm
Q8Z3L5	ALX_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z3M9	CCA_SALTI		cytoplasm
Q8Z3N1	GLNE_SALTI		cytoplasm
Q8Z3P2	QSEC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z3Q8	YQHA_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z3R7	UXAC_SALTI		cytoplasm
Q8Z3T9	MLTC_SALTI	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q8Z3U1	TRMB_SALTI		cytoplasm
Q8Z3U6	HAM1_SALTI		extracellular
Q8Z3U7	YGGU_SALTI		inner membrane
Q8Z3V1	RUVX_SALTI	cytoplasm (potential).	cytoplasm
Q8Z3V2	YQGE_SALTI		inner membrane
Q8Z3W2	ARGO_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z3W5	Y3215_SALTI		inner membrane
Q8Z3X0	GCSP_SALTI		cytoplasm
Q8Z3X3	YQFB_SALTI		inner membrane
Q8Z3X9	IDI_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z3Z9	KDUI_SALTI		cytoplasm
Q8Z406	AAS_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q8Z410	MUTH_SALTI	cytoplasm (by similarity).	extracellular
Q8Z412	TYSY_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z418	PTRA_SALTI	periplasm (by similarity).	extracellular
Q8Z421	ARGA_SALTI	cytoplasm (probable).	cytoplasm
Q8Z428	FUCK_SALTI		cytoplasm
Q8Z429	FUCI_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z437	QUEF_SALTI	cytoplasm (probable).	cytoplasm
Q8Z439	TRUC_SALTI		cytoplasm
Q8Z446	RUMA_SALTI		cytoplasm
Q8Z458	CYSJ_SALTI		cytoplasm
Q8Z459	CYSI_SALTI		cytoplasm
Q8Z460	CYSH_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z470	CYSN_SALTI		cytoplasm
Q8Z471	ISPD_SALTI		cytoplasm
Q8Z472	ISPF_SALTI		extracellular
Q8Z473	TRUD_SALTI		cytoplasm
Q8Z474	PIMT_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z495	IACP_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z497	SICP_SALTI	cytoplasm (probable).	cytoplasm
Q8Z4A6	SIRC_SALTI		cytoplasm
Q8Z4C4	NORW_SALTI	cytoplasm (probable).	cytoplasm
Q8Z4C5	NORV_SALTI	cytoplasm (probable).	cytoplasm
Q8Z4C6	NORR_SALTI		cytoplasm
Q8Z4D4	RECX_SALTI	cytoplasm (potential).	cytoplasm
Q8Z4D6	GSH1_SALTI		cytoplasm
Q8Z4D7	LUXS_SALTI		extracellular
Q8Z4E6	NRDI_SALTI		inner membrane
Q8Z4G0	CSID_SALTI		cytoplasm
Q8Z4I5	TRMD_SALTI	cytoplasm (potential).	cytoplasm
Q8Z4K0	NADB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z4K4	RNC_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z4K5	ERA_SALTI		cytoplasm
Q8Z4K6	PDXJ_SALTI	cytoplasm (probable).	cytoplasm
Q8Z4L2	YFEU_SALTI		cytoplasm
Q8Z4M3	HMP_SALTI		cytoplasm
Q8Z4M8	YHFR_SALTI		inner membrane
Q8Z4N0	ISCS_SALTI		cytoplasm
Q8Z4N1	HSCB_SALTI		inner membrane
Q8Z4N2	HSCA_SALTI		cytoplasm
Q8Z4N3	PEPB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z4P6	ENGA_SALTI		cytoplasm
Q8Z4Q1	EX7L_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z4Q3	GUAA_SALTI		cytoplasm
Q8Z4R0	PPK_SALTI		cytoplasm
Q8Z4R2	PUR5_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z4R8	DAPA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z4S7	YFEW_SALTI		extracellular
Q8Z4T0	TALA_SALTI	cytoplasm (probable).	cytoplasm
Q8Z4T3	EUTS_SALTI		cytoplasm
Q8Z4U3	EUTC_SALTI		cytoplasm
Q8Z4U8	HEM6_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z4V6	CYSA_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z4W1	PDXK_SALTI		cytoplasm

Q8Z4W3	CYSZ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z4X5	MNTH_SALTI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8Z4X8	YFEO_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z4Y8	FADL_SALTI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z4Y9	FADI_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z4Z0	FADJ_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z4Z2	MEPA_SALTI	periplasm (by similarity).	extracellular
Q8Z4Z3	YFCK_SALTI		cytoplasm
Q8Z4Z9	YFCJ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z500	TRUA_SALTI		cytoplasm
Q8Z521	YFBV_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z524	YFBR_SALTI		cytoplasm
Q8Z537	YFBJ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z538	ARNT_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z540	ARNA_SALTI		cytoplasm
Q8Z541	ARNC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z542	ARNB_SALTI		cytoplasm
Q8Z560	UBIG_SALTI		cytoplasm
Q8Z568	ECOT_SALTI	periplasm (by similarity).	periplasm
Q8Z570	NAPA_SALTI	periplasm (by similarity).	periplasm
Q8Z579	YEJM_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z591	PTFAH_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z593	END4_SALTI		extracellular
Q8Z5A6	YEIT_SALTI		inner membrane
Q8Z5A8	CDD_SALTI		cytoplasm
Q8Z5B2	DUSC_SALTI		cytoplasm
Q8Z5C1	YEHT_SALTI		cytoplasm
Q8Z5C8	THIM_SALTI		cytoplasm
Q8Z5F6	MDTC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z5F7	MDTB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane

Q8Z5F8	MDTA_SALTI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8Z5G4	WZA_SALTI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8Z5G5	WZB_SALTI		extracellular
Q8Z5G6	WZC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z5H2	NUDD_SALTI		cytoplasm
Q8Z5J6	HIS2_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z5J7	HIS4_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z5J8	HIS7_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z5J9	HIS8_SALTI		cytoplasm
Q8Z5K0	HISX_SALTI		cytoplasm
Q8Z5K1	HIS1_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z5M7	CBIB_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z5M8	CBID_SALTI		cytoplasm
Q8Z5M9	CBIT_SALTI		cytoplasm
Q8Z5N3	CBIN_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z5N5	CBIO_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z5N6	COBQ_SALTI		cytoplasm
Q8Z5N8	COBS_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z5N9	COBT_SALTI		cytoplasm
Q8Z5R0	YEDQ_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z5R1	DSRB_SALTI		inner membrane
Q8Z5S1	FLIE_SALTI		flagellar
Q8Z5S9	DCYD_SALTI		cytoplasm
Q8Z5T3	UVRC_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z5U4	USPC_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z5V9	CUTC_SALTI	cytoplasm (probable).	periplasm
Q8Z5X5	YEBF_SALTI	secreted protein (by similarity).	inner membrane
Q8Z670	YEBN_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z671	YOBD_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z682	YCGN_SALTI		inner membrane
Q8Z685	FADR_SALTI	cytoplasm (potential).	cytoplasm
Q8Z687	DADA_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z688	ALR2_SALTI		cytoplasm

Q8Z699	ISPE_SALTI		cytoplasm
Q8Z6A0	LOLB_SALTI	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q8Z6E7	YEAH_SALTI		inner membrane
Q8Z6F4	SELD_SALTI		cytoplasm
Q8Z6F5	TOP3_SALTI		cytoplasm
Q8Z6F6	DHE4_SALTI		cytoplasm
Q8Z6F9	ARGM_SALTI		cytoplasm
Q8Z6G3	ASTE_SALTI		periplasm
Q8Z6G5	CHO_SALTI		cytoplasm
Q8Z6G6	NADE_SALTI		cytoplasm
Q8Z6H0	CHBG_SALTI		inner membrane
Q8Z6I5	BTUC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z6I8	YDIU_SALTI		inner membrane
Q8Z6I9	AROH_SALTI		cytoplasm
Q8Z6K0	SUFE_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z6K2	KPYK1_SALTI		cytoplasm
Q8Z6L0	SPIC_SALTI		extracellular
Q8Z6N7	MDTK_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z6P7	YDHK_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z6Q2	PDXH_SALTI		cytoplasm
Q8Z6Q5	TPPB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z6Q7	RNFG_SALTI	bacterial cell inner membrane (potential).	inner membrane
Q8Z6Q8	RNFD_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z6R2	ADD_SALTI		extracellular
Q8Z6R6	FUMC_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z6R7	TUS_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z6X9	BIOD2_SALTI		cytoplasm
Q8Z6Y0	CLCB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z6Y5	DMSD_SALTI		cytoplasm
Q8Z6Z0	YNFC_SALTI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8Z6Z2	YNFA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z706	YNEE_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z728	MAO1_SALTI		cytoplasm
Q8Z740	YNCE_SALTI		cytoplasm
Q8Z765	OPGD_SALTI	periplasm (by similarity).	periplasm

Q8Z784	ZNTB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z787	FNR_SALTI	cytoplasm (probable).	cytoplasm
Q8Z788	USPE_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z7A8	TPX_SALTI		cytoplasm
Q8Z7A9	YCJF_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z7C9	RNB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z7D5	RLUB_SALTI		cytoplasm
Q8Z7D9	TRPC_SALTI		cytoplasm
Q8Z7E0	TRPA_SALTI		cytoplasm
Q8Z7E3	YCIC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z7E5	YCIU_SALTI		inner membrane
Q8Z7G6	RLUE_SALTI		cytoplasm
Q8Z7G9	TRMU_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z7H0	YCFC_SALTI		inner membrane
Q8Z7H2	PHOP_SALTI	cytoplasm (probable).	cytoplasm
Q8Z7H3	PHOQ_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z7H6	PEPT_SALTI	cytoplasm (by similarity).	extracellular
Q8Z7I6	NAGZ_SALTI	cytoplasm (by similarity).	extracellular
Q8Z7I7	THIK_SALTI		cytoplasm
Q8Z7J6	FABH_SALTI	cytoplasm (probable).	cytoplasm
Q8Z7J7	RLUC_SALTI		cytoplasm
Q8Z7J8	YCEQ_SALTI		outer membrane
Q8Z7K2	FLGI_SALTI	periplasm (by similarity).	periplasm
Q8Z7L0	MDTH_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z7L2	PYRC_SALTI		cytoplasm
Q8Z7L5	YCEA_SALTI		inner membrane
Q8Z7L7	MDTG_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z7L8	OPGH_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z7L9	OPGC_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z7M3	CSGB_SALTI		extracellular
Q8Z7N9	WRBA_SALTI		cytoplasm
Q8Z7Q5	HPAC_SALTI		cytoplasm
Q8Z7Q6	YEDX_SALTI	periplasm (potential).	periplasm
Q8Z7R2	SIGE_SALTI	cytoplasm (probable).	cytoplasm
Q8Z7S1	YCBG_SALTI		inner membrane
Q8Z7S9	PYRD_SALTI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z7Y9	PNCB_SALTI	cytoplasm (by similarity).	cytoplasm

Q8Z7Z7	MUKF_SALTI	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q8Z800	KDSB_SALTI	cytoplasm (potential).	cytoplasm
Q8Z801	LPXK_SALTI		cytoplasm
Q8Z803	KCY_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z810	YCAD_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z813	LOLA_SALTI	periplasm (by similarity).	periplasm
Q8Z829	HCP_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z841	RUMB_SALTI		cytoplasm
Q8Z851	YBJL_SALTI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z879	YBHG_SALTI	periplasm (potential).	inner membrane
Q8Z883	YBHO_SALTI	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z887	MOAC_SALTI		cytoplasm
Q8Z889	UVRB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z890	BIOD_SALTI		cytoplasm
Q8Z896	HUTH_SALTI	cytoplasm (potential).	cytoplasm
Q8Z897	HUTU_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z899	HUTG_SALTI		cytoplasm
Q8Z8A0	HUTI_SALTI	cytoplasm (potential).	cytoplasm
Q8Z8A2	6PGL_SALTI		cytoplasm
Q8Z8A4	MODC_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z8B0	GAL1_SALTI	cytoplasm (potential).	cytoplasm
Q8Z8B2	GPMA_SALTI		cytoplasm
Q8Z8B6	ZITB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z8B8	NADA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z8C0	TOLB_SALTI	periplasm (potential).	periplasm
Q8Z8D2	END8_SALTI		cytoplasm
Q8Z8D9	YBGL_SALTI		inner membrane
Q8Z8E4	ATKA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8Z8E5	ATKB_SALTI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8Z8E6	ATKC_SALTI	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8Z8G0	NAGB_SALTI		extracellular
Q8Z8G6	YBEY_SALTI		inner membrane
Q8Z8G7	LNT_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z8H0	RIHA_SALTI		extracellular

Q8Z8H3	KDGT2_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z8H7	NADD_SALTI		cytoplasm
Q8Z8H8	COBD_SALTI		cytoplasm
Q8Z8I0	RLPA_SALTI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8Z8I2	LIPB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z8I3	LIPA_SALTI	cytoplasm (potential).	cytoplasm
Q8Z8L4	ENTS_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z8L9	ENTD_SALTI		periplasm
Q8Z8M0	CAAL_SALTI		cytoplasm
Q8Z8P7	LPXH_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z8R4	SELU_SALTI		cytoplasm
Q8Z8S3	CUER_SALTI	cytoplasm (probable).	cytoplasm
Q8Z8S4	ATCU_SALTI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8Z8T2	HEMH_SALTI	cytoplasm (by similarity).	extracellular
Q8Z8V1	CLPX_SALTI		cytoplasm
Q8Z8W2	YAJQ_SALTI		inner membrane
Q8Z8W3	PANE_SALTI	cytoplasm (potential).	cytoplasm
Q8Z8X1	THII_SALTI	cytoplasm (potential).	cytoplasm
Q8Z8X3	DXS_SALTI		cytoplasm
Q8Z8X6	NUSB_SALTI		cytoplasm
Q8Z8Y0	TGT_SALTI		cytoplasm
Q8Z8Y1	QUEA_SALTI	cytoplasm (potential).	cytoplasm
Q8Z8Y9	YAIE_SALTI		inner membrane
Q8Z903	PRPD_SALTI		cytoplasm
Q8Z932	PROA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z933	Yafa_SALTI		cytoplasm
Q8Z937	FADE_SALTI		cytoplasm
Q8Z985	YAFD_SALTI	cytoplasm (potential).	inner membrane
Q8Z988	DKGB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z989	GMHB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z990	METN_SALTI	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8Z991	METI_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z992	METQ_SALTI	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8Z996	TILS_SALTI	cytoplasm (probable).	cytoplasm
Q8Z9A1	LPXB_SALTI		cytoplasm
Q8Z9A2	LPXA_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z9A3	YAET_SALTI	bacterial cell outer membrane (by similarity).	outer membrane
Q8Z9A4	ECFE_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8Z9A5	UPPS_SALTI		inner membrane
Q8Z9A6	DXR_SALTI		cytoplasm
Q8Z9A7	GLND_SALTI		cytoplasm

Q8Z9A8	DAPD_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z9B1	DGTP_SALTI		cytoplasm
Q8Z9B2	BTUF_SALTI	periplasm (by similarity).	periplasm
Q8Z9B3	CLCA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8Z9B4	GSA_SALTI	cytoplasm (potential).	cytoplasm
Q8Z9C1	SFSA_SALTI		cytoplasm
Q8Z9C2	GLUQ_SALTI		cytoplasm
Q8Z9C3	PCNB_SALTI		cytoplasm
Q8Z9D2	PANB_SALTI		cytoplasm
Q8Z9D3	PANC_SALTI		cytoplasm
Q8Z9E2	SPEE_SALTI		cytoplasm
Q8Z9E3	SPED_SALTI		cytoplasm
Q8Z9E5	YACL_SALTI		inner membrane
Q8Z9F9	GUAC_SALTI		cytoplasm
Q8Z9G5	LPXC_SALTI		cytoplasm
Q8Z9G7	DDLB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z9G8	MURC_SALTI	cytoplasm (probable).	cytoplasm
Q8Z9G9	MURG_SALTI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z9H0	MURD_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z9H1	MRAY_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z9H3	MURE_SALTI	cytoplasm (probable).	cytoplasm
Q8Z9H4	MRAW_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z9H5	MRAZ_SALTI		inner membrane
Q8Z9I0	LEU1_SALTI		cytoplasm
Q8Z9I1	LEU3_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z9I2	LEU2_SALTI		cytoplasm
Q8Z9I3	LEUD_SALTI		cytoplasm
Q8Z9J4	RAPA_SALTI		cytoplasm
Q8Z9J5	RLUA_SALTI		cytoplasm
Q8Z9J6	OSTA_SALTI	bacterial cell outer membrane (by similarity).	outer membrane
Q8Z9J7	KSGA_SALTI		cytoplasm
Q8Z9J8	APAG_SALTI		inner membrane
Q8Z9K0	KEFC_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z9K1	KEFF_SALTI		cytoplasm
Q8Z9K8	FIXX_SALTI		periplasm
Q8Z9K9	FIXC_SALTI		periplasm
Q8Z9L0	FIXA_SALTI		cytoplasm
Q8Z9L1	CAIT_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z9L2	CAIA_SALTI		cytoplasm
Q8Z9L3	CAIB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z9L4	CAIC_SALTI		cytoplasm
Q8Z9L5	CAID_SALTI		cytoplasm
Q8Z9L6	CAIE_SALTI		cytoplasm
Q8Z9L7	CARB_SALTI		cytoplasm
Q8Z9L8	CARA_SALTI		cytoplasm

Q8Z9L9	DAPB_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z9M5	OADG1_SALTI	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q8Z9M9	RIHC_SALTI		extracellular
Q8Z9N1	LSPA_SALTI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8Z9N2	SYI_SALTI	cytoplasm (by similarity).	cytoplasm
Q8Z9R1	DNAK_SALTI		cytoplasm
Q8Z9R5	YAAA_SALTI		inner membrane
Q8Z9R8	GIDA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8Z9R9	GIDB_YERPE		cytoplasm
Q8Z9S4	ATPA_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z9S5	ATPG_YERPE		inner membrane
Q8Z9S7	GLMU_YERPE	cytoplasm (by similarity).	cytoplasm
Q8Z9S8	GLMS_YERPE	cytoplasm (by similarity).	cytoplasm
Q8Z9T1	PSTB2_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8Z9U1	Y4104_YERPE		cytoplasm
Q8Z9U2	TRME_YERPE		cytoplasm
Q8Z9U3	OXAA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8Z9U4	RNPA_YERPE		extracellular
Q8Z9U7	DNAA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8Z9U9	RECF_YERPE	cytoplasm (by similarity).	cytoplasm
Q8Z9V5	IBPA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8Z9V6	IBPB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8Z9V7	Y4083_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8Z9X0	MTLD_YERPE		cytoplasm
Q8Z9X7	FDHD_YERPE	cytoplasm (potential).	cytoplasm
Q8Z9Y0	FDHE_YERPE	cytoplasm (by similarity).	cytoplasm
Q8Z9Y1	SELA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8Z9Z1	XYLA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZA09	HUTU_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZA10	HUTH_YERPE	cytoplasm (potential).	cytoplasm
Q8ZA28	DCTA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZA34	CDH_YERPE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8ZA46	Y3974_YERPE		inner membrane
Q8ZA49	USPA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZA50	USPB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZA75	GLGB_YERPE		extracellular
Q8ZA76	GLGX_YERPE		extracellular
Q8ZA77	GLGC_YERPE		cytoplasm

Q8ZA78	GLGA_YERPE		cytoplasm
Q8ZA84	CAPP_YERPE		cytoplasm
Q8ZA85	ARGE_YERPE	cytoplasm (probable).	extracellular
Q8ZA86	ARGC_YERPE	cytoplasm (probable).	cytoplasm
Q8ZA87	ARGB_YERPE	cytoplasm (probable).	cytoplasm
Q8ZA88	ARLY_YERPE	cytoplasm (probable).	cytoplasm
Q8ZA97	STHA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAA0	TRMA_YERPE		cytoplasm
Q8ZAA1	BTUB_YERPE	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8ZAA2	MURI_YERPE		cytoplasm
Q8ZAA7	HDFR_YERPE		cytoplasm
Q8ZAB3	ILVD_YERPE		cytoplasm
Q8ZAC2	ILVC_YERPE		cytoplasm
Q8ZAD8	RHLB_YERPE		cytoplasm
Q8ZAE1	WECA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZAE3	WECB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAE4	WECC_YERPE		cytoplasm
Q8ZAF0	WECF_YERPE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8ZAF1	WZYE_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZAF2	WECG_YERPE		cytoplasm
Q8ZAG3	Y3839_YERPE		inner membrane
Q8ZAG5	CORA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZAH6	GLPB_YERPE		cytoplasm
Q8ZAI0	TUSA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAL3	METE_YERPE		cytoplasm
Q8ZAL4	DLHH_YERPE		cytoplasm
Q8ZAL8	RMUC_YERPE		cytoplasm
Q8ZAM1	UBIB_YERPE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8ZAM2	TATA_YERPE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8ZAM3	TATB_YERPE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q8ZAM9	FADA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAN0	FADB_YERPE		cytoplasm
Q8ZAN4	MURB_YERPE	cytoplasm (probable).	inner membrane
Q8ZAN6	COAA_YERPE	cytoplasm (probable).	cytoplasm
Q8ZAP0	NUSG_YERPE		cytoplasm
Q8ZAP1	RL11_YERPE		cytoplasm
Q8ZAP2	RL1_YERPE		cytoplasm
Q8ZAP3	RL10_YERPE		cytoplasm
Q8ZAP4	RL7_YERPE		cytoplasm
Q8ZAP5	RPOB_YERPE		cytoplasm

Q8ZAP9	THIG_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAQ1	THIE_YERPE		cytoplasm
Q8ZAQ2	THIC_YERPE		cytoplasm
Q8ZAQ5	NUDC_YERPE		cytoplasm
Q8ZAQ7	DCUP_YERPE	cytoplasm (probable).	cytoplasm
Q8ZAQ8	NFI_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAR2	PUR2_YERPE		cytoplasm
Q8ZAR3	PUR9_YERPE		cytoplasm
Q8ZAR4	META_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAR7	ACEK_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAS2	G6PI_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAS3	PSIE_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZAS8	MALK_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZAS9	LAMB1_YERPE	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8ZAU2	C5622_YERPE	periplasm (by similarity).	periplasm
Q8ZAU5	Y3691_YERPE		cytoplasm
Q8ZAU8	AAEB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZAU9	AAEA_YERPE	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q8ZAV0	AAEX_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZAW9	Y3662_YERPE		cytoplasm
Q8ZAX0	Y3661_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZAX1	AROQ_YERPE		cytoplasm
Q8ZAX6	PRMA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZAX7	DUSB_YERPE		cytoplasm
Q8ZAX8	FIS_YERPE		cytoplasm
Q8ZB04	SSUD_YERPE		cytoplasm
Q8ZB38	PYRI_YERPE		cytoplasm
Q8ZB39	PYRB_YERPE		cytoplasm
Q8ZB41	Y3586_YERPE		cytoplasm
Q8ZB47	KDSC_YERPE		cytoplasm
Q8ZB56	MURA_YERPE	cytoplasm (probable).	inner membrane
Q8ZB62	RS9_YERPE		cytoplasm
Q8ZB72	MTGA_YERPE	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8ZB74	DIAA_YERPE		cytoplasm
Q8ZB75	Y3549_YERPE		inner membrane
Q8ZB81	RS6_YERPE		cytoplasm
Q8ZB82	PRIB_YERPE		cytoplasm
Q8ZB83	RS18_YERPE		cytoplasm
Q8ZB84	RL9_YERPE		cytoplasm
Q8ZB88	RCMNS_YERPE	cytoplasm (by similarity).	cytoplasm

Q8ZB94	MSRA_YERPE		cytoplasm
Q8ZB98	IPYR_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZBA2	ARGR_YERPE	cytoplasm (probable).	cytoplasm
Q8ZBA7	RL27_YERPE		cytoplasm
Q8ZBB3	GREA_YERPE		cytoplasm
Q8ZBB5	RRMJ_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZBB8	GLMM_YERPE		cytoplasm
Q8ZBC0	Y3498_YERPE		inner membrane
Q8ZBC3	RBFA_YERPE	cytoplasm (potential).	cytoplasm
Q8ZBC4	TRUB_YERPE		cytoplasm
Q8ZBC5	RS15_YERPE		cytoplasm
Q8ZBE1	Y3475_YERPE		cytoplasm
Q8ZBE9	Y3467_YERPE		inner membrane
Q8ZBG8	OTC_YERPE	cytoplasm (probable).	cytoplasm
Q8ZBH1	SYV_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZBH3	AMPA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZBH8	Y3432_YERPE		cytoplasm
Q8ZBH9	Y3431_YERPE		inner membrane
Q8ZBI0	COAE_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZBI2	GUAC_YERPE		cytoplasm
Q8ZBJ5	Y3414_YERPE		inner membrane
Q8ZBJ7	SPED_YERPE		cytoplasm
Q8ZBJ8	SPEE_YERPE		cytoplasm
Q8ZBK6	PAND_YERPE		cytoplasm
Q8ZBK7	PANC_YERPE		cytoplasm
Q8ZBK8	PANB_YERPE		cytoplasm
Q8ZBL1	GLUQ_YERPE		cytoplasm
Q8ZBL9	GSA_YERPE	cytoplasm (potential).	cytoplasm
Q8ZBM0	CLCA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8ZBM3	BTUF_YERPE	periplasm (by similarity).	periplasm
Q8ZBM5	DGTP_YERPE		cytoplasm
Q8ZBN1	PYRG_YERPE		cytoplasm
Q8ZBN2	ENO_YERPE	cytoplasm (by similarity).	extracellular
Q8ZBN6	CYSJ_YERPE		cytoplasm
Q8ZBN7	CYSI_YERPE		cytoplasm
Q8ZBP1	CYSD_YERPE		cytoplasm
Q8ZBP2	CYSN_YERPE		cytoplasm
Q8ZBP3	CYSC_YERPE		cytoplasm
Q8ZBP5	FTSB_YERPE	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q8ZBP6	ISPD_YERPE		cytoplasm
Q8ZBP7	ISPF_YERPE		extracellular
Q8ZBP8	TRUD_YERPE		cytoplasm
Q8ZBP9	SURE_YERPE	cytoplasm (potential).	periplasm
Q8ZBQ0	PIMT_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZBQ3	MUTS_YERPE		cytoplasm
Q8ZBU2	GSH1_YERPE		cytoplasm
Q8ZBU3	LUXS_YERPE		extracellular
Q8ZBU7	RS16_YERPE		cytoplasm
Q8ZBU8	RIMM_YERPE	cytoplasm (potential).	cytoplasm
Q8ZBU9	TRMD_YERPE	cytoplasm (potential).	cytoplasm

Q8ZBV0	RL19_YERPE		cytoplasm
Q8ZBV2	Y3287_YERPE		cytoplasm
Q8ZBV7	RLUD_YERPE		cytoplasm
Q8ZBY6	FADE_YERPE		cytoplasm
Q8ZBY7	GMHA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZBZ0	NQRA_YERPE		inner membrane
Q8ZBZ1	NQRB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZBZ2	NQRC_YERPE	bacterial cell inner membrane (potential).	inner membrane
Q8ZBZ3	NQRD_YERPE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZBZ5	NQRF_YERPE	bacterial cell inner membrane (potential).	inner membrane
Q8ZBZ9	DPO4_YERPE	cytoplasm (probable).	cytoplasm
Q8ZC05	XGPT_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZC08	PROB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZC09	PROA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZC15	AROL_YERPE	cytoplasm (probable).	cytoplasm
Q8ZC17	Y3213_YERPE		inner membrane
Q8ZC18	RDGC_YERPE	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q8ZC31	ACPH_YERPE		cytoplasm
Q8ZC32	QUEA_YERPE	cytoplasm (potential).	cytoplasm
Q8ZC33	TGT_YERPE		cytoplasm
Q8ZC39	NRDR_YERPE		cytoplasm
Q8ZC41	RISB_YERPE		cytoplasm
Q8ZC42	NUSB_YERPE		cytoplasm
Q8ZC45	DXS_YERPE		cytoplasm
Q8ZC47	EX7S_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZC49	THII_YERPE	cytoplasm (potential).	cytoplasm
Q8ZC51	PANE_YERPE	cytoplasm (potential).	cytoplasm
Q8ZC52	Y3170_YERPE		inner membrane
Q8ZC64	TIG_YERPE		cytoplasm
Q8ZC65	CLPP_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZC66	CLPX_YERPE		cytoplasm
Q8ZC96	Y3121_YERPE		inner membrane
Q8ZC97	RECR_YERPE		cytoplasm
Q8ZC98	HEMH_YERPE	cytoplasm (by similarity).	extracellular
Q8ZCA7	ATCU_YERPE	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8ZCA8	CUER_YERPE	cytoplasm (probable).	cytoplasm
Q8ZCB8	LPXH_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZCC1	HDA_YERPE		nucleoid
Q8ZCC3	Y3069_YERPE		inner membrane
Q8ZCD0	DAPA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZCD2	PUR7_YERPE		cytoplasm
Q8ZCF3	NAPA_YERPE	periplasm (by similarity).	periplasm
Q8ZCF8	Y3033_YERPE		inner membrane
Q8ZCF9	HEM6_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZCG0	Y3031_YERPE		cytoplasm
Q8ZCG8	NANE_YERPE		cytoplasm

Q8ZCG9	NANK_YERPE		cytoplasm
Q8ZCH3	NANT_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZCK2	MNTH_YERPE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8ZCM2	FBPC_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZCP4	PDXJ_YERPE	cytoplasm (probable).	cytoplasm
Q8ZCP5	ACPS_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZCP8	Y2925_YERPE		cytoplasm
Q8ZCR0	HMP_YERPE		cytoplasm
Q8ZCR1	GLYA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZCS3	ISCA_YERPE		cytoplasm
Q8ZCS4	HSCB_YERPE		inner membrane
Q8ZCS5	HSCA_YERPE		cytoplasm
Q8ZCT2	NDK_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZCT9	ENGA_YERPE		cytoplasm
Q8ZCU2	EX7L_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZCU4	GUAA_YERPE		cytoplasm
Q8ZCV6	YEGP_YERPE		extracellular
Q8ZCV9	MDTC_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZCW0	MDTB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZCW1	MDTA_YERPE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8ZCX2	PPK_YERPE		cytoplasm
Q8ZCX5	PSTB1_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZCX8	PUR5_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZCX9	UPP_YERPE		cytoplasm
Q8ZD12	Y2781_YERPE		cytoplasm
Q8ZD27	TRUA_YERPE		cytoplasm
Q8ZD36	Y2756_YERPE		cytoplasm
Q8ZD40	MEPA_YERPE	periplasm (by similarity).	extracellular
Q8ZD41	AROC_YERPE		cytoplasm
Q8ZD43	Y2749_YERPE		inner membrane
Q8ZD45	FADJ_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZD46	FADI_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZD52	DSBE_YERPE	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q8ZD54	CCME_YERPE	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane

Q8ZD58	CCMA_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZD70	RECO_YERPE		cytoplasm
Q8ZD71	ERA_YERPE		cytoplasm
Q8ZD72	RNC_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZD74	LEPA_YERPE		cytoplasm
Q8ZD80	NADB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZD84	GRCA_YERPE		cytoplasm
Q8ZD85	UNG_YERPE	cytoplasm (by similarity).	extracellular
Q8ZD86	PCP_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZD89	Y2700_YERPE		inner membrane
Q8ZD91	Y2697_YERPE		cytoplasm
Q8ZD92	CPDP_YERPE	periplasm (by similarity).	periplasm
Q8ZD96	ATKA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8ZD97	ATKB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8ZD98	ATKC_YERPE	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8ZDA6	CHBG_YERPE		inner membrane
Q8ZDB1	CRCB3_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZDB2	CRCB2_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZDC4	ASR2_YERPE		fimbrium
Q8ZDC6	NRDI_YERPE		inner membrane
Q8ZDE1	NAGB_YERPE		extracellular
Q8ZDF0	Y2618_YERPE		inner membrane
Q8ZDG1	NADD_YERPE		cytoplasm
Q8ZDG3	Y2605_YERPE		inner membrane
Q8ZDG6	RLPA_YERPE	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8ZDG8	Y2600_YERPE		inner membrane
Q8ZDG9	LIPB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZDH0	LIPA_YERPE	cytoplasm (potential).	cytoplasm
Q8ZDH1	TATE_YERPE	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8ZDH2	CRCB1_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZDJ2	Y2573_YERPE		cytoplasm
Q8ZDJ6	ACKA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZDJ8	Y2564_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZDJ9	Y2563_YERPE		inner membrane
Q8ZDK3	Y2559_YERPE		cytoplasm
Q8ZDL2	NUOG_YERPE		cytoplasm
Q8ZDM8	Y2534_YERPE		inner membrane

Q8ZDP7	TAM_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZDV4	Y2451_YERPE		inner membrane
Q8ZDW7	RL35_YERPE		cytoplasm
Q8ZDW8	RL20_YERPE		cytoplasm
Q8ZDX2	IHFA_YERPE		cytoplasm
Q8ZDX4	BTUC_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZDX6	BTUD_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZDX7	ARNC_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZDX8	ARNA_YERPE		cytoplasm
Q8ZDX9	ARNT_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZDY0	Y2416_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZDY2	LPLA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZDY4	Y2410_YERPE		cytoplasm
Q8ZDZ8	MDTK_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZE08	RNT_YERPE		cytoplasm
Q8ZE19	PDXH_YERPE		cytoplasm
Q8ZE20	SYI_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZE38	Y2347_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZE42	TPX_YERPE		cytoplasm
Q8ZE46	ZNTB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZE60	ACPD_YERPE		extracellular
Q8ZE65	ASR1_YERPE		fimbrium
Q8ZE68	Y2315_YERPE	periplasm (potential).	periplasm
Q8ZE81	USPE_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZE82	FNR_YERPE	cytoplasm (probable).	cytoplasm
Q8ZEB3	CLCB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZEB6	FUMC_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZEC8	RNFA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZED2	RNFD_YERPE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZED3	RNFG_YERPE	bacterial cell inner membrane (potential).	inner membrane
Q8ZED4	RNFE_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q8ZED8	RNB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZEF0	GCH2_YERPE		extracellular
Q8ZEG0	RLUB_YERPE		cytoplasm
Q8ZEG6	TRPG_YERPE		cytoplasm
Q8ZEG7	TRPD_YERPE		cytoplasm
Q8ZEG8	TRPC_YERPE		cytoplasm
Q8ZEG9	TRPB_YERPE		cytoplasm
Q8ZEH0	TRPA_YERPE		cytoplasm
Q8ZEH4	Y2199_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZEH5	ISPZ_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZEI2	CLS_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZEI3	Y2187_YERPE		inner membrane
Q8ZEJ1	KITH_YERPE	cytoplasm (potential).	cytoplasm
Q8ZEJ5	Y2172_YERPE		inner membrane
Q8ZEK1	SELD_YERPE		cytoplasm
Q8ZEK7	MSRB_YERPE		cytoplasm
Q8ZEL2	Y2152_YERPE		inner membrane
Q8ZEL7	DADA_YERPE		cytoplasm
Q8ZEL9	FADR_YERPE	cytoplasm (potential).	cytoplasm
Q8ZEM1	DSBB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZES0	Y2083_YERPE		inner membrane
Q8ZES5	MINC_YERPE		inner membrane
Q8ZES7	MINE_YERPE		inner membrane
Q8ZES9	LCFA_YERPE	membrane; peripheral membrane protein (potential). partially membrane-associated (potential).	cytoplasm
Q8ZEU2	ZNUA_YERPE	periplasm (by similarity).	periplasm
Q8ZEU5	RUVB_YERPE		cytoplasm
Q8ZEU6	RUVA_YERPE		cytoplasm
Q8ZEU7	RUVC_YERPE		cytoplasm
Q8ZEU8	Y2055_YERPE		inner membrane
Q8ZEV5	CUTC_YERPE	cytoplasm (probable).	periplasm
Q8ZEW3	MDTH_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZEW6	Y2037_YERPE		cytoplasm
Q8ZEX4	KDSA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZEX9	HEM1_YERPE		cytoplasm
Q8ZEY0	LOLB_YERPE	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q8ZEY1	ISPE_YERPE		cytoplasm
Q8ZEY2	KPRS_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZEY4	PTH_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZF51	PGSA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZF52	UVRC_YERPE	cytoplasm (by similarity).	cytoplasm

Q8ZF61	WRBA_YERPE		cytoplasm
Q8ZF73	DCYD_YERPE		cytoplasm
Q8ZF85	FLIE_YERPE		flagellar
Q8ZFB1	FLGI2_YERPE	periplasm (by similarity).	periplasm
Q8ZFB2	FLGH2_YERPE	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8ZFD5	YEBF_YERPE	secreted protein (by similarity).	inner membrane
Q8ZFD6	Y1778_YERPE		inner membrane
Q8ZFE0	Y1774_YERPE		inner membrane
Q8ZFF7	Y1755_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZFF8	Y1754_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZFG9	Y1740_YERPE	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8ZFH8	KDUI_YERPE		cytoplasm
Q8ZFJ7	HTPX_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8ZFJ9	PROQ_YERPE	cytoplasm (potential).	inner membrane
Q8ZFL4	ALR2_YERPE		cytoplasm
Q8ZFN8	DSRB_YERPE		inner membrane
Q8ZFQ3	RLUE_YERPE		cytoplasm
Q8ZFQ5	TRMU_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZFQ6	Y1637_YERPE		inner membrane
Q8ZFR0	PEPT_YERPE	cytoplasm (by similarity).	extracellular
Q8ZFR1	NPD_YERPE	cytoplasm (probable).	cytoplasm
Q8ZFR4	LOLD_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZFS2	Y1616_YERPE		inner membrane
Q8ZFS3	NAGZ_YERPE	cytoplasm (by similarity).	extracellular
Q8ZFT4	ACP_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZFT7	FABH_YERPE	cytoplasm (probable).	cytoplasm
Q8ZFT8	PLSX_YERPE		cytoplasm
Q8ZFT9	RL32_YERPE		cytoplasm
Q8ZFU1	RLUC_YERPE		cytoplasm
Q8ZFU4	PYRC_YERPE		cytoplasm
Q8ZFV7	ASSY_YERPE	cytoplasm (probable).	cytoplasm
Q8ZFV8	LLDD_YERPE		cytoplasm
Q8ZFW5	Y1560_YERPE		inner membrane
Q8ZFX4	HIS1_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZFX5	HISX_YERPE		cytoplasm
Q8ZFX6	HIS8_YERPE		cytoplasm
Q8ZFX7	HIS7_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZFX8	HIS5_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZFX9	HIS4_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZFY0	HIS6_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZFY1	HIS2_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZFZ8	DCD_YERPE		extracellular
Q8ZFZ9	URK_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZG08	CDD_YERPE		cytoplasm

Q8ZG09	MAO1_YERPE		cytoplasm
Q8ZG15	GCH1_YERPE		cytoplasm
Q8ZG65	TUSE_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZG70	Y1442_YERPE		inner membrane
Q8ZG71	MGSA_YERPE		cytoplasm
Q8ZG78	Y1433_YERPE		inner membrane
Q8ZG80	FABA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZG91	PYRD_YERPE	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZG93	PNCB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZGA1	MUKF_YERPE	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q8ZGA4	KDSB_YERPE	cytoplasm (potential).	cytoplasm
Q8ZGA8	LPXK_YERPE		cytoplasm
Q8ZGA9	MSBA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZGB1	IHFB_YERPE		cytoplasm
Q8ZGB3	KCY_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZGB4	SERC_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZGC3	Y1380_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZGC6	LOLA_YERPE	periplasm (by similarity).	periplasm
Q8ZGD2	LFTR_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZGD5	CLPS_YERPE		cytoplasm
Q8ZGE1	HCP_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZGG1	RUMB_YERPE		cytoplasm
Q8ZGH1	Y1326_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZGH4	DEOC1_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZGJ0	Y1307_YERPE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZGJ1	END4_YERPE		extracellular
Q8ZGJ9	RPIA2_YERPE		cytoplasm
Q8ZGK7	EFPL_YERPE		cytoplasm
Q8ZGK8	UXUA_YERPE		cytoplasm
Q8ZGM2	RSUA_YERPE		cytoplasm
Q8ZGM4	RL25_YERPE		cytoplasm
Q8ZGM5	NDPA_YERPE	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q8ZGM6	Y1261_YERPE		inner membrane
Q8ZGR6	UBIG_YERPE		cytoplasm
Q8ZGS0	ECOT_YERPE	periplasm (by similarity).	periplasm
Q8ZGS9	ADIC_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZGU5	PHNC_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZGV2	DUSC_YERPE		cytoplasm

Q8ZGV8	BETI_YERPE		cytoplasm
Q8ZGV9	BETB_YERPE		cytoplasm
Q8ZGW0	BETA_YERPE		cytoplasm
Q8ZGW2	MOAE_YERPE		cytoplasm
Q8ZGW4	MOAC_YERPE		cytoplasm
Q8ZGW5	MOAA_YERPE		cytoplasm
Q8ZGW7	UVRB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZGW9	BIOD_YERPE		cytoplasm
Q8ZGX4	6PGL_YERPE		cytoplasm
Q8ZGX6	MODC_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZGY3	GAL1_YERPE	cytoplasm (potential).	cytoplasm
Q8ZGY5	GPMA_YERPE		cytoplasm
Q8ZGY6	ZITB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZGY8	NADA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZGZ1	TOLB_YERPE	periplasm (potential).	periplasm
Q8ZH00	SUCC_YERPE		cytoplasm
Q8ZH09	PPNK_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZH12	Y1103_YERPE		inner membrane
Q8ZH14	SSRP_YERPE	cytoplasm (potential).	cytoplasm
Q8ZH30	RNH_YERPE	cytoplasm (potential).	extracellular
Q8ZH34	Y1077_YERPE	cytoplasm (potential).	inner membrane
Q8ZH36	DKGB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZH37	GMHB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZH38	METN_YERPE	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8ZH39	METI_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZH40	METQ_YERPE	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8ZH47	Y1064_YERPE		inner membrane
Q8ZH50	TILS_YERPE	cytoplasm (probable).	cytoplasm
Q8ZH52	ACCA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZH54	RNH2_YERPE	cytoplasm (potential).	extracellular
Q8ZH55	LPXB_YERPE		cytoplasm
Q8ZH56	LPXA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZH57	FABZ_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZH58	YAET_YERPE	bacterial cell outer membrane (by similarity).	outer membrane
Q8ZH59	ECFE_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8ZH61	UPPS_YERPE		inner membrane
Q8ZH62	DXR_YERPE		cytoplasm
Q8ZH63	RRF_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZH64	PYRH_YERPE		cytoplasm
Q8ZH66	RS2_YERPE		cytoplasm
Q8ZH68	GLND_YERPE		cytoplasm
Q8ZH69	DAPD_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZH72	TRUC_YERPE		cytoplasm
Q8ZH75	QUEF_YERPE	cytoplasm (probable).	cytoplasm

Q8ZH78	MTFA_YERPE		cytoplasm
Q8ZH86	ARGA_YERPE	cytoplasm (probable).	cytoplasm
Q8ZH96	Y1009_YERPE		extracellular
Q8ZHE6	MLTC_YERPE	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q8ZHE7	FETP_YERPE		cytoplasm
Q8ZHE9	TRMB_YERPE		cytoplasm
Q8ZHF1	GLSA1_YERPE		extracellular
Q8ZHF4	HAM1_YERPE		extracellular
Q8ZHF5	Y944_YERPE		inner membrane
Q8ZHG0	AGUA_YERPE		extracellular
Q8ZHG2	RUVX_YERPE	cytoplasm (potential).	cytoplasm
Q8ZHG3	Y936_YERPE		inner membrane
Q8ZHG6	SPRT_YERPE	cytoplasm (potential).	extracellular
Q8ZHG8	SPEA_YERPE		cytoplasm
Q8ZHH3	PGK_YERPE	cytoplasm (potential).	cytoplasm
Q8ZHH6	ARGO_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZHH8	RPIA1_YERPE		cytoplasm
Q8ZHI2	Y911_YERPE		inner membrane
Q8ZHI6	GCST_YERPE		cytoplasm
Q8ZHI7	GCSH_YERPE		cytoplasm
Q8ZHI8	GCSP_YERPE		cytoplasm
Q8ZHJ5	Y897_YERPE		cytoplasm
Q8ZHK1	XERD_YERPE	cytoplasm (by similarity).	nucleoid
Q8ZHP0	LAMB2_YERPE	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8ZHU5	MUTH_YERPE	cytoplasm (by similarity).	extracellular
Q8ZHU8	NUDH_YERPE		cytoplasm
Q8ZHV0	LGT_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZHV1	TYSY_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZHZ4	FLG11_YERPE	periplasm (by similarity).	periplasm
Q8ZHZ5	FLGH1_YERPE	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q8ZI40	DKGA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZI56	RIBB_YERPE		cytoplasm
Q8ZI60	HLDE_YERPE		cytoplasm
Q8ZI61	GLNE_YERPE		cytoplasm
Q8ZI64	CCA_YERPE		cytoplasm
Q8ZI65	UPPP_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZI67	Y647_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZIC5	UXAB_YERPE		cytoplasm
Q8ZIC6	UXAC_YERPE		cytoplasm
Q8ZIE3	LPXC_YERPE		cytoplasm
Q8ZIE7	DDL_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIE8	MURC_YERPE	cytoplasm (probable).	cytoplasm

Q8ZIE9	MURG_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZIF1	MURD_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIF2	MRAY_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZIF4	MURE_YERPE	cytoplasm (probable).	cytoplasm
Q8ZIF7	MRAW_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIF8	MRAZ_YERPE		inner membrane
Q8ZIG8	LEU1_YERPE		cytoplasm
Q8ZIG9	LEU3_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIH0	LEU2_YERPE		cytoplasm
Q8ZIH1	LEUD_YERPE		cytoplasm
Q8ZII0	RAPA_YERPE		cytoplasm
Q8ZIK1	RLUA_YERPE		cytoplasm
Q8ZIK3	OSTA_YERPE	bacterial cell outer membrane (by similarity).	outer membrane
Q8ZIK5	KSGA_YERPE		cytoplasm
Q8ZIK6	APAG_YERPE		inner membrane
Q8ZIK7	APAH_YERPE		extracellular
Q8ZIL4	CARB_YERPE		cytoplasm
Q8ZIL5	CARA_YERPE		cytoplasm
Q8ZIL6	DAPB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIL9	LSPA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8ZIM0	SYI_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIM3	RS20_YERPE		cytoplasm
Q8ZIM6	DNAJ_YERPE	cytoplasm (probable).	cytoplasm
Q8ZIM7	DNAK_YERPE		cytoplasm
Q8ZIN2	TAL_YERPE	cytoplasm (probable).	cytoplasm
Q8ZIN3	Y462_YERPE		inner membrane
Q8ZIN5	KHSE_YERPE	cytoplasm (potential).	cytoplasm
Q8ZIP0	GPMB_YERPE		cytoplasm
Q8ZIP1	Y454_YERPE		inner membrane
Q8ZIQ1	SMP_YERPE	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZIQ2	DEOD_YERPE		cytoplasm
Q8ZIQ3	DEOB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIQ4	DEOC2_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIV4	RLMB_YERPE		cytoplasm
Q8ZIV7	PURA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIW2	HFQ_YERPE		cytoplasm
Q8ZIW3	MIAA_YERPE		cytoplasm
Q8ZIW4	MUTL_YERPE		cytoplasm
Q8ZIW9	ORN_YERPE	cytoplasm (potential).	cytoplasm
Q8ZIX0	ENGC_YERPE		cytoplasm
Q8ZIX1	PSD_YERPE		cytoplasm
Q8ZIX3	SYK3_YERPE		cytoplasm
Q8ZIX7	FRDC_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q8ZIX8	FRDD_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZIY3	CH60_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIY4	CH10_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZIY9	DSBD_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZIZ8	RHAT_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZIZ9	RHAR_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJ00	RHAS_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJ02	RHAB_YERPE		cytoplasm
Q8ZJ03	RHAD_YERPE	cytoplasm (by similarity).	extracellular
Q8ZJ06	SSB_YERPE		cytoplasm
Q8ZJ07	UVRA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJ10	ALR1_YERPE		cytoplasm
Q8ZJ14	DUSA_YERPE		cytoplasm
Q8ZJ16	LEXA_YERPE		cytoplasm
Q8ZJ18	PLSB_YERPE	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZJ73	ACTP_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZJ74	AROE_YERPE		cytoplasm
Q8ZJ77	SMG_YERPE		cytoplasm
Q8ZJ79	DEF_YERPE		cytoplasm
Q8ZJ80	FMT_YERPE		cytoplasm
Q8ZJ81	RSMB_YERPE		cytoplasm
Q8ZJ83	MSCL_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZJ87	RPOA_YERPE		cytoplasm
Q8ZJ88	RS4_YERPE		cytoplasm
Q8ZJ89	RS11_YERPE		cytoplasm
Q8ZJ90	RS13_YERPE		cytoplasm
Q8ZJ93	RL15_YERPE		cytoplasm
Q8ZJ95	RS5_YERPE		cytoplasm
Q8ZJ96	RL18_YERPE		cytoplasm
Q8ZJ98	RS8_YERPE		cytoplasm
Q8ZJ99	RS14_YERPE		cytoplasm
Q8ZJA0	RL5_YERPE		cytoplasm
Q8ZJA1	RL24_YERPE		cytoplasm
Q8ZJA3	RS17_YERPE		cytoplasm
Q8ZJA4	RL29_YERPE		cytoplasm
Q8ZJA5	RL16_YERPE		cytoplasm
Q8ZJA6	RS3_YERPE		cytoplasm
Q8ZJA7	RL22_YERPE		cytoplasm
Q8ZJA8	RS19_YERPE		cytoplasm
Q8ZJA9	RL3_YERPE		cytoplasm
Q8ZJB4	RS7_YERPE		cytoplasm
Q8ZJB5	RS12_YERPE		cytoplasm
Q8ZJB6	TUSB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJB7	TUSC_YERPE	cytoplasm (by similarity).	cytoplasm

Q8ZJB8	TUSD_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJC1	SLYX_YERPE		inner membrane
Q8ZJC4	KEFB_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZJC5	KEFG_YERPE		cytoplasm
Q8ZJD0	TAUB_YERPE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZJD4	Y179_YERPE		inner membrane
Q8ZJE6	TSGA_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZJF3	GPH_YERPE		cytoplasm
Q8ZJF6	AROB_YERPE	cytoplasm (probable).	cytoplasm
Q8ZJF7	AROK_YERPE	cytoplasm (probable).	cytoplasm
Q8ZJG8	HSLO_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJG9	PPCK_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJH2	GREB_YERPE		cytoplasm
Q8ZJI0	Y127_YERPE		inner membrane
Q8ZJI2	MALT_YERPE		cytoplasm
Q8ZJI3	GLPE_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJI8	METJ_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJJ4	HSLV_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJJ5	HSLU_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJJ7	RRAA_YERPE		cytoplasm
Q8ZJK9	TPIS_YERPE		cytoplasm
Q8ZJL6	K6PF_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJL7	FIEF_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZJM6	GPDA_YERPE	cytoplasm (probable).	cytoplasm
Q8ZJM7	SECB_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJN0	GPMI_YERPE		cytoplasm
Q8ZJN2	TDH_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJN4	HLDD_YERPE		cytoplasm
Q8ZJN9	COAD_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJP0	FPG_YERPE		cytoplasm
Q8ZJP1	RL33_YERPE		cytoplasm
Q8ZJP2	RL28_YERPE		cytoplasm
Q8ZJP3	RADC_YERPE		cytoplasm
Q8ZJP5	DUT_YERPE		extracellular
Q8ZJP6	SLMA_YERPE	nucleoid-associated (by similarity).	cytoplasm
Q8ZJP7	PYRE_YERPE		cytoplasm
Q8ZJP8	RNPH_YERPE		cytoplasm
Q8ZJQ2	KGUA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJQ3	RPOZ_YERPE		cytoplasm
Q8ZJR1	RBN_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZJR9	Y020_YERPE		inner membrane
Q8ZJS0	ENGB_YERPE		cytoplasm
Q8ZJS4	MOBA_YERPE	cytoplasm (by similarity).	cytoplasm

Q8ZJT0	KUP_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZJT1	RAVA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJT2	VIAA_YERPE		cytoplasm
Q8ZJT3	ASNA_YERPE	cytoplasm (by similarity).	cytoplasm
Q8ZJU8	GPMB_SALTY		cytoplasm
Q8ZJV0	SMP_SALTY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZJV1	LPLA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZJV7	DEOD_SALTY		cytoplasm
Q8ZJV8	DEOC_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZJX6	OPGB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZK08	YEEN_SALTY		inner membrane
Q8ZK29	AMPA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZK31	SYV_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZK33	ARCA_SALTY	cytoplasm (by similarity).	extracellular
Q8ZK35	OTCC_SALTY	cytoplasm (probable).	cytoplasm
Q8ZK55	Y4428_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZK71	MSRA_SALTY		cytoplasm
Q8ZK76	RCMNS_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZK80	RL9_SALTY		cytoplasm
Q8ZK81	RS18_SALTY		cytoplasm
Q8ZK82	PRIB_SALTY		cytoplasm
Q8ZK85	ULAF_SALTY		cytoplasm
Q8ZK86	ULAE_SALTY		cytoplasm
Q8ZK87	ULAD_SALTY		cytoplasm
Q8ZK88	ULAC_SALTY	cytoplasm (probable).	cytoplasm
Q8ZK89	ULAB_SALTY	cytoplasm (potential).	cytoplasm
Q8ZK90	ULAA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZKA2	RLMB_SALTY		cytoplasm
Q8ZKB0	ENGC_SALTY		cytoplasm
Q8ZKB1	PSD_SALTY		cytoplasm
Q8ZKC3	DSBD_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZKF5	NRFA_SALTY	periplasm (by similarity).	periplasm
Q8ZKF6	ACSA_SALTY		cytoplasm
Q8ZKF8	ACTP_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZKH4	DUSA_SALTY		cytoplasm
Q8ZKH9	PLSB_SALTY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZKI4	G6PI_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKL1	RLUF_SALTY		cytoplasm
Q8ZKL6	ARLY_SALTY	cytoplasm (probable).	cytoplasm
Q8ZKL8	ARGC_SALTY	cytoplasm (probable).	cytoplasm

Q8ZKL9	ARGE_SALTY	cytoplasm (probable).	extracellular
Q8ZKM0	CAPP_SALTY		cytoplasm
Q8ZKM8	FSA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKP3	GLPK_SALTY		cytoplasm
Q8ZKP7	TPIS_SALTY		cytoplasm
Q8ZKR4	FIEF_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZKS1	RHAD_SALTY	cytoplasm (by similarity).	extracellular
Q8ZKS7	FDHD_SALTY	cytoplasm (potential).	cytoplasm
Q8ZKT0	FDHE_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKT5	RBN_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZKU8	RDOA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKU9	MOBA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKW1	KUP_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZKW2	RAVA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKW3	VIAA_SALTY		cytoplasm
Q8ZKW4	ASNA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKW6	GIDA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKW8	ATPG_SALTY		inner membrane
Q8ZKX0	GLMU_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKX1	GLMS_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKY0	YIDZ_SALTY		cytoplasm
Q8ZKY1	MDTL_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZKY3	TRME_SALTY		cytoplasm
Q8ZKY4	OXAA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZKZ7	TORA_SALTY	periplasm (by similarity).	periplasm
Q8ZKZ8	TORD_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZKZ9	CCMA1_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZL03	IBPB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZL04	YIDE_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZL08	SDHD_SALTY		cytoplasm
Q8ZL46	SLMA_SALTY	nucleoid-associated (by similarity).	cytoplasm
Q8ZL48	COAD_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZL52	TDH_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZL56	GPMI_SALTY		cytoplasm
Q8ZL61	LLDD_SALTY		cytoplasm
Q8ZL63	LLDP_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZL67	MTLD_SALTY		cytoplasm
Q8ZL69	SELA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZL83	DLGD_SALTY	cytoplasm (potential).	cytoplasm

Q8ZL90	XYLA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZLB6	YHJQ_SALTY		inner membrane
Q8ZLB7	GUN_SALTY	secreted protein (by similarity).	extracellular
Q8ZLB8	BCSC_SALTY		outer membrane
Q8ZLC8	TREF_SALTY	cytoplasm (by similarity).	extracellular
Q8ZLD7	USPA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZLE2	ACPT_SALTY		cytoplasm
Q8ZLE4	YHHS_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZLG5	GLGB_SALTY		extracellular
Q8ZLG6	GLGX_SALTY		extracellular
Q8ZLI0	RTCA_SALTY	cytoplasm (potential).	cytoplasm
Q8ZLI3	MALT_SALTY		cytoplasm
Q8ZLI7	GNTY_SALTY		inner membrane
Q8ZLI8	GNTX_SALTY		cytoplasm
Q8ZLI9	BIOH_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZLJ2	GREB_SALTY		cytoplasm
Q8ZLJ4	HSLO_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZLK5	GPH_SALTY		cytoplasm
Q8ZLK7	TSGA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZLL3	KEFB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZLL5	SLYX_SALTY		inner membrane
Q8ZLL7	TUSD_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZLL8	TUSC_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZLL9	TUSB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZLM1	RS13_SALTY		cytoplasm
Q8ZLM5	RSMB_SALTY		cytoplasm
Q8ZLM6	FMT_SALTY		cytoplasm
Q8ZLM7	DEF_SALTY		cytoplasm
Q8ZLN1	AROE_SALTY		cytoplasm
Q8ZLP1	YEDZ_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZLP5	AAEB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZLQ6	NANA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZLQ7	NANE2_SALTY		cytoplasm
Q8ZLQ8	NANK_SALTY		cytoplasm
Q8ZLR7	MTGA_SALTY	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q8ZLR8	YHBJ_SALTY		cytoplasm
Q8ZLS0	KDSC_SALTY		cytoplasm
Q8ZLS1	KDSD_SALTY		cytoplasm
Q8ZLS7	RRMJ_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZLT0	ASSY_SALTY	cytoplasm (probable).	cytoplasm
Q8ZLT1	RBFA_SALTY	cytoplasm (potential).	cytoplasm
Q8ZLT2	TRUB_SALTY		cytoplasm
Q8ZLU2	DIAA_SALTY		cytoplasm

Q8ZLU3	YRAN_SALTY		inner membrane
Q8ZLX2	ALX_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZLY4	CCA_SALTY		cytoplasm
Q8ZLY6	GLNE_SALTY		cytoplasm
Q8ZLZ9	QSEC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZM03	YGIQ_SALTY		cytoplasm
Q8ZM06	DKGA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZM13	YQHA_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZM23	UXAC_SALTY		cytoplasm
Q8ZM39	MLTC_SALTY	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q8ZM40	TRMB_SALTY		cytoplasm
Q8ZM45	HAM1_SALTY		extracellular
Q8ZM46	YGGU_SALTY		inner membrane
Q8ZM50	RUVX_SALTY	cytoplasm (potential).	cytoplasm
Q8ZM51	YQGE_SALTY		inner membrane
Q8ZM68	ARGO_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZM71	YGFB_SALTY		inner membrane
Q8ZM75	GCSH_SALTY		cytoplasm
Q8ZM76	GCSP_SALTY		cytoplasm
Q8ZM79	YQFB_SALTY		inner membrane
Q8ZM82	IDI_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZM94	RCNA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZM98	KDUI_SALTY		cytoplasm
Q8ZMA4	AAS_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZMA8	MUTH_SALTY	cytoplasm (by similarity).	extracellular
Q8ZMA9	TYSY_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZMB5	PTRA_SALTY	periplasm (by similarity).	extracellular
Q8ZMB8	ARGA_SALTY	cytoplasm (probable).	cytoplasm
Q8ZMC5	FUCK_SALTY		cytoplasm
Q8ZMC6	FUCI_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZMD3	QUEF_SALTY	cytoplasm (probable).	cytoplasm
Q8ZMD5	TRUC_SALTY		cytoplasm
Q8ZME1	RUMA_SALTY		cytoplasm
Q8ZMF5	CYSN_SALTY		cytoplasm
Q8ZMF6	ISPD_SALTY		cytoplasm
Q8ZMF7	ISPF_SALTY		extracellular
Q8ZMF8	TRUD_SALTY		cytoplasm
Q8ZMF9	PIMT_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZMH3	PRP2_SALTY		extracellular
Q8ZMJ6	NORW_SALTY	cytoplasm (probable).	cytoplasm
Q8ZMJ7	NORV_SALTY	cytoplasm (probable).	cytoplasm
Q8ZMJ8	NORR_SALTY		cytoplasm
Q8ZMK5	RECX_SALTY	cytoplasm (potential).	cytoplasm

Q8ZMX9	NADB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZN19	PDXJ_SALTY	cytoplasm (probable).	cytoplasm
Q8ZN25	YFEU_SALTY		cytoplasm
Q8ZN39	YFHR_SALTY		inner membrane
Q8ZN40	ISCS_SALTY		cytoplasm
Q8ZN41	HSCB_SALTY		inner membrane
Q8ZN42	HSCA_SALTY		cytoplasm
Q8ZN58	EX7L_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZN60	GUAA_SALTY		cytoplasm
Q8ZN71	DAPA_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZN80	YFEW_SALTY		extracellular
Q8ZN83	TALA_SALTY	cytoplasm (probable).	cytoplasm
Q8ZNA5	FADL_SALTY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZNA6	FADI_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZNA7	FADJ_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZNB0	MEPA_SALTY	periplasm (by similarity).	extracellular
Q8ZNB2	YFCK_SALTY		cytoplasm
Q8ZNB8	YFCJ_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZNB9	TRUA_SALTY		cytoplasm
Q8ZND7	YFBV_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZNE0	YFBR_SALTY		cytoplasm
Q8ZNF0	YFBJ_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZNF3	ARNB_SALTY		cytoplasm
Q8ZNH4	ECOT_SALTY	periplasm (by similarity).	periplasm
Q8ZNH6	NAPA_SALTY	periplasm (by similarity).	periplasm
Q8ZNK6	END4_SALTY		extracellular
Q8ZNK7	YEIH_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZNL8	YEIT_SALTY		inner membrane
Q8ZNM0	CDD_SALTY		cytoplasm
Q8ZNM4	DUSC_SALTY		cytoplasm
Q8ZNN2	YEHT_SALTY		cytoplasm
Q8ZNQ1	MDTC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZNQ2	MDTB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZNQ3	MDTA_SALTY	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8ZNQ9	WZA_SALTY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q8ZNT5	YEDQ_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q8ZNT7	DCYD_SALTY		cytoplasm
Q8ZNV0	CUTC_SALTY	cytoplasm (probable).	periplasm
Q8ZNW0	KPYK2_SALTY		cytoplasm
Q8ZNW7	YEBF_SALTY	secreted protein (by similarity).	inner membrane
Q8ZNY9	PRP1_SALTY		extracellular
Q8ZP00	YEBN_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZP01	YOBD_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZP12	YCGN_SALTY		inner membrane
Q8ZP15	FADR_SALTY	cytoplasm (potential).	cytoplasm
Q8ZP17	DADA_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZP20	TREA_SALTY	periplasm (by similarity).	periplasm
Q8ZP43	YCHJ_SALTY		inner membrane
Q8ZP48	YCIU_SALTY		inner membrane
Q8ZP49	YCIC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZP51	RLUB_SALTY		cytoplasm
Q8ZP54	RNB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZP64	YCJF_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZP65	TPX_SALTY		cytoplasm
Q8ZP84	USPE_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZPB3	OPGD_SALTY	periplasm (by similarity).	periplasm
Q8ZPD6	YNCE_SALTY		cytoplasm
Q8ZPE8	MAO1_SALTY		cytoplasm
Q8ZPI0	YNEE_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZPI2	GLSA_SALTY		extracellular
Q8ZPJ2	YNFA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZPJ4	YNFC_SALTY	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8ZPK0	DMSD_SALTY		cytoplasm
Q8ZPK5	CLCB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZPK6	BIOD2_SALTY		cytoplasm
Q8ZPK9	ASR_SALTY		fimbrium
Q8ZPL7	FUMC_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZPL9	ADD_SALTY		extracellular
Q8ZPM2	RNFC_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	cytoplasm
Q8ZPM3	RNFD_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane

Q8ZPM4	RNFG_SALTY	bacterial cell inner membrane (potential).	inner membrane
Q8ZPM6	TPPB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZPM9	PDXH_SALTY		cytoplasm
Q8ZPN4	YDHK_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZPP2	MDTK_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZPQ1	SUFE_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZPR4	YDIB_SALTY		cytoplasm
Q8ZPS4	AROH_SALTY		cytoplasm
Q8ZPS5	YDIU_SALTY		inner membrane
Q8ZPS8	BTUC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZPU1	CHBG_SALTY		inner membrane
Q8ZPU5	NADE_SALTY		cytoplasm
Q8ZPU6	CHO_SALTY		cytoplasm
Q8ZPU8	ASTE_SALTY		periplasm
Q8ZPV2	ARGM_SALTY		cytoplasm
Q8ZPV5	SELD_SALTY		cytoplasm
Q8ZPW3	YEAH_SALTY		inner membrane
Q8ZPZ1	RLUE_SALTY		cytoplasm
Q8ZPZ4	TRMU_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZPZ5	YCFC_SALTY		inner membrane
Q8ZQ06	NAGZ_SALTY	cytoplasm (by similarity).	extracellular
Q8ZQ07	THIK_SALTY		cytoplasm
Q8ZQ16	RLUC_SALTY		cytoplasm
Q8ZQ19	MDTH_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZQ23	YCEA_SALTY		inner membrane
Q8ZQ25	MDTG_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZQ26	OPGH_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZQ27	OPGC_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZQ40	WRBA_SALTY		cytoplasm
Q8ZQ51	HPAC_SALTY		cytoplasm
Q8ZQ52	YEDX_SALTY	periplasm (potential).	periplasm
Q8ZQ68	YCBG_SALTY		inner membrane
Q8ZQB7	MUKF_SALTY	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q8ZQC0	KDSB_SALTY	cytoplasm (potential).	cytoplasm
Q8ZQC2	LPXK_SALTY		cytoplasm
Q8ZQC4	KCY_SALTY	cytoplasm (by similarity).	cytoplasm

Q8ZQD2	YCAD_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZQD4	LOLA_SALTY	periplasm (by similarity).	periplasm
Q8ZQE8	HCP_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZQJ5	RUMB_SALTY		cytoplasm
Q8ZQK4	YBJL_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q8ZQN7	DING_SALTY		cytoplasm
Q8ZQP0	YBHG_SALTY	periplasm (potential).	inner membrane
Q8ZQP4	YBHO_SALTY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZQQ1	MOAC_SALTY		cytoplasm
Q8ZQQ4	UVRB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZQQ5	BIOD_SALTY		cytoplasm
Q8ZQQ9	HUTH_SALTY	cytoplasm (potential).	cytoplasm
Q8ZQR1	HUTG_SALTY		cytoplasm
Q8ZQR2	HUTI_SALTY	cytoplasm (potential).	cytoplasm
Q8ZQR4	6PGL_SALTY		cytoplasm
Q8ZQR6	MODC_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZQS2	GPMA_SALTY		cytoplasm
Q8ZQT3	ZITB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZQT5	TOLB_SALTY	periplasm (potential).	periplasm
Q8ZQU2	DHSB_SALTY		cytoplasm
Q8ZQU3	DHSA_SALTY		inner membrane
Q8ZQU6	END8_SALTY		cytoplasm
Q8ZQV6	YBGL_SALTY		inner membrane
Q8ZQW1	ATKA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8ZQW2	ATKB_SALTY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8ZQW3	ATKC_SALTY	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q8ZQX1	FLAV_SALTY		cytoplasm
Q8ZQX7	NAGB_SALTY		extracellular
Q8ZQY4	RIHA_SALTY		extracellular
Q8ZQZ4	KDGT2_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZQZ8	NADD_SALTY		cytoplasm
Q8ZR01	RLPA_SALTY	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8ZR03	LIPB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZR04	LIPA_SALTY	cytoplasm (potential).	cytoplasm
Q8ZR35	ENTS_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q8ZR41	CAAL_SALTY		cytoplasm
Q8ZR69	LPXH_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZR87	ALLA_SALTY		extracellular
Q8ZR88	SELU_SALTY		cytoplasm
Q8ZR95	ATCU_SALTY	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q8ZRC0	CLPX_SALTY		cytoplasm
Q8ZRC9	YAJQ_SALTY		inner membrane
Q8ZRD1	DXS_SALTY		cytoplasm
Q8ZRD3	NUSB_SALTY		cytoplasm
Q8ZRD8	TGT_SALTY		cytoplasm
Q8ZRD9	QUEA_SALTY	cytoplasm (potential).	cytoplasm
Q8ZRE7	YAIE_SALTY		inner membrane
Q8ZRF8	YAIT_SALTY		extracellular
Q8ZRI9	LEUD2_SALTY		cytoplasm
Q8ZRJ0	LEU22_SALTY		cytoplasm
Q8ZRJ7	FADE_SALTY		cytoplasm
Q8ZRM4	YAFD_SALTY	cytoplasm (potential).	inner membrane
Q8ZRM7	DKGB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZRM8	GMHB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZRM9	METN_SALTY	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q8ZRN0	METI_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZRN1	METQ_SALTY	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q8ZRN5	TILS_SALTY	cytoplasm (probable).	cytoplasm
Q8ZRN9	LPXB_SALTY		cytoplasm
Q8ZRP0	YAET_SALTY	bacterial cell outer membrane (by similarity).	outer membrane
Q8ZRP1	ECFE_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8ZRP2	UPPS_SALTY		inner membrane
Q8ZRP3	DXR_SALTY		cytoplasm
Q8ZRP4	DAPD_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZRP7	BTUF_SALTY	periplasm (by similarity).	periplasm
Q8ZRP8	CLCA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q8ZRQ6	SFSA_SALTY		cytoplasm
Q8ZRQ7	GLUQ_SALTY		cytoplasm
Q8ZRQ8	PCNB_SALTY		cytoplasm
Q8ZRR0	PANB_SALTY		cytoplasm
Q8ZRR1	PANC_SALTY		cytoplasm
Q8ZRS3	SPEE_SALTY		cytoplasm
Q8ZRS4	SPED_SALTY		cytoplasm
Q8ZRS6	YACL_SALTY		inner membrane
Q8ZRT5	GUAC_SALTY		cytoplasm
Q8ZRT9	LPXC_SALTY		cytoplasm
Q8ZRU1	DDLB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZRU2	MURC_SALTY	cytoplasm (probable).	cytoplasm

Q8ZRU3	MURG_SALTY	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q8ZRU4	MURD_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZRU5	MRAY_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZRU7	MURE_SALTY	cytoplasm (probable).	cytoplasm
Q8ZRU8	MRAW_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZRU9	MRAZ_SALTY		inner membrane
Q8ZRV8	RAPA_SALTY		cytoplasm
Q8ZRV9	RLUA_SALTY		cytoplasm
Q8ZRW0	OSTA_SALTY	bacterial cell outer membrane (by similarity).	outer membrane
Q8ZRW2	KEFC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZRW3	KEFF_SALTY		cytoplasm
Q8ZRW8	FIXX_SALTY		periplasm
Q8ZRW9	FIXC_SALTY		periplasm
Q8ZRX0	FIXA_SALTY		cytoplasm
Q8ZRX1	CAIT_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZRX2	CAIA_SALTY		cytoplasm
Q8ZRX3	CAIB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZRX4	CAIC_SALTY		cytoplasm
Q8ZRX5	CAID_SALTY		cytoplasm
Q8ZRX6	CAIE_SALTY		cytoplasm
Q8ZRX8	DAPB_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZRY4	OADB1_SALTY	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q8ZRY7	RIHC_SALTY		extracellular
Q8ZRY9	LSPA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q8ZRZ0	SYI_SALTY	cytoplasm (by similarity).	cytoplasm
Q8ZS17	YAAA_SALTY		inner membrane
Q91UN2	MERB_RHIME		inner membrane
Q925W7	RS11_RHIME		cytoplasm
Q925X7	RS13_RHIME		cytoplasm
Q925Z2	RPOA_RHIME		cytoplasm
Q926A5	RL17_RHIME		cytoplasm
Q926C0	OMP19_RHIME	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q926C2	TOLB_RHIME	periplasm (potential).	periplasm
Q926C3	OMP16_RHIME	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q92FV3	DCUP_RICCN	cytoplasm (probable).	cytoplasm
Q92FV4	HEMH_RICCN	cytoplasm (by similarity).	extracellular
Q92FV8	HEM6_RICCN	cytoplasm (by similarity).	cytoplasm
Q92FW6	MUTL_RICCN		cytoplasm
Q92FW7	RL33_RICCN		cytoplasm
Q92FW8	RS16_RICCN		cytoplasm
Q92FX0	LIPB_RICCN	cytoplasm (by similarity).	cytoplasm

Q92FX1	Y1356_RICCN		inner membrane
Q92FZ0	Y1337_RICCN		inner membrane
Q92FZ5	GREA_RICCN		cytoplasm
Q92FZ8	RPOD_RICCN		cytoplasm
Q92FZ9	TRUA_RICCN		cytoplasm
Q92G13	HEMK_RICCN		cytoplasm
Q92G20	TIG_RICCN		cytoplasm
Q92G24	Y1301_RICCN		inner membrane
Q92G30	SSB_RICCN		cytoplasm
Q92G31	UVRA_RICCN	cytoplasm (by similarity).	cytoplasm
Q92G55	XERC_RICCN	cytoplasm (by similarity).	nucleoid
Q92G57	Y1266_RICCN	cytoplasm (potential).	inner membrane
Q92G85	ATPD_RICCN		inner membrane
Q92G86	ATPA_RICCN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92G88	ATPB_RICCN		inner membrane
Q92G89	ATPE_RICCN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92G90	ACON_RICCN		cytoplasm
Q92G92	NUOG_RICCN		cytoplasm
Q92G94	NUOI_RICCN		cytoplasm
Q92G95	CCMA_RICCN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92G96	NUOM_RICCN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92G97	NUOL_RICCN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92G98	NUOK_RICCN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92GB1	UDG_RICCN		cytoplasm
Q92GC0	RL32_RICCN		cytoplasm
Q92GC1	FABH_RICCN	cytoplasm (probable).	cytoplasm
Q92GC8	Y1195_RICCN		cytoplasm
Q92GC9	KGUA_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GD8	ACP_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GE1	RECA_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GE8	TRME_RICCN		cytoplasm
Q92GG0	RL27_RICCN		cytoplasm
Q92GG3	TATA_RICCN	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q92GH4	END3_RICCN		cytoplasm
Q92GH5	GLRXA_RICCN		cytoplasm
Q92GH6	Y1147_RICCN		cytoplasm
Q92GH7	GLYA_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GH8	LIPA_RICCN	cytoplasm (potential).	cytoplasm
Q92GI1	Y1142_RICCN		inner membrane
Q92GL0	SUR1_RICCN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q92GL1	DPO3E_RICCN		cytoplasm
Q92GL2	COAE_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GL5	RNH_RICCN	cytoplasm (potential).	extracellular
Q92GM6	TGT_RICCN		cytoplasm
Q92GN1	LPXK_RICCN		cytoplasm
Q92GP5	LOLD_RICCN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92GQ4	CLPX_RICCN		cytoplasm
Q92GR5	RMUC_RICCN		cytoplasm
Q92GR9	SYV_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GS5	KTHY_RICCN		cytoplasm
Q92GT4	FPG_RICCN		cytoplasm
Q92GT5	UBIE_RICCN		cytoplasm
Q92GU2	ENGA_RICCN		cytoplasm
Q92GU6	EX7L_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GV0	KSGA_RICCN		cytoplasm
Q92GV1	CSPA_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GV5	Y1017_RICCN		cytoplasm
Q92GW0	FUMC_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GW5	RS10_RICCN		cytoplasm
Q92GW6	RL3_RICCN		cytoplasm
Q92GW7	RL4_RICCN		cytoplasm
Q92GW9	RL2_RICCN		cytoplasm
Q92GX0	RS19_RICCN		cytoplasm
Q92GX1	RL22_RICCN		cytoplasm
Q92GX2	RS3_RICCN		cytoplasm
Q92GX3	RL16_RICCN		cytoplasm
Q92GX4	RL29_RICCN		cytoplasm
Q92GX5	RS17_RICCN		cytoplasm
Q92GX7	RL24_RICCN		cytoplasm
Q92GX8	RL5_RICCN		cytoplasm
Q92GX9	RS14_RICCN		cytoplasm
Q92GY0	RS8_RICCN		cytoplasm
Q92GY2	RL18_RICCN		cytoplasm
Q92GY3	RS5_RICCN		cytoplasm
Q92GY5	RL15_RICCN		cytoplasm
Q92GY7	KAD_RICCN	cytoplasm (by similarity).	cytoplasm
Q92GY8	RS13_RICCN		cytoplasm
Q92GY9	RS11_RICCN		cytoplasm
Q92GZ0	RPOA_RICCN		cytoplasm
Q92GZ1	RL17_RICCN		cytoplasm
Q92GZ2	RS20_RICCN		cytoplasm
Q92GZ5	GRPE_RICCN	cytoplasm (probable).	cytoplasm
Q92GZ7	RNPH_RICCN		cytoplasm
Q92H03	CH10_RICCN	cytoplasm (by similarity).	cytoplasm
Q92H04	CH60_RICCN	cytoplasm (by similarity).	cytoplasm
Q92H07	UBIG_RICCN		cytoplasm
Q92H19	SYI_RICCN	cytoplasm (by similarity).	cytoplasm
Q92H23	RS21_RICCN		cytoplasm
Q92H35	RNPA_RICCN		extracellular
Q92H37	RL20_RICCN		cytoplasm
Q92H38	RL35_RICCN		cytoplasm
Q92H40	RL25_RICCN		cytoplasm
Q92H41	PTH_RICCN	cytoplasm (by similarity).	cytoplasm
Q92H56	DNAA_RICCN	cytoplasm (by similarity).	cytoplasm

Q92H59	MURE_RICCN	cytoplasm (probable).	cytoplasm
Q92H60	MURF_RICCN	cytoplasm (probable).	cytoplasm
Q92H61	MRAY_RICCN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92H74	IPYR_RICCN	cytoplasm (by similarity).	cytoplasm
Q92H75	CCME_RICCN	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q92H76	SCO22_RICCN		cytoplasm
Q92H87	GYRB_RICCN		cytoplasm
Q92H88	MURA_RICCN	cytoplasm (probable).	inner membrane
Q92H89	RPOZ_RICCN		cytoplasm
Q92H90	ACPS_RICCN	cytoplasm (by similarity).	cytoplasm
Q92H91	PLP_RICCN	bacterial cell outer membrane; peripheral membrane protein (by similarity).	cytoplasm
Q92HA7	UVRC_RICCN	cytoplasm (by similarity).	cytoplasm
Q92HB3	MRAZ_RICCN		inner membrane
Q92HB4	MRAW_RICCN		cytoplasm
Q92HD6	Y835_RICCN		inner membrane
Q92HE8	SYU_RICCN	cytoplasm (by similarity).	cytoplasm
Q92HF3	Y818_RICCN		inner membrane
Q92HF4	NUSA_RICCN		cytoplasm
Q92HG1	RADA_RICCN		cytoplasm
Q92HG4	Y807_RICCN		cytoplasm
Q92HG8	DNAB_RICCN		nucleoid
Q92HH1	PRIA_RICCN		cytoplasm
Q92HI8	PPDK_RICCN		cytoplasm
Q92HJ3	SODF_RICCN		periplasm
Q92HK7	ODP2_RICCN		inner membrane
Q92HL4	DBHL_RICCN		cytoplasm
Q92HL7	Y754_RICCN		inner membrane
Q92HM3	KCY_RICCN	cytoplasm (by similarity).	cytoplasm
Q92HM5	CLPP_RICCN	cytoplasm (by similarity).	cytoplasm
Q92HM7	IDI2_RICCN	cytoplasm (by similarity).	cytoplasm
Q92HP1	ISCS_RICCN		cytoplasm
Q92HR5	HEM3_RICCN		cytoplasm
Q92HU0	Y681_RICCN		inner membrane
Q92HU7	DEFL_RICCN		cytoplasm
Q92HU9	Y672_RICCN		cytoplasm
Q92HV3	Y668_RICCN		cytoplasm
Q92HV5	TRUB_RICCN		cytoplasm
Q92HV6	RS15_RICCN		cytoplasm
Q92HW0	Y660_RICCN		inner membrane
Q92HW4	MIAA_RICCN		cytoplasm
Q92HY4	FOLD_RICCN		cytoplasm
Q92I02	TRXB_RICCN	cytoplasm (by similarity).	cytoplasm
Q92I05	GPDA_RICCN	cytoplasm (probable).	cytoplasm
Q92I08	PPNK_RICCN	cytoplasm (by similarity).	cytoplasm
Q92I11	RECR_RICCN		cytoplasm
Q92I14	RBFA_RICCN	cytoplasm (potential).	cytoplasm
Q92I21	SUCC_RICCN		cytoplasm
Q92I24	SSRP_RICCN	cytoplasm (potential).	cytoplasm
Q92I25	DAPA_RICCN	cytoplasm (by similarity).	cytoplasm

Q92I26	ILVE_RICCN		cytoplasm
Q92I30	UPPS_RICCN		inner membrane
Q92I37	DPO3B_RICCN	cytoplasm (by similarity).	cytoplasm
Q92I41	DAFP_RICCN	cytoplasm (by similarity).	cytoplasm
Q92I58	MURG_RICCN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92I60	MURD_RICCN	cytoplasm (by similarity).	cytoplasm
Q92I62	LSPA_RICCN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q92I65	COX2_RICCN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92I71	Y549_RICCN		extracellular
Q92I74	DUT_RICCN		extracellular
Q92I78	Y542_RICCN		cytoplasm
Q92I81	ISPZ_RICCN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92I83	RLPA_RICCN		cytoplasm
Q92I87	RUVB_RICCN		cytoplasm
Q92I89	RUVA_RICCN		cytoplasm
Q92I93	GCH1_RICCN		cytoplasm
Q92I96	KDSB_RICCN	cytoplasm (potential).	cytoplasm
Q92I97	PYRG_RICCN		cytoplasm
Q92IA0	MDH_RICCN		cytoplasm
Q92IC5	LNT_RICCN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92IC6	FABI_RICCN	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92IC9	XERD_RICCN	cytoplasm (by similarity).	nucleoid
Q92ID6	NUOB_RICCN		cytoplasm
Q92ID9	NUOE_RICCN		cytoplasm
Q92IE0	3MGH_RICCN		cytoplasm
Q92IE5	EX7S_RICCN	cytoplasm (by similarity).	cytoplasm
Q92IE7	RIMM_RICCN	cytoplasm (potential).	cytoplasm
Q92IF1	COXX_RICCN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q92IF2	RS4_RICCN		cytoplasm
Q92IG5	RUVX_RICCN	cytoplasm (potential).	cytoplasm
Q92IH1	TOP1_RICCN		cytoplasm
Q92IH7	CYAY_RICCN		inner membrane
Q92II0	LPXB_RICCN		cytoplasm
Q92II6	HSLU_RICCN	cytoplasm (by similarity).	cytoplasm
Q92II7	HSLV_RICCN	cytoplasm (by similarity).	cytoplasm
Q92IL0	TRMU_RICCN	cytoplasm (by similarity).	cytoplasm
Q92IL2	COXZ_RICCN		inner membrane
Q92IL4	TOLB_RICCN	periplasm (potential).	periplasm
Q92IL5	THYX_RICCN		cytoplasm
Q92IL9	MUTS_RICCN		cytoplasm
Q92IQ1	LEPA_RICCN		cytoplasm
Q92IQ7	HSP19_RICCN		cytoplasm

Q92IR4	Y356_RICCN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92IR6	Y354_RICCN		inner membrane
Q92IR7	IDH_RICCN		cytoplasm
Q92IS2	ODPB_RICCN		cytoplasm
Q92IS3	ODPA_RICCN		cytoplasm
Q92IS6	RLUC_RICCN		cytoplasm
Q92IT1	LPXC_RICCN		cytoplasm
Q92IT7	DDL_RICCN	cytoplasm (by similarity).	cytoplasm
Q92IT8	MURB_RICCN	cytoplasm (probable).	inner membrane
Q92IT9	MURC_RICCN	cytoplasm (probable).	cytoplasm
Q92IU5	PSD_RICCN		cytoplasm
Q92IU6	Y324_RICCN		cytoplasm
Q92IV0	NUDH_RICCN		cytoplasm
Q92IV4	RS9_RICCN		cytoplasm
Q92IV6	Y314_RICCN		cytoplasm
Q92IX6	PUR7_RICCN		cytoplasm
Q92IX7	Y293_RICCN		extracellular
Q92IY7	QUEA_RICCN	cytoplasm (potential).	cytoplasm
Q92IZ1	DEF1_RICCN		cytoplasm
Q92IZ6	GYRA_RICCN		cytoplasm
Q92J03	UVRB_RICCN	cytoplasm (by similarity).	cytoplasm
Q92J05	RNH2_RICCN	cytoplasm (potential).	extracellular
Q92J06	HSCB_RICCN		inner membrane
Q92J07	HSCA_RICCN		cytoplasm
Q92J08	FER2_RICCN		cytoplasm
Q92J26	DAPD_RICCN	cytoplasm (by similarity).	cytoplasm
Q92J36	DNAK_RICCN		cytoplasm
Q92J37	DNAJ_RICCN	cytoplasm (probable).	cytoplasm
Q92J40	REC_N_RICCN		cytoplasm
Q92J42	ODO1_RICCN		cytoplasm
Q92J43	ODO2_RICCN		inner membrane
Q92J55	SRP54_RICCN		cytoplasm
Q92J57	DBH_RICCN		cytoplasm
Q92J59	Y210_RICCN		cytoplasm
Q92J64	RRMJ_RICCN	cytoplasm (by similarity).	cytoplasm
Q92J65	NUSB_RICCN		cytoplasm
Q92J66	Y203_RICCN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q92J71	PYRH_RICCN		cytoplasm
Q92J72	RRF_RICCN	cytoplasm (by similarity).	cytoplasm
Q92J74	GATC_RICCN		cytoplasm
Q92J75	GATA_RICCN		cytoplasm
Q92J76	GATB_RICCN		cytoplasm
Q92J79	DAPB_RICCN	cytoplasm (by similarity).	cytoplasm
Q92J81	Y188_RICCN		cytoplasm
Q92J85	AMPA_RICCN	cytoplasm (by similarity).	cytoplasm
Q92J87	RL7_RICCN		cytoplasm
Q92J88	RL10_RICCN		cytoplasm
Q92J89	RL1_RICCN		cytoplasm
Q92J90	RL11_RICCN		cytoplasm
Q92J91	NUSG_RICCN		cytoplasm
Q92J94	RS7_RICCN		cytoplasm
Q92J95	RS12_RICCN		cytoplasm

Q92JA0	Y167_RICCN		inner membrane
Q92JA1	DEGP_RICCN		cytoplasm
Q92JA4	MRP_RICCN		cytoplasm
Q92JA8	RUVC_RICCN		cytoplasm
Q92JA9	ERA_RICCN		cytoplasm
Q92JB0	RNC_RICCN	cytoplasm (by similarity).	cytoplasm
Q92JB1	LEP_RICCN	bacterial cell membrane; single-pass type ii membrane protein (potential).	inner membrane
Q92JB2	NUOF_RICCN		periplasm
Q92JB4	Y153_RICCN		inner membrane
Q92JB5	RL19_RICCN		cytoplasm
Q92JB6	TRMD_RICCN	cytoplasm (potential).	cytoplasm
Q92JC9	ENGB_RICCN		cytoplasm
Q92JD0	RL31_RICCN		cytoplasm
Q92JD1	RL28_RICCN		cytoplasm
Q92JD2	Y135_RICCN		extracellular
Q92JD3	Y134_RICCN		cytoplasm
Q92JD4	Y133_RICCN		cytoplasm
Q92JD5	Y132_RICCN		cytoplasm
Q92JD6	Y131_RICCN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92JD9	ALR_RICCN		cytoplasm
Q92JE7	AAT_RICCN	cytoplasm (by similarity).	cytoplasm
Q92JF1	Y116_RICCN		cytoplasm
Q92JF2	Y115_RICCN		extracellular
Q92JF5	RS2_RICCN		cytoplasm
Q92JF9	Y108_RICCN		cytoplasm
Q92JG5	QUEF_RICCN	cytoplasm (probable).	cytoplasm
Q92JG7	SECB_RICCN	cytoplasm (by similarity).	cytoplasm
Q92JG8	DCD_RICCN		extracellular
Q92JH0	PARC_RICCN	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q92JH1	Y096_RICCN		inner membrane
Q92JH3	DGTL1_RICCN		cytoplasm
Q92JH6	Y091_RICCN		cytoplasm
Q92JH7	KDSA_RICCN	cytoplasm (by similarity).	cytoplasm
Q92JH8	Y089_RICCN		inner membrane
Q92JI2	GIDB_RICCN		cytoplasm
Q92JI3	GIDA_RICCN	cytoplasm (by similarity).	cytoplasm
Q92JI4	NDK_RICCN	cytoplasm (by similarity).	cytoplasm
Q92JI7	DEF2_RICCN		cytoplasm
Q92JI8	Y079_RICCN		cytoplasm
Q92JJ2	PGSA_RICCN	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92JJ3	OXAA_RICCN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92JJ5	LGT_RICCN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92JJ7	Y070_RICCN		cytoplasm
Q92JJ8	DHSB_RICCN		cytoplasm

Q92JK0	TILS_RICCN	cytoplasm (probable).	cytoplasm
Q92JK1	RL9_RICCN		cytoplasm
Q92JK2	RS18_RICCN		cytoplasm
Q92JK3	RS6_RICCN		cytoplasm
Q92JK7	Y060_RICCN		cytoplasm
Q92JK8	CLPB_RICCN	cytoplasm (probable).	cytoplasm
Q92JK9	Y058_RICCN		cytoplasm
Q92JL0	Y057_RICCN		cytoplasm
Q92JL2	Y055_RICCN		cytoplasm
Q92JL4	Y053_RICCN		cytoplasm
Q92JL5	Y052_RICCN		cytoplasm
Q92JL6	Y051_RICCN		inner membrane
Q92JL7	Y050_RICCN		cytoplasm
Q92JL8	Y049_RICCN		cytoplasm
Q92JL9	Y048_RICCN		cytoplasm
Q92JM0	Y047_RICCN		extracellular
Q92JM2	Y045_RICCN		inner membrane
Q92JM4	Y043_RICCN		inner membrane
Q92JM5	SCO21_RICCN		cytoplasm
Q92JM6	Y041_RICCN		cytoplasm
Q92JM7	Y040_RICCN		extracellular
Q92JM8	Y039_RICCN		cytoplasm
Q92JN5	RECF_RICCN	cytoplasm (by similarity).	cytoplasm
Q92JP1	ATPL_RICCN	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92JP3	ATPF_RICCN		inner membrane
Q92JP4	Y023_RICCN		cytoplasm
Q92JP5	Y022_RICCN		extracellular
Q92JQ1	Y016_RICCN		cytoplasm
Q92JQ4	Y013_RICCN		cytoplasm
Q92JQ5	Y012_RICCN		cytoplasm
Q92JQ6	DUS_RICCN		cytoplasm
Q92JQ7	LPXD_RICCN		cytoplasm
Q92JQ8	FABZ_RICCN	cytoplasm (by similarity).	cytoplasm
Q92JQ9	LPXA_RICCN	cytoplasm (by similarity).	cytoplasm
Q92JR0	Y007_RICCN		cytoplasm
Q92JR5	THIO_RICCN		cytoplasm
Q92JR6	Y001_RICCN		cytoplasm
Q92JY3	TILS_RHIME	cytoplasm (probable).	cytoplasm
Q92KF8	ATE_RHIME	cytoplasm (potential).	cytoplasm
Q92KH5	RECO_RHIME		cytoplasm
Q92KL6	HISZ_RHIME	cytoplasm (by similarity).	cytoplasm
Q92KT8	SLYX_RHIME		inner membrane
Q92KW2	GIDA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92KW3	GIDB_RHIME		cytoplasm
Q92KX2	ACSA2_RHIME		cytoplasm
Q92KX3	HTPX_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q92KX6	PUR9_RHIME		cytoplasm
Q92KY8	LEU3_RHIME	cytoplasm (by similarity).	cytoplasm
Q92KZ7	SP39_RHIME	periplasm (probable).	periplasm
Q92L19	ISPG_RHIME		cytoplasm
Q92L21	HIS82_RHIME		cytoplasm

Q92L23	MTGA_RHIME	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q92L25	RL32_RHIME		cytoplasm
Q92L39	RL19_RHIME		cytoplasm
Q92L41	TRMD_RHIME	cytoplasm (potential).	cytoplasm
Q92L42	RIMM_RHIME	cytoplasm (potential).	cytoplasm
Q92L43	RS16_RHIME		cytoplasm
Q92L46	DAPF_RHIME	cytoplasm (by similarity).	cytoplasm
Q92L49	ISPZ_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92L50	Y3235_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92L55	CCMA_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92L66	PHS_RHIME		cytoplasm
Q92L73	ASSY_RHIME	cytoplasm (probable).	cytoplasm
Q92L76	LEU2_RHIME		cytoplasm
Q92L99	META_RHIME	cytoplasm (by similarity).	cytoplasm
Q92LA1	LEUD_RHIME		cytoplasm
Q92LA6	DNE21_RHIME	cytoplasm (probable).	cytoplasm
Q92LA8	NUDH_RHIME		cytoplasm
Q92LB0	Y3164_RHIME		inner membrane
Q92LB2	PROA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92LB3	PROB1_RHIME	cytoplasm (by similarity).	cytoplasm
Q92LB7	RL27_RHIME		cytoplasm
Q92LF7	PROB2_RHIME	cytoplasm (by similarity).	cytoplasm
Q92LH1	IPYR_RHIME	cytoplasm (by similarity).	cytoplasm
Q92LH7	HEM3_RHIME		cytoplasm
Q92LI5	MSRA2_RHIME		cytoplasm
Q92LK1	XERC_RHIME	cytoplasm (by similarity).	nucleoid
Q92LK3	TAL_RHIME	cytoplasm (probable).	cytoplasm
Q92LK6	ATPA_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92LK7	ATPG_RHIME		inner membrane
Q92LK9	ATPE_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92LP6	DXR_RHIME		cytoplasm
Q92LU2	MODC_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92LU5	UNG_RHIME	cytoplasm (by similarity).	extracellular
Q92LW9	XYLA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92LX6	Y2887_RHIME		inner membrane
Q92M03	RS21B_RHIME		cytoplasm
Q92M13	GLGC_RHIME		cytoplasm
Q92M14	GLGB_RHIME		extracellular
Q92M28	ILVD_RHIME		cytoplasm
Q92M52	HEMH_RHIME	cytoplasm (by similarity).	extracellular
Q92M53	OMP10_RHIME	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane

Q92M59	Y2793_RHIME		inner membrane
Q92M71	SUHB_RHIME		cytoplasm
Q92M79	PGK_RHIME	cytoplasm (potential).	cytoplasm
Q92M88	Y2753_RHIME		inner membrane
Q92M90	RUVC_RHIME		cytoplasm
Q92M91	RUVA_RHIME		cytoplasm
Q92M92	RUVB_RHIME		cytoplasm
Q92M99	GLMM_RHIME		cytoplasm
Q92MA5	PURA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92MB5	TRMU_RHIME	cytoplasm (by similarity).	cytoplasm
Q92ME2	ACCA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92ME3	XERD_RHIME	cytoplasm (by similarity).	nucleoid
Q92ME6	AROK_RHIME	cytoplasm (probable).	cytoplasm
Q92ME7	AROB_RHIME	cytoplasm (probable).	cytoplasm
Q92MF4	RL28_RHIME		cytoplasm
Q92MG0	HIS81_RHIME		cytoplasm
Q92MH1	ARLY1_RHIME	cytoplasm (probable).	cytoplasm
Q92MJ1	ARGJ_RHIME	cytoplasm (probable).	cytoplasm
Q92MK1	UBIG_RHIME		cytoplasm
Q92MK7	CLPB_RHIME	cytoplasm (probable).	cytoplasm
Q92MQ3	DEOC_RHIME	cytoplasm (by similarity).	cytoplasm
Q92MQ8	G6P11_RHIME	cytoplasm (by similarity).	cytoplasm
Q92MT4	XYLD_RHIME		cytoplasm
Q92MY3	UREE_RHIME	cytoplasm (by similarity).	cytoplasm
Q92N33	ADEC1_RHIME		cytoplasm
Q92N38	KITH_RHIME	cytoplasm (potential).	cytoplasm
Q92N42	FTHS_RHIME		cytoplasm
Q92N45	QUEF_RHIME	cytoplasm (probable).	cytoplasm
Q92N66	CLPS2_RHIME		cytoplasm
Q92N67	PTH_RHIME	cytoplasm (by similarity).	cytoplasm
Q92N68	RL25_RHIME		cytoplasm
Q92N73	KPRS_RHIME	cytoplasm (by similarity).	cytoplasm
Q92N77	LGT_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92N95	CARA_RHIME		cytoplasm
Q92NH8	TPIS2_RHIME		cytoplasm
Q92NI1	RISB2_RHIME		cytoplasm
Q92NL3	MRAZ_RHIME		inner membrane
Q92NL4	MRAW_RHIME		cytoplasm
Q92NL6	MURE_RHIME	cytoplasm (probable).	cytoplasm
Q92NL9	MURG_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92NM0	MURC_RHIME	cytoplasm (probable).	cytoplasm
Q92NM1	MURB_RHIME	cytoplasm (probable).	inner membrane
Q92NM3	AQPZ1_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92NM4	DDL_RHIME	cytoplasm (by similarity).	cytoplasm
Q92NN0	PANC_RHIME		cytoplasm
Q92NN1	PANB_RHIME		cytoplasm
Q92NN5	PRMA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92NQ5	TYSY_RHIME	cytoplasm (by similarity).	cytoplasm
Q92NR2	MIAA_RHIME		cytoplasm
Q92NT6	PDXJ_RHIME		cytoplasm

Q92P31	COBQ_RHIME		cytoplasm
Q92P48	COBB_RHIME		cytoplasm
Q92P51	ACPXL_RHIME	cytoplasm (by similarity).	cytoplasm
Q92P97	COBS_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92P98	COBT_RHIME		cytoplasm
Q92PB2	MOBA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PB4	MOAA_RHIME		cytoplasm
Q92PB6	FUMC_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PB8	RPIA_RHIME		cytoplasm
Q92PD8	HEM6_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PF5	Y1807_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92PG8	MURI_RHIME		cytoplasm
Q92PG9	RS4_RHIME		cytoplasm
Q92PH0	GLSA_RHIME		extracellular
Q92PH7	PURL_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PI1	PURQ_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PI3	PUR7_RHIME		cytoplasm
Q92PK5	SYV_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PL6	RADC_RHIME		cytoplasm
Q92PR8	MOAC_RHIME		cytoplasm
Q92PR9	TRPC_RHIME		cytoplasm
Q92PS0	TRPD_RHIME		cytoplasm
Q92PS3	GLMU_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PS4	GLMS_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PU1	XGPT_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PU3	WRBA1_RHIME		cytoplasm
Q92PW3	LEXA_RHIME		cytoplasm
Q92PW7	KTHY_RHIME		cytoplasm
Q92PY4	TGT_RHIME		cytoplasm
Q92PY5	QUEA_RHIME	cytoplasm (potential).	cytoplasm
Q92PY8	COAD_RHIME	cytoplasm (by similarity).	cytoplasm
Q92PZ0	NORM_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92PZ4	CARB_RHIME		cytoplasm
Q92Q10	GCSH_RHIME		cytoplasm
Q92Q11	GCSP_RHIME		cytoplasm
Q92Q12	TIG_RHIME		cytoplasm
Q92Q15	GID_RHIME	cytoplasm (by similarity).	cytoplasm
Q92Q24	TATB_RHIME	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q92Q25	TATA_RHIME	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q92Q32	DGTL1_RHIME		cytoplasm
Q92Q37	SYV_RHIME	cytoplasm (by similarity).	cytoplasm
Q92Q45	LPXA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92Q46	FABZ_RHIME	cytoplasm (by similarity).	cytoplasm
Q92Q47	LPXD_RHIME		cytoplasm

Q92Q49	Y1501_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q92Q51	UPPS_RHIME		inner membrane
Q92Q52	RRF_RHIME	cytoplasm (by similarity).	cytoplasm
Q92Q53	PYRH_RHIME		cytoplasm
Q92Q55	RS2_RHIME		cytoplasm
Q92Q63	CLPS1_RHIME		cytoplasm
Q92Q90	ISPDF_RHIME		cytoplasm
Q92Q94	LIPA_RHIME	cytoplasm (potential).	cytoplasm
Q92Q98	ENO_RHIME	cytoplasm (by similarity).	extracellular
Q92Q99	KDSA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92QA0	PYRG_RHIME		cytoplasm
Q92QA1	TPIS1_RHIME		cytoplasm
Q92QB2	Y1422_RHIME		inner membrane
Q92QB4	GCH1_RHIME		cytoplasm
Q92QB5	HIS3_RHIME	cytoplasm (by similarity).	cytoplasm
Q92QD0	DEFL_RHIME		cytoplasm
Q92QD3	Y1398_RHIME		inner membrane
Q92QD5	LIPB_RHIME	cytoplasm (by similarity).	cytoplasm
Q92QD8	Y1393_RHIME		inner membrane
Q92QE1	CRCB_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92QE7	Y1383_RHIME		cytoplasm
Q92QE8	Y1382_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92QF1	RL15_RHIME		cytoplasm
Q92QF3	RS5_RHIME		cytoplasm
Q92QF4	RL18_RHIME		cytoplasm
Q92QF6	RS8_RHIME		cytoplasm
Q92QF7	RS14_RHIME		cytoplasm
Q92QF8	RL5_RHIME		cytoplasm
Q92QF9	RL24_RHIME		cytoplasm
Q92QG1	RS17_RHIME		cytoplasm
Q92QG2	RL29_RHIME		cytoplasm
Q92QG3	RL16_RHIME		cytoplasm
Q92QG4	RS3_RHIME		cytoplasm
Q92QG5	RL22_RHIME		cytoplasm
Q92QG7	RL2_RHIME		cytoplasm
Q92QG9	RL4_RHIME		cytoplasm
Q92QH0	RL3_RHIME		cytoplasm
Q92QH1	RS10_RHIME		cytoplasm
Q92QH3	RS7_RHIME		cytoplasm
Q92QH4	RS12_RHIME		cytoplasm
Q92QH6	RPOC_RHIME		cytoplasm
Q92QH7	RPOB_RHIME		cytoplasm
Q92QH8	RL7_RHIME		cytoplasm
Q92QH9	RL10_RHIME		cytoplasm
Q92QI0	RL1_RHIME		cytoplasm
Q92QI1	RL11_RHIME		cytoplasm
Q92QI7	ACPD_RHIME		extracellular
Q92QJ0	PPNK_RHIME	cytoplasm (by similarity).	cytoplasm
Q92QJ9	AROQ_RHIME		cytoplasm
Q92QK2	LFTR_RHIME	cytoplasm (by similarity).	cytoplasm

Q92QK5	GATB_RHIME		cytoplasm
Q92QK7	GATA_RHIME		cytoplasm
Q92QK8	GATC_RHIME		cytoplasm
Q92QL0	RUVX_RHIME	cytoplasm (potential).	cytoplasm
Q92QL5	PYRB_RHIME		cytoplasm
Q92QL7	Y1302_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92QM4	RL33_RHIME		cytoplasm
Q92QM8	DPO41_RHIME	cytoplasm (probable).	cytoplasm
Q92QN0	LOLD_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92QQ2	CLPX_RHIME		cytoplasm
Q92QR5	RS9_RHIME		cytoplasm
Q92QR7	ARGC_RHIME	cytoplasm (probable).	cytoplasm
Q92QT3	IHFA_RHIME		cytoplasm
Q92QT4	FABH_RHIME	cytoplasm (probable).	cytoplasm
Q92QT5	PLSX_RHIME		cytoplasm
Q92QT9	NUSB_RHIME		cytoplasm
Q92QU0	RISB1_RHIME		cytoplasm
Q92QU6	GLYA1_RHIME	cytoplasm (by similarity).	cytoplasm
Q92QW2	PUR5_RHIME	cytoplasm (by similarity).	cytoplasm
Q92QX5	MOAE_RHIME		cytoplasm
Q92QX9	NDK_RHIME	cytoplasm (by similarity).	cytoplasm
Q92QY7	AMPA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92QZ0	PDXA1_RHIME	cytoplasm (probable).	cytoplasm
Q92QZ1	KSGA_RHIME		cytoplasm
Q92QZ2	KGUA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92QZ7	RS6_RHIME		cytoplasm
Q92QZ8	RS18_RHIME		cytoplasm
Q92QZ9	RL9_RHIME		cytoplasm
Q92R07	ALR_RHIME		cytoplasm
Q92R32	NADB_RHIME	cytoplasm (by similarity).	cytoplasm
Q92R33	NADA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92R46	ERA_RHIME		cytoplasm
Q92R47	RNC_RHIME	cytoplasm (by similarity).	cytoplasm
Q92R49	ACPS_RHIME	cytoplasm (by similarity).	cytoplasm
Q92R52	RPOZ_RHIME		cytoplasm
Q92R54	SSRP_RHIME	cytoplasm (potential).	cytoplasm
Q92R55	DAPA1_RHIME	cytoplasm (by similarity).	cytoplasm
Q92R68	Y1043_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92R86	GLNE_RHIME		cytoplasm
Q92R94	Y1011_RHIME		inner membrane
Q92RA4	MSRB1_RHIME		cytoplasm
Q92RB2	Y982_RHIME		inner membrane
Q92RC5	TAM_RHIME	cytoplasm (by similarity).	cytoplasm
Q92RF8	Y917_RHIME		inner membrane
Q92RG0	RNH_RHIME	cytoplasm (potential).	extracellular
Q92RG1	KHSE_RHIME		cytoplasm
Q92RG2	ISPH_RHIME		cytoplasm
Q92RG6	COXZ_RHIME		inner membrane
Q92RH8	PDXH_RHIME		cytoplasm
Q92RI9	EX7S_RHIME	cytoplasm (by similarity).	cytoplasm

Q92RJ1	DXS_RHIME		cytoplasm
Q92RJ3	Y878_RHIME		cytoplasm
Q92RL0	PUR2_RHIME		cytoplasm
Q92RM1	ISPE_RHIME		cytoplasm
Q92RM4	RNH2_RHIME	cytoplasm (potential).	extracellular
Q92RN0	KUP1_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92RP4	MUTL_RHIME		cytoplasm
Q92RP7	LPXK_RHIME		cytoplasm
Q92RR6	HIS1_RHIME	cytoplasm (by similarity).	cytoplasm
Q92RT9	RRMJ_RHIME	cytoplasm (by similarity).	cytoplasm
Q92S14	RHAL_RHIME		cytoplasm
Q92S21	Y616_RHIME		cytoplasm
Q92S22	Y615_RHIME	cytoplasm (potential).	inner membrane
Q92S25	Y612_RHIME		inner membrane
Q92S26	HISX1_RHIME		cytoplasm
Q92S27	MURA_RHIME	cytoplasm (probable).	inner membrane
Q92S49	PNCB_RHIME	cytoplasm (by similarity).	cytoplasm
Q92S72	MSCL_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92S97	APAG_RHIME		inner membrane
Q92S99	OTC_RHIME	cytoplasm (probable).	cytoplasm
Q92SA0	ARGD_RHIME	cytoplasm (probable).	cytoplasm
Q92SA1	PSTB_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92SC4	G6PI_RHIME	cytoplasm (by similarity).	cytoplasm
Q92SC6	PYRE_RHIME		cytoplasm
Q92SC7	PYRC_RHIME		cytoplasm
Q92SF4	RNPA_RHIME		extracellular
Q92SF5	OXAA_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q92SF6	ENGB_RHIME		cytoplasm
Q92SG0	ARGB_RHIME	cytoplasm (probable).	cytoplasm
Q92SG5	DAPD_RHIME	cytoplasm (by similarity).	cytoplasm
Q92SH4	TRUA_RHIME		cytoplasm
Q92SH5	FMT_RHIME		cytoplasm
Q92SH6	DEF_RHIME		cytoplasm
Q92SI3	TRMB_RHIME		cytoplasm
Q92SI5	Y407_RHIME		inner membrane
Q92SJ7	IHFB_RHIME		cytoplasm
Q92SK0	GRPE_RHIME	cytoplasm (probable).	cytoplasm
Q92SK1	HRCA_RHIME		cytoplasm
Q92SK2	RNPH_RHIME		cytoplasm
Q92SK4	HAM1_RHIME		extracellular
Q92SK7	UBIE_RHIME		cytoplasm
Q92SM6	DUT_RHIME		extracellular
Q92SN3	GSHB_RHIME		cytoplasm
Q92SN4	Y337_RHIME		inner membrane
Q92SN8	PYRF_RHIME		cytoplasm
Q92SP2	UPPP_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q92SQ3	GUAA_RHIME		cytoplasm
Q92SR7	EX7L_RHIME	cytoplasm (by similarity).	cytoplasm
Q92ST1	RL20_RHIME		cytoplasm
Q92ST2	RL35_RHIME		cytoplasm
Q92SU3	LEPA_RHIME		cytoplasm
Q92SV4	KCY_RHIME	cytoplasm (by similarity).	cytoplasm
Q92SV5	AROA_RHIME	cytoplasm (probable).	cytoplasm
Q92SV8	FABA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92SW1	RS15_RHIME		cytoplasm
Q92SW2	TRUB_RHIME		cytoplasm
Q92SW3	RBFA_RHIME	cytoplasm (potential).	cytoplasm
Q92SW6	Y236_RHIME		inner membrane
Q92SW9	RECR_RHIME		cytoplasm
Q92SX1	Y231_RHIME		inner membrane
Q92SY7	MSRA1_RHIME		cytoplasm
Q92SZ1	CUTC_RHIME	cytoplasm (probable).	periplasm
Q92T07	DNAJ_RHIME	cytoplasm (probable).	cytoplasm
Q92T25	GPMA_RHIME		cytoplasm
Q92T27	GLK_RHIME	cytoplasm (by similarity).	cytoplasm
Q92T28	MGSA_RHIME		cytoplasm
Q92T47	DEOB_RHIME	cytoplasm (by similarity).	cytoplasm
Q92T48	ADD_RHIME		extracellular
Q92T49	UPP_RHIME		cytoplasm
Q92T50	TYPH_RHIME		cytoplasm
Q92T74	PHK1_RHIME		cytoplasm
Q92TA4	MASZ_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TA7	HSLU_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TA9	HSLV_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TB0	HIS7_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TB1	HIS5_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TB2	HIS4_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TB3	HIS6_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TB4	HIS2_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TB5	COAA_RHIME	cytoplasm (probable).	cytoplasm
Q92TC1	SAHH_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TC8	TRPA_RHIME		cytoplasm
Q92TC9	TRPB_RHIME		cytoplasm
Q92TD0	TRPF_RHIME		cytoplasm
Q92TE7	SECB_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TE9	COAE_RHIME	cytoplasm (by similarity).	cytoplasm
Q92TF0	AROE_RHIME		cytoplasm
Q92TF1	Y002_RHIME	cytoplasm (potential).	cytoplasm
Q92TF2	Y001_RHIME		cytoplasm
Q92TI9	THIC_RHIME		cytoplasm
Q92TP3	PDXA3_RHIME	cytoplasm (probable).	cytoplasm
Q92TT1	3MGH_RHIME		cytoplasm
Q92TW1	ALLA2_RHIME		extracellular
Q92TY6	NADE_RHIME		cytoplasm
Q92TY9	MINC_RHIME		inner membrane
Q92UG4	ALLA1_RHIME		extracellular
Q92UG5	Y5866_RHIME		cytoplasm
Q92UI1	G6PI2_RHIME	cytoplasm (by similarity).	cytoplasm
Q92UK6	ENGA_RHIME		cytoplasm
Q92UV5	FBPC2_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane

Q92UW5	Y5671_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92UW6	XSC_RHIME	cytoplasm (by similarity).	cytoplasm
Q92UX0	TAUB_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92V04	UXAC_RHIME		cytoplasm
Q92V09	KDUI2_RHIME		cytoplasm
Q92V27	Y5606_RHIME		cytoplasm
Q92V44	Y5573_RHIME	periplasm (potential).	inner membrane
Q92V71	PHNC_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92V80	HUTU_RHIME	cytoplasm (by similarity).	cytoplasm
Q92VB9	Y5489_RHIME		cytoplasm
Q92VC1	ALLC_RHIME		extracellular
Q92VC6	ADEC2_RHIME		cytoplasm
Q92VJ2	CYSA2_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92VM6	ARLY2_RHIME	cytoplasm (probable).	cytoplasm
Q92VT5	Y614_RHIME		extracellular
Q92W37	Y5208_RHIME		inner membrane
Q92WB5	UXUA_RHIME		cytoplasm
Q92WJ0	FBPC1_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92WP0	NANA_RHIME	cytoplasm (by similarity).	cytoplasm
Q92WX6	GFA_RHIME		extracellular
Q92X16	PDXA2_RHIME	cytoplasm (probable).	cytoplasm
Q92XH8	DPO42_RHIME	cytoplasm (probable).	cytoplasm
Q92XI9	ATKA_RHIME	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q92XJ0	ATKB_RHIME	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q92XJ1	ATKC_RHIME	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q92XP4	OOXA1_RHIME		cytoplasm
Q92XP5	OOXB1_RHIME		cytoplasm
Q92XS8	GLYA2_RHIME	cytoplasm (by similarity).	cytoplasm
Q92XV9	Y4531_RHIME		periplasm
Q92XW1	CYSA1_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q92Y27	WRBA3_RHIME		cytoplasm
Q92Y45	MSRA3_RHIME		cytoplasm
Q92Y46	MSRB2_RHIME		cytoplasm
Q92Y93	KUP2_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q92YC7	Y4357_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q92YD2	BETB2_RHIME		cytoplasm
Q92YE0	ADEC4_RHIME		cytoplasm
Q92YE2	ADEC3_RHIME		cytoplasm
Q92YG5	ARCA2_RHIME	cytoplasm (potential).	extracellular
Q92YS6	KDGD_RHIME		cytoplasm
Q92YY3	Y4129_RHIME		inner membrane
Q92Z29	NIRK_RHIME	periplasm (by similarity).	periplasm
Q92Z36	NAPA_RHIME	periplasm (by similarity).	periplasm
Q92ZA4	PHK2_RHIME		cytoplasm
Q92ZJ6	DNE22_RHIME	cytoplasm (probable).	cytoplasm
Q92ZK3	NODM1_RHIME	cytoplasm (by similarity).	cytoplasm
Q92ZL2	NIFE_RHIME		cytoplasm
Q92ZQ3	CH104_RHIME	cytoplasm (by similarity).	cytoplasm
Q92ZQ4	CH604_RHIME	cytoplasm (by similarity).	cytoplasm
Q92ZT0	ARCC_RHIME	cytoplasm (potential).	cytoplasm
Q92ZT1	OTCC_RHIME	cytoplasm (probable).	cytoplasm
Q92ZT2	ARCA1_RHIME	cytoplasm (potential).	extracellular
Q92ZW9	AQPZ2_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q930I4	HISX2_RHIME		cytoplasm
Q930J0	HIS83_RHIME		cytoplasm
Q930L2	WRBA2_RHIME		cytoplasm
Q930T2	KDU11_RHIME		cytoplasm
Q930X9	CH103_RHIME	cytoplasm (by similarity).	cytoplasm
Q930Y0	CH603_RHIME	cytoplasm (by similarity).	cytoplasm
Q931D0	SELD_RHIME		cytoplasm
Q931D7	FDHE_RHIME	cytoplasm (by similarity).	cytoplasm
Q932V0	ALR1_SHIDY		cytoplasm
Q933E0	LKTB_PASTR	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q933I3	LKTB_MANGL	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q934G6	TOLB_DICD3	periplasm (potential).	periplasm
Q934J6	PHON_SALTI	periplasm (by similarity).	periplasm
Q934T0	PYRB_PSYT1		cytoplasm
Q934T6	REPA_SALTI		cytoplasm
Q936V7	OTCC_PSEME	cytoplasm (probable).	cytoplasm
Q937H8	FLHD_ENTS2	cytoplasm (by similarity).	cytoplasm
Q937L4	CPNA_COMTE		cytoplasm
Q937L5	CPNB_COMTE		cytoplasm
Q937W7	MASZ_RHILV	cytoplasm (by similarity).	cytoplasm
Q938W0	HIS22_BRAJA	cytoplasm (by similarity).	cytoplasm
Q939I2	NANE_KLEAE		cytoplasm
Q93A90	RBL_NITS4		extracellular
Q93AA7	YBHG_YERPE	periplasm (potential).	inner membrane
Q93AG0	1A1D_RHILV		cytoplasm
Q93CC3	HOSA_ECOLI		cytoplasm
Q93CH6	CUER_SALTY	cytoplasm (probable).	cytoplasm
Q93CP9	NADA_SHIFL	cytoplasm (by similarity).	cytoplasm
Q93E92	RIB2_PHOLE		cytoplasm

Q93ED4	AROA_YERRU	cytoplasm (probable).	cytoplasm
Q93EJ3	ARGJ_HELHP	cytoplasm (probable).	cytoplasm
Q93FE6	KAD_RHIME	cytoplasm (by similarity).	cytoplasm
Q93FG6	LKB2B_PASHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q93FH0	LKB2A_PASHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q93FH2	LKB13_PASHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q93FH3	LKB16_PASHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q93FH6	LKB1B_PASHA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q93FQ6	3MGH_EHRRU		cytoplasm
Q93FR9	GPDA_EHRRU	cytoplasm (probable).	cytoplasm
Q93FU7	CH10_ENTAE	cytoplasm (by similarity).	cytoplasm
Q93GP7	SSB2_SALTY		cytoplasm
Q93GT8	CH60_COLMA	cytoplasm (by similarity).	cytoplasm
Q93GT9	CH10_COLMA	cytoplasm (by similarity).	cytoplasm
Q93GW2	CH60_WOLRE	cytoplasm (by similarity).	cytoplasm
Q93HP9	ALR1_SHISO		cytoplasm
Q93HX3	NAPA_MAGMG	periplasm (by similarity).	periplasm
Q93IM5	GLK_SALTY	cytoplasm (by similarity).	cytoplasm
Q93IM6	YFEO_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q93IN1	BCSB_SALTY	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q93IN2	BCSA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q93JR5	NIFW_RHIET		cytoplasm
Q93JT2	TPMT_PSEST	cytoplasm (by similarity).	cytoplasm
Q93JW0	4OT4_PSEPU		cytoplasm
Q93JY3	GGPS_PSEAG		cytoplasm
Q93K97	ADPP_ECOLI		periplasm
Q93KF3	MSRAB_CAMFE		cytoplasm
Q93KU0	YSCY_YEREN	cytoplasm (by similarity).	cytoplasm
Q93L23	MDCC_BURCE	cytoplasm (by similarity).	cytoplasm
Q93LC7	LUXS_PROMI		extracellular
Q93LP2	RUVB_BURCE		cytoplasm
Q93MH1	CH60_RHOPA	cytoplasm (by similarity).	cytoplasm
Q93MH2	CH10_RHOPA	cytoplasm (by similarity).	cytoplasm
Q93MH4	ASR_SALTI		fimbrium
Q93MH5	ASR_KLEPN		inner membrane
Q93MH6	ASR_ENTCL		inner membrane
Q93MI1	CSRA_PROMI		cytoplasm
Q93MK7	RPOB_WOLPI		cytoplasm
Q93MK8	RPOB_EHRR1		cytoplasm
Q93MK9	RPOB_EHRSE		cytoplasm

Q93MQ9	LEXA_XANCA		cytoplasm
Q93MV4	ATKC_MYXXA	bacterial cell membrane; single-pass membrane protein (probable).	inner membrane
Q93MV5	ATKB_MYXXA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q93N34	CH60_BUCRP	cytoplasm (by similarity).	cytoplasm
Q93N35	CH60_BUCRM	cytoplasm (by similarity).	cytoplasm
Q93NE2	TRUA_MYXXA		cytoplasm
Q93P00	CHEY_YEREN	cytoplasm (by similarity).	cytoplasm
Q93PD4	Y2324_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q93PD8	ARNA_YERPS		cytoplasm
Q93PD9	ARNC_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q93PE0	ARNB_YERPE		cytoplasm
Q93PJ4	URE1_HELHP	cytoplasm (by similarity).	cytoplasm
Q93PJ5	URE23_HELHP	cytoplasm (by similarity).	cytoplasm
Q93PP8	RUBR2_DESDE		cytoplasm
Q93PP9	BFR_DESDE		cytoplasm
Q93PU3	TTGI_PSEPU	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
Q93PU4	TTGH_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q93PU5	TTGG_PSEPU	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
Q93PU6	TTGV_PSEPU		cytoplasm
Q93PU7	TTGW_PSEPU		cytoplasm
Q93Q11	FADA_PSEOL	cytoplasm (by similarity).	cytoplasm
Q93Q12	FADB_PSEOL		cytoplasm
Q93Q64	DLGD_KLEOX	cytoplasm (potential).	cytoplasm
Q93R37	RECA_CAMLA	cytoplasm (by similarity).	cytoplasm
Q93S39	MSRA_OCHAN		cytoplasm
Q93SE0	BTUB2_ECOLI	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q93SM8	Y2729_BURPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q93SX7	Y189_ACISE		inner membrane
Q93T12	TFDF_COMAC		cytoplasm
Q93T81	URE11_BRUAB	cytoplasm (by similarity).	cytoplasm
Q93TE2	CRCB_PSEYM	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q93TF4	SECB_PSEFL	cytoplasm (by similarity).	cytoplasm
Q93TG4	TOLB_BRUME	periplasm (potential).	periplasm
Q93TX3	HUTH_STIAU	cytoplasm (potential).	cytoplasm
Q93U24	CSGA_ECO57		extracellular
Q93UJ2	GMHA_BURPS	cytoplasm (by similarity).	cytoplasm
Q93UZ0	RBLL_CHRVI		cytoplasm
Q981G1	Y9388_RHILO		inner membrane

Q981H7	ALR3_RHILO		cytoplasm
Q981J9	CH605_RHILO	cytoplasm (by similarity).	cytoplasm
Q981K0	CH105_RHILO	cytoplasm (by similarity).	cytoplasm
Q981V0	ARLY2_RHILO	cytoplasm (probable).	cytoplasm
Q981X2	DADA3_RHILO		cytoplasm
Q982F2	NADA2_RHILO	cytoplasm (by similarity).	cytoplasm
Q982G0	PAND2_RHILO		cytoplasm
Q982V5	CLPX_RHILO		cytoplasm
Q982V6	CLPP2_RHILO	cytoplasm (by similarity).	cytoplasm
Q982W9	RS9_RHILO		cytoplasm
Q982X3	ARGC_RHILO	cytoplasm (probable).	cytoplasm
Q982Z7	IHFA_RHILO		cytoplasm
Q982Z8	FABH_RHILO	cytoplasm (probable).	cytoplasm
Q982Z9	PLSX_RHILO		cytoplasm
Q983A3	HPPA_RHILO	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q983A9	NUSB_RHILO		cytoplasm
Q983B0	RISB2_RHILO		cytoplasm
Q983B3	NRDR_RHILO		cytoplasm
Q983B6	GLYA1_RHILO	cytoplasm (by similarity).	cytoplasm
Q983E4	ATE_RHILO	cytoplasm (potential).	cytoplasm
Q983F8	RRMJ_RHILO	cytoplasm (by similarity).	cytoplasm
Q983H8	HUTU_RHILO	cytoplasm (by similarity).	cytoplasm
Q983I0	HUTH_RHILO	cytoplasm (potential).	cytoplasm
Q983I1	HUTI_RHILO	cytoplasm (potential).	cytoplasm
Q983J4	HGD_RHILO		cytoplasm
Q983J7	TDH_RHILO	cytoplasm (by similarity).	cytoplasm
Q983K4	QUEF_RHILO	cytoplasm (probable).	cytoplasm
Q983L2	MUTL_RHILO		cytoplasm
Q983N4	SYI_RHILO	cytoplasm (by similarity).	cytoplasm
Q983S3	CH104_RHILO	cytoplasm (by similarity).	cytoplasm
Q983S4	CH604_RHILO	cytoplasm (by similarity).	cytoplasm
Q983T9	Y8179_RHILO		periplasm
Q983U5	FUMC_RHILO	cytoplasm (by similarity).	cytoplasm
Q984K6	PUR5_RHILO	cytoplasm (by similarity).	cytoplasm
Q984N8	NDK_RHILO	cytoplasm (by similarity).	cytoplasm
Q984P0	MOAE_RHILO		cytoplasm
Q984S1	AMPA_RHILO	cytoplasm (by similarity).	cytoplasm
Q984S6	PDXA_RHILO	cytoplasm (probable).	cytoplasm
Q984S7	KSGA_RHILO		cytoplasm
Q984S9	KGUA_RHILO	cytoplasm (by similarity).	cytoplasm
Q984T3	ACP_RHILO	cytoplasm (by similarity).	cytoplasm
Q984T8	RS6_RHILO		cytoplasm
Q984T9	RS18_RHILO		cytoplasm
Q984U0	RL9_RHILO		cytoplasm
Q984U2	Y7841_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q984W0	PSD_RHILO		cytoplasm
Q985A3	RECO_RHILO		cytoplasm
Q985A5	ERA_RHILO		cytoplasm
Q985A6	RNC_RHILO	cytoplasm (by similarity).	cytoplasm
Q985A8	ACPS_RHILO	cytoplasm (by similarity).	cytoplasm
Q985B1	PYRE2_RHILO		cytoplasm
Q985B3	RPOZ_RHILO		cytoplasm

Q985B9	SSRP_RHILO	cytoplasm (potential).	cytoplasm
Q985D2	Y7730_RHILO		inner membrane
Q985F6	GLNE_RHILO		cytoplasm
Q985G2	CCME_RHILO	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q985M2	BETI_RHILO		cytoplasm
Q985M5	BETA_RHILO		cytoplasm
Q985M6	BETB_RHILO		cytoplasm
Q985P2	GLGA_RHILO		cytoplasm
Q985P3	GLGC_RHILO		cytoplasm
Q985P4	GLGB_RHILO		extracellular
Q985V6	Y7511_RHILO		inner membrane
Q985W1	RNH_RHILO	cytoplasm (potential).	extracellular
Q985W2	KHSE_RHILO		cytoplasm
Q985W3	ISPH_RHILO		cytoplasm
Q985W8	COXZ_RHILO		inner membrane
Q985X9	Y7480_RHILO		cytoplasm
Q985Y3	DXS_RHILO		cytoplasm
Q985Y6	EX7S_RHILO	cytoplasm (by similarity).	cytoplasm
Q986A0	PDXH_RHILO		cytoplasm
Q986A5	PUR2_RHILO		cytoplasm
Q986C6	ISPE_RHILO		cytoplasm
Q986D0	RNH2_RHILO	cytoplasm (potential).	extracellular
Q986D9	ENGA_RHILO		cytoplasm
Q986N2	RISB1_RHILO		cytoplasm
Q986N6	TPIS2_RHILO		cytoplasm
Q986Q6	MURA2_RHILO	cytoplasm (probable).	inner membrane
Q986R8	DCTA2_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q987C6	HISX3_RHILO		cytoplasm
Q987C8	HIS81_RHILO		cytoplasm
Q987N5	PANE_RHILO	cytoplasm (potential).	cytoplasm
Q987R3	GUAA_RHILO		cytoplasm
Q987S8	HIS1_RHILO	cytoplasm (by similarity).	cytoplasm
Q987S9	HISZ_RHILO	cytoplasm (by similarity).	cytoplasm
Q987V3	EX7L_RHILO	cytoplasm (by similarity).	cytoplasm
Q987X5	UXUA_RHILO		cytoplasm
Q988H0	NADE_RHILO		cytoplasm
Q988I1	ARCA_RHILO	cytoplasm (potential).	extracellular
Q988P7	HISX2_RHILO		cytoplasm
Q988V7	PHK_RHILO		cytoplasm
Q989A3	6PGL_RHILO		extracellular
Q989E5	MURA1_RHILO	cytoplasm (probable).	inner membrane
Q989E7	HISX1_RHILO		cytoplasm
Q989E8	Y6455_RHILO		inner membrane
Q989F1	YSE2_RHILO	cytoplasm (potential).	inner membrane
Q989F2	Y6451_RHILO		cytoplasm
Q989G8	DGT1B_RHILO		cytoplasm
Q989L5	IDI2_RHILO	cytoplasm (by similarity).	cytoplasm
Q989S7	PROB2_RHILO	cytoplasm (by similarity).	cytoplasm
Q98A05	ALR1_RHILO		cytoplasm
Q98A07	DCHS_RHILO		cytoplasm
Q98A73	METE_RHILO		cytoplasm

Q98A81	GLYA2_RHILO	cytoplasm (by similarity).	cytoplasm
Q98AM7	1A1D_RHILO		cytoplasm
Q98AN6	ADC1_RHILO		cytoplasm
Q98AN7	PYRE1_RHILO		cytoplasm
Q98AP3	NIFN_RHILO		periplasm
Q98AP4	NIFE_RHILO		cytoplasm
Q98AP7	NIFH_RHILO		cytoplasm
Q98AT3	NIFW_RHILO		cytoplasm
Q98AV2	DCTA1_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98AV7	NADA1_RHILO	cytoplasm (by similarity).	inner membrane
Q98AV8	NADB_RHILO	cytoplasm (by similarity).	cytoplasm
Q98AX8	CH103_RHILO	cytoplasm (by similarity).	cytoplasm
Q98AX9	CH603_RHILO	cytoplasm (by similarity).	cytoplasm
Q98AZ3	THIC_RHILO		cytoplasm
Q98AZ6	THIG_RHILO	cytoplasm (by similarity).	cytoplasm
Q98B00	HIS83_RHILO		cytoplasm
Q98B65	RHAL_RHILO		cytoplasm
Q98B75	DADA2_RHILO		cytoplasm
Q98B79	MSCL3_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98BB4	APAG_RHILO		inner membrane
Q98BB6	OTC_RHILO	cytoplasm (probable).	cytoplasm
Q98BB7	ARGD_RHILO	cytoplasm (probable).	cytoplasm
Q98BG5	DEOB_RHILO	cytoplasm (by similarity).	cytoplasm
Q98BG6	FPG_RHILO		cytoplasm
Q98BG8	RS20_RHILO		cytoplasm
Q98BG9	DNAA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98BH1	RECF_RHILO	cytoplasm (by similarity).	cytoplasm
Q98BH8	FABA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98BI4	RS15_RHILO		cytoplasm
Q98BI6	TRUB_RHILO		cytoplasm
Q98BI7	RBFA_RHILO	cytoplasm (potential).	cytoplasm
Q98BJ1	Y5550_RHILO		inner membrane
Q98BJ3	TRMB_RHILO		cytoplasm
Q98BJ6	LNT_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98BK1	Y5536_RHILO		inner membrane
Q98BM4	RECR_RHILO		cytoplasm
Q98BM7	Y5504_RHILO		inner membrane
Q98BV5	G6PI_RHILO	cytoplasm (by similarity).	cytoplasm
Q98BZ8	ILVD1_RHILO		cytoplasm
Q98C03	DHAA_RHILO		cytoplasm
Q98C10	DUT_RHILO		extracellular
Q98C21	MUTS_RHILO		cytoplasm
Q98C27	GLND_RHILO		cytoplasm
Q98CB9	SSUD1_RHILO		cytoplasm
Q98CC1	AROA_RHILO	cytoplasm (probable).	cytoplasm
Q98CC2	KCY_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CH8	Y5144_RHILO		inner membrane
Q98CI7	Y5133_RHILO		cytoplasm
Q98CJ0	ALLA_RHILO		extracellular
Q98CL7	PPCK_RHILO	cytoplasm (by similarity).	cytoplasm

Q98CM3	SAHH_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CN6	TRPA_RHILO		cytoplasm
Q98CN7	TRPB_RHILO		cytoplasm
Q98CN8	TRPF_RHILO		cytoplasm
Q98CP7	RL35_RHILO		cytoplasm
Q98CP8	RL20_RHILO		cytoplasm
Q98CR8	XYLA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CS9	COAA_RHILO	cytoplasm (probable).	cytoplasm
Q98CT0	HIS2_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CT1	HIS6_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CT3	HIS4_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CT5	HIS5_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CT7	HIS7_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CT8	HSLV_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CU0	Y5005_RHILO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q98CU1	HSLU_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CY3	URE3_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CY6	URE2_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CY9	URE1_RHILO	cytoplasm (by similarity).	cytoplasm
Q98CZ2	UREE_RHILO	cytoplasm (by similarity).	cytoplasm
Q98D10	XYLD_RHILO		cytoplasm
Q98D15	NORM_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98D24	PNCB_RHILO	cytoplasm (by similarity).	cytoplasm
Q98D52	DEF_RHILO		cytoplasm
Q98D53	FMT_RHILO		cytoplasm
Q98D55	TRUA_RHILO		cytoplasm
Q98D61	DAPD_RHILO	cytoplasm (by similarity).	cytoplasm
Q98D72	PH4H_RHILO		cytoplasm
Q98D76	ARGB_RHILO	cytoplasm (probable).	cytoplasm
Q98D85	ENGB_RHILO		cytoplasm
Q98D88	OXAA_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98D89	RNPA_RHILO		extracellular
Q98DD1	DNAK_RHILO		cytoplasm
Q98DD2	DNAJ_RHILO	cytoplasm (probable).	cytoplasm
Q98DD5	PYRF_RHILO		cytoplasm
Q98DE8	GSHB_RHILO		cytoplasm
Q98DG5	MSCL2_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98DH7	MSCL1_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98DI0	YAG6_RHILO		cytoplasm
Q98DK4	MASZ_RHILO	cytoplasm (by similarity).	cytoplasm
Q98DM0	GPMA_RHILO		cytoplasm
Q98DM2	GLK_RHILO	cytoplasm (by similarity).	cytoplasm
Q98DM7	UPPP1_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98DM8	Y4633_RHILO		inner membrane

Q98DM9	Y4632_RHILO		cytoplasm
Q98DN4	HAM1_RHILO		extracellular
Q98DN6	RNPH_RHILO		cytoplasm
Q98DN7	HRCA_RHILO		cytoplasm
Q98DR6	3MGH_RHILO		cytoplasm
Q98DT4	SSUD2_RHILO		cytoplasm
Q98DU8	SECB_RHILO	cytoplasm (by similarity).	cytoplasm
Q98DV1	LEPA_RHILO		cytoplasm
Q98DV6	MSRA2_RHILO		cytoplasm
Q98DW6	TAUB_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98DY2	COAE_RHILO	cytoplasm (by similarity).	cytoplasm
Q98DY3	AROE_RHILO		cytoplasm
Q98DY4	Y4491_RHILO	cytoplasm (potential).	inner membrane
Q98DY5	Y4488_RHILO		cytoplasm
Q98DY6	DCUP_RHILO	cytoplasm (probable).	cytoplasm
Q98DZ1	GIDA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98DZ2	GIDB_RHILO		cytoplasm
Q98E34	DNAE2_RHILO	cytoplasm (probable).	cytoplasm
Q98E51	LEUD_RHILO		cytoplasm
Q98E57	LEU3_RHILO	cytoplasm (by similarity).	cytoplasm
Q98E66	Y4386_RHILO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q98E70	RS16_RHILO		cytoplasm
Q98E81	ASSY_RHILO	cytoplasm (probable).	cytoplasm
Q98E92	PHS_RHILO		cytoplasm
Q98EA4	CCMA_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98EB3	ISPZ_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98EB6	DAPF_RHILO	cytoplasm (by similarity).	cytoplasm
Q98EC1	OMP19_RHILO	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q98EC4	MDH_RHILO		cytoplasm
Q98EC5	SUCC_RHILO		cytoplasm
Q98ED9	XERC_RHILO	cytoplasm (by similarity).	nucleoid
Q98EE0	RIMM_RHILO	cytoplasm (potential).	cytoplasm
Q98EE1	TRMD_RHILO	cytoplasm (potential).	cytoplasm
Q98EF1	LEU2_RHILO		cytoplasm
Q98EI7	HEM3_RHILO		cytoplasm
Q98EP2	Y4154_RHILO		inner membrane
Q98ER2	IPYR_RHILO	cytoplasm (by similarity).	cytoplasm
Q98ES7	PUR9_RHILO		cytoplasm
Q98ET0	HTPX_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q98ET8	ACSA_RHILO		cytoplasm
Q98EV0	TAL_RHILO	cytoplasm (probable).	cytoplasm
Q98EV6	ATPA_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98EV7	ATPG_RHILO		inner membrane

Q98EV9	ATPE_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98EW1	UXAC_RHILO		cytoplasm
Q98EZ0	RL27_RHILO		cytoplasm
Q98EZ4	PROB1_RHILO	cytoplasm (by similarity).	cytoplasm
Q98EZ5	PROA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98F00	Y4004_RHILO		inner membrane
Q98F04	NUDH_RHILO		cytoplasm
Q98F05	ALR2_RHILO		cytoplasm
Q98F08	DADA1_RHILO		cytoplasm
Q98F28	Y3968_RHILO		inner membrane
Q98F44	Y3945_RHILO		inner membrane
Q98F50	RL31_RHILO		cytoplasm
Q98F59	SUHB_RHILO		cytoplasm
Q98F72	RUVC_RHILO		cytoplasm
Q98F75	RUVA_RHILO		cytoplasm
Q98F76	RUVB_RHILO		cytoplasm
Q98F84	TOLB_RHILO	periplasm (potential).	periplasm
Q98F85	OMP16_RHILO	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q98F87	TILS_RHILO	cytoplasm (probable).	cytoplasm
Q98F91	GLMM_RHILO		cytoplasm
Q98F97	PURA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98FF1	Y3804_RHILO		cytoplasm
Q98FG0	ISPG_RHILO		cytoplasm
Q98FG8	MTGA_RHILO	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q98FG9	RL32_RHILO		cytoplasm
Q98FJ1	PGK_RHILO	cytoplasm (potential).	cytoplasm
Q98FL2	PSTS_RHILO	periplasm (potential).	periplasm
Q98FL3	PSTC_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98FL4	PSTA_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98FL5	PSTB_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98FS9	ADC3_RHILO		cytoplasm
Q98FX7	ACCA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98FX8	XERD_RHILO	cytoplasm (by similarity).	nucleoid
Q98FY0	AROK_RHILO	cytoplasm (probable).	cytoplasm
Q98FY1	AROB_RHILO	cytoplasm (probable).	cytoplasm
Q98FZ7	RL28_RHILO		cytoplasm
Q98G09	METX_RHILO	cytoplasm (by similarity).	cytoplasm
Q98G10	HIS82_RHILO		cytoplasm
Q98G36	ARLY1_RHILO	cytoplasm (probable).	cytoplasm
Q98G43	FBPC_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98G72	ARGJ_RHILO	cytoplasm (probable).	cytoplasm
Q98G87	UBIG_RHILO		cytoplasm
Q98G94	HEMK_RHILO		cytoplasm

Q98G96	CLPB_RHILO	cytoplasm (probable).	cytoplasm
Q98GC6	Y3387_RHILO		inner membrane
Q98GF5	PHNC_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98GG1	FOSX_RHILO	cytoplasm (by similarity).	cytoplasm
Q98GP9	CYSH_RHILO	cytoplasm (by similarity).	cytoplasm
Q98GQ5	GRPE_RHILO	cytoplasm (probable).	cytoplasm
Q98GR6	Y8638_RHILO		inner membrane
Q98GR7	Y3206_RHILO		inner membrane
Q98GS0	IHFB_RHILO		cytoplasm
Q98GV1	UBIE_RHILO		cytoplasm
Q98GV2	ADD_RHILO		extracellular
Q98GV3	UPP_RHILO		cytoplasm
Q98GV5	TYPH_RHILO		cytoplasm
Q98GW1	SLYX_RHILO		inner membrane
Q98GX5	ATKA_RHILO	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q98GX6	ATKB_RHILO	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q98GX7	ATKC_RHILO	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q98GY4	RS21_RHILO		cytoplasm
Q98H33	SP39_RHILO	periplasm (probable).	periplasm
Q98H61	HEMH_RHILO	cytoplasm (by similarity).	extracellular
Q98H63	OMP10_RHILO	bacterial cell outer membrane; lipid-anchor (by similarity).	outer membrane
Q98H64	HSS_RHILO		cytoplasm
Q98H78	Y2992_RHILO		inner membrane
Q98HB7	FLBT_RHILO		cytoplasm
Q98HD3	FLGH_RHILO	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q98HD8	FLIE_RHILO		flagellar
Q98HE3	FLGF_RHILO		flagellar
Q98HL0	TRMU_RHILO	cytoplasm (by similarity).	cytoplasm
Q98HN3	LEU1_RHILO		cytoplasm
Q98HQ4	FTHS_RHILO		cytoplasm
Q98HR4	KITH_RHILO	cytoplasm (potential).	cytoplasm
Q98HS2	CLPS2_RHILO		cytoplasm
Q98HV6	PTH_RHILO	cytoplasm (by similarity).	cytoplasm
Q98HV7	RL25_RHILO		cytoplasm
Q98HW3	KPRS_RHILO	cytoplasm (by similarity).	cytoplasm
Q98HW8	LGT_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98I01	UVRB_RHILO	cytoplasm (by similarity).	cytoplasm
Q98I23	PUR72_RHILO		cytoplasm
Q98I49	GREA_RHILO		cytoplasm
Q98I52	HLDD_RHILO		cytoplasm
Q98I54	HLDE_RHILO		cytoplasm
Q98I87	CARB_RHILO		cytoplasm

Q98I99	MNTH_RHILO	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q98IA7	CARA_RHILO		cytoplasm
Q98IF8	ACPD_RHILO		extracellular
Q98IG7	Y2411_RHILO		inner membrane
Q98IH6	ADC2_RHILO		cytoplasm
Q98IH9	CH602_RHILO	cytoplasm (by similarity).	cytoplasm
Q98II0	CH102_RHILO	cytoplasm (by similarity).	cytoplasm
Q98IV4	CH101_RHILO	cytoplasm (by similarity).	cytoplasm
Q98IV5	CH601_RHILO	cytoplasm (by similarity).	cytoplasm
Q98IV9	Y2225_RHILO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q98J52	Y2098_RHILO		cytoplasm
Q98JB6	HTPG_RHILO	cytoplasm (by similarity).	cytoplasm
Q98JG8	MINC_RHILO		inner membrane
Q98JM4	ENGC_RHILO		cytoplasm
Q98JM5	DPO42_RHILO	cytoplasm (probable).	cytoplasm
Q98JS0	DDLA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98JV5	MSRA1_RHILO		cytoplasm
Q98K23	CYSA_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98K73	TAM_RHILO	cytoplasm (by similarity).	cytoplasm
Q98KA5	MRAW_RHILO		cytoplasm
Q98KA8	MURE_RHILO	cytoplasm (probable).	cytoplasm
Q98KB0	MRAY_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98KB1	MURD_RHILO	cytoplasm (by similarity).	cytoplasm
Q98KB3	MURG_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98KB4	MURC_RHILO	cytoplasm (probable).	cytoplasm
Q98KB5	MURB_RHILO	cytoplasm (probable).	inner membrane
Q98KB6	DDLB_RHILO	cytoplasm (by similarity).	cytoplasm
Q98KC1	Y1543_RHILO	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q98KD0	PRMA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98KH9	TYSY_RHILO	cytoplasm (by similarity).	cytoplasm
Q98KI3	MODC_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98KJ4	MIAA_RHILO		cytoplasm
Q98KK0	BODG_RHILO	cytoplasm (by similarity).	cytoplasm
Q98KL6	PDXJ_RHILO		cytoplasm
Q98KL7	KUP2_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98KL8	KUP1_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98KM7	ILVC_RHILO		cytoplasm
Q98KN6	Y1393_RHILO		cytoplasm
Q98KN9	COBT_RHILO		cytoplasm

Q98KP0	COBS_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98KP1	COBB_RHILO		cytoplasm
Q98KS7	LOLD_RHILO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q98KT0	LIPB_RHILO	cytoplasm (by similarity).	cytoplasm
Q98KT9	MTKA_RHILO		cytoplasm
Q98L33	COBQ_RHILO		cytoplasm
Q98L53	ACPXL_RHILO	cytoplasm (by similarity).	cytoplasm
Q98L96	CUTC_RHILO	cytoplasm (probable).	periplasm
Q98LB1	SYV_RHILO	cytoplasm (by similarity).	cytoplasm
Q98LB3	ILVD2_RHILO		cytoplasm
Q98LB9	DGT1A_RHILO		cytoplasm
Q98LC5	TATA_RHILO	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q98LC9	SURE_RHILO	cytoplasm (potential).	periplasm
Q98LE1	GID_RHILO	cytoplasm (by similarity).	cytoplasm
Q98LE8	TIG_RHILO		cytoplasm
Q98LQ5	HIS3_RHILO	cytoplasm (by similarity).	cytoplasm
Q98LQ6	GCH1_RHILO		cytoplasm
Q98LQ9	Y918_RHILO		inner membrane
Q98LT6	GCSP_RHILO		cytoplasm
Q98LT7	GCSH_RHILO		cytoplasm
Q98LU3	Y875_RHILO		inner membrane
Q98LU4	GFA_RHILO		extracellular
Q98LV1	DPO41_RHILO	cytoplasm (probable).	cytoplasm
Q98LW4	RL33_RHILO		cytoplasm
Q98LX2	GLMU_RHILO	cytoplasm (by similarity).	cytoplasm
Q98LX5	GLMS_RHILO	cytoplasm (by similarity).	cytoplasm
Q98M18	Y776_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98M36	UVRA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98M38	CLPP1_RHILO	cytoplasm (by similarity).	cytoplasm
Q98M41	SSB_RHILO		cytoplasm
Q98M51	COAD_RHILO	cytoplasm (by similarity).	cytoplasm
Q98M55	QUEA_RHILO	cytoplasm (potential).	cytoplasm
Q98M57	TGT_RHILO		cytoplasm
Q98M73	GLPK_RHILO		cytoplasm
Q98M84	Y688_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98M86	PYRB_RHILO		cytoplasm
Q98M90	RUVX_RHILO	cytoplasm (potential).	cytoplasm
Q98M91	Y680_RHILO		extracellular
Q98M93	GATC_RHILO		cytoplasm
Q98M95	GATA_RHILO		cytoplasm
Q98MA3	CLPS1_RHILO		cytoplasm
Q98MB2	RS2_RHILO		cytoplasm
Q98MB7	PYRH_RHILO		cytoplasm
Q98MB8	RRF_RHILO	cytoplasm (by similarity).	cytoplasm
Q98MB9	UPPS_RHILO		inner membrane

Q98MC1	Y638_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q98MC4	LPXD_RHILO		cytoplasm
Q98MC5	FABZ_RHILO	cytoplasm (by similarity).	cytoplasm
Q98MC6	LPXA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98MD2	LEXA_RHILO		cytoplasm
Q98ME2	MOAC_RHILO		cytoplasm
Q98ME3	TRPC_RHILO		cytoplasm
Q98ME4	TRPD_RHILO		cytoplasm
Q98ME7	TPIS1_RHILO		cytoplasm
Q98MF0	PYRG_RHILO		cytoplasm
Q98MK2	MOBA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98MK6	MOAA_RHILO		cytoplasm
Q98ML8	GPH_RHILO		cytoplasm
Q98ML9	RPIA_RHILO		cytoplasm
Q98MV4	KTHY_RHILO		cytoplasm
Q98MX2	HFQ_RHILO		cytoplasm
Q98MX9	ISPDF_RHILO		cytoplasm
Q98MY2	LIPA_RHILO	cytoplasm (potential).	cytoplasm
Q98MZ3	ENO_RHILO	cytoplasm (by similarity).	extracellular
Q98MZ6	KDSA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98N18	GLNB_RHILO		cytoplasm
Q98N26	CRCB_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98N33	RPOA_RHILO		cytoplasm
Q98N34	RS11_RHILO		cytoplasm
Q98N35	RS13_RHILO		cytoplasm
Q98N36	KAD_RHILO	cytoplasm (by similarity).	cytoplasm
Q98N38	RL15_RHILO		cytoplasm
Q98N40	RS5_RHILO		cytoplasm
Q98N41	RL18_RHILO		cytoplasm
Q98N43	RS8_RHILO		cytoplasm
Q98N44	RS14_RHILO		cytoplasm
Q98N45	RL5_RHILO		cytoplasm
Q98N46	RL24_RHILO		cytoplasm
Q98N48	RS17_RHILO		cytoplasm
Q98N49	RL29_RHILO		cytoplasm
Q98N50	RL16_RHILO		cytoplasm
Q98N51	RS3_RHILO		cytoplasm
Q98N52	RL22_RHILO		cytoplasm
Q98N53	RS19_RHILO		cytoplasm
Q98N54	RL2_RHILO		cytoplasm
Q98N56	RL4_RHILO		cytoplasm
Q98N57	RL3_RHILO		cytoplasm
Q98N58	RS10_RHILO		cytoplasm
Q98N60	RS7_RHILO		cytoplasm
Q98N61	RS12_RHILO		cytoplasm
Q98N65	RPOC_RHILO		cytoplasm
Q98N66	RPOB_RHILO		cytoplasm
Q98N67	RL7_RHILO		cytoplasm
Q98N68	RL10_RHILO		cytoplasm
Q98N69	RL1_RHILO		cytoplasm
Q98N70	RL11_RHILO		cytoplasm
Q98NA6	PPNK_RHILO	cytoplasm (by similarity).	cytoplasm

Q98NB7	GLSA_RHILO		extracellular
Q98NC1	AROQ_RHILO		cytoplasm
Q98NC4	LFTR_RHILO	cytoplasm (by similarity).	cytoplasm
Q98NC9	PANB_RHILO		cytoplasm
Q98ND0	PANC_RHILO		cytoplasm
Q98ND3	GATB_RHILO		cytoplasm
Q98NF9	ADEC_RHILO		cytoplasm
Q98NH3	XGPT_RHILO	cytoplasm (by similarity).	cytoplasm
Q98NJ1	UPPP2_RHILO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q98NM8	PUR71_RHILO		cytoplasm
Q98NN0	PURQ_RHILO	cytoplasm (by similarity).	cytoplasm
Q98NN7	PURL_RHILO	cytoplasm (by similarity).	cytoplasm
Q98NP5	RS4_RHILO		cytoplasm
Q98NP9	MURI_RHILO		cytoplasm
Q98NQ6	RECA_RHILO	cytoplasm (by similarity).	cytoplasm
Q98NS5	SYU_RHILO	cytoplasm (by similarity).	cytoplasm
Q98NW7	DPO43_RHILO	cytoplasm (probable).	cytoplasm
Q98P28	Y9637_RHILO		inner membrane
Q98PA0	Y9551_RHILO		cytoplasm
Q99191	RFBX_SALTI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q99342	YFI3_ECOLI		extracellular
Q99390	YPT2_ECOLI		cytoplasm
Q9A224	SECB_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A228	HIS2_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A229	HIS6_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A230	HIS4_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A231	HIS5_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A232	HIS7_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A238	HSLU_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A239	HSLV_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A246	FABA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A253	DUT_CAUCR		extracellular
Q9A258	UBIE_CAUCR		cytoplasm
Q9A259	FPG_CAUCR		cytoplasm
Q9A280	DAPF_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A288	ISPZ_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9A298	CCMA_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9A2A3	Y3663_CAUCR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9A2B1	MDH_CAUCR		cytoplasm
Q9A2B4	RIMM_CAUCR	cytoplasm (potential).	cytoplasm
Q9A2C5	HLDE_CAUCR		cytoplasm
Q9A2E3	Y3622_CAUCR		inner membrane
Q9A2F1	TAL_CAUCR	cytoplasm (probable).	cytoplasm
Q9A2G0	UPPP_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q9A2H2	AROA_CAUCR	cytoplasm (probable).	cytoplasm
Q9A2H3	KCY_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A2H5	IHFB_CAUCR		cytoplasm
Q9A2H6	MSCL_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9A2I0	ACSA_CAUCR		cytoplasm
Q9A2L1	DAPB_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A2P5	HIS1_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A2S6	NPD_CAUCR	cytoplasm (probable).	cytoplasm
Q9A2V7	ATPA_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9A2V8	ATPG_CAUCR		inner membrane
Q9A2W1	ATPE_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9A2W6	NUDH_CAUCR		cytoplasm
Q9A2X3	Y3433_CAUCR		inner membrane
Q9A2X6	PROA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A311	Y3395_CAUCR		inner membrane
Q9A321	Y3385_CAUCR		inner membrane
Q9A341	RNH_CAUCR	cytoplasm (potential).	extracellular
Q9A342	KHSE_CAUCR		cytoplasm
Q9A345	ISPH_CAUCR		cytoplasm
Q9A352	GCSH_CAUCR		cytoplasm
Q9A353	GCSPA_CAUCR		cytoplasm
Q9A354	GCSPB_CAUCR		cytoplasm
Q9A385	Y3319_CAUCR		inner membrane
Q9A3A7	RS21_CAUCR		cytoplasm
Q9A3C8	Y3276_CAUCR		inner membrane
Q9A3C9	RL31_CAUCR		cytoplasm
Q9A3D5	SUHB_CAUCR		cytoplasm
Q9A3E9	Y3255_CAUCR		inner membrane
Q9A3F5	PGK_CAUCR	cytoplasm (potential).	cytoplasm
Q9A3G1	Y3243_CAUCR		inner membrane
Q9A3G2	PUR72_CAUCR		cytoplasm
Q9A3G6	RUVC_CAUCR		cytoplasm
Q9A3G7	RUVA_CAUCR		cytoplasm
Q9A3G8	RUVB_CAUCR		cytoplasm
Q9A3H4	TOLB_CAUCR	periplasm (potential).	periplasm
Q9A3H7	TILS_CAUCR	cytoplasm (probable).	cytoplasm
Q9A3I3	YX21_CAUCR		cytoplasm
Q9A3J1	IMUA_CAUCR		cytoplasm
Q9A3J2	IMUB_CAUCR		cytoplasm
Q9A3J3	DNAE2_CAUCR	cytoplasm (probable).	cytoplasm
Q9A3K2	RS12_CAUCR		cytoplasm
Q9A3K3	RS7_CAUCR		cytoplasm
Q9A3M3	ADD_CAUCR		extracellular
Q9A3N5	Y3167_CAUCR		cytoplasm
Q9A3P8	AROC_CAUCR		cytoplasm
Q9A3U9	PURA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A3Y4	ARGJ_CAUCR	cytoplasm (probable).	cytoplasm
Q9A434	AROB_CAUCR	cytoplasm (probable).	cytoplasm
Q9A435	AROK_CAUCR	cytoplasm (probable).	cytoplasm
Q9A437	XERD_CAUCR	cytoplasm (by similarity).	nucleoid

Q9A448	ACCA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A478	Y2965_CAUCR		cytoplasm
Q9A4C3	NADB_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A4C4	NADA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A4D0	Y2906_CAUCR		inner membrane
Q9A4D6	CARB_CAUCR		cytoplasm
Q9A4F3	UVRC_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A4I3	GREA_CAUCR		cytoplasm
Q9A4J7	CARA_CAUCR		cytoplasm
Q9A4N2	Y2798_CAUCR		cytoplasm
Q9A4N5	Y2795_CAUCR		cytoplasm
Q9A4R9	CCME_CAUCR	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q9A4S7	GLNE_CAUCR		cytoplasm
Q9A4T2	Y2748_CAUCR		cytoplasm
Q9A4T3	Y2747_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9A4Z6	SFSA_CAUCR		cytoplasm
Q9A515	QUEF_CAUCR	cytoplasm (probable).	cytoplasm
Q9A545	Y2620_CAUCR		cytoplasm
Q9A566	GID_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A594	MRAZ_CAUCR		inner membrane
Q9A595	MURE_CAUCR	cytoplasm (probable).	cytoplasm
Q9A597	MURD_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A5A1	MURG_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9A5A6	MURC_CAUCR	cytoplasm (probable).	cytoplasm
Q9A5A7	MURB_CAUCR	cytoplasm (probable).	inner membrane
Q9A5A9	DDL_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A5B6	HIS82_CAUCR		cytoplasm
Q9A5B8	HGD_CAUCR		cytoplasm
Q9A5D9	RS4_CAUCR		cytoplasm
Q9A5E1	HTPX_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9A5F0	PURL_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A5F3	PURQ_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A5F9	PUR71_CAUCR		cytoplasm
Q9A5I0	CLPS_CAUCR		cytoplasm
Q9A5I1	DPO4_CAUCR	cytoplasm (probable).	cytoplasm
Q9A5I8	RL33_CAUCR		cytoplasm
Q9A5K1	Y2446_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9A5K4	PYRB_CAUCR		cytoplasm
Q9A5K8	RUVX_CAUCR	cytoplasm (potential).	cytoplasm
Q9A5K9	GATC_CAUCR		cytoplasm
Q9A5L0	GATA_CAUCR		cytoplasm
Q9A5L1	GATB_CAUCR		cytoplasm
Q9A5S0	Y2377_CAUCR		inner membrane
Q9A5T7	CUTC_CAUCR	cytoplasm (probable).	periplasm
Q9A5U7	MURA_CAUCR	cytoplasm (probable).	inner membrane

Q9A5V1	HISX_CAUCR		cytoplasm
Q9A5V2	Y2345_CAUCR		inner membrane
Q9A5V5	Y2342_CAUCR	cytoplasm (potential).	inner membrane
Q9A5V7	Y2340_CAUCR		cytoplasm
Q9A5Z2	GPH_CAUCR		cytoplasm
Q9A5Z3	GLMU_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A5Z4	RPIA_CAUCR		cytoplasm
Q9A602	PAND_CAUCR		cytoplasm
Q9A627	UPP_CAUCR		cytoplasm
Q9A634	GPMA_CAUCR		cytoplasm
Q9A649	EX7L_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A652	ARGD_CAUCR	cytoplasm (probable).	cytoplasm
Q9A653	OTC_CAUCR	cytoplasm (probable).	cytoplasm
Q9A655	APAG_CAUCR		inner membrane
Q9A671	HIS81_CAUCR		cytoplasm
Q9A683	ARLY_CAUCR	cytoplasm (probable).	cytoplasm
Q9A6B1	MSRB_CAUCR		cytoplasm
Q9A6B8	LIPB_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A6C8	PANC_CAUCR		cytoplasm
Q9A6H0	TYSY_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A6H4	ILVC_CAUCR		cytoplasm
Q9A6I5	FUMC_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A6J5	MIAA_CAUCR		cytoplasm
Q9A6M3	EX7S_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A6M5	DXS_CAUCR		cytoplasm
Q9A6N1	6PGL_CAUCR		extracellular
Q9A6N3	GLK_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A6Q5	THIC_CAUCR		cytoplasm
Q9A6R5	OPGH_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9A6S5	DGTL1_CAUCR		cytoplasm
Q9A6T1	TATB_CAUCR	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9A6T5	SURE_CAUCR	cytoplasm (potential).	periplasm
Q9A6T6	PIMT_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A6U9	Y1984_CAUCR	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q9A6V2	CRCB_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9A6V7	Y1976_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9A6Z7	LOLD2_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9A700	DPO3A_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A703	RS2_CAUCR		cytoplasm
Q9A705	PYRH_CAUCR		cytoplasm
Q9A706	RRF_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A707	UPPS_CAUCR		inner membrane
Q9A709	DXR_CAUCR		cytoplasm

Q9A710	MMPA_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9A713	LPXD_CAUCR		cytoplasm
Q9A714	FABZ_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A715	LPXA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A724	LEXA_CAUCR		cytoplasm
Q9A727	TRPC_CAUCR		cytoplasm
Q9A728	TRPD_CAUCR		cytoplasm
Q9A733	TPIS_CAUCR		cytoplasm
Q9A741	LFTR_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A744	AROQ_CAUCR		cytoplasm
Q9A746	THIG_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A756	SYI_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A777	CALB_CAUCR		cytoplasm
Q9A7H8	HFQ_CAUCR		cytoplasm
Q9A7I5	ISPDF_CAUCR		cytoplasm
Q9A7I8	LIPA_CAUCR	cytoplasm (potential).	cytoplasm
Q9A7J9	ENO_CAUCR	cytoplasm (by similarity).	extracellular
Q9A7K2	RL32_CAUCR		cytoplasm
Q9A7K3	PYRG_CAUCR		cytoplasm
Q9A7M2	NDK_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A7M9	AMPA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A7N4	PDXA_CAUCR	cytoplasm (probable).	cytoplasm
Q9A7N5	KSGA_CAUCR		cytoplasm
Q9A7N9	KGUA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A7P3	ACP_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A7Q1	RS6_CAUCR		cytoplasm
Q9A7Q2	RS18_CAUCR		cytoplasm
Q9A7Q3	RL9_CAUCR		cytoplasm
Q9A7Q9	ALR_CAUCR		cytoplasm
Q9A7U9	GUAA_CAUCR		cytoplasm
Q9A7V4	RRMJ_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A7V7	PH4H_CAUCR		cytoplasm
Q9A7X1	CYSA_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9A7X6	ATKC_CAUCR	bacterial cell inner membrane; single-pass membrane protein (probable).	inner membrane
Q9A7X7	ATKB_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9A7X8	ATKA_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9A7Y1	TGT_CAUCR		cytoplasm
Q9A7Y2	QUEA_CAUCR	cytoplasm (potential).	cytoplasm
Q9A7Z2	BIOD_CAUCR		cytoplasm
Q9A7Z5	ATE_CAUCR	cytoplasm (potential).	cytoplasm
Q9A802	RECO_CAUCR		cytoplasm
Q9A805	RNC_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A807	ACPS_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A808	PDXJ_CAUCR	cytoplasm (probable).	cytoplasm
Q9A810	PYRE_CAUCR		cytoplasm
Q9A823	LEU1_CAUCR		cytoplasm

Q9A838	PRMA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A871	CAPP_CAUCR		cytoplasm
Q9A873	KDUI_CAUCR		cytoplasm
Q9A874	UXAC_CAUCR		cytoplasm
Q9A894	SSB_CAUCR		cytoplasm
Q9A8C5	KDSA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A8H5	ARGC_CAUCR	cytoplasm (probable).	cytoplasm
Q9A8H6	RS9_CAUCR		cytoplasm
Q9A8I4	FABH_CAUCR	cytoplasm (probable).	cytoplasm
Q9A8I5	PLSX_CAUCR		cytoplasm
Q9A8J0	HPPA_CAUCR	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9A8J3	NUSB_CAUCR		cytoplasm
Q9A8J4	RISB2_CAUCR		cytoplasm
Q9A8J5	NRDR_CAUCR		cytoplasm
Q9A8J6	GLYA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A8L7	ISPE_CAUCR		cytoplasm
Q9A8M7	Y1326_CAUCR		cytoplasm
Q9A8N3	SYV_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A8S9	RPOA_CAUCR		cytoplasm
Q9A8T0	RS11_CAUCR		cytoplasm
Q9A8T1	RS13_CAUCR		cytoplasm
Q9A8T2	KAD_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A8T4	RL15_CAUCR		cytoplasm
Q9A8T6	RS5_CAUCR		cytoplasm
Q9A8T7	RL18_CAUCR		cytoplasm
Q9A8T9	RS8_CAUCR		cytoplasm
Q9A8U0	RS14_CAUCR		cytoplasm
Q9A8U1	RL5_CAUCR		cytoplasm
Q9A8U2	RL24_CAUCR		cytoplasm
Q9A8U4	RS17_CAUCR		cytoplasm
Q9A8U5	RL29_CAUCR		cytoplasm
Q9A8U6	RL16_CAUCR		cytoplasm
Q9A8U7	RS3_CAUCR		cytoplasm
Q9A8U8	RL22_CAUCR		cytoplasm
Q9A8U9	RS19_CAUCR		cytoplasm
Q9A8V0	RL2_CAUCR		cytoplasm
Q9A8V2	RL4_CAUCR		cytoplasm
Q9A8V3	RL3_CAUCR		cytoplasm
Q9A8V4	RS10_CAUCR		cytoplasm
Q9A8Y2	Y1215_CAUCR		inner membrane
Q9A8Z4	Y1201_CAUCR		cytoplasm
Q9A8Z9	SSRP_CAUCR	cytoplasm (potential).	cytoplasm
Q9A900	DAPA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A919	DHMA_CAUCR		cytoplasm
Q9A943	LLDD_CAUCR		cytoplasm
Q9A9A7	RECA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A9C8	Y1060_CAUCR		inner membrane
Q9A9E2	RL35_CAUCR		cytoplasm
Q9A9E3	RL20_CAUCR		cytoplasm
Q9A9E9	MSRA2_CAUCR		cytoplasm
Q9A9F4	LEPA_CAUCR		cytoplasm
Q9A9I6	MSRA1_CAUCR		cytoplasm
Q9A9M1	HUTU_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9A9M3	FLIE_CAUCR		flagellar

Q9A9N8	Y938_CAUCR		cytoplasm
Q9A9P4	LOLD1_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9A9Q6	PDXH_CAUCR		cytoplasm
Q9A9S4	RISB1_CAUCR		cytoplasm
Q9A9T4	CLPB_CAUCR	cytoplasm (probable).	cytoplasm
Q9A9W0	ISPG_CAUCR		cytoplasm
Q9A9X1	UBIG_CAUCR		cytoplasm
Q9AA17	WRBA_CAUCR		cytoplasm
Q9AA39	RNPA_CAUCR		extracellular
Q9AA40	OXAA_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9AA58	Y748_CAUCR		cytoplasm
Q9AAA5	SYI_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9AAE1	RL28_CAUCR		cytoplasm
Q9AAF9	RL11_CAUCR		cytoplasm
Q9AAG0	RL1_CAUCR		cytoplasm
Q9AAH2	DCTA_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9AAL3	ARGM_CAUCR		cytoplasm
Q9AAS1	METX_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9AAT8	HEM6_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9AAU1	RPOC_CAUCR		cytoplasm
Q9AAU2	RPOB_CAUCR		cytoplasm
Q9AAU7	RL7_CAUCR		cytoplasm
Q9AAV1	LGT_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9AAV6	KPRS_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9AAV8	RL25_CAUCR		cytoplasm
Q9AAV9	PTH_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9AAW1	METE_CAUCR		cytoplasm
Q9AAX9	Y463_CAUCR		inner membrane
Q9AAY3	GCH1_CAUCR		cytoplasm
Q9AAY5	HIS3_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9AB09	Y425_CAUCR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9AB70	PHNC_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9AB72	DHSL_CAUCR		cytoplasm
Q9AB80	Y351_CAUCR	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
Q9AB94	SUCC_CAUCR		cytoplasm
Q9ABA6	MTGA_CAUCR	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q9ABB3	RL27_CAUCR		cytoplasm
Q9ABB6	PROB_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9ABD2	PUR2_CAUCR		cytoplasm

Q9ABD6	PSTB_CAUCR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ABE5	ARGB_CAUCR	cytoplasm (probable).	cytoplasm
Q9ABE7	DAPD_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9ABE9	FMT_CAUCR		cytoplasm
Q9ABF0	TRUA_CAUCR		cytoplasm
Q9ABF2	GLSA_CAUCR		extracellular
Q9ABF5	DEF_CAUCR		cytoplasm
Q9ABF8	RECR_CAUCR		cytoplasm
Q9ABF9	Y268_CAUCR		inner membrane
Q9ABG6	PANE_CAUCR	cytoplasm (potential).	cytoplasm
Q9ABH0	SAHH_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9ABI2	PHS_CAUCR		cytoplasm
Q9ABK5	G6PI_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9ABM9	TRMD_CAUCR	cytoplasm (potential).	cytoplasm
Q9ABN0	LEU2_CAUCR		cytoplasm
Q9ABN1	LEUD_CAUCR		cytoplasm
Q9ABN3	LEU3_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9ABS4	HAM1_CAUCR		extracellular
Q9ABS7	Y143_CAUCR		inner membrane
Q9ABS9	GSHB_CAUCR		cytoplasm
Q9ABT9	KUP_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ABU1	ASSY_CAUCR	cytoplasm (probable).	cytoplasm
Q9ABV2	GLMS_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9ABV3	GLMM_CAUCR		cytoplasm
Q9ABW5	PYRF_CAUCR		cytoplasm
Q9ABX9	Y091_CAUCR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ABY4	PUR9_CAUCR		cytoplasm
Q9ABZ8	HEM3_CAUCR		cytoplasm
Q9AC08	Y061_CAUCR		cytoplasm
Q9AC14	Y054_CAUCR		inner membrane
Q9AC16	LNT_CAUCR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9AC20	IPYR_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9AC22	Y045_CAUCR		inner membrane
Q9AC30	RBFA_CAUCR	cytoplasm (potential).	cytoplasm
Q9AC31	RS15_CAUCR		cytoplasm
Q9AC39	Y027_CAUCR		cytoplasm
Q9AC48	MOAA_CAUCR		cytoplasm
Q9AC50	MOAE_CAUCR		cytoplasm
Q9AC52	MOAC_CAUCR		cytoplasm
Q9AC53	GLND_CAUCR		cytoplasm
Q9AC54	MUTS_CAUCR		cytoplasm
Q9AC57	AROE_CAUCR		cytoplasm
Q9AC58	Y002_CAUCR	cytoplasm (potential).	inner membrane
Q9AC59	Y001_CAUCR		cytoplasm
Q9ACP0	IGAA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ADT1	FTR_METCA	cytoplasm (by similarity).	cytoplasm

Q9AEQ5	PTH_BRUAB	cytoplasm (by similarity).	cytoplasm
Q9AES0	RNPH_PASHA		cytoplasm
Q9AEV8	APAH_BURMA		extracellular
Q9AF21	COMA_XANP2		cytoplasm
Q9AFF7	PRRB_PRB02	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9AG20	MINC_NEIGO		inner membrane
Q9AGA6	AGLB_KLEPN		cytoplasm
Q9AGA7	PTUCB_KLEPN	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9AGD5	CLCB_YERPS	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9AGM5	LEXA_RALEU		cytoplasm
Q9AGU4	HUTU_PSESX	cytoplasm (by similarity).	cytoplasm
Q9AHF0	1A1D_AGRTU		cytoplasm
Q9AHK1	LUXS_VIBVU		extracellular
Q9AHY3	FADB_PSEPU		cytoplasm
Q9AI34	GMHB_BURMA	cytoplasm (by similarity).	cytoplasm
Q9AI36	GMHA_BURMA	cytoplasm (by similarity).	cytoplasm
Q9AIF4	RS14_CARRU		cytoplasm
Q9AIF9	RS3_CARRU		cytoplasm
Q9AIG8	RS7_CARRU		cytoplasm
Q9AIG9	RS12_CARRU		cytoplasm
Q9AIN8	RL3_CARRU		cytoplasm
Q9AIP8	LPXD_METCA		cytoplasm
Q9AIU0	TTGR_PSEPU		cytoplasm
Q9AIU3	RPOB_ANAPH		cytoplasm
Q9AJ37	SCA4_RICFE	cytoplasm (probable).	cytoplasm
Q9AJ63	SCA4_RICTY	cytoplasm (probable).	cytoplasm
Q9AJ79	SCA4_RICJA	cytoplasm (probable).	cytoplasm
Q9AJA7	LUXD_SHEHA		cytoplasm
Q9AJG9	MRAW_VIBPR		cytoplasm
Q9AJH1	MRAW_VIBPA		cytoplasm
Q9AJQ8	METH_VIBFI		cytoplasm
Q9AKC4	FER2_RICTY		cytoplasm
Q9AKD8	MRAY_RICTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9AKE3	SSRP_RICTY	cytoplasm (potential).	cytoplasm
Q9AKE4	DAPA_RICTY	cytoplasm (by similarity).	cytoplasm
Q9AKH1	FER2_RICRI		cytoplasm
Q9AKI4	Y600_RICRI		cytoplasm
Q9AKI7	MURE_RICRI	cytoplasm (probable).	cytoplasm
Q9AKI9	MRAY_RICRI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9AKJ8	SSRP_RICRI	cytoplasm (potential).	cytoplasm
Q9AKJ9	DAPA_RICRI	cytoplasm (by similarity).	cytoplasm
Q9AKM6	FER2_RICMO		cytoplasm
Q9AKP0	MURE_RICMO	cytoplasm (probable).	cytoplasm
Q9AKP2	MRAY_RICMO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q9AKQ2	SSRP_RICMO	cytoplasm (potential).	cytoplasm
Q9AKQ3	DAPA_RICMO	cytoplasm (by similarity).	cytoplasm
Q9AKR8	ARGC_RHIET	cytoplasm (probable).	cytoplasm
Q9AKT2	CH10_PSEHT	cytoplasm (by similarity).	cytoplasm
Q9AL83	NIKR_BRUSU		cytoplasm
Q9AL99	MSRAB_ACTAC		cytoplasm
Q9ALA9	CH601_VIBVU	cytoplasm (by similarity).	cytoplasm
Q9ALJ1	RL20_VIBME		cytoplasm
Q9ALJ2	RL35_VIBME		cytoplasm
Q9AM27	UXAC_XANOR		cytoplasm
Q9AML4	PSTB_EDWTA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9AMN9	DUSC_PSEAC		cytoplasm
Q9AMR9	PANC_BRAJA		cytoplasm
Q9AMS0	PANB1_BRAJA		cytoplasm
Q9AMS6	PAND_BRAJA		cytoplasm
Q9AMV8	METE_BRAJA		cytoplasm
Q9AMW4	Y2058_BRAJA		cytoplasm
Q9ANL1	NIFW_BRAJA		cytoplasm
Q9ANP1	HYPD1_BRAJA		extracellular
Q9ANP8	HYP1_BRAJA		cytoplasm
Q9ANR2	DCTA_BRAJA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ANR9	CH60_WIGBR	cytoplasm (by similarity).	cytoplasm
Q9ANS0	CH10_WIGBR	cytoplasm (by similarity).	cytoplasm
Q9AP01	ALLA_BURCE		extracellular
Q9AP31	RECX_XANOR	cytoplasm (potential).	cytoplasm
Q9APB2	RECA_RALEU	cytoplasm (by similarity).	cytoplasm
Q9APD9	ZRAR_KLEOX	cytoplasm (probable).	cytoplasm
Q9APE0	ZRAS_KLEOX	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9APJ2	RL32_METEX		cytoplasm
Q9AQ50	FNR_KLEOX	cytoplasm (probable).	cytoplasm
Q9AQI4	SELD_PSEOC		cytoplasm
Q9AQT2	UREE_RHOCA	cytoplasm (by similarity).	cytoplasm
Q9AQT6	URE3_RHOCA	cytoplasm (by similarity).	cytoplasm
Q9CJK1	GLNB_PASMU		cytoplasm
Q9CJK7	LPXB_PASMU		cytoplasm
Q9CJK8	LPXA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CJK9	FABZ_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CJL0	LPXD_PASMU		cytoplasm
Q9CJL2	Y1991_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9CJL4	UPPS_PASMU		inner membrane
Q9CJL5	PYRH_PASMU		cytoplasm
Q9CJM2	CLPP_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CJM5	Y1972_PASMU		inner membrane
Q9CJN5	LEU1_PASMU		cytoplasm
Q9CJN6	LEU3_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CJN7	LEU2_PASMU		cytoplasm
Q9CJN8	LEUD_PASMU		cytoplasm
Q9CJN9	Y1958_PASMU		inner membrane

Q9CJP3	RLMB_PASMU		cytoplasm
Q9CJP4	SSB_PASMU		cytoplasm
Q9CJR0	Y1934_PASMU		inner membrane
Q9CJR2	Y1932_PASMU		cytoplasm
Q9CJR4	Y1928_PASMU		inner membrane
Q9CJR6	RLPA_PASMU		cytoplasm
Q9CJR9	Y1923_PASMU		inner membrane
Q9CJS1	RHLB_PASMU		cytoplasm
Q9CJS5	ACP_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CJS8	FABH_PASMU	cytoplasm (probable).	cytoplasm
Q9CJS9	RL32_PASMU		cytoplasm
Q9CJT7	BIOD2_PASMU		cytoplasm
Q9CJU2	PSD_PASMU		cytoplasm
Q9CJU4	Y1897_PASMU	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q9CJU5	PROB_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CJV6	NDPA_PASMU	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q9CJV7	Y1884_PASMU		inner membrane
Q9CJW1	DUSC_PASMU		cytoplasm
Q9CJW3	DJLA_PASMU	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q9CJW4	PYRE_PASMU		cytoplasm
Q9CJW5	RNPH_PASMU		cytoplasm
Q9CJW9	PYRG_PASMU		cytoplasm
Q9CJX0	RUVX_PASMU	cytoplasm (potential).	cytoplasm
Q9CJX1	Y1869_PASMU		inner membrane
Q9CJX3	RUMA_PASMU		cytoplasm
Q9CJY8	Y1850_PASMU		inner membrane
Q9CJZ2	PTSBC_PASMU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9CK01	RAPA_PASMU		cytoplasm
Q9CK02	RLUA_PASMU		cytoplasm
Q9CK03	Y1835_PASMU		inner membrane
Q9CK20	RECX_PASMU	cytoplasm (potential).	cytoplasm
Q9CK24	HEM3_PASMU		cytoplasm
Q9CK29	GLMU_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CK30	Y1805_PASMU		cytoplasm
Q9CK31	Y1804_PASMU		inner membrane
Q9CK32	TRMA_PASMU		cytoplasm
Q9CK40	TORD_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CK41	TORA_PASMU	periplasm (by similarity).	periplasm
Q9CK44	Y1790_PASMU		cytoplasm
Q9CK54	Y1780_PASMU		cytoplasm
Q9CK55	Y1779_PASMU		inner membrane
Q9CK59	Y1775_PASMU		cytoplasm
Q9CK61	Y1773_PASMU		cytoplasm
Q9CK62	Y1772_PASMU		cytoplasm
Q9CK63	Y1771_PASMU		cytoplasm
Q9CK66	SELA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CK84	NUSG_PASMU		cytoplasm
Q9CK85	RL11_PASMU		cytoplasm
Q9CK86	RL1_PASMU		cytoplasm

Q9CK89	RL10_PASMU		cytoplasm
Q9CK90	RL7_PASMU		cytoplasm
Q9CK91	RPOB_PASMU		cytoplasm
Q9CK92	RPOC_PASMU		cytoplasm
Q9CK94	DBH_PASMU		cytoplasm
Q9CK95	METQ_PASMU	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q9CK96	METI_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CK97	METN_PASMU	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q9CK98	GMHB_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CKA2	Y1723_PASMU		inner membrane
Q9CKA5	Y1720_PASMU	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q9CKA6	RLUD_PASMU		cytoplasm
Q9CKB0	NANA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CKB3	NANK_PASMU		cytoplasm
Q9CKB7	Y1707_PASMU		outer membrane
Q9CKC0	CLPB_PASMU	cytoplasm (probable).	cytoplasm
Q9CKC2	XERC_PASMU	cytoplasm (by similarity).	nucleoid
Q9CKC7	Y1696_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CKC8	ZIPA_PASMU	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q9CKC9	CYSZ_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CKD3	TATA_PASMU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9CKD4	UBIB_PASMU	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9CKD6	UBIE_PASMU		cytoplasm
Q9CKD7	Y1685_PASMU		cytoplasm
Q9CKE9	KTHY_PASMU		cytoplasm
Q9CKF3	Y1668_PASMU		inner membrane
Q9CKF5	HAM1_PASMU		extracellular
Q9CKF7	SYI_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CKG0	RS20_PASMU		cytoplasm
Q9CKG2	Y1656_PASMU		inner membrane
Q9CKG3	Y1655_PASMU		extracellular
Q9CKG4	Y1654_PASMU		cytoplasm
Q9CKG8	PDXA_PASMU	cytoplasm (probable).	cytoplasm
Q9CKH9	TALB_PASMU	cytoplasm (probable).	cytoplasm
Q9CKI7	CRCB_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CKJ2	THD1_PASMU		cytoplasm
Q9CKJ5	GPH_PASMU		cytoplasm
Q9CKJ8	Y1617_PASMU		inner membrane

Q9CKJ9	RBN_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CKK0	DTD_PASMU	cytoplasm (probable).	cytoplasm
Q9CKK4	TRUD_PASMU		cytoplasm
Q9CKK5	FTSB_PASMU	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q9CKK7	RLUE_PASMU		cytoplasm
Q9CKL0	TALA_PASMU	cytoplasm (probable).	cytoplasm
Q9CKL2	OSTA_PASMU	bacterial cell outer membrane (by similarity).	outer membrane
Q9CKL8	NAPA_PASMU	periplasm (by similarity).	periplasm
Q9CKM2	Y1590_PASMU		cytoplasm
Q9CKM8	Y1583_PASMU		cytoplasm
Q9CKN6	ASNA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CKN9	Y1571_PASMU		cytoplasm
Q9CKP7	RSMB_PASMU		cytoplasm
Q9CKP9	Y1557_PASMU		inner membrane
Q9CKQ9	HSLO_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CKR3	Y1543_PASMU		outer membrane
Q9CKR4	PPCK_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CKU6	Y1509_PASMU	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q9CKU9	GPMA_PASMU		cytoplasm
Q9CKV0	CARB_PASMU		cytoplasm
Q9CKV2	Y1503_PASMU		cytoplasm
Q9CKV3	CARA_PASMU		cytoplasm
Q9CKV5	Y1500_PASMU		inner membrane
Q9CKV9	TUSA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CKW0	ATPE_PASMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9CKW1	ATPB_PASMU		inner membrane
Q9CKW2	ATPA_PASMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9CKX3	Y1478_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CKY2	Y1469_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CKY6	Y1464_PASMU		cytoplasm
Q9CKY9	Y1461_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CL00	Y1449_PASMU		outer membrane
Q9CL02	Y1447_PASMU		inner membrane
Q9CL07	GLPB_PASMU		cytoplasm
Q9CL11	Y1437_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CL12	GLPE_PASMU	cytoplasm (by similarity).	cytoplasm

Q9CL16	SECB_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CL17	GPDA_PASMU	cytoplasm (probable).	cytoplasm
Q9CL21	RECQ_PASMU		cytoplasm
Q9CL24	OADB_PASMU	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CL26	OADG_PASMU	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q9CL27	TNAA_PASMU		cytoplasm
Q9CL29	DUSA_PASMU		cytoplasm
Q9CL32	RL3_PASMU		cytoplasm
Q9CL33	RL4_PASMU		cytoplasm
Q9CL35	RL2_PASMU		cytoplasm
Q9CL36	RL22_PASMU		cytoplasm
Q9CL37	RS3_PASMU		cytoplasm
Q9CL38	RL16_PASMU		cytoplasm
Q9CL39	RL29_PASMU		cytoplasm
Q9CL40	RS17_PASMU		cytoplasm
Q9CL41	RL24_PASMU		cytoplasm
Q9CL42	RL5_PASMU		cytoplasm
Q9CL43	RS14_PASMU		cytoplasm
Q9CL44	RS8_PASMU		cytoplasm
Q9CL46	RL18_PASMU		cytoplasm
Q9CL47	RS5_PASMU		cytoplasm
Q9CL49	RL15_PASMU		cytoplasm
Q9CL51	RS13_PASMU		cytoplasm
Q9CL52	RS11_PASMU		cytoplasm
Q9CL53	RS4_PASMU		cytoplasm
Q9CL57	Y1386_PASMU		inner membrane
Q9CL60	SPEA_PASMU		cytoplasm
Q9CL73	XGPT_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CL85	RS7_PASMU		cytoplasm
Q9CL86	RS12_PASMU		cytoplasm
Q9CL88	TUSC_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CL89	TUSD_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CL92	SLYX_PASMU		inner membrane
Q9CL97	HLDD_PASMU		cytoplasm
Q9CLA3	TRMU_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLA6	NQRF_PASMU	bacterial cell inner membrane (potential).	inner membrane
Q9CLA7	NQRE_PASMU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9CLA8	NQRD_PASMU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9CLA9	NQRC_PASMU	bacterial cell inner membrane (potential).	inner membrane
Q9CLB0	NQRB_PASMU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9CLB1	NQRA_PASMU		inner membrane
Q9CLB8	MLTC_PASMU	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q9CLB9	FETP_PASMU		cytoplasm

Q9CLC2	TRMB_PASMU		cytoplasm
Q9CLC4	CORA_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CLC6	Y1313_PASMU		inner membrane
Q9CLD4	COAD_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLD5	KDKA_PASMU	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q9CLE0	RL19_PASMU		cytoplasm
Q9CLE1	TRMD_PASMU	cytoplasm (potential).	cytoplasm
Q9CLE2	Y1295_PASMU		outer membrane
Q9CLE6	DEOD_PASMU		cytoplasm
Q9CLE7	METJ_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLE9	USPA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLF1	ILVC_PASMU		cytoplasm
Q9CLF4	Y1281_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CLF7	Y1278_PASMU		cytoplasm
Q9CLG6	Y1268_PASMU	cytoplasm (potential).	inner membrane
Q9CLG7	RMUC_PASMU		cytoplasm
Q9CLH5	DLGD_PASMU	cytoplasm (potential).	cytoplasm
Q9CLI9	PRIM_PASMU		cytoplasm
Q9CLJ0	RS21_PASMU		cytoplasm
Q9CLJ2	Y1237_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CLJ5	PDXT_PASMU		cytoplasm
Q9CLJ6	PDXS_PASMU		extracellular
Q9CLL5	KSGA_PASMU		cytoplasm
Q9CLL7	PYRR_PASMU		cytoplasm
Q9CLL8	HIS2_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLL9	Y1205_PASMU		cytoplasm
Q9CLM0	HIS6_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLM1	HIS4_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLM2	Y1201_PASMU		cytoplasm
Q9CLM3	HIS81_PASMU		cytoplasm
Q9CLM4	HISX_PASMU		cytoplasm
Q9CLN1	Y1189_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CLN7	PLSB_PASMU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9CLN8	RS6_PASMU		cytoplasm
Q9CLN9	PRIB_PASMU		cytoplasm
Q9CLP0	RL9_PASMU		cytoplasm
Q9CLP1	Y1176_PASMU	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q9CLP9	RRAA_PASMU		cytoplasm
Q9CLQ1	TRME_PASMU		cytoplasm
Q9CLQ2	OXAA_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CLQ3	Y1164_PASMU		inner membrane

Q9CLQ4	DNAA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLQ6	RECF_PASMU	cytoplasm (potential).	cytoplasm
Q9CLQ8	Y1156_PASMU		inner membrane
Q9CLQ9	SLMA_PASMU	nucleoid-associated (by similarity).	cytoplasm
Q9CLR1	RL28_PASMU		cytoplasm
Q9CLR7	RL31_PASMU		cytoplasm
Q9CLT5	Y1124_PASMU		inner membrane
Q9CLT6	Y1123_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CLT7	Y1122_PASMU		cytoplasm
Q9CLT8	Y1121_PASMU		cytoplasm
Q9CLT9	ARGE_PASMU	cytoplasm (probable).	extracellular
Q9CLU4	MODD_PASMU		cytoplasm
Q9CLV1	Y1101_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CLV5	MENB_PASMU		cytoplasm
Q9CLV6	Y1095_PASMU		inner membrane
Q9CLV7	MENC_PASMU		cytoplasm
Q9CLW2	PRMA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLW3	DUSB_PASMU		cytoplasm
Q9CLW4	PUR4_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CLW6	Y1083_PASMU		cytoplasm
Q9CLW7	Y1082_PASMU		cytoplasm
Q9CLX3	Y1076_PASMU		cytoplasm
Q9CLX8	Y1071_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CLY2	MUTH_PASMU	cytoplasm (by similarity).	extracellular
Q9CLY7	MTLD_PASMU		cytoplasm
Q9CLY8	PTM3C_PASMU	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CLZ7	DAPA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CM00	GSHAB_PASMU		cytoplasm
Q9CM13	CORC_PASMU		inner membrane
Q9CM14	LNT_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CM16	PEPB_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CM17	NDK_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CM18	SPRT_PASMU	cytoplasm (potential).	extracellular
Q9CM20	HTPG_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CM49	THIO_PASMU		cytoplasm
Q9CM50	NPD_PASMU	cytoplasm (probable).	cytoplasm
Q9CM51	RLUC_PASMU		cytoplasm
Q9CM61	Y980_PASMU		inner membrane
Q9CM71	TOLB_PASMU	periplasm (potential).	periplasm
Q9CM80	FBPC_PASMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9CM85	URK_PASMU	cytoplasm (by similarity).	cytoplasm

Q9CM87	SOTB_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CM94	NRDG_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CM98	PROA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CMB1	MSRB_PASMU		cytoplasm
Q9CMB2	RPOZ_PASMU		cytoplasm
Q9CMB4	RECG_PASMU		cytoplasm
Q9CMC3	Y909_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CMC6	HFQ_PASMU		cytoplasm
Q9CMC7	MIAA_PASMU		cytoplasm
Q9CMD0	Y901_PASMU		inner membrane
Q9CMD1	ENGC_PASMU		cytoplasm
Q9CME6	HLDE_PASMU		cytoplasm
Q9CME9	Y880_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CMF1	DGTL1_PASMU		cytoplasm
Q9CMF4	NAGB_PASMU		extracellular
Q9CMG7	MSBA_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CMG8	LPXK_PASMU		cytoplasm
Q9CMI6	UBIG_PASMU		cytoplasm
Q9CMI7	HIS82_PASMU		cytoplasm
Q9CMI8	Y836_PASMU		inner membrane
Q9CMJ4	Y830_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CMJ6	ARGA_PASMU	cytoplasm (probable).	cytoplasm
Q9CMJ8	RIMK_PASMU		cytoplasm
Q9CMJ9	Y825_PASMU		inner membrane
Q9CMK1	FUMC_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CMK4	Y819_PASMU		cytoplasm
Q9CMK5	SYV_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CMK9	TPL_PASMU		cytoplasm
Q9CML8	IHFB_PASMU		cytoplasm
Q9CMM1	PYRF_PASMU		cytoplasm
Q9CMM8	SELD_PASMU		cytoplasm
Q9CMM9	Y788_PASMU		cytoplasm
Q9CMP0	CHS_PASMU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9CMP7	Y768_PASMU		cytoplasm
Q9CMQ4	Y761_PASMU		inner membrane
Q9CMQ6	Y758_PASMU		cytoplasm
Q9CMQ7	TRUB_PASMU		cytoplasm
Q9CMQ8	SYV_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CMR2	NRDR_PASMU		cytoplasm
Q9CMR5	Y747_PASMU		cytoplasm
Q9CMS0	MODC_PASMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9CMS2	DNAJ_PASMU	cytoplasm (probable).	cytoplasm

Q9CMS3	Y739_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CMV2	Y703_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CMW1	ACSA_PASMU		cytoplasm
Q9CMX0	Y682_PASMU		inner membrane
Q9CMX6	Y674_PASMU		periplasm
Q9CMY0	Y670_PASMU		cytoplasm
Q9CMY2	FNR_PASMU	cytoplasm (probable).	cytoplasm
Q9CMY9	RLUB_PASMU		cytoplasm
Q9CMZ1	Y659_PASMU		cytoplasm
Q9CMZ2	DAPD_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CMZ3	Y657_PASMU		cytoplasm
Q9CMZ6	Y654_PASMU		inner membrane
Q9CMZ9	NORM_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CN04	Y645_PASMU		cytoplasm
Q9CN08	BIOD1_PASMU		cytoplasm
Q9CN10	RL25_PASMU		cytoplasm
Q9CN17	Y630_PASMU		cytoplasm
Q9CN18	IHFA_PASMU		cytoplasm
Q9CN20	Y626_PASMU		cytoplasm
Q9CN21	MOAA_PASMU		cytoplasm
Q9CN22	MOAC_PASMU		cytoplasm
Q9CN24	MOAE_PASMU		cytoplasm
Q9CN30	TOP1_PASMU		cytoplasm
Q9CN31	Y614_PASMU		cytoplasm
Q9CN32	Y613_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CN38	MUKF_PASMU	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q9CN40	MSRA_PASMU		cytoplasm
Q9CN41	RL20_PASMU		cytoplasm
Q9CN56	Y587_PASMU		cytoplasm
Q9CN71	MTFA_PASMU		cytoplasm
Q9CN78	LOLD_PASMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9CN81	Y557_PASMU		extracellular
Q9CN82	HEMK_PASMU		cytoplasm
Q9CN84	Y553_PASMU	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q9CN86	MDH_PASMU		cytoplasm
Q9CN88	PURR_PASMU		cytoplasm
Q9CN89	CAPP_PASMU		cytoplasm
Q9CN90	PHSG_PASMU		cytoplasm
Q9CN91	GLGA_PASMU		cytoplasm
Q9CN92	GLGC_PASMU		cytoplasm
Q9CN94	GLGB_PASMU		extracellular
Q9CN96	Y539_PASMU		inner membrane

Q9CN97	Y538_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CN98	Y537_PASMU		cytoplasm
Q9CN99	Y536_PASMU		inner membrane
Q9CNA0	EX7S_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CNA2	Y531_PASMU		cytoplasm
Q9CNA6	CUTC_PASMU	cytoplasm (probable).	periplasm
Q9CNB0	SSPA_PASMU		cytoplasm
Q9CNB1	RS9_PASMU		cytoplasm
Q9CNB3	Y519_PASMU		periplasm
Q9CNC4	LST_PASMU		cytoplasm
Q9CNC5	Y507_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CNC7	Y505_PASMU		cytoplasm
Q9CNC8	Y504_PASMU		cytoplasm
Q9CND0	Y502_PASMU		cytoplasm
Q9CND4	Y498_PASMU		cytoplasm
Q9CND8	Y494_PASMU		cytoplasm
Q9CND9	Y493_PASMU		cytoplasm
Q9CNE0	Y492_PASMU		cytoplasm
Q9CNE1	Y491_PASMU		cytoplasm
Q9CNE3	Y489_PASMU		cytoplasm
Q9CNE8	FABA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CNF0	Y482_PASMU		inner membrane
Q9CNF2	Y480_PASMU		inner membrane
Q9CNF3	TRUC_PASMU		cytoplasm
Q9CNF6	QUEF_PASMU	cytoplasm (probable).	cytoplasm
Q9CNG4	DPO4_PASMU	cytoplasm (probable).	cytoplasm
Q9CNG8	RFE_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CNG9	GSA_PASMU	cytoplasm (potential).	cytoplasm
Q9CNH1	GLND_PASMU		cytoplasm
Q9CNH4	Y457_PASMU		inner membrane
Q9CNH7	PEPE_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CNJ0	GLMM_PASMU		cytoplasm
Q9CNJ3	RRMJ_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CNJ4	PSTS_PASMU	periplasm (potential).	periplasm
Q9CNJ5	PSTC_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CNJ6	PSTA_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CNJ7	PSTB_PASMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9CNL2	G6PI_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CNL3	Y415_PASMU		inner membrane
Q9CNL5	ALR_PASMU		cytoplasm
Q9CNL6	DNAB_PASMU		cytoplasm
Q9CNM2	FDHE_PASMU	cytoplasm (by similarity).	cytoplasm

Q9CNM5	Y402_PASMU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9CNN2	Y395_PASMU		inner membrane
Q9CNN3	MGSA_PASMU		cytoplasm
Q9CNN6	Y391_PASMU		inner membrane
Q9CNN7	Y390_PASMU		cytoplasm
Q9CNP0	RNFA_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CNP2	RNFC_PASMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	cytoplasm
Q9CNP3	RNFD_PASMU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9CNP4	RNFG_PASMU	bacterial cell inner membrane (potential).	inner membrane
Q9CNP5	RNFE_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CNQ2	GLNE_PASMU		cytoplasm
Q9CNS7	RL27_PASMU		cytoplasm
Q9CNT1	ARGD_PASMU	cytoplasm (probable).	cytoplasm
Q9CNT7	Y338_PASMU		cytoplasm
Q9CNU1	GRPE_PASMU	cytoplasm (probable).	cytoplasm
Q9CNU2	PPNK_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CNU9	TRPR_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CNV0	MTGA_PASMU	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q9CNV2	HSCA_PASMU		cytoplasm
Q9CNV3	HSCB_PASMU		inner membrane
Q9CNV8	SUHB_PASMU		cytoplasm
Q9CNV9	Y314_PASMU		cytoplasm
Q9CNW3	Y310_PASMU		cytoplasm
Q9CNW4	Y309_PASMU		cytoplasm
Q9CNX1	RS15_PASMU		cytoplasm
Q9CNX7	Y294_PASMU		cytoplasm
Q9CNX8	GUAA_PASMU		cytoplasm
Q9CNX9	ACCA_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CNY0	Y291_PASMU		cytoplasm
Q9CNY2	TILS_PASMU	cytoplasm (probable).	cytoplasm
Q9CNY5	GALE_PASMU		cytoplasm
Q9CNZ0	SUCC_PASMU		cytoplasm
Q9CP02	PROQ_PASMU	cytoplasm (potential).	inner membrane
Q9CP05	PEPT_PASMU	cytoplasm (by similarity).	extracellular
Q9CP11	CDD_PASMU		cytoplasm
Q9CP15	Y253_PASMU		cytoplasm
Q9CP21	CCA_PASMU		cytoplasm
Q9CP22	KPRS_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CP40	DSBD_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CP41	RL31B_PASMU		cytoplasm
Q9CP43	UBID_PASMU		cytoplasm

Q9CP49	Y211_PASMU		inner membrane
Q9CP52	SECG_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CP53	TOP3_PASMU		cytoplasm
Q9CP58	FRDC_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CP59	FRDD_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CP65	DSBC_PASMU	periplasm (by similarity).	periplasm
Q9CP72	Y183_PASMU		cytoplasm
Q9CP74	RNB_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CP85	Y169_PASMU		cytoplasm
Q9CP86	EX7L_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CP88	Y166_PASMU		inner membrane
Q9CP90	ENGD_PASMU		cytoplasm
Q9CP96	Y157_PASMU		cytoplasm
Q9CPA2	RBSR_PASMU		cytoplasm
Q9CPA5	LPXC_PASMU		cytoplasm
Q9CPB0	MURD_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CPB4	MRAW_PASMU		cytoplasm
Q9CPB5	MRAZ_PASMU		inner membrane
Q9CPC8	Y119_PASMU		cytoplasm
Q9CPD1	Y116_PASMU		cytoplasm
Q9CPD3	KHSE_PASMU	cytoplasm (potential).	cytoplasm
Q9CPD5	Y112_PASMU		cytoplasm
Q9CPD6	Y111_PASMU		cytoplasm
Q9CPD7	Y110_PASMU		cytoplasm
Q9CPD8	Y109_PASMU		cytoplasm
Q9CPE0	DPO3E_PASMU		cytoplasm
Q9CPE3	LPXH_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CPE8	PROC_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CPF0	XERD_PASMU	cytoplasm (by similarity).	nucleoid
Q9CPF4	Y089_PASMU		cytoplasm
Q9CPF5	COAE_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CPG2	LGT_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CPH0	NAGZ_PASMU	cytoplasm (by similarity).	extracellular
Q9CPH1	RUMB_PASMU		cytoplasm
Q9CPH2	K6PF_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CPH5	Y066_PASMU		inner membrane
Q9CPH6	GRCA_PASMU		cytoplasm
Q9CPH8	ERA_PASMU		cytoplasm
Q9CPI4	MEND_PASMU		cytoplasm
Q9CPI5	MENF_PASMU		cytoplasm
Q9CPJ0	FADR_PASMU	cytoplasm (potential).	cytoplasm
Q9CPK3	DPO3A_PASMU	cytoplasm (by similarity).	cytoplasm
Q9CPK9	NRF_X_PASMU	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (potential).	periplasm
Q9CPL4	NRFA_PASMU	periplasm (by similarity).	periplasm
Q9CPL6	PUR5_PASMU	cytoplasm (by similarity).	cytoplasm

Q9CPL8	UPP_PASMU		cytoplasm
Q9CPL9	URAA_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9CPM2	Y015_PASMU		cytoplasm
Q9CPM6	DSBE_PASMU	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q9CPM8	CCME_PASMU	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q9CPN2	CCMA_PASMU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9CPN4	RSUA_PASMU		cytoplasm
Q9CPN6	SODM_PASMU		periplasm
Q9ETX2	LKTA_MANGL	secreted protein (by similarity).	extracellular
Q9EUT5	GLGA_RHITR		cytoplasm
Q9EUT6	GLGC_RHITR		cytoplasm
Q9EV04	PSD_DICD3		cytoplasm
Q9EV27	LKA2E_PASHA	secreted protein (by similarity).	extracellular
Q9EV29	LKA2D_PASHA	secreted protein (by similarity).	extracellular
Q9EV30	LKA2A_PASHA	secreted protein (by similarity).	extracellular
Q9EV31	LKA13_PASHA	secreted protein (by similarity).	extracellular
Q9EV32	LKA16_PASHA	secreted protein (by similarity).	extracellular
Q9EV33	LKA14_PASHA	secreted protein (by similarity).	extracellular
Q9EV34	LKTA8_PASHA	secreted protein (by similarity).	extracellular
Q9EV47	LPXC_SHEVI		cytoplasm
Q9EV79	6PGL_PSEPU		extracellular
Q9EVE1	LEU3_BUCUD	cytoplasm (by similarity).	cytoplasm
Q9EVE4	DPO3B_BUCRP	cytoplasm (by similarity).	cytoplasm
Q9EVG3	LEU3_BUCML	cytoplasm (by similarity).	cytoplasm
Q9EVG6	LEU3_BUCUA	cytoplasm (by similarity).	cytoplasm
Q9EVG9	LEU3_BUCUM	cytoplasm (by similarity).	cytoplasm
Q9EVH5	LEU3_BUCUE	cytoplasm (by similarity).	cytoplasm
Q9EVH8	LEU3_BUCUH	cytoplasm (by similarity).	cytoplasm
Q9EVI1	LEU3_BUCUO	cytoplasm (by similarity).	cytoplasm
Q9EVI4	LEU3_BUCUS	cytoplasm (by similarity).	cytoplasm
Q9EVI7	LEU3_BUCUN	cytoplasm (by similarity).	cytoplasm
Q9EVN2	RNFE_PSEST	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9EVN3	RNFG_PSEST	bacterial cell inner membrane (potential).	inner membrane

Q9EVN4	RNFD_PSEST	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9EXN5	MCHF_ECOLI	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9EXN6	MCHE_ECOLI	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9EXQ1	FNR_ACTAC		cytoplasm
Q9EXU8	PQQE_RHIME		cytoplasm
Q9EXU9	PQQD_RHIME		cytoplasm
Q9EXV0	PQQC_RHIME		cytoplasm
Q9EXV1	PQQB_RHIME		cytoplasm
Q9EXV4	F16P1_RHIME		cytoplasm
Q9EYF1	DINI1_ECO57		cytoplasm
Q9EYG9	SUCC_RHIME		cytoplasm
Q9EYJ6	MDH_RHIME		cytoplasm
Q9EYM2	LOLD_ACIAD	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9EYP3	LIPA_BURPS	cytoplasm (potential).	cytoplasm
Q9EYT4	FLIE_PSEFL		flagellar
Q9EYU1	PAND1_RHILO		cytoplasm
Q9EYV3	PYRF_RHIFR		cytoplasm
Q9EYV5	GADX_ECO27		cytoplasm
Q9EYY3	FTSB_KLEAE	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q9EYY5	GPH_KLEAE		cytoplasm
Q9EYZ5	HFQ_PHOPR		cytoplasm
Q9EZ97	SECB_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9EZI1	ACP_XANAL	cytoplasm (by similarity).	cytoplasm
Q9F0D4	HYBF_KLEPN		cytoplasm
Q9F0H3	HRCQB_PSESM	cytoplasm (by similarity). or loosely associated with a membrane component, probably hrcqa (by similarity).	inner membrane
Q9F0J6	ROO_DESGI		inner membrane
Q9F0W4	NOSZ_PSEFL	periplasm (by similarity).	periplasm
Q9F0Y6	FADA_ENTCL	cytoplasm (by similarity).	cytoplasm
Q9F0Y7	FADB_ENTCL		cytoplasm
Q9F173	DPO1_SALTY		cytoplasm
Q9F187	METE_RALEU		cytoplasm
Q9F1M9	MURC_SHEVI	cytoplasm (probable).	cytoplasm
Q9F1N0	MURG_SHEVI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9F1N2	MURD_SHEVI	cytoplasm (by similarity).	cytoplasm
Q9F1N3	MRAY_SHEVI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9F1N8	MRAW_SHEVI		cytoplasm
Q9F1N9	MRAZ_SHEVI		inner membrane

Q9F1U7	PYRD_FLALU	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9F3T4	AVRP2_PSESJ		cytoplasm
Q9F495	OPGH_DICD3	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9F496	OPGG_DICD3	periplasm (by similarity).	periplasm
Q9F4E2	YREP_BUCGU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9F4E4	CH10_BUCTS	cytoplasm (by similarity).	cytoplasm
Q9F4E6	CH10_BUCTT	cytoplasm (by similarity).	cytoplasm
Q9F4E8	CH10_BUCTC	cytoplasm (by similarity).	cytoplasm
Q9F4F0	CH10_BUCPS	cytoplasm (by similarity).	cytoplasm
Q9F4F2	CH10_BUCPP	cytoplasm (by similarity).	cytoplasm
Q9F4F4	CH10_BUCGU	cytoplasm (by similarity).	cytoplasm
Q9F4G9	YCJF_YEREN	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9F4I7	PYRR_PSEFL		cytoplasm
Q9F4I8	RUVX_PSEFL	cytoplasm (potential).	cytoplasm
Q9F4K4	MODE_HERSE		cytoplasm
Q9F5A5	KUP_AGRRH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9F5M1	RL7_NEIPE		cytoplasm
Q9F5N7	NORM_BURVI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9F5X1	GIDA_PSEU2	cytoplasm (by similarity).	cytoplasm
Q9F5Y0	RNFH_AZOVI		inner membrane
Q9F5Y1	RNFE_AZOVI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9F618	FLBT_RHIME		cytoplasm
Q9F664	GALU_HAEDU		cytoplasm
Q9F665	CSRA_HAEDU		cytoplasm
Q9F672	RECA_HERSE	cytoplasm (by similarity).	cytoplasm
Q9F6B2	SLYA_EDWTA		cytoplasm
Q9F712	CH601_BURPS	cytoplasm (by similarity).	cytoplasm
Q9F746	CLPB_YEREN	cytoplasm (probable).	cytoplasm
Q9F771	XERCL_PSEAE		cytoplasm
Q9F7A2	NUDD_SALTY		cytoplasm
Q9F7B1	WZC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9F7B2	WZB_SALTY		extracellular
Q9F7D4	GALE_YERPE		cytoplasm
Q9F7L5	COAE_PRB01	cytoplasm (by similarity).	cytoplasm
Q9F7L6	ILVC_PRB01		cytoplasm
Q9F7P4	PRRG_PRB01	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9F7R5	ACSA_PRB01		cytoplasm
Q9F7S4	DUT_PRB01		extracellular

Q9F812	THIG_ERWAM	cytoplasm (by similarity).	cytoplasm
Q9F854	HISX_VIBCH		cytoplasm
Q9F855	RLUE_VIBCH		cytoplasm
Q9F8R8	HOSA_ECO11		cytoplasm
Q9F9K6	RL9_RHILT		cytoplasm
Q9F9U0	PQQB_PSEST		cytoplasm
Q9FA11	NIFW_ACEDI		cytoplasm
Q9FA32	IMPC2_SALTY		cytoplasm
Q9FA43	CSID_SALTY		cytoplasm
Q9FA52	OPGH_RHOS4	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9FA54	OPGG_RHOS4	periplasm (by similarity).	periplasm
Q9FAD7	CHEY_ENTCL	cytoplasm (by similarity).	cytoplasm
Q9FAS4	UREE_VIBPA	cytoplasm (by similarity).	cytoplasm
Q9FAS7	URE3_VIBPA	cytoplasm (by similarity).	cytoplasm
Q9FCZ8	HRPA_ERWST	extracellular and secreted via type iii secretion system (by similarity).	extracellular
Q9FD13	GLNE_PASHA		cytoplasm
Q9FD28	RL27_BURPS		cytoplasm
Q9FDC7	RPOA_BARHE		cytoplasm
Q9FDI9	PSD_RHIME		cytoplasm
Q9FDL4	EX7L_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9FDL5	PUR2_ZYMMO		cytoplasm
Q9FDL9	RL28_ZYMMO		cytoplasm
Q9FDM1	LGT_ZYMMO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HT05	RNPA_PSEAE		extracellular
Q9HT06	OXAA_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HT07	TRME_PSEAE		cytoplasm
Q9HT09	GIDA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HT10	GIDB_PSEAE		cytoplasm
Q9HT18	ATPA_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9HT19	ATPG_PSEAE		inner membrane
Q9HT21	ATPE_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9HT22	GLMU_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HT25	GLMS_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HT35	Y5539_PSEAE		inner membrane
Q9HT81	ENGB_PSEAE		cytoplasm
Q9HT89	YID1_PSEAE	periplasm (potential).	periplasm
Q9HTE9	GLYA1_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HTF1	LTAE_PSEAE		cytoplasm
Q9HTH0	CLS_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HTI1	Y5383_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q9HTJ0	BETI_PSEAE		cytoplasm
Q9HTJ1	BETB_PSEAE		cytoplasm
Q9HTJ2	BETA_PSEAE		cytoplasm
Q9HTL0	DBHA_PSEAE		cytoplasm
Q9HTM1	RPOZ_PSEAE		cytoplasm
Q9HTM2	KGUA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HTN2	ARGB_PSEAE	cytoplasm (probable).	cytoplasm
Q9HTN3	DUT_PSEAE		extracellular
Q9HTN5	RADC_PSEAE		cytoplasm
Q9HTN8	RL28_PSEAE		cytoplasm
Q9HTN9	RL33_PSEAE		cytoplasm
Q9HTQ0	DADA1_PSEAE		cytoplasm
Q9HTQ2	ALR2_PSEAE		cytoplasm
Q9HTR0	NORM_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HTS5	CYAY_PSEAE		inner membrane
Q9HTU7	Y5247_PSEAE		inner membrane
Q9HTV3	UBID_PSEAE		cytoplasm
Q9HTW5	YGFB_PSEAE		inner membrane
Q9HTX3	P5217_PSEAE		inner membrane
Q9HTX5	GCST_PSEAE		cytoplasm
Q9HTX6	GCSH2_PSEAE		cytoplasm
Q9HTX7	GCSP2_PSEAE		cytoplasm
Q9HTY6	GSH1_PSEAE		cytoplasm
Q9HTZ2	RIMK_PSEAE		cytoplasm
Q9HTZ6	HSLO_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HTZ7	PPCK_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HU36	FETP_PSEAE		cytoplasm
Q9HU41	HIS7_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HU42	HIS51_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HU43	HIS4_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HU44	HIS61_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HU53	GPMI_PSEAE		cytoplasm
Q9HU55	GLRX_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HU56	SECB_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HU65	GLNA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HU66	THII_PSEAE	cytoplasm (potential).	cytoplasm
Q9HU83	HUTU_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HU85	HUTH_PSEAE	cytoplasm (potential).	cytoplasm
Q9HU91	HUTI_PSEAE	cytoplasm (potential).	cytoplasm
Q9HU99	DADA2_PSEAE		cytoplasm
Q9HUA4	DTD_PSEAE	cytoplasm (probable).	cytoplasm
Q9HUA5	OPGG_PSEAE	periplasm (by similarity).	periplasm
Q9HUA6	OPGH_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HUB0	Y5073_PSEAE		inner membrane
Q9HUB4	TATB_PSEAE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9HUB5	TATA_PSEAE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9HUB6	HIS2_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUB7	HIS3_PSEAE	cytoplasm (by similarity).	cytoplasm

Q9HUB8	UBIB_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HUC0	UBIE_PSEAE		cytoplasm
Q9HUC5	HSLU_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUC6	HSLV_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUD0	RL31_PSEAE		cytoplasm
Q9HUE6	Y5023_PSEAE		inner membrane
Q9HUE8	CVRA_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9HUF1	MSRA_PSEAE		cytoplasm
Q9HUF4	GLNE_PSEAE		cytoplasm
Q9HUG6	PA99_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9HUG8	MSBA_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HUG9	HLDE_PSEAE		cytoplasm
Q9HUJ2	THIC_PSEAE		cytoplasm
Q9HUK8	PSD_PSEAE		cytoplasm
Q9HUL3	ENGC_PSEAE		cytoplasm
Q9HUL8	MUTL_PSEAE		cytoplasm
Q9HUL9	MIAA_PSEAE		cytoplasm
Q9HUM0	HFQ_PSEAE		cytoplasm
Q9HUM5	HISZ_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUM6	PURA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUM8	RLMB_PSEAE		cytoplasm
Q9HUM9	RS6_PSEAE		cytoplasm
Q9HUN0	RS18_PSEAE		cytoplasm
Q9HUN2	RL9_PSEAE		cytoplasm
Q9HUN4	ALR1_PSEAE		cytoplasm
Q9HUN6	Y4928_PSEAE		cytoplasm
Q9HUP3	NADE_PSEAE		cytoplasm
Q9HUP4	PNCB1_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUS2	UREE_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUU5	URE1_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUU6	URE2_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUU8	URE3_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUV8	PUR2_PSEAE		cytoplasm
Q9HUV9	PUR9_PSEAE		cytoplasm
Q9HUW0	FISL_PSEAE		cytoplasm
Q9HUW1	DUSB_PSEAE		cytoplasm
Q9HUW3	PRMA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HUW5	DSBD1_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HUW9	Y4841_PSEAE		extracellular
Q9HUX1	SPEA_PSEAE		cytoplasm
Q9HV00	FDHE_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HV01	SELA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HV11	Y4798_PSEAE		cytoplasm
Q9HV30	Y4778_PSEAE	cytoplasm (probable).	cytoplasm
Q9HV34	SPEE2_PSEAE		cytoplasm
Q9HV35	SPEH_PSEAE		cytoplasm

Q9HV37	LLDD_PSEAE		cytoplasm
Q9HV40	SSRP_PSEAE	cytoplasm (potential).	cytoplasm
Q9HV42	GRPE_PSEAE	cytoplasm (probable).	cytoplasm
Q9HV43	DNAK_PSEAE		cytoplasm
Q9HV44	DNAJ_PSEAE	cytoplasm (probable).	cytoplasm
Q9HV46	GREA_PSEAE		cytoplasm
Q9HV50	GLMM_PSEAE		cytoplasm
Q9HV51	TPIS_PSEAE		cytoplasm
Q9HV52	SECG_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HV53	Y4746_PSEAE		inner membrane
Q9HV56	RBFA_PSEAE	cytoplasm (potential).	cytoplasm
Q9HV58	RS15_PSEAE		cytoplasm
Q9HV61	Y4738_PSEAE		cytoplasm
Q9HV66	ACSA2_PSEAE		cytoplasm
Q9HV67	G6PI_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HV68	PAND_PSEAE		cytoplasm
Q9HV69	PANC_PSEAE		cytoplasm
Q9HV70	PANB2_PSEAE		cytoplasm
Q9HV75	GLUQ_PSEAE		cytoplasm
Q9HV77	SFSA_PSEAE		cytoplasm
Q9HV78	TRMA_PSEAE		cytoplasm
Q9HVA2	ILVC_PSEAE		cytoplasm
Q9HVA4	Y4692_PSEAE		cytoplasm
Q9HVA5	Y4691_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HVC3	PTH_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HVC4	RL25_PSEAE		cytoplasm
Q9HVC5	KPRS_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HVD0	MURI_PSEAE		cytoplasm
Q9HVD7	HEMH_PSEAE	cytoplasm (by similarity).	extracellular
Q9HVE6	UPP_PSEAE		cytoplasm
Q9HVF1	MQO2_PSEAE		cytoplasm
Q9H VH7	MSCL_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HVI7	GLYA3_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HVJ9	RTCA_PSEAE	cytoplasm (potential).	cytoplasm
Q9HVL0	Y4574_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9HVL7	RL27_PSEAE		cytoplasm
Q9HVL9	PROB_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HVM1	RS20_PSEAE		cytoplasm
Q9HVM4	SYI_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HVM5	LSPA_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9HVM7	ISPH_PSEAE		cytoplasm
Q9HVN5	CLPB_PSEAE	cytoplasm (probable).	cytoplasm
Q9HVP7	Y4530_PSEAE		cytoplasm
Q9HVP8	COAE_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HVQ7	PIUC_PSEAE		cytoplasm
Q9HVR0	Y4511_PSEAE		inner membrane

Q9HVT1	Y4490_PSEAE		cytoplasm
Q9HVT2	Y4489_PSEAE		cytoplasm
Q9HVT7	GATB_PSEAE		cytoplasm
Q9HVT8	GATA_PSEAE		cytoplasm
Q9HVT9	GATC_PSEAE		cytoplasm
Q9HVU3	Y4478_PSEAE	cytoplasm (potential).	inner membrane
Q9HVU8	Y4473_PSEAE		inner membrane
Q9HVV2	PTHP_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HVV3	Y4465_PSEAE		cytoplasm
Q9HVW7	MURA_PSEAE	cytoplasm (probable).	inner membrane
Q9HVW8	HIS1_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HVW9	HISX_PSEAE		cytoplasm
Q9HVX0	HIS81_PSEAE		cytoplasm
Q9HVX2	Y4445_PSEAE		cytoplasm
Q9HVV3	RS9_PSEAE		cytoplasm
Q9HVZ0	GMHA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HVZ1	Y4424_PSEAE		inner membrane
Q9HVZ3	Y4422_PSEAE		cytoplasm
Q9HVZ4	MRAZ_PSEAE		inner membrane
Q9HVZ5	MRAW_PSEAE		cytoplasm
Q9HVZ8	MRAY_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HVZ9	MURD_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HW01	MURG_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9HW02	MURC_PSEAE	cytoplasm (probable).	cytoplasm
Q9HW04	ARGJ_PSEAE	cytoplasm (probable).	cytoplasm
Q9HW09	PANE_PSEAE	cytoplasm (potential).	cytoplasm
Q9HW11	Y4395_PSEAE		inner membrane
Q9HW19	CRCB_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HW26	PNCB2_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HW38	ICIA_PSEAE		cytoplasm
Q9HWB8	SBCC_PSEAE		cytoplasm
Q9HWC4	NUSG_PSEAE		cytoplasm
Q9HWC5	RL11_PSEAE		cytoplasm
Q9HWC6	RL1_PSEAE		cytoplasm
Q9HWC7	RL10_PSEAE		cytoplasm
Q9HWC8	RL7_PSEAE		cytoplasm
Q9HWC9	RPOC_PSEAE		cytoplasm
Q9HWD0	RS12_PSEAE		cytoplasm
Q9HWD1	RS7_PSEAE		cytoplasm
Q9HWD4	RS10_PSEAE		cytoplasm
Q9HWD5	RL3_PSEAE		cytoplasm
Q9HWD6	RL4_PSEAE		cytoplasm
Q9HWD8	RL2_PSEAE		cytoplasm
Q9HWD9	RS19_PSEAE		cytoplasm
Q9HWE0	RL22_PSEAE		cytoplasm
Q9HWE1	RS3_PSEAE		cytoplasm
Q9HWE2	RL16_PSEAE		cytoplasm
Q9HWE3	RL29_PSEAE		cytoplasm
Q9HWE4	RS17_PSEAE		cytoplasm
Q9HWE6	RL24_PSEAE		cytoplasm

Q9HWE7	RL5_PSEAE		cytoplasm
Q9HWE8	RS14_PSEAE		cytoplasm
Q9HWE9	RS8_PSEAE		cytoplasm
Q9HWF1	RL18_PSEAE		cytoplasm
Q9HWF2	RS5_PSEAE		cytoplasm
Q9HWF3	RL30_PSEAE		cytoplasm
Q9HWF4	RL15_PSEAE		cytoplasm
Q9HWF7	RS13_PSEAE		cytoplasm
Q9HWF8	RS11_PSEAE		cytoplasm
Q9HWG0	UVRA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HWI0	DDLA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HWP3	SYI1_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HWR3	BPHY_PSEAE		inner membrane
Q9HWR7	SOTB_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HWS2	Y4106_PSEAE		inner membrane
Q9HWX1	NRDR_PSEAE		cytoplasm
Q9HWX4	RIBAB_PSEAE		cytoplasm
Q9HWX5	RISB_PSEAE		cytoplasm
Q9HWX6	NUSB_PSEAE		cytoplasm
Q9HWY1	GCH2_PSEAE		extracellular
Q9HWY5	EX7S_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HWZ3	AQPZ_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HWZ6	IPYR_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HX02	EUTC_PSEAE		cytoplasm
Q9HX08	PAAD_PSEAE		cytoplasm
Q9HX17	3MGH_PSEAE		cytoplasm
Q9HX20	PROA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HX21	NADD_PSEAE		cytoplasm
Q9HX23	Y4004_PSEAE		inner membrane
Q9HX25	LIPA_PSEAE	cytoplasm (potential).	cytoplasm
Q9HX37	Y3982_PSEAE		inner membrane
Q9HX40	THIE_PSEAE		cytoplasm
Q9HX48	RLUE_PSEAE		cytoplasm
Q9HX79	TAUB_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9HX91	P3922_PSEAE		periplasm
Q9HX95	MOAC_PSEAE		cytoplasm
Q9HX97	MOAE_PSEAE		cytoplasm
Q9HXD6	MOAA1_PSEAE		cytoplasm
Q9HXE5	RHLB_PSEAE		cytoplasm
Q9HXF2	Y3854_PSEAE		cytoplasm
Q9HXF7	NDPA_PSEAE	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q9HXH0	SYV_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HXH8	QUEA_PSEAE	cytoplasm (potential).	cytoplasm
Q9HXH9	TGT_PSEAE		cytoplasm
Q9HXI4	SUHB_PSEAE		cytoplasm
Q9HXI8	ISCS_PSEAE		cytoplasm
Q9HXJ1	HSCB_PSEAE		inner membrane
Q9HXJ4	ISPG_PSEAE		cytoplasm

Q9HXJ8	ENGA_PSEAE		cytoplasm
Q9HXK5	LEU1_PSEAE		cytoplasm
Q9HXL8	EX7L_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HXM6	GUAA_PSEAE		cytoplasm
Q9HXN2	PUR4_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HXP9	RS16_PSEAE		cytoplasm
Q9HXQ0	RIMM_PSEAE	cytoplasm (potential).	cytoplasm
Q9HXQ1	TRMD_PSEAE	cytoplasm (potential).	cytoplasm
Q9HXQ2	RL19_PSEAE		cytoplasm
Q9HXQ6	XERD_PSEAE	cytoplasm (by similarity).	nucleoid
Q9HXU7	Y3693_PSEAE		inner membrane
Q9HXV3	CAPP_PSEAE		cytoplasm
Q9HXV4	KAD_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HXW0	Y3680_PSEAE		inner membrane
Q9HXW7	PLSB_PSEAE	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9HXX3	CSD_PSEAE		cytoplasm
Q9HXY2	UPPS_PSEAE		inner membrane
Q9HXY3	Y3649_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9HXY5	SKPL_PSEAE		cytoplasm
Q9HXY6	LPXD_PSEAE		cytoplasm
Q9HXY7	FABZ_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HXY8	LPXB_PSEAE		cytoplasm
Q9HXY9	RNH2_PSEAE	cytoplasm (potential).	extracellular
Q9HXZ1	DPO3A_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HXZ2	ACCA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HXZ3	TILS_PSEAE	cytoplasm (probable).	cytoplasm
Q9HXZ4	PYRG_PSEAE		cytoplasm
Q9HXZ5	ENO_PSEAE	cytoplasm (by similarity).	extracellular
Q9HY04	TRUD_PSEAE		cytoplasm
Q9HY05	SURE_PSEAE	cytoplasm (potential).	periplasm
Q9HY08	MUTS_PSEAE		cytoplasm
Q9HY25	RL31B_PSEAE		cytoplasm
Q9HY26	RL362_PSEAE		cytoplasm
Q9HY41	GLPK2_PSEAE		cytoplasm
Q9HY42	Y3578_PSEAE		cytoplasm
Q9HY59	Y3558_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HY63	ARNA_PSEAE		cytoplasm
Q9HY65	ARNB_PSEAE		cytoplasm
Q9HY69	ALG44_PSEAE		inner membrane
Q9HY72	Y3539_PSEAE		inner membrane
Q9HY82	RNT_PSEAE		cytoplasm
Q9HY84	ASSY_PSEAE	cytoplasm (probable).	cytoplasm
Q9HYA4	ASPD_PSEAE		cytoplasm
Q9HYB5	RNFE_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HYB6	RNFG_PSEAE	bacterial cell inner membrane (potential).	inner membrane

Q9HYB7	RNFD_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9HYB8	RNFC_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	cytoplasm
Q9HYC0	RNFA_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HYC9	DCD_PSEAE		extracellular
Q9HYD5	MAO1_PSEAE		cytoplasm
Q9HYE3	Y3463_PSEAE		inner membrane
Q9HYF0	Y3456_PSEAE		cytoplasm
Q9HYF4	MQO1_PSEAE		cytoplasm
Q9HYG2	SSUD_PSEAE		cytoplasm
Q9HYG8	GCH11_PSEAE		cytoplasm
Q9HYH1	Y3435_PSEAE		cytoplasm
Q9HYH5	Y3430_PSEAE		cytoplasm
Q9HYI6	Y3419_PSEAE		cytoplasm
Q9HYL2	NOSZ1_PSEAE	periplasm (by similarity).	periplasm
Q9HYL3	NSR1_PSEAE	bacterial cell membrane; peripheral membrane protein (potential).	cytoplasm
Q9HYL7	PHNC2_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9HYN9	SDHD_PSEAE		cytoplasm
Q9HYQ8	HLDD_PSEAE		cytoplasm
Q9HYR2	FABH_PSEAE	cytoplasm (probable).	cytoplasm
Q9HYR9	CLPP2_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HYT0	PHNC1_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9HYT3	Y3311_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9HYT6	RAPA_PSEAE		cytoplasm
Q9HYV6	Y3287_PSEAE		cytoplasm
Q9HYW0	Y3283_PSEAE		inner membrane
Q9HYW6	Y3275_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HYX7	RDGC_PSEAE	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q9HYZ3	APEB_PSEAE		extracellular
Q9HYZ5	MINE_PSEAE		inner membrane
Q9HYZ7	MINC_PSEAE		inner membrane
Q9HZ00	Y3240_PSEAE		cytoplasm
Q9HZ17	ACPD3_PSEAE		extracellular
Q9HZ39	ISPZ_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HZ46	GLK_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HZ55	RLUB_PSEAE		cytoplasm
Q9HZ59	HUTG_PSEAE		cytoplasm

Q9HZ62	GPH2_PSEAE		cytoplasm
Q9HZ63	UBIG_PSEAE		cytoplasm
Q9HZ65	MTNA_PSEAE		cytoplasm
Q9HZ66	SERC_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HZ68	HIS82_PSEAE		cytoplasm
Q9HZ70	KCY_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HZ95	DUSC_PSEAE		cytoplasm
Q9HZA3	LEU2_PSEAE		cytoplasm
Q9HZA4	LEUD_PSEAE		cytoplasm
Q9HZC0	PPNK_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HZF8	PYRD_PSEAE	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9HZG5	DGTL2_PSEAE		cytoplasm
Q9HZJ2	FADB_PSEAE		cytoplasm
Q9HZJ3	FADA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9HZJ5	TOP1_PSEAE		cytoplasm
Q9HZK0	NAGZ_PSEAE	cytoplasm (by similarity).	extracellular
Q9HZK1	MTNP_PSEAE		cytoplasm
Q9HZK6	NQRA_PSEAE		inner membrane
Q9HZK7	NQRB_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9HZK8	NQRC_PSEAE	bacterial cell inner membrane (potential).	inner membrane
Q9HZK9	NQRD_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9HZL0	NQRE_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9HZL1	NQRF_PSEAE	bacterial cell inner membrane (potential).	inner membrane
Q9HZL7	LOLD_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9HZM3	LPXK_PSEAE		cytoplasm
Q9HZM5	KDSB_PSEAE	cytoplasm (potential).	cytoplasm
Q9HZM7	MURB_PSEAE	cytoplasm (probable).	inner membrane
Q9HZM9	RLUC_PSEAE		cytoplasm
Q9HZN2	Y2972_PSEAE	cytoplasm (potential).	inner membrane
Q9HZN4	RL32_PSEAE		cytoplasm
Q9HZN5	PLSX_PSEAE		cytoplasm
Q9HZN8	KTHY_PSEAE		cytoplasm
Q9HZP5	ETFD_PSEAE		cytoplasm
Q9HZP8	Y2950_PSEAE		cytoplasm
Q9HZT7	Y2910_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9HZT9	CBID_PSEAE		cytoplasm
Q9HZY5	GREB_PSEAE		cytoplasm
Q9I011	TPMT_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I013	HTPX_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9I016	MSRB_PSEAE		cytoplasm

Q9I037	QUEF_PSEAE	cytoplasm (probable).	cytoplasm
Q9I047	TAL_PSEAE	cytoplasm (probable).	cytoplasm
Q9I048	DUSA_PSEAE		cytoplasm
Q9I073	Y2770_PSEAE		cytoplasm
Q9I088	ECOT_PSEAE	periplasm (by similarity).	periplasm
Q9I0A1	RL35_PSEAE		cytoplasm
Q9I0A2	RL20_PSEAE		cytoplasm
Q9I0H4	HMP_PSEAE		cytoplasm
Q9I0J6	NUOG_PSEAE		cytoplasm
Q9I0K4	ACEA_PSEAE		cytoplasm
Q9I0L1	Y2627_PSEAE		inner membrane
Q9I0L2	TRMU_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I0L7	CLPS_PSEAE		cytoplasm
Q9I0M0	ATE_PSEAE	cytoplasm (potential).	cytoplasm
Q9I0M1	LFTR_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I0M4	LOLA_PSEAE	periplasm (by similarity).	periplasm
Q9I0N2	TUSC_PSEAE	cytoplasm (by similarity).	inner membrane
Q9I0N3	TUSD_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I0Q1	UVRC_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I0S1	TAM_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I104	DSBD2_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I106	DSBG_PSEAE	periplasm (by similarity).	periplasm
Q9I136	GCSH1_PSEAE		cytoplasm
Q9I137	GCSP1_PSEAE		cytoplasm
Q9I138	GLYA2_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I163	Y2418_PSEAE		cytoplasm
Q9I165	TREA_PSEAE	periplasm (by similarity).	periplasm
Q9I1C2	MSUD_PSEAE		cytoplasm
Q9I1L9	DLDH1_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I1Q5	PDXA2_PSEAE	cytoplasm (probable).	cytoplasm
Q9I1T4	CAAL_PSEAE		cytoplasm
Q9I1V0	GLGA_PSEAE		cytoplasm
Q9I1W2	GLGB_PSEAE		extracellular
Q9I1Z9	Y2116_PSEAE		inner membrane
Q9I203	Y2112_PSEAE		inner membrane
Q9I263	CYNS_PSEAE		cytoplasm
Q9I270	Y2045_PSEAE		inner membrane
Q9I291	GALU1_PSEAE		cytoplasm
Q9I2B6	Y1993_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I2C0	PQQE_PSEAE		cytoplasm
Q9I2C1	PQQD_PSEAE		cytoplasm
Q9I2C2	PQQC_PSEAE		cytoplasm
Q9I2C3	PQQB_PSEAE		cytoplasm
Q9I2D2	PQQF_PSEAE		extracellular
Q9I2E2	ACPD2_PSEAE		extracellular
Q9I2E5	UPPP_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I2N4	MODC_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9I2P8	Y1847_PSEAE		inner membrane

Q9I2S9	RNH_PSEAE	cytoplasm (potential).	extracellular
Q9I2U0	CLPX_PSEAE		cytoplasm
Q9I2U1	CLPP1_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I2U2	TIG_PSEAE		cytoplasm
Q9I2V0	LPXH_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I2W7	RRAA_PSEAE		cytoplasm
Q9I2X0	Y1769_PSEAE		cytoplasm
Q9I310	Y1727_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9I344	AROC_PSEAE		cytoplasm
Q9I347	Y1678_PSEAE		cytoplasm
Q9I351	GCH12_PSEAE		cytoplasm
Q9I352	HEMTL_PSEAE		cytoplasm
Q9I382	SELU_PSEAE		cytoplasm
Q9I383	SELD_PSEAE		cytoplasm
Q9I387	GLSA_PSEAE		extracellular
Q9I3A4	Y1618_PSEAE		cytoplasm
Q9I3A8	GPDA_PSEAE	cytoplasm (probable).	cytoplasm
Q9I3C3	PANB1_PSEAE		cytoplasm
Q9I3C5	HTPG_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I3D8	Y1579_PSEAE		cytoplasm
Q9I3E3	Y1574_PSEAE		inner membrane
Q9I3F3	TUSA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I3F4	MTFA_PSEAE		cytoplasm
Q9I3H7	Y1536_PSEAE		inner membrane
Q9I3H9	RECR_PSEAE		cytoplasm
Q9I3I0	Y1533_PSEAE		inner membrane
Q9I3I5	ZIPA_PSEAE	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q9I3J5	Y1518_PSEAE		cytoplasm
Q9I3J8	ALLC_PSEAE		extracellular
Q9I3J9	ALLA_PSEAE		extracellular
Q9I3K7	MOAA2_PSEAE		cytoplasm
Q9I3N1	DSBE_PSEAE	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q9I3N3	CCME_PSEAE	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q9I3N7	CCMA_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9I3W8	ACEK_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I3W9	PDXB_PSEAE		cytoplasm
Q9I3Y3	PMPM_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I407	ASPQ_PSEAE	periplasm (by similarity).	periplasm
Q9I445	Y1299_PSEAE		inner membrane
Q9I452	THTM_PSEAE	cytoplasm (probable).	cytoplasm

Q9I463	COBS_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I465	COBT_PSEAE		cytoplasm
Q9I467	COBQ_PSEAE		cytoplasm
Q9I471	COBB_PSEAE		cytoplasm
Q9I492	Y1252_PSEAE	cytoplasm (potential).	cytoplasm
Q9I4C8	Y1210_PSEAE		cytoplasm
Q9I4D3	Y1205_PSEAE		cytoplasm
Q9I4E1	NPD2_PSEAE	cytoplasm (probable).	cytoplasm
Q9I4E9	SPRT_PSEAE	cytoplasm (potential).	extracellular
Q9I4F5	DCTA2_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I4G3	NAPA_PSEAE	periplasm (by similarity).	periplasm
Q9I4K0	HCHA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I4K6	FOSA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I4L0	NPD1_PSEAE	cytoplasm (probable).	cytoplasm
Q9I4L1	DGTP_PSEAE		cytoplasm
Q9I4N6	FLISB_PSEAE		flagellar
Q9I4P4	FLGJ_PSEAE	periplasm (by similarity).	extracellular
Q9I4P5	FLGI_PSEAE	periplasm (by similarity).	periplasm
Q9I4P6	FLGH_PSEAE	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q9I4S5	PDXH_PSEAE		cytoplasm
Q9I4T5	Y1039_PSEAE		inner membrane
Q9I4U3	RMUC_PSEAE		cytoplasm
Q9I4W0	PUR7_PSEAE		cytoplasm
Q9I4W3	DAPA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I4W9	NADA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I4Z9	SLYX_PSEAE		inner membrane
Q9I507	RBN_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I513	PUR5_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I525	RUMA_PSEAE		cytoplasm
Q9I534	DPO4_PSEAE	cytoplasm (probable).	cytoplasm
Q9I540	KUP_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I558	ACSA1_PSEAE		cytoplasm
Q9I583	Y858_PSEAE		inner membrane
Q9I587	FUMC2_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I595	CYSZ_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I5A4	ACKA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I5F3	ACPD1_PSEAE		extracellular
Q9I5G5	PDXJ_PSEAE	cytoplasm (probable).	cytoplasm
Q9I5G8	LEPA_PSEAE		cytoplasm
Q9I5G9	Y760_PSEAE		cytoplasm
Q9I5H9	UNG_PSEAE	cytoplasm (by similarity).	extracellular
Q9I5J6	RSUA_PSEAE		cytoplasm
Q9I5Q2	DNAE2_PSEAE	cytoplasm (probable).	cytoplasm
Q9I5Q3	SYI2_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I5Q9	ARGC_PSEAE	cytoplasm (probable).	cytoplasm

Q9I5R7	SPED_PSEAE		cytoplasm
Q9I5U2	OSTA_PSEAE	bacterial cell outer membrane (by similarity).	outer membrane
Q9I5U4	PDXA1_PSEAE	cytoplasm (probable).	cytoplasm
Q9I5U5	KSGA_PSEAE		cytoplasm
Q9I5U6	APAG_PSEAE		inner membrane
Q9I5U7	APAH_PSEAE		extracellular
Q9I5U8	GLPE_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I5V0	Y587_PSEAE		inner membrane
Q9I5V3	CCA_PSEAE		cytoplasm
Q9I5V6	Y581_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I5V8	RS21_PSEAE		cytoplasm
Q9I5W0	PRIM_PSEAE		cytoplasm
Q9I5W9	Y567_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9I5Y1	ALF_PSEAE		cytoplasm
Q9I5Y4	PGK_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I614	BIOD_PSEAE		cytoplasm
Q9I626	Y492_PSEAE		inner membrane
Q9I633	Y485_PSEAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9I636	MASZ_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I644	Y474_PSEAE		cytoplasm
Q9I685	SAHH_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I690	Y423_PSEAE	periplasm (potential).	periplasm
Q9I697	GSHB_PSEAE		cytoplasm
Q9I699	RUVX_PSEAE	cytoplasm (potential).	cytoplasm
Q9I6A8	HAM1_PSEAE		extracellular
Q9I6B3	TRMB_PSEAE		cytoplasm
Q9I6B4	THIG_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I6B7	MTGA_PSEAE	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q9I6C8	CALB_PSEAE		cytoplasm
Q9I6D1	COAD_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I6E0	ILVD_PSEAE		cytoplasm
Q9I6F1	TYSY_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I6F2	LGT_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I6G1	RPIA_PSEAE		cytoplasm
Q9I6G2	Y329_PSEAE		outer membrane
Q9I6J9	AGUA_PSEAE		extracellular
Q9I6L0	CYSA_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9I6P3	AROQ2_PSEAE		cytoplasm
Q9I6S5	MDCG_PSEAE		cytoplasm
Q9I6S8	MDCC_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I6S9	MDCB_PSEAE		cytoplasm
Q9I6V7	CHER2_PSEAE		cytoplasm
Q9I6Y4	ADD_PSEAE		extracellular

Q9I713	DCTA1_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9I767	5NTD_PSEAE		periplasm
Q9I778	KPTA_PSEAE		cytoplasm
Q9I783	P049_PSEAE		cytoplasm
Q9I7A8	DEF_PSEAE		cytoplasm
Q9I7A9	RSMB_PSEAE		cytoplasm
Q9I7C0	GMHB_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I7C3	RECF_PSEAE	cytoplasm (potential).	cytoplasm
Q9I7C4	DPO3B_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9I7C5	DNAA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9JMT3	INF7_ECOLI		cytoplasm
Q9JMT9	YUAA_ECOLI		inner membrane
Q9JN05	NPD_CAMJE	cytoplasm (probable).	cytoplasm
Q9JP87	CARA_RHOGE		cytoplasm
Q9JP88	RUMA_RHOGE		cytoplasm
Q9JP92	SURE_RHOGE	cytoplasm (potential).	periplasm
Q9JPA2	BCHN_RHOGE		cytoplasm
Q9JPA3	BCHB_RHOGE		cytoplasm
Q9JPA5	BCHL_RHOGE		cytoplasm
Q9JPB9	BCHZ_RHOGE		cytoplasm
Q9JPD1	UBIE_RHOGE		cytoplasm
Q9JQL7	RS17_NEIMA		cytoplasm
Q9JQQ6	RL18_NEIMA		cytoplasm
Q9JQT2	RS15_NEIMA		cytoplasm
Q9JQX5	RRMJ_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JQX8	HLDD_NEIMA		cytoplasm
Q9JR00	GRPE_NEIMA	cytoplasm (probable).	cytoplasm
Q9JR26	RL16_NEIMA		cytoplasm
Q9JR30	DBHC_NEIMA		cytoplasm
Q9JRD8	RL22_NEIMA		cytoplasm
Q9JRE9	Y1193_NEIMA		cytoplasm
Q9JRG1	DPO4_NEIMA	cytoplasm (probable).	cytoplasm
Q9JRM6	COAE_ACTAC	cytoplasm (by similarity).	cytoplasm
Q9JRP9	RECA_ACTAC	cytoplasm (by similarity).	cytoplasm
Q9JRS8	YORX_ACTAC		inner membrane
Q9JRT1	CYSH_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JRV8	ACCA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JRZ6	Y1133_NEIMB		cytoplasm
Q9JS04	HSCA_NEIMB		cytoplasm
Q9JS33	CYSI_NEIMB		cytoplasm
Q9JS45	CYSJ_NEIMB		cytoplasm
Q9JS61	ILVD_NEIMB		cytoplasm
Q9JSM1	EX7S_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JSM2	ENGC_NEIMA		cytoplasm
Q9JSM5	RUVA_NEIMA		cytoplasm
Q9JSM7	RLPA_NEIMA		outer membrane
Q9JSN0	BIOH_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JSN4	TRPC_NEIMA		cytoplasm
Q9JSQ9	DYR_NEIMA		cytoplasm
Q9JSR7	QUEF_NEIMA	cytoplasm (probable).	cytoplasm
Q9JSS3	Y2158_NEIMA		cytoplasm
Q9JSS4	COAE_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JSS6	G6PI2_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JSS7	DAPD_NEIMA	cytoplasm (by similarity).	cytoplasm

Q9JST7	DTD_NEIMA	cytoplasm (probable).	cytoplasm
Q9JST9	GLUQ_NEIMA		cytoplasm
Q9JSU1	TAL_NEIMA	cytoplasm (probable).	cytoplasm
Q9JSU8	Y2126_NEIMA		cytoplasm
Q9JSX4	NADB_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JSX6	NADA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JSY6	UPPP_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JSY8	MRAZ_NEIMA		inner membrane
Q9JSY9	MRAW_NEIMA		cytoplasm
Q9JSZ0	MURE_NEIMA	cytoplasm (probable).	cytoplasm
Q9JSZ3	MRAY_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JSZ5	MURD_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JSZ7	MURG_NEIMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JSZ8	MURC_NEIMA	cytoplasm (probable).	cytoplasm
Q9JSZ9	DDL_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JT07	ACKA2_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JT15	RECO_NEIMA		cytoplasm
Q9JT25	SPEA_NEIMA		cytoplasm
Q9JT57	TYSY_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JT68	UBID_NEIMA		cytoplasm
Q9JT70	DHPS_NEIMA		cytoplasm
Q9JT71	GLMM_NEIMA		cytoplasm
Q9JT78	NUDH_NEIMA		cytoplasm
Q9JT81	AROC_NEIMA		cytoplasm
Q9JT82	TRMA_NEIMA		cytoplasm
Q9JT86	GCSP_NEIMA		cytoplasm
Q9JT96	KGUA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JTA1	Y1912_NEIMA		cytoplasm
Q9JTA3	Y1909_NEIMA		cytoplasm
Q9JTA4	Y1908_NEIMA		inner membrane
Q9JTA6	ALR_NEIMA		cytoplasm
Q9JTA9	DSBB_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JTB0	Y1902_NEIMA		inner membrane
Q9JTB8	ANIA_NEIMA	bacterial cell outer membrane; lipid-anchor (probable).	periplasm
Q9JTD9	RNH_NEIMA	cytoplasm (potential).	extracellular
Q9JTE9	Y1806_NEIMA		inner membrane
Q9JTF2	GPMA_NEIMA		cytoplasm
Q9JTH8	HIS8_NEIMA		cytoplasm
Q9JTH9	HISX_NEIMA		cytoplasm
Q9JTI3	ILVC_NEIMA		cytoplasm
Q9JTI4	OTC_NEIMA	cytoplasm (probable).	cytoplasm
Q9JTI8	AMPA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JTJ6	GSHB_NEIMA		cytoplasm
Q9JTJ9	TRMU_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JTK1	PYRG_NEIMA		cytoplasm

Q9JTL9	DSBD_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JTM0	ACKA1_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JTM3	ISPD_NEIMA		cytoplasm
Q9JTM4	ISPF_NEIMA		extracellular
Q9JTM5	RPIA_NEIMA		cytoplasm
Q9JTM8	PNCB_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JTN2	RNPH_NEIMA		cytoplasm
Q9JTP0	SURE_NEIMA	cytoplasm (potential).	periplasm
Q9JTP4	RECX_NEIMA	cytoplasm (potential).	cytoplasm
Q9JTP5	GPH_NEIMA		cytoplasm
Q9JTP9	CLPB_NEIMA	cytoplasm (probable).	cytoplasm
Q9JTR0	FUMC_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JTR9	AROD_NEIMA		cytoplasm
Q9JTS0	Y1657_NEIMA		inner membrane
Q9JTS2	MUTL_NEIMA		cytoplasm
Q9JTT3	AROA_NEIMA	cytoplasm (probable).	cytoplasm
Q9JTT4	GREA_NEIMA		cytoplasm
Q9JTU3	RUVC_NEIMA		cytoplasm
Q9JTW1	G6P11_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JTW6	HSCB_NEIMA		inner membrane
Q9JTX0	ISCS_NEIMA		cytoplasm
Q9JTX5	TRUB_NEIMA		cytoplasm
Q9JTX7	RBFA_NEIMA	cytoplasm (potential).	cytoplasm
Q9JTX8	CLPX_NEIMA		cytoplasm
Q9JTX9	ARGD_NEIMA	cytoplasm (probable).	cytoplasm
Q9JTY8	EX7L_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JTZ1	PDXH_NEIMA		cytoplasm
Q9JTZ3	GATB_NEIMA		cytoplasm
Q9JTZ5	GATA_NEIMA		cytoplasm
Q9JTZ6	GATC_NEIMA		cytoplasm
Q9JU03	SUHB_NEIMA		cytoplasm
Q9JU11	RUVX_NEIMA	cytoplasm (potential).	cytoplasm
Q9JU12	Y1550_NEIMA		inner membrane
Q9JU19	TRMB_NEIMA		cytoplasm
Q9JU21	UVRC_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JU24	RS6_NEIMA		cytoplasm
Q9JU25	PRIB_NEIMA		cytoplasm
Q9JU26	RL9_NEIMA		cytoplasm
Q9JU32	TIG_NEIMA		cytoplasm
Q9JU33	CLPP_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JU34	ISPG_NEIMA		cytoplasm
Q9JU41	PLSC_NEIMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JU46	ENO_NEIMA	cytoplasm (by similarity).	extracellular
Q9JU48	KDSA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JU49	PAND_NEIMA		cytoplasm
Q9JU79	LEU3_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JU81	LEUD_NEIMA		cytoplasm
Q9JU82	LEU2_NEIMA		cytoplasm
Q9JU84	RADC_NEIMA		cytoplasm
Q9JU97	RIBAB_NEIMA		cytoplasm
Q9JU99	GCH2_NEIMA		extracellular
Q9JUA2	PUR5_NEIMA	cytoplasm (by similarity).	cytoplasm

Q9JUA5	MOBA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUB0	RUVB_NEIMA		cytoplasm
Q9JUB2	CCA_NEIMA		cytoplasm
Q9JUB6	RECR_NEIMA		cytoplasm
Q9JUC4	UNG_NEIMA	cytoplasm (by similarity).	extracellular
Q9JUC7	LIPB_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUC8	LIPA_NEIMA	cytoplasm (potential).	cytoplasm
Q9JUC9	GLND_NEIMA		cytoplasm
Q9JUD5	CYSH_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUD8	CYSJ_NEIMA		cytoplasm
Q9JUD9	CYSI_NEIMA		cytoplasm
Q9JUE0	ILVD_NEIMA		cytoplasm
Q9JUE9	TILS_NEIMA	cytoplasm (probable).	cytoplasm
Q9JUF0	ACCA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUF2	Y1343_NEIMA		cytoplasm
Q9JUF4	HSCA_NEIMA		cytoplasm
Q9JUK2	ARGB_NEIMA	cytoplasm (probable).	cytoplasm
Q9JUK4	LGT_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JUK6	LEU1_NEIMA		cytoplasm
Q9JUK7	PROB_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUK8	PROA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUL1	CRCB_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JUL4	Y1261_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JUP6	DUSC_NEIMA		cytoplasm
Q9JUQ8	PUR9_NEIMA		cytoplasm
Q9JUS3	PSD_NEIMA		cytoplasm
Q9JUS4	UVRA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUT0	SUCC_NEIMA		cytoplasm
Q9JUT6	METE_NEIMA		cytoplasm
Q9JUT8	RL31B_NEIMA		cytoplasm
Q9JUT9	METX_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUU5	MIAA_NEIMA		cytoplasm
Q9JUU8	RLMB_NEIMA		cytoplasm
Q9JUU9	DAPA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUV1	PIP_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JUV5	LST_NEIMA		cytoplasm
Q9JUV9	Y1114_NEIMA		inner membrane
Q9JUW1	DUT_NEIMA		extracellular
Q9JUX4	CYSA_NEIMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JUX7	RL25_NEIMA		cytoplasm
Q9JUX8	ISPE_NEIMA		cytoplasm
Q9JUY0	PANB_NEIMA		cytoplasm
Q9JV01	ENGA_NEIMA		cytoplasm
Q9JV02	RDGC_NEIMA	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q9JV11	Y1049_NEIMA		inner membrane
Q9JV14	CLPS_NEIMA		cytoplasm

Q9JV18	PYRF_NEIMA		cytoplasm
Q9JV19	HTPX_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9JV25	PURA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JV26	HISZ_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JV27	NORM_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JV28	MURB_NEIMA	cytoplasm (probable).	inner membrane
Q9JV35	Y1013_NEIMA		inner membrane
Q9JV42	PTH_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JV52	DCUP_NEIMA	cytoplasm (probable).	cytoplasm
Q9JV58	UPP_NEIMA		cytoplasm
Q9JV65	LEPA_NEIMA		cytoplasm
Q9JV69	DAPF_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JV73	PUR7_NEIMA		cytoplasm
Q9JV76	XERD_NEIMA	cytoplasm (by similarity).	nucleoid
Q9JV83	UBIE_NEIMA		cytoplasm
Q9JV90	Y948_NEIMA		cytoplasm
Q9JV91	HPRK_NEIMA		cytoplasm
Q9JV95	BIOD_NEIMA		cytoplasm
Q9JVA1	RL20_NEIMA		cytoplasm
Q9JVA4	TGT_NEIMA		cytoplasm
Q9JVA5	HEMH_NEIMA	cytoplasm (by similarity).	extracellular
Q9JVA8	LNT_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JVB6	RLUD_NEIMA		cytoplasm
Q9JVB7	COML_NEIMA	bacterial cell membrane; lipid-anchor (by similarity).	outer membrane
Q9JVC0	TRPB_NEIMA		cytoplasm
Q9JVC2	KSGA_NEIMA		cytoplasm
Q9JVD0	GREB_NEIMA		cytoplasm
Q9JVD1	TRPF_NEIMA		cytoplasm
Q9JVD2	ERA_NEIMA		cytoplasm
Q9JVD3	RNC_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVD5	NUSB_NEIMA		cytoplasm
Q9JVD6	PYRC_NEIMA		cytoplasm
Q9JVE0	TRPA_NEIMA		cytoplasm
Q9JVE4	LPXK_NEIMA		cytoplasm
Q9JVE7	KTHY_NEIMA		cytoplasm
Q9JVF4	APAH_NEIMA		extracellular
Q9JVG3	IPYR_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVG5	HAM1_NEIMA		extracellular
Q9JVG7	ARLY_NEIMA	cytoplasm (probable).	cytoplasm
Q9JVH1	FBPC_NEIMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JVH3	HIS5_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVH4	HIS4_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVH5	HIS6_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVH6	HIS3_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVI4	Y827_NEIMA		cytoplasm
Q9JVJ9	HIS2_NEIMA	cytoplasm (by similarity).	cytoplasm

Q9JVK1	TATB_NEIMA	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9JVK3	Y802_NEIMA	cytoplasm (potential).	inner membrane
Q9JVK9	RIMM_NEIMA	cytoplasm (potential).	cytoplasm
Q9JVL0	TRMD_NEIMA	cytoplasm (potential).	cytoplasm
Q9JVL1	RL19_NEIMA		cytoplasm
Q9JVP1	GCSH_NEIMA		cytoplasm
Q9JVP2	GCST_NEIMA		cytoplasm
Q9JVP8	NQRA_NEIMA		inner membrane
Q9JVP9	NQRB_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9JVQ0	NQRC_NEIMA	bacterial cell inner membrane (potential).	inner membrane
Q9JVQ1	NQRD_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9JVQ2	NQRE_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9JVQ3	NQRF_NEIMA	bacterial cell inner membrane (potential).	inner membrane
Q9JVQ5	UBIB_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JVQ9	DNAK_NEIMA		cytoplasm
Q9JVS0	LPXH_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVS4	HEM3_NEIMA		cytoplasm
Q9JVS5	Y717_NEIMA		inner membrane
Q9JVT3	NAGZ_NEIMA	cytoplasm (by similarity).	extracellular
Q9JVT9	RBN_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JVU6	ARGC_NEIMA	cytoplasm (probable).	cytoplasm
Q9JVU8	SECB_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVU9	GLRX_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVW1	ENGB_NEIMA		cytoplasm
Q9JVW4	PILQ_NEIMA	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
Q9JVW5	AROB_NEIMA	cytoplasm (probable).	cytoplasm
Q9JVW7	NRDR_NEIMA		cytoplasm
Q9JVX8	DPO3A_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVV4	SYI_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVV6	SYI_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JVV8	FTHS_NEIMA		cytoplasm
Q9JVZ6	CARA_NEIMA		cytoplasm
Q9JW02	CARB_NEIMA		cytoplasm
Q9JW05	QUEA_NEIMA	cytoplasm (potential).	cytoplasm
Q9JW08	PRMA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JW09	ORN_NEIMA	cytoplasm (potential).	cytoplasm
Q9JW10	GSA_NEIMA	cytoplasm (potential).	cytoplasm
Q9JW13	DXS_NEIMA		cytoplasm
Q9JW14	XERC_NEIMA	cytoplasm (by similarity).	nucleoid
Q9JW21	ARGA_NEIMA	cytoplasm (probable).	cytoplasm

Q9JW31	TPIS_NEIMA		cytoplasm
Q9JW42	PPK_NEIMA		cytoplasm
Q9JW45	DNAA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JW46	RNPA_NEIMA		extracellular
Q9JW48	OXAA_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JW50	Y546_NEIMA	cytoplasm (potential).	inner membrane
Q9JW52	RL32_NEIMA		cytoplasm
Q9JW54	PLSX_NEIMA		cytoplasm
Q9JW56	FABH_NEIMA	cytoplasm (probable).	cytoplasm
Q9JW59	MSBA_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JW60	GUAA_NEIMA		cytoplasm
Q9JW69	ATPE_NEIMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JW71	ATPG_NEIMA		inner membrane
Q9JW72	ATPA_NEIMA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JW77	PARB_NEIMA		cytoplasm
Q9JW78	PAAD_NEIMA		cytoplasm
Q9JWA2	HPUB_NEIMA	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
Q9JWA3	CH10_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JWA6	DCDA_NEIMA		cytoplasm
Q9JWA8	Y465_NEIMA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9JWB0	LUXS_NEIMA		extracellular
Q9JWB7	TRME_NEIMA		cytoplasm
Q9JWC5	PUR4_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JWC8	HSLO_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JWD6	VSR_NEIMA		extracellular
Q9JWE5	KHSE_NEIMA		cytoplasm
Q9JWE6	UBIG_NEIMA		cytoplasm
Q9JWE9	GMHB_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JWF1	TRUA_NEIMA		cytoplasm
Q9JWF3	THIC_NEIMA		cytoplasm
Q9JWG3	RMUC_NEIMA		inner membrane
Q9JWG6	Y382_NEIMA		cytoplasm
Q9JWH0	GPDA_NEIMA	cytoplasm (probable).	cytoplasm
Q9JWH1	CAPP_NEIMA		cytoplasm
Q9JWH4	SLYX_NEIMA		inner membrane
Q9JWI2	THIE_NEIMA		cytoplasm
Q9JWI4	THIG_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JWJ7	Y342_NEIMA		cytoplasm
Q9JWJ8	Y341_NEIMA		inner membrane
Q9JWK3	MQO_NEIMA		cytoplasm
Q9JWM1	ASSY_NEIMA	cytoplasm (probable).	cytoplasm
Q9JWM8	MSRAB_NEIMA		cytoplasm
Q9JWN3	GLMU_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JWN9	GLMS_NEIMA	cytoplasm (by similarity).	cytoplasm

Q9JWS2	LPXC_NEIMA		cytoplasm
Q9JWS7	MURA_NEIMA	cytoplasm (probable).	inner membrane
Q9JWS8	PGK_NEIMA	cytoplasm (potential).	cytoplasm
Q9JWT7	MUTS_NEIMA		cytoplasm
Q9JWU6	PUR2_NEIMA		cytoplasm
Q9JWV1	Y228_NEIMA		inner membrane
Q9JWW8	SACB1_NEIMA		cytoplasm
Q9JWY6	PYRI_NEIMA		cytoplasm
Q9JWY9	FMT_NEIMA		cytoplasm
Q9JX02	RL11_NEIMA		cytoplasm
Q9JX03	RPOC_NEIMA		cytoplasm
Q9JX12	RL2_NEIMA		cytoplasm
Q9JX13	RL5_NEIMA		cytoplasm
Q9JX15	RL15_NEIMA		cytoplasm
Q9JX17	MINC_NEIMA		inner membrane
Q9JX18	MINE_NEIMA		inner membrane
Q9JX22	SYV_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JX23	ZUPT_NEIMA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9JX24	DADA_NEIMA		cytoplasm
Q9JX26	LPXA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JX28	FABZ_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JX29	LPXD_NEIMA		cytoplasm
Q9JX32	Y084_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9JX33	DXR_NEIMA		cytoplasm
Q9JX35	UPPS_NEIMA		inner membrane
Q9JX38	GIDB_NEIMA		cytoplasm
Q9JX40	RNH2_NEIMA	cytoplasm (potential).	extracellular
Q9JX41	GIDA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JX42	PDXA_NEIMA	cytoplasm (probable).	cytoplasm
Q9JX45	LPXB_NEIMA		cytoplasm
Q9JX48	DAPB_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JX50	LFTR_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9JX59	Y052_NEIMA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JX66	PYRD_NEIMA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JX72	GLNE_NEIMA		cytoplasm
Q9JX80	NUOC_NEIMA		inner membrane
Q9JX92	NUOL_NEIMA	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9JX94	MUTS_NEIMB		cytoplasm
Q9JXA3	PUR2_NEIMB		cytoplasm
Q9JXB0	Y2142_NEIMB		inner membrane
Q9JXC1	ASSY_NEIMB	cytoplasm (probable).	cytoplasm
Q9JXD7	MQO_NEIMB		cytoplasm
Q9JXE2	Y2089_NEIMB		inner membrane
Q9JXE3	Y2088_NEIMB		cytoplasm
Q9JXF5	THIG_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXF7	THIE_NEIMB		cytoplasm

Q9JXG3	SLYX_NEIMB		inner membrane
Q9JXG5	CAPP_NEIMB		cytoplasm
Q9JXG6	GPDA_NEIMB	cytoplasm (probable).	cytoplasm
Q9JXG9	Y2054_NEIMB		cytoplasm
Q9JXH2	RMUC_NEIMB		inner membrane
Q9JXI0	THIC_NEIMB		cytoplasm
Q9JXI2	TRUA_NEIMB		cytoplasm
Q9JXI5	GMHB_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXI7	UBIG_NEIMB		cytoplasm
Q9JXK1	HSLO_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXK5	PUR4_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXL4	TRME_NEIMB		cytoplasm
Q9JXL8	LUXS_NEIMB		extracellular
Q9JXM0	Y1979_NEIMB	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9JXM2	DCDA_NEIMB		cytoplasm
Q9JXM4	CH10_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXP4	PAAD_NEIMB		cytoplasm
Q9JXP5	PARB_NEIMB		cytoplasm
Q9JXQ0	ATPA_NEIMB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JXQ1	ATPG_NEIMB		inner membrane
Q9JXQ3	ATPE_NEIMB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JXR2	GUAA_NEIMB		cytoplasm
Q9JXR3	MSBA_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JXR6	FABH_NEIMB	cytoplasm (probable).	cytoplasm
Q9JXR8	PLSX_NEIMB		cytoplasm
Q9JXS0	RL32_NEIMB		cytoplasm
Q9JXS2	Y1909_NEIMB	cytoplasm (potential).	inner membrane
Q9JXS4	OXAA_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JXS6	RNPA_NEIMB		extracellular
Q9JXS7	DNAA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXS9	PPK_NEIMB		cytoplasm
Q9JXT8	TPIS_NEIMB		cytoplasm
Q9JXU9	ARGA_NEIMB	cytoplasm (probable).	cytoplasm
Q9JXV6	XERC_NEIMB	cytoplasm (by similarity).	nucleoid
Q9JXV7	DXS_NEIMB		cytoplasm
Q9JXW0	GSA_NEIMB	cytoplasm (potential).	cytoplasm
Q9JXW1	ORN_NEIMB	cytoplasm (potential).	cytoplasm
Q9JXW2	PRMA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXW4	QUEA_NEIMB	cytoplasm (potential).	cytoplasm
Q9JXW8	CARB_NEIMB		cytoplasm
Q9JXX4	CARA_NEIMB		cytoplasm
Q9JXY2	FTHS_NEIMB		cytoplasm
Q9JXY5	SYI_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXY7	SYI_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXZ2	DPO3A_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JXZ9	NRDR_NEIMB		cytoplasm

Q9JY01	AROB_NEIMB	cytoplasm (probable).	cytoplasm
Q9JY03	ENGB_NEIMB		cytoplasm
Q9JY15	GLRX_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JY16	SECB_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JY18	ARGC_NEIMB	cytoplasm (probable).	cytoplasm
Q9JY72	TYSY_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JY86	UBID_NEIMB		cytoplasm
Q9JY89	GLMM_NEIMB		cytoplasm
Q9JY96	NUDH_NEIMB		cytoplasm
Q9JY99	AROC_NEIMB		cytoplasm
Q9JYA0	TRMA_NEIMB		cytoplasm
Q9JYB5	KGUA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JYC0	Y1655_NEIMB		cytoplasm
Q9JYC2	Y1653_NEIMB		cytoplasm
Q9JYC3	Y1652_NEIMB		inner membrane
Q9JYC4	ALR_NEIMB		cytoplasm
Q9JYC6	DSBB_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JYC7	Y1648_NEIMB		inner membrane
Q9JYE1	ANIA_NEIMB	bacterial cell outer membrane; lipid-anchor (probable).	periplasm
Q9JYE5	RNH_NEIMB	cytoplasm (potential).	extracellular
Q9JYF4	Y1608_NEIMB		inner membrane
Q9JYF7	GPMA_NEIMB		cytoplasm
Q9JYH7	HIS8_NEIMB		cytoplasm
Q9JYH8	HISX_NEIMB		cytoplasm
Q9JYI2	ILVC_NEIMB		cytoplasm
Q9JYI3	OTC_NEIMB	cytoplasm (probable).	cytoplasm
Q9JYJ3	GSHB_NEIMB		cytoplasm
Q9JYJ6	TRMU_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JYJ8	PYRG_NEIMB		cytoplasm
Q9JYM0	DSBD_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JYM1	ACKA1_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JYM4	ISPD_NEIMB		cytoplasm
Q9JYM5	ISPF_NEIMB		extracellular
Q9JYM6	RPIA_NEIMB		cytoplasm
Q9JYM9	PNCB_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JYN5	RNPH_NEIMB		cytoplasm
Q9JYP8	SURE_NEIMB	cytoplasm (potential).	periplasm
Q9JYQ3	RECX_NEIMB	cytoplasm (potential).	cytoplasm
Q9JYQ8	CLPB_NEIMB	cytoplasm (probable).	cytoplasm
Q9JYS8	DPO4_NEIMB	cytoplasm (probable).	cytoplasm
Q9JYT0	AROD_NEIMB		cytoplasm
Q9JYT1	Y1444_NEIMB		inner membrane
Q9JYT2	MUTL_NEIMB		cytoplasm
Q9JYU1	AROA_NEIMB	cytoplasm (probable).	cytoplasm
Q9JYU3	GREA_NEIMB		cytoplasm
Q9JYV1	RUVC_NEIMB		cytoplasm
Q9JYV5	FRPC_NEIMB	bacterial cell outer membrane; peripheral membrane protein (by similarity). secreted protein (by similarity).	extracellular
Q9JYX3	G6P11_NEIMB	cytoplasm (by similarity).	cytoplasm

Q9JYX6	HSCB_NEIMB		inner membrane
Q9JYY0	ISCS_NEIMB		cytoplasm
Q9JYY1	TRUB_NEIMB		cytoplasm
Q9JYY2	RBFA_NEIMB	cytoplasm (potential).	cytoplasm
Q9JYY3	CLPX_NEIMB		cytoplasm
Q9JYY4	ARGD_NEIMB	cytoplasm (probable).	cytoplasm
Q9JYZ2	EX7L_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JYZ5	PDXH_NEIMB		cytoplasm
Q9JYZ7	GATB_NEIMB		cytoplasm
Q9JYZ9	GATA_NEIMB		cytoplasm
Q9JZ00	GATC_NEIMB		cytoplasm
Q9JZ07	SUHB_NEIMB		cytoplasm
Q9JZ16	RUVX_NEIMB	cytoplasm (potential).	cytoplasm
Q9JZ17	Y1336_NEIMB		inner membrane
Q9JZ24	TRMB_NEIMB		cytoplasm
Q9JZ26	UVRC_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZ29	RS6_NEIMB		cytoplasm
Q9JZ30	PRIB_NEIMB		cytoplasm
Q9JZ31	RL9_NEIMB		cytoplasm
Q9JZ37	TIG_NEIMB		cytoplasm
Q9JZ38	CLPP_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZ40	ISPG_NEIMB		cytoplasm
Q9JZ47	PLSC_NEIMB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JZ53	ENO_NEIMB	cytoplasm (by similarity).	extracellular
Q9JZ55	KDSA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZ56	PAND_NEIMB		cytoplasm
Q9JZ77	RIBAB_NEIMB		cytoplasm
Q9JZ78	GCH2_NEIMB		extracellular
Q9JZ80	PUR5_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZ86	RUVB_NEIMB		cytoplasm
Q9JZ88	CCA_NEIMB		cytoplasm
Q9JZ92	RECR_NEIMB		cytoplasm
Q9JZA1	UNG_NEIMB	cytoplasm (by similarity).	extracellular
Q9JZA4	LIPB_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZA5	LIPA_NEIMB	cytoplasm (potential).	cytoplasm
Q9JZB4	GLND_NEIMB		cytoplasm
Q9JZF7	ARGB_NEIMB	cytoplasm (probable).	cytoplasm
Q9JZF9	LGT_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JZG1	LEU1_NEIMB		cytoplasm
Q9JZG2	PROB_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZG3	PROA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZG6	CRCB_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JZG9	Y1062_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9JZI3	RADC_NEIMB		cytoplasm
Q9JZI5	LEU2_NEIMB		cytoplasm
Q9JZI6	LEUD_NEIMB		cytoplasm
Q9JZI9	LEU3_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZL5	DUSC_NEIMB		cytoplasm

Q9JZM7	PUR9_NEIMB		cytoplasm
Q9JZP0	PSD_NEIMB		cytoplasm
Q9JZP1	UVRA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZP4	SUCC_NEIMB		cytoplasm
Q9JZQ2	METE_NEIMB		cytoplasm
Q9JZQ4	RL31B_NEIMB		cytoplasm
Q9JZQ5	METX_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZR0	MIAA_NEIMB		cytoplasm
Q9JZR3	RLMB_NEIMB		cytoplasm
Q9JZR4	DAPA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZR6	PIP_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9JZU5	Y895_NEIMB		inner membrane
Q9JZU7	DUT_NEIMB		extracellular
Q9JZW0	CYSA_NEIMB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9JZW3	RL25_NEIMB		cytoplasm
Q9JZW4	ISPE_NEIMB		cytoplasm
Q9JZW6	PANB_NEIMB		cytoplasm
Q9JZY1	ENGA_NEIMB		cytoplasm
Q9JZY2	RDGC_NEIMB	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q9JZZ2	Y840_NEIMB		inner membrane
Q9JZZ5	CLPS_NEIMB		cytoplasm
Q9K002	HLDD_NEIMB		cytoplasm
Q9K005	PYRF_NEIMB		cytoplasm
Q9K006	HTPX_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9K012	PURA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K013	HISZ_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K015	NORM_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9K016	MURB_NEIMB	cytoplasm (probable).	inner membrane
Q9K023	Y803_NEIMB		inner membrane
Q9K029	PTH_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K041	DCUP_NEIMB	cytoplasm (probable).	cytoplasm
Q9K048	UPP_NEIMB		cytoplasm
Q9K055	LEPA_NEIMB		cytoplasm
Q9K060	DAPF_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K063	PUR7_NEIMB		cytoplasm
Q9K068	XERD_NEIMB	cytoplasm (by similarity).	nucleoid
Q9K075	UBIE_NEIMB		cytoplasm
Q9K080	Y738_NEIMB		cytoplasm
Q9K081	HPRK_NEIMB		cytoplasm
Q9K085	BIOD_NEIMB		cytoplasm
Q9K093	RL20_NEIMB		cytoplasm
Q9K096	TGT_NEIMB		cytoplasm
Q9K097	HEMH_NEIMB	cytoplasm (by similarity).	extracellular
Q9K0A2	LNT_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9K0B0	RLUD_NEIMB		cytoplasm

Q9K0B1	COML_NEIMB	bacterial cell membrane; lipid-anchor (by similarity).	outer membrane
Q9K0B5	TRPB_NEIMB		cytoplasm
Q9K0B7	KSGA_NEIMB		cytoplasm
Q9K0C5	GREB_NEIMB		cytoplasm
Q9K0C6	TRPF_NEIMB		cytoplasm
Q9K0C7	ERA_NEIMB		cytoplasm
Q9K0C8	RNC_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0D0	NUSB_NEIMB		cytoplasm
Q9K0D1	PYRC_NEIMB		cytoplasm
Q9K0D4	TRPA_NEIMB		cytoplasm
Q9K0D7	LPXK_NEIMB		cytoplasm
Q9K0D9	KTHY_NEIMB		cytoplasm
Q9K0G4	IPYR_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0G6	HAM1_NEIMB		extracellular
Q9K0G8	ARLY_NEIMB	cytoplasm (probable).	cytoplasm
Q9K0H2	HIS5_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0H3	HIS4_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0H4	HIS6_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0H5	HIS3_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0I1	Y619_NEIMB		cytoplasm
Q9K0J4	HIS2_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0J6	TATB_NEIMB	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9K0J8	Y598_NEIMB	cytoplasm (potential).	inner membrane
Q9K0K3	RIMM_NEIMB	cytoplasm (potential).	cytoplasm
Q9K0K4	TRMD_NEIMB	cytoplasm (potential).	cytoplasm
Q9K0K5	RL19_NEIMB		cytoplasm
Q9K0K9	FRPA_NEIMB	bacterial cell outer membrane; peripheral membrane protein (by similarity). secreted protein (by similarity).	extracellular
Q9K0L7	GCSH_NEIMB		cytoplasm
Q9K0L8	GCST_NEIMB		cytoplasm
Q9K0M3	NQRA_NEIMB		inner membrane
Q9K0M4	NQRB_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9K0M5	NQRC_NEIMB	bacterial cell inner membrane (potential).	inner membrane
Q9K0M6	NQRD_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9K0M7	NQRE_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9K0M8	NQRF_NEIMB	bacterial cell inner membrane (potential).	inner membrane
Q9K0N0	UBIB_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9K0N4	DNAK_NEIMB		cytoplasm
Q9K0P2	LPXH_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0P6	HEM3_NEIMB		cytoplasm
Q9K0P7	Y538_NEIMB		inner membrane

Q9K0Q4	NAGZ_NEIMB	cytoplasm (by similarity).	extracellular
Q9K0R0	RBN_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9K0U3	SPEA_NEIMB		cytoplasm
Q9K0V0	TBPB_NEIMB	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q9K0V9	PDXJ_NEIMB	cytoplasm (probable).	cytoplasm
Q9K0W0	RECO_NEIMB		cytoplasm
Q9K0X1	ACKA2_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0Y0	DDL_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0Y1	MURC_NEIMB	cytoplasm (probable).	cytoplasm
Q9K0Y2	MURG_NEIMB	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9K0Y4	MURD_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K0Y6	MRAY_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9K0Y9	MURE_NEIMB	cytoplasm (probable).	cytoplasm
Q9K0Z0	MRAW_NEIMB		cytoplasm
Q9K0Z1	MRAZ_NEIMB		inner membrane
Q9K0Z3	UPPP_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9K105	NADA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K107	NADB_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K132	Y361_NEIMB		cytoplasm
Q9K139	TAL_NEIMB	cytoplasm (probable).	cytoplasm
Q9K141	GLUQ_NEIMB		cytoplasm
Q9K143	DTD_NEIMB	cytoplasm (probable).	cytoplasm
Q9K152	DAPD_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K153	G6PI2_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K155	Y330_NEIMB		cytoplasm
Q9K161	QUEF_NEIMB	cytoplasm (probable).	cytoplasm
Q9K168	DYR_NEIMB		cytoplasm
Q9K192	TRPC_NEIMB		cytoplasm
Q9K197	BIOH_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K1A0	RLPA_NEIMB		outer membrane
Q9K1A2	RUVA_NEIMB		cytoplasm
Q9K1A4	ENG_C_NEIMB		cytoplasm
Q9K1A5	EX7S_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K1B0	NUOL_NEIMB	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9K1C1	NUOC_NEIMB		inner membrane
Q9K1D5	GLNE_NEIMB		cytoplasm
Q9K1D7	PYRD_NEIMB	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9K1E1	Y215_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9K1E9	LFTR_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K1F1	DAPB_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K1F5	LPXB_NEIMB		cytoplasm

Q9K1F9	PDXA_NEIMB	cytoplasm (probable).	cytoplasm
Q9K1G0	GIDA_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K1G1	RNH2_NEIMB	cytoplasm (potential).	extracellular
Q9K1G3	GIDB_NEIMB		cytoplasm
Q9K1G6	UPPS_NEIMB		inner membrane
Q9K1G8	DXR_NEIMB		cytoplasm
Q9K1G9	Y183_NEIMB	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9K1H5	DADA_NEIMB		cytoplasm
Q9K1H6	ZUPT_NEIMB	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9K1H7	SYV_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K1H9	MINE_NEIMB		inner membrane
Q9K1I0	MINC_NEIMB		inner membrane
Q9K1I2	RL15_NEIMB		cytoplasm
Q9K1I4	RL5_NEIMB		cytoplasm
Q9K1I5	RL2_NEIMB		cytoplasm
Q9K1J1	RPOC_NEIMB		cytoplasm
Q9K1J3	RL11_NEIMB		cytoplasm
Q9K1K6	FMT_NEIMB		cytoplasm
Q9K1K9	PYRI_NEIMB		cytoplasm
Q9K1N8	MSRAB_NEIMB		cytoplasm
Q9K1P3	GLMU_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K1P9	GLMS_NEIMB	cytoplasm (by similarity).	cytoplasm
Q9K1Q3	LPXC_NEIMB		cytoplasm
Q9K1Q9	MURA_NEIMB	cytoplasm (probable).	inner membrane
Q9K1R0	PGK_NEIMB	cytoplasm (potential).	cytoplasm
Q9K2W5	AMPA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9K2Y1	TOLC_VIBCH	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
Q9K323	CEP_VIBCH	secreted protein (probable).	inner membrane
Q9K3D6	ARGHA_MORAB	cytoplasm (probable).	cytoplasm
Q9K4Q3	IHFA_MYXXA		cytoplasm
Q9K4U8	NORR2_RALEU		cytoplasm
Q9K4V0	NORR1_RALEU		cytoplasm
Q9K4V1	GSTI_RHILE		cytoplasm
Q9K4Y8	ASSY_MORAB	cytoplasm (probable).	cytoplasm
Q9K4Y9	OTC_MORAB	cytoplasm (probable).	cytoplasm
Q9K4Z0	ARGB_MORAB	cytoplasm (probable).	cytoplasm
Q9K4Z1	ARGC_MORAB	cytoplasm (probable).	cytoplasm
Q9K4Z2	ARGE_MORAB	cytoplasm (probable).	extracellular
Q9K4Z3	ASSY_MORPR	cytoplasm (probable).	cytoplasm
Q9K4Z4	OTC_MORPR	cytoplasm (probable).	cytoplasm
Q9K4Z5	ARGB_MORPR	cytoplasm (probable).	cytoplasm
Q9K4Z6	ARGC_MORPR	cytoplasm (probable).	cytoplasm
Q9K4Z7	ARGE_MORPR	cytoplasm (probable).	extracellular
Q9KGS7	INH_PSEBR	periplasm (by similarity).	periplasm
Q9KGU6	DXR_PSEAE		cytoplasm
Q9KGU7	DXS_PSEAE		cytoplasm
Q9KGW3	NUOG_PSEFL		cytoplasm
Q9KH12	PURL_RHIFR	cytoplasm (by similarity).	cytoplasm
Q9KH33	IMDH_RHITR		cytoplasm
Q9KHL6	LPXC_RALSO		cytoplasm

Q9KHR3	SSUD_BUTSP		cytoplasm
Q9KHS2	SSUD_PSESP		cytoplasm
Q9KHS6	DBHB_PSEF5		cytoplasm
Q9KHV6	T2A1_RUEGE		extracellular
Q9KI21	SURE_COXBU	cytoplasm (potential).	periplasm
Q9KIQ6	YPTO_VIBHA		cytoplasm
Q9KIT1	FLR_DESGI		cytoplasm
Q9KIZ4	C167_POLCB		cytoplasm
Q9KJC1	ARPC_PSEPU	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
Q9KJC2	ARPB_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9KJC3	ARPA_PSEPU	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
Q9KJC4	ARPR_PSEPU		cytoplasm
Q9KJE8	BBSG_THAAR		cytoplasm
Q9KJG4	RPOB_BARHE		cytoplasm
Q9KJH8	PHAG_PSEOL		cytoplasm
Q9KJM5	RPOB_BARQU		cytoplasm
Q9KJV4	FLIB_SALCH		cytoplasm
Q9KJZ3	DSBD_PSEUK	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KK72	NUDH_BARCL		cytoplasm
Q9KKL3	CHER3_VIBCH		cytoplasm
Q9KKM4	PDXH_VIBCH		cytoplasm
Q9KKP2	RIBB_VIBCH		cytoplasm
Q9KKP3	Y3859_VIBCH		inner membrane
Q9KKQ6	MTLD_VIBCH		cytoplasm
Q9KKQ7	PTM3C_VIBCH	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KKS5	NAGB_VIBCH		extracellular
Q9KKW4	Y3786_VIBCH		cytoplasm
Q9KKW6	LLDD_VIBCH		cytoplasm
Q9KKY0	Y3770_VIBCH		inner membrane
Q9KKY1	Y3769_VIBCH		cytoplasm
Q9KKZ9	Y3751_VIBCH		inner membrane
Q9KL04	MALK_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KL06	MALF_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KL07	MALG_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KL16	CSPV_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KL20	RUMB_VIBCH		cytoplasm
Q9KL24	PYRC_VIBCH		cytoplasm
Q9KL54	Y3694_VIBCH		cytoplasm
Q9KL62	TDH_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KL72	SDHD_VIBCH		cytoplasm
Q9KL96	Y0G0_VIBCH		cytoplasm
Q9KLB8	PH4H_VIBCH		cytoplasm

Q9KLB9	PHS_VIBCH		cytoplasm
Q9KLC1	ECTA_VIBCH		cytoplasm
Q9KLC2	ECTB_VIBCH		cytoplasm
Q9KLC3	ECTC_VIBCH		cytoplasm
Q9KLC6	CH602_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KLC7	CH102_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KLD1	SPEA_VIBCH		cytoplasm
Q9KLE1	RNB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KLG0	Y3584_VIBCH		cytoplasm
Q9KLG8	Y3576_VIBCH		inner membrane
Q9KLJ3	FABH2_VIBCH	cytoplasm (probable).	cytoplasm
Q9KLJ6	GLPB_VIBCH		cytoplasm
Q9KLJ9	GLPK_VIBCH		cytoplasm
Q9KLK2	Y3541_VIBCH		inner membrane
Q9KLK4	Y3539_VIBCH		cytoplasm
Q9KLK6	LUXP_VIBCH	periplasm (probable).	periplasm
Q9KLK7	LUXQ_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9KLL6	COBQ_VIBCH		cytoplasm
Q9KLL9	MODC_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KLN2	MGSA_VIBCH		cytoplasm
Q9KLP4	GLGC2_VIBCH		cytoplasm
Q9KLQ5	FBPC_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KLR4	NAPA_VIBCH	periplasm (by similarity).	periplasm
Q9KLS6	DCUC_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KLW7	TKT2_VIBCH		extracellular
Q9KLW8	TAL_VIBCH	cytoplasm (probable).	cytoplasm
Q9K LX5	GCH1_VIBCH		cytoplasm
Q9K LX6	MSRAB_VIBCH		cytoplasm
Q9K LX7	FTHS_VIBCH		cytoplasm
Q9K LX9	MSCL_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KM09	Y3380_VIBCH		cytoplasm
Q9KM17	DDL_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KM20	Y3369_VIBCH		inner membrane
Q9KM50	Y3339_VIBCH	periplasm (potential).	periplasm
Q9KM59	KUP_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KM62	CLCA_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9KM67	SBCC_VIBCH		cytoplasm
Q9KM76	NRDG_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KMG3	Y3195_VIBCH		cytoplasm
Q9KMI9	Y3160_VIBCH		inner membrane
Q9KMJ4	Y3154_VIBCH		cytoplasm
Q9KMM3	DGT1B_VIBCH		cytoplasm

Q9KMP4	GLYA2_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KMP5	GCSH_VIBCH		cytoplasm
Q9KMT5	ACKA2_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KMW1	NADE_VIBCH		cytoplasm
Q9KMW9	GUAC_VIBCH		cytoplasm
Q9KMY3	HMP_VIBCH		cytoplasm
Q9KMY6	PEPT_VIBCH	cytoplasm (by similarity).	extracellular
Q9KN00	CSPA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KN05	TNAA_VIBCH		cytoplasm
Q9KN16	DEF2_VIBCH		cytoplasm
Q9KN67	PNCB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KN74	Y2891_VIBCH		inner membrane
Q9KN92	PSTB1_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KNA8	PHR_VIBCH		cytoplasm
Q9KNB2	DEOD2_VIBCH		cytoplasm
Q9KND9	Y2826_VIBCH		inner membrane
Q9KNE8	GLGB_VIBCH		extracellular
Q9KNF3	MALT_VIBCH		cytoplasm
Q9KNF8	Y2806_VIBCH		inner membrane
Q9KNG4	GIDA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNG5	GIDB_VIBCH		cytoplasm
Q9KNG7	PARB_VIBCH		cytoplasm
Q9KNG8	ATPZ_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9KNH1	ATPF_VIBCH		inner membrane
Q9KNH3	ATPA_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KNH4	ATPG_VIBCH		inner membrane
Q9KNH5	ATPB_VIBCH		inner membrane
Q9KNH6	ATPE_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KNH7	GLMU_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNI0	FADA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNI7	ADD_VIBCH		extracellular
Q9KNJ6	RBN_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KNJ7	DTD_VIBCH	cytoplasm (probable).	cytoplasm
Q9KNK0	PPCK_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNK2	HSLO_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNL2	Y2720_VIBCH		inner membrane
Q9KNL4	BIOH_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNL7	GREB_VIBCH		cytoplasm
Q9KNM3	RPOZ_VIBCH		cytoplasm
Q9KNM4	KGUA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNM9	CVRA_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9KNN1	DSBD_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane

Q9KNP2	K6PF_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNP9	METJ_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNQ2	RL31_VIBCH		cytoplasm
Q9KNQ6	HSLV_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNQ7	HSLU_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNQ9	RRAA2_VIBCH		cytoplasm
Q9KNR1	TPIS_VIBCH		cytoplasm
Q9KNR6	CH101_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNR7	CH601_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNS2	FRDD_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KNS3	FRDC_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KNS6	SYK3_VIBCH		cytoplasm
Q9KNS8	SECB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNT0	GPDA_VIBCH	cytoplasm (probable).	cytoplasm
Q9KNT4	CAPP_VIBCH		cytoplasm
Q9KNT5	ARGE_VIBCH	cytoplasm (probable).	extracellular
Q9KNT6	ARGC_VIBCH	cytoplasm (probable).	cytoplasm
Q9KNT7	ARGB_VIBCH	cytoplasm (probable).	cytoplasm
Q9KNT8	ASSY_VIBCH	cytoplasm (probable).	cytoplasm
Q9KNT9	ARLY_VIBCH	cytoplasm (probable).	cytoplasm
Q9KNU5	PBPA_VIBCH	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q9KNV1	AROK_VIBCH	cytoplasm (probable).	cytoplasm
Q9KNV2	AROB_VIBCH	cytoplasm (probable).	cytoplasm
Q9KNV6	GPH_VIBCH		cytoplasm
Q9KNW2	ARGD_VIBCH	cytoplasm (probable).	cytoplasm
Q9KNW8	Y2612_VIBCH		inner membrane
Q9KNX8	PURA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KNY1	RNR_VIBCH		cytoplasm
Q9KNY2	RLMB_VIBCH		cytoplasm
Q9KNY3	RS10_VIBCH		cytoplasm
Q9KNY4	RL3_VIBCH		cytoplasm
Q9KNY5	RL4_VIBCH		cytoplasm
Q9KNY7	RL2_VIBCH		cytoplasm
Q9KNY8	RS19_VIBCH		cytoplasm
Q9KNY9	RL22_VIBCH		cytoplasm
Q9KNZ0	RS3_VIBCH		cytoplasm
Q9KNZ1	RL16_VIBCH		cytoplasm
Q9KNZ2	RL29_VIBCH		cytoplasm
Q9KNZ3	RS17_VIBCH		cytoplasm
Q9KNZ5	RL24_VIBCH		cytoplasm
Q9KNZ6	RL5_VIBCH		cytoplasm
Q9KNZ7	RS14_VIBCH		cytoplasm
Q9KNZ8	RS8_VIBCH		cytoplasm
Q9KNZ9	RL6_VIBCH		cytoplasm
Q9KP00	RL18_VIBCH		cytoplasm
Q9KP01	RS5_VIBCH		cytoplasm
Q9KP03	RL15_VIBCH		cytoplasm
Q9KP05	RS13_VIBCH		cytoplasm
Q9KP07	RS4_VIBCH		cytoplasm
Q9KP08	RPOA_VIBCH		cytoplasm

Q9KP19	CYSD_VIBCH		cytoplasm
Q9KP20	CYSN_VIBCH		cytoplasm
Q9KP21	CYSC_VIBCH		cytoplasm
Q9KP30	MSRA_VIBCH		cytoplasm
Q9KP33	Y2546_VIBCH		inner membrane
Q9KP34	IPYR_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KP38	PAAD_VIBCH		cytoplasm
Q9KP43	Y2536_VIBCH		cytoplasm
Q9KP46	PTSO_VIBCH	cytoplasm (probable).	cytoplasm
Q9KP47	Y2532_VIBCH		cytoplasm
Q9KP62	MURA_VIBCH	cytoplasm (probable).	inner membrane
Q9KP65	PYRI_VIBCH		cytoplasm
Q9KP66	PYRB_VIBCH		cytoplasm
Q9KP68	OTC_VIBCH	cytoplasm (probable).	cytoplasm
Q9KP70	RAPA_VIBCH		cytoplasm
Q9KP71	RLUA_VIBCH		cytoplasm
Q9KP73	SYV_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KP80	LEUD_VIBCH		cytoplasm
Q9KP81	LEU2_VIBCH		cytoplasm
Q9KP82	LEU3_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KP83	LEU1_VIBCH		cytoplasm
Q9KP93	RPIA_VIBCH		cytoplasm
Q9KP97	Y2476_VIBCH		inner membrane
Q9KPA2	Y2471_VIBCH		cytoplasm
Q9KPA4	NADB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPB0	LEPA_VIBCH		cytoplasm
Q9KPB2	RNC_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPB3	ERA_VIBCH		cytoplasm
Q9KPB4	RECO_VIBCH		cytoplasm
Q9KPB5	PDXJ_VIBCH	cytoplasm (probable).	cytoplasm
Q9KPB6	ACPS_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPC1	RUMA_VIBCH		cytoplasm
Q9KPC4	PYRG_VIBCH		cytoplasm
Q9KPC5	ENO_VIBCH	cytoplasm (by similarity).	extracellular
Q9KPC6	CCA_VIBCH		cytoplasm
Q9KPD4	GLNE_VIBCH		cytoplasm
Q9KPE1	Y2429_VIBCH		cytoplasm
Q9KPE2	Y2428_VIBCH		inner membrane
Q9KPE3	COAE_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPE9	XERD_VIBCH	cytoplasm (by similarity).	nucleoid
Q9KPF6	DLDH_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPF9	MRAW_VIBCH		cytoplasm
Q9KPG4	MRAY_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KPG5	MURD_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPG7	MURG_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KPG8	MURC_VIBCH	cytoplasm (probable).	cytoplasm
Q9KPH2	LPXC_VIBCH		cytoplasm
Q9KPH7	DAPB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPH8	CARA_VIBCH		cytoplasm
Q9KPH9	CARB_VIBCH		cytoplasm
Q9KPI4	Y2383_VIBCH		cytoplasm

Q9KPI5	Y2382_VIBCH	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9KPI6	BTUF_VIBCH	periplasm (by similarity).	periplasm
Q9KPK1	RRAA1_VIBCH		cytoplasm
Q9KPK4	KHSE_VIBCH	cytoplasm (potential).	cytoplasm
Q9KPK6	GRCA_VIBCH		cytoplasm
Q9KPK7	END4_VIBCH		extracellular
Q9KPK8	UNG_VIBCH	cytoplasm (by similarity).	extracellular
Q9KPL2	Y2355_VIBCH		inner membrane
Q9KPL7	DEOC_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPL8	TYPH_VIBCH		cytoplasm
Q9KPL9	DEOB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPM0	DEOD1_VIBCH		cytoplasm
Q9KPQ0	ARGA_VIBCH	cytoplasm (probable).	cytoplasm
Q9KPQ4	MLTA_VIBCH	bacterial cell outer membrane; lipid-anchor (by similarity).	extracellular
Q9KPQ7	CSD_VIBCH		cytoplasm
Q9KPQ9	PANE_VIBCH	cytoplasm (potential).	cytoplasm
Q9KPS0	BOLA_VIBCH		cytoplasm
Q9KPS1	NQRA_VIBCH		inner membrane
Q9KPS2	NQRB_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9KPS5	DPO4_VIBCH	cytoplasm (probable).	cytoplasm
Q9KPT1	RIMK_VIBCH		cytoplasm
Q9KPT5	XGPT_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPT6	Y2276_VIBCH		cytoplasm
Q9KPT8	PROB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPT9	PROA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPU0	NRDR_VIBCH		cytoplasm
Q9KPU4	RISB_VIBCH		cytoplasm
Q9KPU5	NUSB_VIBCH		cytoplasm
Q9KPU8	Y2264_VIBCH		inner membrane
Q9KPV0	GLND_VIBCH		cytoplasm
Q9KPV2	RS2_VIBCH		cytoplasm
Q9KPV4	PYRH_VIBCH		cytoplasm
Q9KPV5	RRF_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPV6	UPPS_VIBCH		inner membrane
Q9KPV8	DXR_VIBCH		cytoplasm
Q9KPV9	Y2253_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9KPW1	SKP_VIBCH	periplasm (by similarity).	periplasm
Q9KPW2	LPXD_VIBCH		cytoplasm
Q9KPW3	FABZ_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPW4	LPXA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPW5	LPXB_VIBCH		cytoplasm
Q9KPW8	ACCA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPX0	TILS_VIBCH	cytoplasm (probable).	cytoplasm
Q9KPX2	PADC_VIBCH		cytoplasm
Q9KPX4	Y2238_VIBCH	cytoplasm (potential).	inner membrane
Q9KPX8	RNH_VIBCH	cytoplasm (potential).	extracellular
Q9KPY2	GMHA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPY6	PUR5_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KPY7	UPP_VIBCH		cytoplasm

Q9KPZ7	ATCU_VIBCH	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9KQ06	CHER1_VIBCH		cytoplasm
Q9KQ13	FLGH_VIBCH	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q9KQ14	FLGI_VIBCH	periplasm (by similarity).	periplasm
Q9KQ15	FLGJ_VIBCH	periplasm (by similarity).	extracellular
Q9KQ21	PTH_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KQ22	KPRS_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KQ23	ISPE_VIBCH		cytoplasm
Q9KQ24	HEM1_VIBCH		cytoplasm
Q9KQ26	HEMK_VIBCH		cytoplasm
Q9KQ28	Y2176_VIBCH		extracellular
Q9KQ29	KDSA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KQ30	5NTD_VIBCH	bacterial cell outer membrane; lipid-anchor (by similarity).	periplasm
Q9KQ40	Y2164_VIBCH		inner membrane
Q9KQ47	DAPA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KQ62	FLAG_VIBCH		cytoplasm
Q9KQ65	FLIS_VIBCH		flagellar
Q9KQ82	Y2119_VIBCH		inner membrane
Q9KQ83	Y2118_VIBCH		cytoplasm
Q9KQ85	AROC_VIBCH		cytoplasm
Q9KQ91	Y2110_VIBCH		cytoplasm
Q9KQ92	PDXB_VIBCH		cytoplasm
Q9KQA7	Y2093_VIBCH		cytoplasm
Q9KQB5	SUCC_VIBCH		cytoplasm
Q9KQE3	CCMA_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KQE7	CCME_VIBCH	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q9KQE9	DSBE_VIBCH	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q9KQF5	TOP3_VIBCH		cytoplasm
Q9KQF8	Y2040_VIBCH		inner membrane
Q9KQF9	NDPA_VIBCH	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	inner membrane
Q9KQH0	RLUC_VIBCH		cytoplasm
Q9KQH1	Y2027_VIBCH	cytoplasm (potential).	inner membrane
Q9KQH3	RL32_VIBCH		cytoplasm
Q9KQH4	PLSX_VIBCH		cytoplasm
Q9KQH5	FABH1_VIBCH	cytoplasm (probable).	cytoplasm
Q9KQH7	FABG_VIBCH		cytoplasm
Q9KQH8	ACP_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KQH9	FABF_VIBCH		cytoplasm
Q9KQI0	PABC_VIBCH		cytoplasm
Q9KQI2	KTHY_VIBCH		cytoplasm
Q9KQK0	MSRB_VIBCH		cytoplasm
Q9KQL9	DGT1A_VIBCH		cytoplasm

Q9KQM0	Y1978_VIBCH		cytoplasm
Q9KQM6	MENC_VIBCH		extracellular
Q9KQN7	MINE_VIBCH		inner membrane
Q9KQN9	MINC_VIBCH		inner membrane
Q9KQR7	Y1931_VIBCH		inner membrane
Q9KQS5	TIG_VIBCH		cytoplasm
Q9KQS6	CLPP_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KQS7	CLPX_VIBCH		cytoplasm
Q9KQS9	DBHB_VIBCH		cytoplasm
Q9KQT2	KCY_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KQT4	IHFB_VIBCH		cytoplasm
Q9KQT7	PYRF_VIBCH		cytoplasm
Q9KQU6	DSBB_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KQU8	FADR_VIBCH	cytoplasm (potential).	cytoplasm
Q9KQV6	Y1892_VIBCH		inner membrane
Q9KQW9	MSBA_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KQX0	LPXK_VIBCH		cytoplasm
Q9KQX2	KDSB_VIBCH	cytoplasm (potential).	cytoplasm
Q9KQX4	Y1873_VIBCH		inner membrane
Q9KQX6	Y1871_VIBCH		inner membrane
Q9KQZ4	Y1853_VIBCH		inner membrane
Q9KQZ5	Y1852_VIBCH		inner membrane
Q9KQZ7	LPXH_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KR00	RUVC_VIBCH		cytoplasm
Q9KR01	RUVA_VIBCH		cytoplasm
Q9KR02	RUVB_VIBCH		cytoplasm
Q9KR11	TOLB_VIBCH	periplasm (potential).	periplasm
Q9KR14	NADA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KR33	CRYD_VIBCH		cytoplasm
Q9KR57	Y1786_VIBCH		cytoplasm
Q9KR61	NANK_VIBCH		cytoplasm
Q9KR62	NANE_VIBCH		cytoplasm
Q9KR69	Y1774_VIBCH		periplasm
Q9KR70	Y1773_VIBCH		extracellular
Q9KRA3	Y1738_VIBCH		cytoplasm
Q9KRA6	ATE_VIBCH	cytoplasm (potential).	cytoplasm
Q9KRA7	LFTR_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KRB0	AROA_VIBCH	cytoplasm (probable).	cytoplasm
Q9KRB2	TOP1_VIBCH		cytoplasm
Q9KRB5	GLGC1_VIBCH		cytoplasm
Q9KRB6	GLGA_VIBCH		cytoplasm
Q9KRC2	TORD_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KRC6	MUKF_VIBCH	nucleoid (by similarity). restricted to the nucleoid region (by similarity).	nucleoid
Q9KRD1	Y1711_VIBCH		cytoplasm
Q9KRD8	METE_VIBCH		cytoplasm
Q9KRE2	ISPZ_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KRF0	TORA_VIBCH	periplasm (by similarity).	periplasm

Q9KRJ0	COLA_VIBCH	secreted protein (by similarity).	extracellular
Q9KRK0	RL25_VIBCH		cytoplasm
Q9KRK5	RSUA_VIBCH		cytoplasm
Q9KRM5	META_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KRP1	GAL1_VIBCH	cytoplasm (potential).	cytoplasm
Q9KRQ1	CATA_VIBCH	periplasm (probable).	periplasm
Q9KRR0	Y1576_VIBCH		inner membrane
Q9KRU4	NORM_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KRU5	Y1539_VIBCH		cytoplasm
Q9KRV8	MOBA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KRX4	NPD_VIBCH	cytoplasm (probable).	cytoplasm
Q9KRX5	Y1508_VIBCH		inner membrane
Q9KRY6	PROQ_VIBCH	cytoplasm (potential).	inner membrane
Q9KRZ2	PYRD_VIBCH	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KS00	FABA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KS02	Y1481_VIBCH		inner membrane
Q9KS27	FNR_VIBCH	cytoplasm (probable).	cytoplasm
Q9KS61	CHER2_VIBCH		cytoplasm
Q9KS67	CRY2_VIBCH		cytoplasm
Q9KS71	LPLA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSA1	Y1358_VIBCH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9KSB2	MAAI_VIBCH		cytoplasm
Q9KSB4	Y1345_VIBCH		cytoplasm
Q9KSC2	PRPB_VIBCH		cytoplasm
Q9KSE5	ALR2_VIBCH		cytoplasm
Q9KSG8	OPGG_VIBCH	periplasm (by similarity).	periplasm
Q9KSG9	OPGH_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KSH1	Y1285_VIBCH		inner membrane
Q9KSJ3	GCH2_VIBCH		extracellular
Q9KSJ7	Y1259_VIBCH		inner membrane
Q9KSJ9	UBIG_VIBCH		cytoplasm
Q9KSL1	BTUD_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KSL2	BTUC_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KSL4	ASTE_VIBCH		periplasm
Q9KSL8	COBS_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KSL9	COBT_VIBCH		cytoplasm
Q9KSM3	Y1233_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KSM5	CDD_VIBCH		cytoplasm
Q9KSN0	TPMT_VIBCH	cytoplasm (by similarity).	cytoplasm

Q9KSN4	IHFA_VIBCH		cytoplasm
Q9KSP2	UVRB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSP7	EFPL_VIBCH		cytoplasm
Q9KSP8	Y1208_VIBCH		inner membrane
Q9KSQ1	HUTI_VIBCH	cytoplasm (potential).	cytoplasm
Q9KSQ2	HUTG_VIBCH		cytoplasm
Q9KSQ3	HUTU_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSQ4	HUTH_VIBCH	cytoplasm (potential).	cytoplasm
Q9KSR6	PUR7_VIBCH		cytoplasm
Q9KSR8	MAO1_VIBCH		cytoplasm
Q9KSS4	TRXB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSS7	RLUB_VIBCH		cytoplasm
Q9KST2	TRPE_VIBCH		cytoplasm
Q9KST3	TRPG_VIBCH		cytoplasm
Q9KST4	TRPD_VIBCH		cytoplasm
Q9KST5	TRPC_VIBCH		cytoplasm
Q9KST6	TRPB_VIBCH		cytoplasm
Q9KST7	TRPA_VIBCH		cytoplasm
Q9KST9	KITH_VIBCH	cytoplasm (potential).	cytoplasm
Q9KSU7	SERC_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSW0	GLRX_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSW1	Y1145_VIBCH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9KSW3	CLPS_VIBCH		cytoplasm
Q9KSW4	CSPD_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSW7	HIS2_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSW8	HIS6_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSW9	HIS4_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSX0	HIS5_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSX1	HIS7_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSX2	HIS8_VIBCH		cytoplasm
Q9KSX4	HIS1_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSX8	TRMU_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KSX9	Y1127_VIBCH		inner membrane
Q9KSY9	HTPX_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9KSZ1	BIOD_VIBCH		cytoplasm
Q9KT00	DUSC_VIBCH		cytoplasm
Q9KT06	Y1099_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KT07	ACKA1_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KT47	Y1058_VIBCH		inner membrane
Q9KT48	Y1057_VIBCH		inner membrane
Q9KT49	RECR_VIBCH		cytoplasm
Q9KT50	Y1055_VIBCH		inner membrane
Q9KT58	FADJ_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KT59	FADI_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KT67	URK_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KT77	MOAE_VIBCH		cytoplasm
Q9KT79	MOAC_VIBCH		cytoplasm
Q9KT81	MOAA_VIBCH		cytoplasm
Q9KT82	Y1023_VIBCH		cytoplasm
Q9KT83	LUXU_VIBCH		inner membrane

Q9KT84	LUXO1_VIBCH		cytoplasm
Q9KT86	RNFA_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KT88	RNFC_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	cytoplasm
Q9KT89	RNFD_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9KT90	RNFG_VIBCH	bacterial cell inner membrane (potential).	inner membrane
Q9KT91	RNFE_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KT93	LGUL_VIBCH		extracellular
Q9KT97	RNT_VIBCH		cytoplasm
Q9KTA4	TRUA_VIBCH		cytoplasm
Q9KTB6	HEMH_VIBCH	cytoplasm (by similarity).	extracellular
Q9KTC8	CUER_VIBCH	cytoplasm (probable).	cytoplasm
Q9KTD2	ZIPA_VIBCH	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q9KTD3	CYSZ_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KTD6	PTHP_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTE2	Y960_VIBCH		inner membrane
Q9KTE3	CORC_VIBCH		inner membrane
Q9KTE4	LNT_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KTF1	Y951_VIBCH		inner membrane
Q9KTF4	RLPA_VIBCH	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q9KTF7	Y945_VIBCH		inner membrane
Q9KTF8	LIPB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTF9	LIPA_VIBCH	cytoplasm (potential).	cytoplasm
Q9KTG1	GLYA1_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTJ4	GMHB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTJ5	METN_VIBCH	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q9KTJ6	METI_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KTJ7	METQ_VIBCH	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q9KTK0	QUEF_VIBCH	cytoplasm (probable).	cytoplasm
Q9KTK4	EX9_VIBCH		cytoplasm
Q9KTK8	THII_VIBCH	cytoplasm (potential).	cytoplasm
Q9KTL1	EX7S_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTL3	DXS_VIBCH		cytoplasm
Q9KTL4	TRUC_VIBCH		cytoplasm
Q9KTM1	Y881_VIBCH		inner membrane
Q9KTM4	RL31B_VIBCH		cytoplasm

Q9KTN0	Y872_VIBCH		inner membrane
Q9KTN2	PUR4_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTP8	PPNK_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTQ4	ACFD_VIBCH	bacterial cell membrane; lipid-anchor (potential).	extracellular
Q9KTT7	CITG_VIBCH		cytoplasm
Q9KTT8	CITX_VIBCH		cytoplasm
Q9KTU1	CITD_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTU4	OADG2_VIBCH	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q9KTU5	OADB_VIBCH	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KTV1	DADA_VIBCH		cytoplasm
Q9KTW2	GUAA_VIBCH		cytoplasm
Q9KTW4	EX7L_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTW7	ENGA_VIBCH		cytoplasm
Q9KTX1	ISPG_VIBCH		cytoplasm
Q9KTX4	NDK_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTX5	PEPB_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KTX8	HSCA_VIBCH		cytoplasm
Q9KTX9	HSCB_VIBCH		inner membrane
Q9KTY2	ISCS_VIBCH		cytoplasm
Q9KTY5	SUHB_VIBCH		cytoplasm
Q9KTY9	TGT_VIBCH		cytoplasm
Q9KTZ1	QUEA_VIBCH	cytoplasm (potential).	cytoplasm
Q9KU00	CUTC_VIBCH	cytoplasm (probable).	cytoplasm
Q9KU04	PSTB2_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KU07	PPK_VIBCH		cytoplasm
Q9KU08	PPX_VIBCH		cytoplasm
Q9KU12	RDGC_VIBCH	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q9KU18	CLPB_VIBCH	cytoplasm (probable).	cytoplasm
Q9KU20	RLUD_VIBCH		cytoplasm
Q9KU21	Y708_VIBCH	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q9KU25	NSPS_VIBCH	periplasm (probable).	periplasm
Q9KU26	MBAA_VIBCH		inner membrane
Q9KU27	Y702_VIBCH		inner membrane
Q9KU28	TRPR_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KU36	Y693_VIBCH		cytoplasm
Q9KU37	NAGZ_VIBCH	cytoplasm (by similarity).	extracellular
Q9KU39	Y690_VIBCH		cytoplasm
Q9KU44	ISPH_VIBCH		cytoplasm
Q9KU46	LSPA_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9KU47	SYI_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KU53	NUDH_VIBCH		cytoplasm
Q9KU56	MUTH_VIBCH	cytoplasm (by similarity).	extracellular
Q9KU77	RS15_VIBCH		cytoplasm
Q9KU78	TRUB_VIBCH		cytoplasm

Q9KU79	RBFA_VIBCH	cytoplasm (potential).	cytoplasm
Q9KU82	Y641_VIBCH		inner membrane
Q9KU84	GLMM_VIBCH		cytoplasm
Q9KU87	RRMJ_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KU89	GREA_VIBCH		cytoplasm
Q9KU92	SY Y2_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KU97	GSA_VIBCH	cytoplasm (potential).	cytoplasm
Q9KUB7	Y605_VIBCH		inner membrane
Q9KUC0	PBPB_VIBCH	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q9KUC5	SFSA_VIBCH		cytoplasm
Q9KUC7	GLUQ_VIBCH		cytoplasm
Q9KUD0	PANB_VIBCH		cytoplasm
Q9KUD1	PANC_VIBCH		cytoplasm
Q9KUD9	Y582_VIBCH		cytoplasm
Q9KUE1	Y580_VIBCH		inner membrane
Q9KUF0	RS9_VIBCH		cytoplasm
Q9KUF7	RL19_VIBCH		cytoplasm
Q9KUF8	TRMD_VIBCH	cytoplasm (potential).	cytoplasm
Q9KUF9	RIMM_VIBCH	cytoplasm (potential).	cytoplasm
Q9KUG0	RS16_VIBCH		cytoplasm
Q9KUG4	LUXS_VIBCH		extracellular
Q9KUG5	GSH1_VIBCH		cytoplasm
Q9KUH2	OADG1_VIBCH	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q9KUH3	CSRA_VIBCH		cytoplasm
Q9KUI0	CYSA_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KUI6	MUTS_VIBCH		cytoplasm
Q9KUI8	PIMT_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KUI9	SURE_VIBCH	cytoplasm (potential).	periplasm
Q9KUJ0	TRUD_VIBCH		cytoplasm
Q9KUJ1	ISPF_VIBCH		extracellular
Q9KUJ2	ISPD_VIBCH		cytoplasm
Q9KUJ3	FTSB_VIBCH	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q9KUJ4	UPPP_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KUJ7	Y523_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KUK8	Y510_VIBCH		cytoplasm
Q9KUM8	GLMS_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KUN3	ICIA_VIBCH		cytoplasm
Q9KUP2	TKT1_VIBCH		extracellular
Q9KUP4	SPRT_VIBCH	cytoplasm (potential).	extracellular
Q9KUP7	GSHB_VIBCH		cytoplasm
Q9KUP8	Y467_VIBCH		inner membrane
Q9KUP9	RUVX_VIBCH	cytoplasm (potential).	cytoplasm

Q9KUQ0	SY11_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KUQ4	Y461_VIBCH		cytoplasm
Q9KUQ7	Y458_VIBCH		inner membrane
Q9KUQ9	HAM1_VIBCH		extracellular
Q9KUR1	GLSA_VIBCH		extracellular
Q9KUR2	TRMB_VIBCH		cytoplasm
Q9KUR4	FETP_VIBCH		cytoplasm
Q9KUR8	DJLA_VIBCH	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane
Q9KUR9	OSTA_VIBCH	bacterial cell outer membrane (by similarity).	outer membrane
Q9KUS1	PDXA_VIBCH	cytoplasm (probable).	cytoplasm
Q9KUS2	KSGA_VIBCH		cytoplasm
Q9KUS3	APAG_VIBCH		inner membrane
Q9KUS4	APAH_VIBCH		extracellular
Q9KUS9	RL27_VIBCH		cytoplasm
Q9KUT3	MDH_VIBCH		cytoplasm
Q9KUT4	ARGR_VIBCH	cytoplasm (probable).	cytoplasm
Q9KUU7	Y418_VIBCH	cytoplasm (potential).	inner membrane
Q9KUW2	SSB_VIBCH		cytoplasm
Q9KUW5	UVRA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KUW9	METH_VIBCH		cytoplasm
Q9KUX2	CYSH_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KUX3	CYSI_VIBCH		cytoplasm
Q9KUX4	CYSJ_VIBCH		cytoplasm
Q9KUX9	DUSA_VIBCH		cytoplasm
Q9KUY4	G6PI_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KUY6	ALR1_VIBCH		cytoplasm
Q9KUY9	RL9_VIBCH		cytoplasm
Q9KUZ0	RS18_VIBCH		cytoplasm
Q9KUZ1	PRIB_VIBCH		cytoplasm
Q9KUZ2	RS6_VIBCH		cytoplasm
Q9KUZ8	RS7_VIBCH		cytoplasm
Q9KUZ9	RS12_VIBCH		cytoplasm
Q9KV01	TUSC_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KV02	TUSD_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KV06	SLYX_VIBCH		inner membrane
Q9KV08	HFLC_VIBCH		extracellular
Q9KV09	HFLK_VIBCH		inner membrane
Q9KV11	HFQ_VIBCH		cytoplasm
Q9KV12	MIAA_VIBCH		cytoplasm
Q9KV13	MUTL_VIBCH		cytoplasm
Q9KV17	ORN_VIBCH	cytoplasm (potential).	cytoplasm
Q9KV18	ENGC_VIBCH		cytoplasm
Q9KV19	PSD_VIBCH		cytoplasm
Q9KV22	GPMI_VIBCH		cytoplasm
Q9KV26	DCUP_VIBCH	cytoplasm (probable).	cytoplasm
Q9KV27	NUDC_VIBCH		cytoplasm
Q9KV29	RPOC_VIBCH		cytoplasm
Q9KV30	RPOB_VIBCH		cytoplasm
Q9KV31	RL7_VIBCH		cytoplasm
Q9KV32	RL10_VIBCH		cytoplasm
Q9KV33	RL1_VIBCH		cytoplasm
Q9KV34	RL11_VIBCH		cytoplasm
Q9KV35	NUSG_VIBCH		cytoplasm

Q9KV36	SECE_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KV38	COAA_VIBCH	cytoplasm (probable).	cytoplasm
Q9KV40	MURB_VIBCH	cytoplasm (probable).	inner membrane
Q9KV48	UBID_VIBCH		cytoplasm
Q9KV52	RHLB_VIBCH		cytoplasm
Q9KV53	GPPA_VIBCH		extracellular
Q9KV59	ACSA_VIBCH		cytoplasm
Q9KV60	AROQ_VIBCH		cytoplasm
Q9KV64	PRMA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KV66	DUSB_VIBCH		cytoplasm
Q9KV80	PUR9_VIBCH		cytoplasm
Q9KV81	PUR2_VIBCH		cytoplasm
Q9KV83	DBHA_VIBCH		cytoplasm
Q9KVB9	KDKA_VIBCH	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q9KVC4	COAD_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KVC5	FPG_VIBCH		cytoplasm
Q9KVC7	RL33_VIBCH		cytoplasm
Q9KVC8	RL28_VIBCH		cytoplasm
Q9KVC9	RADC_VIBCH		cytoplasm
Q9KVD1	COABC_VIBCH		cytoplasm
Q9KVD2	SLMA_VIBCH	nucleoid-associated (by similarity).	cytoplasm
Q9KVD5	PYRE_VIBCH		cytoplasm
Q9KVD6	RNPH_VIBCH		cytoplasm
Q9KVD8	Y208_VIBCH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9KVD9	PTSBC_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9KVE0	Y206_VIBCH		cytoplasm
Q9KVF1	Y195_VIBCH	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9KVI4	ILVC_VIBCH		cytoplasm
Q9KVI7	MURI_VIBCH		cytoplasm
Q9KVI9	BTUB_VIBCH	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q9KVJ0	TRMA_VIBCH		cytoplasm
Q9KVL4	XERC_VIBCH	cytoplasm (by similarity).	nucleoid
Q9KVL6	DAPF_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KVL7	DCDA_VIBCH		cytoplasm
Q9KVL8	CYAY_VIBCH		inner membrane
Q9KVM1	HEM3_VIBCH		cytoplasm
Q9KVM7	Y114_VIBCH		inner membrane
Q9KVN0	ENGB_VIBCH		cytoplasm
Q9KVP1	GLPE_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KVP8	PLSB_VIBCH	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9KVP9	LEXA_VIBCH		cytoplasm

Q9KVQ4	UBIB_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KVQ6	UBIE_VIBCH		cytoplasm
Q9KVQ7	RMUC_VIBCH		cytoplasm
Q9KVR0	USPB_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KVR6	Y073_VIBCH		inner membrane
Q9KVS4	THIG_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KVS8	THIC_VIBCH		cytoplasm
Q9KVS9	CRCB_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KVT3	AROE_VIBCH		cytoplasm
Q9KVT4	HEM6_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KVT7	PUR6_VIBCH		cytoplasm
Q9KVT8	PURK_VIBCH		cytoplasm
Q9KVU0	SMG_VIBCH		cytoplasm
Q9KVU3	DEF1_VIBCH		cytoplasm
Q9KVU4	FMT_VIBCH		cytoplasm
Q9KVU5	RSMB_VIBCH		cytoplasm
Q9KVV0	ILVD_VIBCH		cytoplasm
Q9KVV4	TUSA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KVX3	GYRB_VIBCH		cytoplasm
Q9KVX4	RECF_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KVX5	DPO3B_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KVX6	DNAA_VIBCH	cytoplasm (by similarity).	cytoplasm
Q9KVY2	RNPA_VIBCH		extracellular
Q9KVY3	Y005_VIBCH		inner membrane
Q9KVY4	OXAA_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9KVY5	TRME_VIBCH		cytoplasm
Q9KVY6	MIOC_VIBCH		cytoplasm
Q9KVZ7	T2X1_XANCO		extracellular
Q9KVZ8	MTX1_XANCO		cytoplasm
Q9KW13	RPOC_SHEVI		cytoplasm
Q9KW14	RPOB_SHEVI		cytoplasm
Q9KW52	HEM1_DESVM		cytoplasm
Q9KWA3	ACSA_AGRRH		cytoplasm
Q9KWD1	IDI_AGRRH	cytoplasm (by similarity).	cytoplasm
Q9KWE4	HUTH_AGRRH	cytoplasm (potential).	cytoplasm
Q9KWE5	HUTI_AGRRH	cytoplasm (potential).	cytoplasm
Q9KWI8	BCHZ_ERYLO		cytoplasm
Q9KWM8	COPD_PSESF	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9KWN1	SDH_AGRTU		cytoplasm
Q9KVV3	TTGF_PSEPU	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
Q9KVV4	TTGE_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9KVV5	TTGD_PSEPU	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane

Q9L3B0	PQQE_GLUOX		cytoplasm
Q9L3B1	PQQD_GLUOX		cytoplasm
Q9L3B2	PQQC_GLUOX		cytoplasm
Q9L3B3	PQQB_GLUOX		cytoplasm
Q9L3I7	RUVX_AERHY	cytoplasm (potential).	cytoplasm
Q9L4D3	GREA_XANCP		cytoplasm
Q9L4K0	BETA_CHRSL		cytoplasm
Q9L4K1	BETB_HALEL		cytoplasm
Q9L4K2	BETI_CHRSL		cytoplasm
Q9L4T0	LUXS_SALTY		extracellular
Q9L516	GRPE_PSYS1	cytoplasm (probable).	cytoplasm
Q9L560	PGK_BRUAB	cytoplasm (by similarity).	cytoplasm
Q9L5D6	G7AC_BREDI	periplasm (potential).	periplasm
Q9L5W4	RL7_LIBAC		cytoplasm
Q9L5W5	RL10_LIBAC		cytoplasm
Q9L645	NRDG_SALTY	cytoplasm (by similarity).	cytoplasm
Q9L646	NRDD_SALTY		cytoplasm
Q9L690	CH603_RHILE	cytoplasm (by similarity).	cytoplasm
Q9L691	CH602_RHILE	cytoplasm (by similarity).	cytoplasm
Q9L699	Y1001_PASMU		cytoplasm
Q9L6A2	MIOC_PASMU		cytoplasm
Q9L6A3	MREB_PASMU		cytoplasm
Q9L6B3	DSBB_PASMU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9L6B4	NANE_PASMU		cytoplasm
Q9L6B5	Y1466_PASMU		cytoplasm
Q9L6B6	ATPG_PASMU		inner membrane
Q9L6B7	IMDH_PASMU		cytoplasm
Q9L6B8	PUR1_PASMU		cytoplasm
Q9L6K8	RIBB_VIBFI		cytoplasm
Q9L6L1	HEMG_SALTY		cytoplasm
Q9L6L2	TRKH_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9L6L5	FADB_SALTY		cytoplasm
Q9L6L9	FRE_SALTY		cytoplasm
Q9L6M0	UBID_SALTY	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9L6M9	DLHH_SALTY		extracellular
Q9L6N1	METE_SALTY		cytoplasm
Q9L6N6	RHTB_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9L6N7	RHTC_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9L6P1	RARD_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9L6Q3	HEM4_SALTY		cytoplasm
Q9L6Q8	WECF_SALTY	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q9L6R4	WECC_SALTY		cytoplasm

Q9L6R5	WECB_SALTY	cytoplasm (by similarity).	cytoplasm
Q9L6R7	WECA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9L6S1	REP_SALTY		cytoplasm
Q9L6W2	QUEA_PSEU2	cytoplasm (potential).	cytoplasm
Q9L6W9	QUEA_PSESM	cytoplasm (potential).	cytoplasm
Q9L6X9	TUS_YERPE	cytoplasm (by similarity).	cytoplasm
Q9L772	AQPZ_BRUME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9L7A3	CCA_HAEDU		cytoplasm
Q9L7A6	Y470_HAEDU	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q9L7A7	RLUD_HAEDU		cytoplasm
Q9L7P5	CH601_VIBPA	cytoplasm (by similarity).	cytoplasm
Q9L7R3	OMPL_SALTY	bacterial cell outer membrane; multi-pass membrane protein (by similarity).	outer membrane
Q9L7R4	YIHO_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9L7R5	YIHP_SALTY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9L7R9	YIHT_SALTY		cytoplasm
Q9L7S0	YIHU_SALTY		cytoplasm
Q9L7T2	FPG_PSEAE		cytoplasm
Q9L7V3	Y584_HAEDU		cytoplasm
Q9L7X5	CLPX_BRUAB		cytoplasm
Q9L7X6	CLPP_BRUAB	cytoplasm (by similarity).	cytoplasm
Q9L7Z1	DNAK_VIBPR		cytoplasm
Q9L7Z3	GRPE_VIBPR	cytoplasm (probable).	cytoplasm
Q9L867	ACPD_AZOBR		extracellular
Q9L894	OTSB_SALTY		cytoplasm
Q9L8J3	AHPA_PASMU		inner membrane
Q9L8J5	BCHL_RHORU		cytoplasm
Q9L9C3	RUVB_RHIET		cytoplasm
Q9L9C4	RUVA_RHIET		cytoplasm
Q9L9C8	RS16_THIFE		cytoplasm
Q9L9I0	ZRAP_SALTY	periplasm (by similarity).	periplasm
Q9L9I4	DCUP_SALTY	cytoplasm (probable).	cytoplasm
Q9L9I5	NUDC_SALTY		cytoplasm
Q9L9I6	RSD_SALTY		cytoplasm
Q9L9I7	THIC_SALTY		cytoplasm
Q9L9I8	THIE_SALTY		cytoplasm
Q9L9J1	THIG_SALTY	cytoplasm (by similarity).	cytoplasm
Q9L9K3	COAA_SALTY	cytoplasm (probable).	cytoplasm
Q9LA01	CCME_RHOCA	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q9LA02	ARGC_RHOCA	cytoplasm (probable).	cytoplasm
Q9LA24	FNR_SHIDY	cytoplasm (probable).	cytoplasm

Q9LA29	YCJF_SHIDY	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9LA96	DBHA_AERHY		cytoplasm
Q9LAG8	TRPA_AZOBR		cytoplasm
Q9LCJ0	NQRF_VIBPA	bacterial cell inner membrane (potential).	inner membrane
Q9LCT0	PYRF_BRAJA		cytoplasm
Q9LCT6	DDLB_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9P9T1	Y1585_XYLFA		cytoplasm
Q9P9T2	Y1581_XYLFA		cytoplasm
Q9P9T3	Y1567_XYLFA		cytoplasm
Q9P9T5	Y1569_XYLFA		cytoplasm
Q9P9T6	Y503_XYLFA		cytoplasm
Q9P9U0	RNPA_XYLFA		extracellular
Q9P9U1	OXAA_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9P9U3	TRME_XYLFA		cytoplasm
Q9PA24	RHLB_XYLFA		cytoplasm
Q9PA28	UNG_XYLFA	cytoplasm (by similarity).	extracellular
Q9PA34	NORM_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PA38	OPGD_XYLFA	periplasm (by similarity).	periplasm
Q9PA72	HEM1_XYLFA		cytoplasm
Q9PA74	LOLB_XYLFA	bacterial cell outer membrane; lipid-anchor (by similarity).	inner membrane
Q9PA75	ISPE_XYLFA		cytoplasm
Q9PA76	KPRS_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PA77	RL25_XYLFA		cytoplasm
Q9PA78	PTH_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PA81	NUSG_XYLFA		cytoplasm
Q9PA82	RL11_XYLFA		cytoplasm
Q9PA83	RL1_XYLFA		cytoplasm
Q9PA84	RL10_XYLFA		cytoplasm
Q9PA85	RL7_XYLFA		cytoplasm
Q9PA86	RPOB_XYLFA		cytoplasm
Q9PA87	RPOC_XYLFA		cytoplasm
Q9PA89	RS7_XYLFA		cytoplasm
Q9PA93	HTPX_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9PA99	Y2619_XYLFA		inner membrane
Q9PAA9	Y2609_XYLFA		inner membrane
Q9PAB4	PHS_XYLFA		cytoplasm
Q9PAB5	Y2603_XYLFA		inner membrane
Q9PAC7	PPK_XYLFA		cytoplasm
Q9PAD0	LPXH_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PAE3	ISPG_XYLFA		cytoplasm
Q9PAE6	MURB_XYLFA	cytoplasm (probable).	inner membrane
Q9PAE7	PYRD_XYLFA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PAF8	RS18_XYLFA		cytoplasm
Q9PAF9	RL9_XYLFA		cytoplasm

Q9PAG1	ZIPA_XYLFA	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q9PAG5	MTNA_XYLFA		cytoplasm
Q9PAH1	SUCC_XYLFA		cytoplasm
Q9PAI2	COAE_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PAL9	BIOD_XYLFA		cytoplasm
Q9PAM0	SUHB_XYLFA		cytoplasm
Q9PAM5	UBIG_XYLFA		cytoplasm
Q9PAM6	GPH_XYLFA		cytoplasm
Q9PAN0	METX_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PAN4	DSBE_XYLFA	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q9PAN6	CCME_XYLFA	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q9PAP0	CCMA_XYLFA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PAP2	RLUB_XYLFA		cytoplasm
Q9PAQ6	KCY_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PAQ8	IHFB_XYLFA		cytoplasm
Q9PAR6	GUAA_XYLFA		cytoplasm
Q9PAR9	UVRA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PAS2	RL27_XYLFA		cytoplasm
Q9PAS7	SYI_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PAS8	LSPA_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9PAW2	QUEF_XYLFA	cytoplasm (probable).	cytoplasm
Q9PAX0	LEU2_XYLFA		cytoplasm
Q9PAX1	LEUD_XYLFA		cytoplasm
Q9PAX3	LEU3_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PAY7	RUMA_XYLFA		cytoplasm
Q9PAZ0	NAGZ_XYLFA	cytoplasm (by similarity).	extracellular
Q9PAZ2	MTNP_XYLFA		cytoplasm
Q9PAZ7	SSRP_XYLFA	cytoplasm (potential).	cytoplasm
Q9PAZ9	Y2346_XYLFA		inner membrane
Q9PB02	RECN_XYLFA		cytoplasm
Q9PB03	HRCA_XYLFA		cytoplasm
Q9PB04	GRPE_XYLFA	cytoplasm (probable).	cytoplasm
Q9PB05	DNAK_XYLFA		cytoplasm
Q9PB06	DNAJ_XYLFA	cytoplasm (probable).	cytoplasm
Q9PB12	LGT_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PB13	TYSY_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PB19	SERC_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PB21	AROA_XYLFA	cytoplasm (probable).	cytoplasm
Q9PB34	UVRC_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PB43	GSA_XYLFA	cytoplasm (potential).	cytoplasm
Q9PB63	PARB_XYLFA		cytoplasm
Q9PB72	METE_XYLFA		cytoplasm

Q9PB76	GLPK_XYLFA		cytoplasm
Q9PB87	Y2257_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PB89	ACSA_XYLFA		cytoplasm
Q9PB95	DXS_XYLFA		cytoplasm
Q9PB96	RECO_XYLFA		cytoplasm
Q9PB97	ERA_XYLFA		cytoplasm
Q9PB98	RNC_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBA1	LEPA_XYLFA		cytoplasm
Q9PBB6	Y2228_XYLFA		inner membrane
Q9PBB8	PYRB_XYLFA		cytoplasm
Q9PBC0	KHSE_XYLFA	cytoplasm (potential).	cytoplasm
Q9PBC4	HIS1_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBC5	HISX_XYLFA		cytoplasm
Q9PBC6	HIS8_XYLFA		cytoplasm
Q9PBC7	HIS7_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBC8	HIS5_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBC9	HIS4_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBD0	HIS6_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBD1	HIS2_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBE1	EFPL_XYLFA		cytoplasm
Q9PBE3	PRMA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBG5	NADD_XYLFA		cytoplasm
Q9PBI6	RNH_XYLFA	cytoplasm (potential).	extracellular
Q9PBJ1	KDKA_XYLFA	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q9PBJ4	APAH_XYLFA		extracellular
Q9PBJ5	APAG_XYLFA		inner membrane
Q9PBJ6	KSGA_XYLFA		cytoplasm
Q9PBJ8	RNT_XYLFA		cytoplasm
Q9PBJ9	PHOU_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBK0	PSTB_XYLFA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PBK1	PSTA_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PBK2	PSTC_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PBK3	PSTS_XYLFA	periplasm (potential).	periplasm
Q9PBN4	GIDA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBP4	Y2096_XYLFA		inner membrane
Q9PBQ0	PPNK_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PBR9	Y2071_XYLFA		inner membrane
Q9PBX0	RPIA_XYLFA		cytoplasm
Q9PBX5	Y2010_XYLFA		inner membrane
Q9PBY7	FISL_XYLFA		cytoplasm
Q9PC00	RLMB_XYLFA		cytoplasm
Q9PC02	GCH1_XYLFA		cytoplasm
Q9PC09	PUR2_XYLFA		cytoplasm
Q9PC10	PUR9_XYLFA		cytoplasm
Q9PC29	GSHB_XYLFA		cytoplasm
Q9PC45	MSRA_XYLFA		cytoplasm

Q9PC50	GIDB_XYLFA		cytoplasm
Q9PC57	NADB_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PC58	NADA_XYLFA	cytoplasm (by similarity).	inner membrane
Q9PC61	TRPR_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PC73	FETP_XYLFA		cytoplasm
Q9PC75	Y1906_XYLFA		inner membrane
Q9PC76	RUVC_XYLFA		cytoplasm
Q9PC77	RUVA_XYLFA		cytoplasm
Q9PC78	KUP_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PC79	RUVB_XYLFA		cytoplasm
Q9PC84	TOLB_XYLFA	periplasm (potential).	periplasm
Q9PC88	GPMA_XYLFA		cytoplasm
Q9PC89	NFI_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PC93	THIC_XYLFA		cytoplasm
Q9PCE0	UPPP_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PCE6	Y1835_XYLFA		cytoplasm
Q9PCE8	UBIB_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PCF9	ILVC_XYLFA		cytoplasm
Q9PCG3	LEU1_XYLFA		cytoplasm
Q9PCG4	FABH_XYLFA	cytoplasm (probable).	cytoplasm
Q9PCG5	RL32_XYLFA		cytoplasm
Q9PCH1	RECR_XYLFA		cytoplasm
Q9PCH2	Y1808_XYLFA		inner membrane
Q9PCH7	GPDA_XYLFA	cytoplasm (probable).	cytoplasm
Q9PCH8	SECB_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PCQ3	Y1725_XYLFA		cytoplasm
Q9PCR2	MUTS_XYLFA		cytoplasm
Q9PCR3	MTGA_XYLFA	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q9PCT9	Y1666_XYLFA		extracellular
Q9PCX7	HEM3_XYLFA		cytoplasm
Q9PCY1	OPGH_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PCZ7	SLYX_XYLFA		inner membrane
Q9PD25	FUMC_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PD40	SPED_XYLFA		cytoplasm
Q9PD43	RS9_XYLFA		cytoplasm
Q9PD45	RL31B_XYLFA		cytoplasm
Q9PD67	HEMK_XYLFA		cytoplasm
Q9PD69	PIP_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PD73	HAM1_XYLFA		extracellular
Q9PD74	RNPH_XYLFA		cytoplasm
Q9PD76	KGUA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PD77	RPOZ_XYLFA		cytoplasm
Q9PD78	CYSNC_XYLFA		cytoplasm
Q9PD79	CYSD_XYLFA		cytoplasm
Q9PD82	CYSH_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PD85	Y1494_XYLFA		extracellular

Q9PD92	UBIE_XYLFA		cytoplasm
Q9PD94	HSLU_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PD96	XERC_XYLFA	cytoplasm (by similarity).	nucleoid
Q9PD98	DAPF_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PDA6	CSD_XYLFA		cytoplasm
Q9PDB1	GLMM_XYLFA		cytoplasm
Q9PDB9	Y1460_XYLFA		cytoplasm
Q9PDC5	CRCB_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PDC7	LOLA_XYLFA	periplasm (by similarity).	periplasm
Q9PDD3	LFTR_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PDD7	CLPS_XYLFA		cytoplasm
Q9PDD9	TRMU_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PDE0	Y1439_XYLFA		inner membrane
Q9PDE9	ENGB_XYLFA		cytoplasm
Q9PDF2	ARGD_XYLFA	cytoplasm (probable).	cytoplasm
Q9PDF4	XERD_XYLFA	cytoplasm (by similarity).	nucleoid
Q9PDF6	PUR4_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PDG4	MURA_XYLFA	cytoplasm (probable).	inner membrane
Q9PDH3	HPRK_XYLFA		cytoplasm
Q9PDH4	Y1405_XYLFA		cytoplasm
Q9PDH6	PTHP_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PDI7	SSB2_XYLFA		cytoplasm
Q9PDJ4	GCSP_XYLFA		cytoplasm
Q9PDK3	TRPA_XYLFA		cytoplasm
Q9PDK4	TRPB_XYLFA		cytoplasm
Q9PDK5	TRPF_XYLFA		cytoplasm
Q9PDK6	TRUA_XYLFA		cytoplasm
Q9PDL0	AROC_XYLFA		cytoplasm
Q9PDL1	Y1368_XYLFA		cytoplasm
Q9PDL4	PSD_XYLFA		cytoplasm
Q9PDL7	CCA_XYLFA		cytoplasm
Q9PDM3	BIOH_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PDM9	RPOD_XYLFA		cytoplasm
Q9PDN2	CYSA_XYLFA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PDN8	CUTC_XYLFA	cytoplasm (probable).	periplasm
Q9PDP2	PDXH_XYLFA		cytoplasm
Q9PDP4	AROK_XYLFA	cytoplasm (probable).	cytoplasm
Q9PDP5	AROB_XYLFA	cytoplasm (probable).	cytoplasm
Q9PDP7	DCUP_XYLFA	cytoplasm (probable).	cytoplasm
Q9PDQ7	MINC_XYLFA		inner membrane
Q9PDR5	QUEA_XYLFA	cytoplasm (potential).	cytoplasm
Q9PDT5	ISPF_XYLFA		extracellular
Q9PDT6	ISPD_XYLFA		cytoplasm
Q9PDT7	FTSB_XYLFA	bacterial cell inner membrane; single-pass type ii membrane protein (potential). colocalizes with ftsl to the division site (by similarity).	inner membrane
Q9PDT8	ENO_XYLFA	cytoplasm (by similarity).	extracellular
Q9PDU0	KDSA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PDU1	PYRG_XYLFA		cytoplasm
Q9PDV8	Y1271_XYLFA		inner membrane

Q9PDV9	LIPB_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PDW0	LIPA_XYLFA	cytoplasm (potential).	cytoplasm
Q9PDW9	Y1260_XYLFA		cytoplasm
Q9PDX2	ORN_XYLFA	cytoplasm (potential).	cytoplasm
Q9PDX7	Y1252_XYLFA		cytoplasm
Q9PE17	MDH_XYLFA		cytoplasm
Q9PE38	DBH_XYLFA		cytoplasm
Q9PE40	CLPX_XYLFA		cytoplasm
Q9PE41	CLPP_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PE42	TIG_XYLFA		cytoplasm
Q9PE46	LIFO_XYLFA	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
Q9PE53	RS4_XYLFA		cytoplasm
Q9PE54	RS11_XYLFA		cytoplasm
Q9PE55	RS13_XYLFA		cytoplasm
Q9PE59	RS5_XYLFA		cytoplasm
Q9PE60	RL18_XYLFA		cytoplasm
Q9PE62	RS8_XYLFA		cytoplasm
Q9PE63	RS14_XYLFA		cytoplasm
Q9PE64	RL5_XYLFA		cytoplasm
Q9PE65	RL24_XYLFA		cytoplasm
Q9PE67	RS17_XYLFA		cytoplasm
Q9PE68	RL29_XYLFA		cytoplasm
Q9PE69	RL16_XYLFA		cytoplasm
Q9PE70	RS3_XYLFA		cytoplasm
Q9PE71	RL22_XYLFA		cytoplasm
Q9PE72	RS19_XYLFA		cytoplasm
Q9PE73	RL2_XYLFA		cytoplasm
Q9PE75	RL4_XYLFA		cytoplasm
Q9PE76	RL3_XYLFA		cytoplasm
Q9PE83	ATPA_XYLFA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PE84	ATPG_XYLFA		inner membrane
Q9PE86	ATPE_XYLFA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PE88	GLMU_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PE98	Y1130_XYLFA		inner membrane
Q9PEA4	Y1124_XYLFA	cytoplasm (potential).	inner membrane
Q9PEB0	MURD_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEC0	GREA_XYLFA		cytoplasm
Q9PEC1	CARB_XYLFA		cytoplasm
Q9PEC2	CARA_XYLFA		cytoplasm
Q9PEC3	DAPB_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PED1	PNCB_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEE6	LPXK_XYLFA		cytoplasm
Q9PEE7	MSBA_XYLFA	bacterial cell inner membrane; multi- pass membrane protein (by similarity).	inner membrane
Q9PEF2	LOLC_XYLFA	bacterial cell inner membrane; multi- pass membrane protein (by similarity).	inner membrane
Q9PEF4	Y1074_XYLFA		cytoplasm
Q9PEG4	GLK_XYLFA	cytoplasm (by similarity).	cytoplasm

Q9PEG5	6PGL_XYLFA		extracellular
Q9PEH0	PYRH_XYLFA		cytoplasm
Q9PEH7	RRF_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEH8	UPPS_XYLFA		inner membrane
Q9PEI0	DXR_XYLFA		cytoplasm
Q9PEI1	Y1047_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9PEI3	LPXD_XYLFA		cytoplasm
Q9PEI4	FABZ_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEI5	LPXA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEI6	LPXB_XYLFA		cytoplasm
Q9PEI7	RNH2_XYLFA	cytoplasm (potential).	extracellular
Q9PEJ1	SAHH_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEJ7	PLSB_XYLFA	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PEL0	ATE_XYLFA	cytoplasm (potential).	cytoplasm
Q9PEL3	MNTH_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9PEM3	PROA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEM4	PROB_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEM5	ARLY_XYLFA	cytoplasm (probable).	cytoplasm
Q9PEM6	ARGC_XYLFA	cytoplasm (probable).	cytoplasm
Q9PEM7	ARGB_XYLFA	cytoplasm (probable).	cytoplasm
Q9PEM9	ASSY_XYLFA	cytoplasm (probable).	cytoplasm
Q9PEN0	AOTC_XYLFA	cytoplasm (probable).	cytoplasm
Q9PEN9	Y989_XYLFA		inner membrane
Q9PEP8	COAD_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEQ0	HTPG_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PEQ2	DCTA_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PER1	UVRB_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PER5	DAPA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PES3	NUSB_XYLFA		cytoplasm
Q9PES4	RISB_XYLFA		cytoplasm
Q9PET0	NRDR_XYLFA		cytoplasm
Q9PET2	GLYA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PET6	MQO_XYLFA		cytoplasm
Q9PET8	Y940_XYLFA		cytoplasm
Q9PET9	RLUD_XYLFA		cytoplasm
Q9PEU0	Y938_XYLFA	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q9PEV0	RSMB_XYLFA		cytoplasm
Q9PEV1	FMT_XYLFA		cytoplasm
Q9PEV5	SMG_XYLFA		cytoplasm
Q9PEV8	TOP1_XYLFA		cytoplasm
Q9PEX5	Y903_XYLFA		inner membrane
Q9PF20	SURE_XYLFA	cytoplasm (potential).	periplasm
Q9PF26	ALR_XYLFA		cytoplasm
Q9PF27	DADA_XYLFA		cytoplasm
Q9PF29	MSRB_XYLFA		cytoplasm
Q9PF39	PDXA_XYLFA	cytoplasm (probable).	cytoplasm

Q9PF41	OSTA_XYLFA	bacterial cell outer membrane (by similarity).	outer membrane
Q9PF52	ALF1_XYLFA		cytoplasm
Q9PF55	PGK_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PF75	LPXC_XYLFA		cytoplasm
Q9PF79	DDL_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PF80	MURC_XYLFA	cytoplasm (probable).	cytoplasm
Q9PF81	MURG_XYLFA	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PF83	MRAY_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PF85	MURE_XYLFA	cytoplasm (probable).	cytoplasm
Q9PF88	MRAW_XYLFA		cytoplasm
Q9PF89	MRAZ_XYLFA		inner membrane
Q9PF94	TRMB_XYLFA		cytoplasm
Q9PF95	THIG_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PFB6	DCD_XYLFA		extracellular
Q9PFB8	MUTL_XYLFA		cytoplasm
Q9PFC3	EX7L_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PFD0	MTFA_XYLFA		cytoplasm
Q9PFD5	IHFA_XYLFA		cytoplasm
Q9PFD8	RL20_XYLFA		cytoplasm
Q9PFJ7	EX7S_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PFJ8	TILS_XYLFA	cytoplasm (probable).	cytoplasm
Q9PFN3	AROE_XYLFA		cytoplasm
Q9PFP2	CH60_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PFQ9	Y598_XYLFA		cytoplasm
Q9PFS0	PUR5_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PFS7	KTHY_XYLFA		cytoplasm
Q9PFT5	FABA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PFT9	RDGC_XYLFA	nucleoid (by similarity). seems to be associated with the nucleoid (by similarity).	cytoplasm
Q9PFU1	HEMH_XYLFA	cytoplasm (by similarity).	extracellular
Q9PFU3	TATA_XYLFA	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9PFU4	TATB_XYLFA	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9PFV1	ENGC_XYLFA		cytoplasm
Q9PFV3	Y554_XYLFA		inner membrane
Q9PFV5	Y552_XYLFA		cytoplasm
Q9PG37	ENGA_XYLFA		cytoplasm
Q9PG44	NDK_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PG47	PURA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PG68	RS21_XYLFA		cytoplasm
Q9PG89	RMUC_XYLFA		cytoplasm
Q9PGA9	NUDH_XYLFA		cytoplasm
Q9PGB5	DUSA_XYLFA		cytoplasm
Q9PGC1	CLPB_XYLFA	cytoplasm (probable).	cytoplasm
Q9PGC4	THIE_XYLFA		cytoplasm
Q9PGC5	C1331_XYLFA		cytoplasm

Q9PGD4	PBPA_XYLFA	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q9PGE6	C1332_XYLFA		cytoplasm
Q9PGG2	DSBB_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PGG4	Y338_XYLFA		inner membrane
Q9PGJ7	TPIS_XYLFA		cytoplasm
Q9PGM3	KAD_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PGR0	RS15_XYLFA		cytoplasm
Q9PGR1	TRUB_XYLFA		cytoplasm
Q9PGR2	RBFA_XYLFA	cytoplasm (potential).	cytoplasm
Q9PGR5	Y233_XYLFA		inner membrane
Q9PGR6	G6PI_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PGR7	PAND_XYLFA		cytoplasm
Q9PGR8	PANC_XYLFA		cytoplasm
Q9PGR9	PANB_XYLFA		cytoplasm
Q9PGS5	TGT_XYLFA		cytoplasm
Q9PGT5	TRPC_XYLFA		cytoplasm
Q9PGT6	TRPD_XYLFA		cytoplasm
Q9PGU3	PUR7_XYLFA		cytoplasm
Q9PGU4	DPO3A_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PGU5	ACCA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PGW5	GCST_XYLFA		cytoplasm
Q9PGW7	GCSH_XYLFA		cytoplasm
Q9PGX7	SYX_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PGZ3	PYRE_XYLFA		cytoplasm
Q9PGZ6	DUT_XYLFA		extracellular
Q9PGZ8	RADC_XYLFA		cytoplasm
Q9PH02	SPEA_XYLFA		cytoplasm
Q9PH03	SPEE_XYLFA		cytoplasm
Q9PH05	GLMS_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PH08	AMPA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PH12	SYV_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PH23	RECA_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PH24	LEXA_XYLFA		cytoplasm
Q9PH32	DAPD_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PH36	RL19_XYLFA		cytoplasm
Q9PH37	TRMD_XYLFA	cytoplasm (potential).	cytoplasm
Q9PH38	RIMM_XYLFA	cytoplasm (potential).	cytoplasm
Q9PH39	RS16_XYLFA		cytoplasm
Q9PH47	ILVD_XYLFA		cytoplasm
Q9PH52	RRMJ_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PH56	MIAA_XYLFA		cytoplasm
Q9PH57	HFQ_XYLFA		cytoplasm
Q9PH84	PDXJ_XYLFA	cytoplasm (probable).	cytoplasm
Q9PH91	HMP_XYLFA		cytoplasm
Q9PH97	AROQ_XYLFA		cytoplasm
Q9PHA5	MSCL_XYLFA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PHB0	PYRF_XYLFA		cytoplasm
Q9PHC7	HEM6_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PHE1	RECF_XYLFA	cytoplasm (by similarity).	cytoplasm
Q9PHE3	DNAA_XYLFA	cytoplasm (by similarity).	cytoplasm

Q9PHE7	SSB1_XYLFA		cytoplasm
Q9PHE9	PARB2_XYLFA		cytoplasm
Q9PHK2	TOP1P_XYLFA		cytoplasm
Q9PHK6	RPP_XYLFA		nucleoid
Q9PHK7	ASSY_CAMJE	cytoplasm (probable).	cytoplasm
Q9PHK8	RL9_CAMJE		cytoplasm
Q9PHK9	HSLV_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PHL0	HSLU_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PHL1	ERA_CAMJE		cytoplasm
Q9PHL7	ENGB_CAMJE		cytoplasm
Q9PHM1	RLPA_CAMJE	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q9PHM5	RECN_CAMJE		cytoplasm
Q9PHM6	PPNK_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PHM9	IPYR_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PHN0	MSRA_CAMJE		cytoplasm
Q9PHN2	RUVX_CAMJE	cytoplasm (potential).	cytoplasm
Q9PHN5	ILVC_CAMJE		cytoplasm
Q9PHP0	HYP_A_CAMJE		cytoplasm
Q9PHQ1	PSTB_CAMJE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PHR3	DSBD_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PHS3	Y593_CAMJE	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9PHT5	NUDH_CAMJE		cytoplasm
Q9PHT7	TATB_CAMJE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9PHT8	TATC_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9PHT9	QUEA_CAMJE	cytoplasm (potential).	cytoplasm
Q9PHU0	LPXD_CAMJE		cytoplasm
Q9PHU4	RIBAB_CAMJE		cytoplasm
Q9PHW9	HEM3_CAMJE		cytoplasm
Q9PHX2	HEM1_CAMJE		cytoplasm
Q9PHY1	SUCC_CAMJE		cytoplasm
Q9PHY2	MDH_CAMJE		cytoplasm
Q9PHY8	FLIE_CAMJE		flagellar
Q9PHZ3	HTPG_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PHZ4	CRCB_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PHZ7	PURQ_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PHZ9	PUR7_CAMJE		cytoplasm
Q9PI02	CLPB_CAMJE	cytoplasm (probable).	cytoplasm
Q9PI04	Y507_CAMJE	cytoplasm (potential).	inner membrane
Q9PI08	HEMH_CAMJE	cytoplasm (by similarity).	extracellular
Q9PI11	TRPC_CAMJE		cytoplasm
Q9PI17	RS7_CAMJE		cytoplasm
Q9PI18	RS12_CAMJE		cytoplasm
Q9PI30	RPOC_CAMJE		cytoplasm

Q9PI32	RL7_CAMJE		cytoplasm
Q9PI33	RL10_CAMJE		cytoplasm
Q9PI34	RL1_CAMJE		cytoplasm
Q9PI35	RL11_CAMJE		cytoplasm
Q9PI36	NUSG_CAMJE		cytoplasm
Q9PI38	RL33_CAMJE		cytoplasm
Q9PI55	THIC_CAMJE		cytoplasm
Q9PI58	RL28_CAMJE		cytoplasm
Q9PI62	ACCA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PI64	ACP_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PI71	GPMI_CAMJE		cytoplasm
Q9PI72	MRAY_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PI73	MURD_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PI98	LGT_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PIA5	GATC_CAMJE		cytoplasm
Q9PIB6	ENGA_CAMJE		cytoplasm
Q9PIB8	KDSA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PIB9	RISB_CAMJE		cytoplasm
Q9PIC0	NUSB_CAMJE		cytoplasm
Q9PIC1	PYRF_CAMJE		cytoplasm
Q9PIC3	Y379_CAMJE		cytoplasm
Q9PIC8	Y374_CAMJE		inner membrane
Q9PID1	Y371_CAMJE	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q9PID2	RS21_CAMJE		cytoplasm
Q9PIE2	GLMM_CAMJE		cytoplasm
Q9PIE4	Y357_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PIF1	TRPA_CAMJE		cytoplasm
Q9PIF2	TRPB_CAMJE		cytoplasm
Q9PIF3	TRPF_CAMJE		cytoplasm
Q9PIG7	NDK_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PIH0	PLSX_CAMJE		cytoplasm
Q9PIH1	FABH_CAMJE	cytoplasm (probable).	cytoplasm
Q9PIH3	SERC_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PIH4	EX7L_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PIH5	UBIE_CAMJE		cytoplasm
Q9PIH8	DXS_CAMJE		cytoplasm
Q9PII2	HIS8_CAMJE		cytoplasm
Q9PII5	DCDA_CAMJE		cytoplasm
Q9PII7	PTH_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PII8	RL25_CAMJE		cytoplasm
Q9PIJ1	BIOD_CAMJE		cytoplasm
Q9PIK1	PANB_CAMJE		cytoplasm
Q9PIK2	PANC_CAMJE		cytoplasm
Q9PIK3	PAND_CAMJE		cytoplasm
Q9PIK6	SURE_CAMJE	cytoplasm (potential).	periplasm
Q9PIK8	LPXB_CAMJE		cytoplasm
Q9PIK9	GREA_CAMJE		cytoplasm
Q9PIL5	TAL_CAMJE	cytoplasm (probable).	cytoplasm
Q9PIL7	CARB_CAMJE		cytoplasm

Q9PIM0	CLPX_CAMJE		cytoplasm
Q9PIM1	LPXA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PIM2	FABZ_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PIM5	Y270_CAMJE		cytoplasm
Q9PIN2	ZUPT_CAMJE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9PIP2	Y253_CAMJE		cytoplasm
Q9PIP3	MOAC_CAMJE		cytoplasm
Q9PIQ0	RL20_CAMJE		cytoplasm
Q9PIQ1	RL35_CAMJE		cytoplasm
Q9PIQ3	HEMTL_CAMJE		cytoplasm
Q9PIQ8	Y236_CAMJE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9PIR0	RRF_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PIR1	PYRE_CAMJE		cytoplasm
Q9PIR7	ARGD_CAMJE	cytoplasm (probable).	cytoplasm
Q9PIR8	ARGB_CAMJE	cytoplasm (probable).	cytoplasm
Q9PIS0	ARGC_CAMJE	cytoplasm (probable).	cytoplasm
Q9PIS4	UPPP_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PIT2	DAPB_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PIT8	DEF_CAMJE		cytoplasm
Q9PIW1	Y167_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PIW2	MIAA_CAMJE		cytoplasm
Q9PIW6	MOAA_CAMJE		cytoplasm
Q9PIX2	RL31_CAMJE		cytoplasm
Q9PIX9	Y148_CAMJE		inner membrane
Q9PIY9	Y138_CAMJE		inner membrane
Q9PIZ0	RBFA_CAMJE	cytoplasm (potential).	cytoplasm
Q9PIZ3	KHSE_CAMJE	cytoplasm (potential).	cytoplasm
Q9PIZ5	LPXC_CAMJE		cytoplasm
Q9PJ01	Y126_CAMJE		inner membrane
Q9PJ06	Y121_CAMJE		inner membrane
Q9PJ14	TOLB_CAMJE	periplasm (potential).	periplasm
Q9PJ18	ATPE_CAMJE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PJ20	ATPG_CAMJE		inner membrane
Q9PJ21	ATPA_CAMJE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PJ25	PARB_CAMJE		cytoplasm
Q9PJ28	FMT_CAMJE		cytoplasm
Q9PJ29	PROB_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PJ31	RL27_CAMJE		cytoplasm
Q9PJ40	UNG_CAMJE	cytoplasm (by similarity).	extracellular
Q9PJ53	AROQ_CAMJE		cytoplasm
Q9PJ54	HPPK_CAMJE		cytoplasm
Q9PJ66	TRMU_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PJ84	PYRG_CAMJE		cytoplasm
Q9PJ85	THYX_CAMJE		cytoplasm

Q9PJ98	ILVD_CAMJE		cytoplasm
Q9PJA1	RNH2_CAMJE	cytoplasm (potential).	extracellular
Q9PJB0	DNAA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PLU8	RUVC_CAMJE		cytoplasm
Q9PLV2	META_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PLV4	QUEF_CAMJE	cytoplasm (probable).	cytoplasm
Q9PLV9	LEU1_CAMJE		cytoplasm
Q9PLW0	LEU3_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PLW1	LEU2_CAMJE		cytoplasm
Q9PLW2	LEUD_CAMJE		cytoplasm
Q9PLW7	KSGA_CAMJE		cytoplasm
Q9PLX0	RS10_CAMJE		cytoplasm
Q9PLX1	RL3_CAMJE		cytoplasm
Q9PLX2	RL4_CAMJE		cytoplasm
Q9PLX4	RL2_CAMJE		cytoplasm
Q9PLX5	RS19_CAMJE		cytoplasm
Q9PLX6	RL22_CAMJE		cytoplasm
Q9PLX7	RS3_CAMJE		cytoplasm
Q9PLX8	RL16_CAMJE		cytoplasm
Q9PLX9	RL29_CAMJE		cytoplasm
Q9PLY0	RS17_CAMJE		cytoplasm
Q9PLY2	RL24_CAMJE		cytoplasm
Q9PLY3	RL5_CAMJE		cytoplasm
Q9PLY4	RS14_CAMJE		cytoplasm
Q9PLY5	RS8_CAMJE		cytoplasm
Q9PLY7	RL18_CAMJE		cytoplasm
Q9PLY8	RS5_CAMJE		cytoplasm
Q9PLZ2	TOP1_CAMJE		cytoplasm
Q9PM01	MURB_CAMJE	cytoplasm (probable).	inner membrane
Q9PM24	MURI_CAMJE		cytoplasm
Q9PM34	Y1642_CAMJE		inner membrane
Q9PM37	PRIM_CAMJE		cytoplasm
Q9PM39	RNH_CAMJE	cytoplasm (potential).	extracellular
Q9PM40	RNC_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PM41	AROC_CAMJE		cytoplasm
Q9PM64	RS20_CAMJE		cytoplasm
Q9PM68	ISPDF_CAMJE		cytoplasm
Q9PM71	HIS2_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PM72	HIS61_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PM74	HIS4_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PM75	HIS52_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PM76	HIS7_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PM77	HISX_CAMJE		cytoplasm
Q9PM78	HIS1_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PM80	RPOA_CAMJE		cytoplasm
Q9PM81	RS4_CAMJE		cytoplasm
Q9PM82	RS11_CAMJE		cytoplasm
Q9PM83	RS13_CAMJE		cytoplasm
Q9PM99	NUOD_CAMJE		cytoplasm
Q9PMA7	NUOL_CAMJE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9PMC8	Y1541_CAMJE		inner membrane
Q9PMD2	ACSA_CAMJE		cytoplasm
Q9PMD4	G6PI_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMD8	DAPF_CAMJE	cytoplasm (by similarity).	cytoplasm

Q9PMD9	COAE_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMF9	SELD_CAMJE		cytoplasm
Q9PMG4	PURA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMG8	CARA_CAMJE		cytoplasm
Q9PMI3	RS9_CAMJE		cytoplasm
Q9PMJ7	Y1463_CAMJE		cytoplasm
Q9PMJ8	FLGI_CAMJE	periplasm (by similarity).	periplasm
Q9PMK3	TRUD_CAMJE		cytoplasm
Q9PMK7	TILS_CAMJE	cytoplasm (probable).	cytoplasm
Q9PML1	Y1449_CAMJE		inner membrane
Q9PMN3	GMHA2_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMP8	ACPS_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMQ3	NADD_CAMJE		cytoplasm
Q9PMQ5	PGK_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMQ6	TPIS_CAMJE		cytoplasm
Q9PMQ9	FEOB_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	cytoplasm
Q9PMR0	Y1397_CAMJE		cytoplasm
Q9PMS2	SELA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMS6	HAM1_CAMJE		extracellular
Q9PMT4	GLMS_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMT7	RUVB_CAMJE		cytoplasm
Q9PMU0	PPK_CAMJE		cytoplasm
Q9PMU9	MOBA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMV3	DXR_CAMJE		cytoplasm
Q9PMY4	HIS51_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMY5	HIS62_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PMZ6	FABHL_CAMJE	cytoplasm (probable).	cytoplasm
Q9PN07	DCD_CAMJE		extracellular
Q9PN13	UPP_CAMJE		cytoplasm
Q9PN20	TRMB_CAMJE		cytoplasm
Q9PN24	PYRH_CAMJE		cytoplasm
Q9PN25	RPOZ_CAMJE		cytoplasm
Q9PN27	SYU_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PN30	Y1268_CAMJE		cytoplasm
Q9PN35	RECR_CAMJE		cytoplasm
Q9PN47	PUR2_CAMJE		cytoplasm
Q9PN49	GUAA_CAMJE		cytoplasm
Q9PN51	UVRC_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PN54	DCUP_CAMJE	cytoplasm (probable).	cytoplasm
Q9PN58	PDXA_CAMJE	cytoplasm (probable).	cytoplasm
Q9PN59	PDXJ_CAMJE	cytoplasm (probable).	cytoplasm
Q9PN78	Y1217_CAMJE		cytoplasm
Q9PN86	Y1209_CAMJE		inner membrane
Q9PN90	RADA_CAMJE		cytoplasm
Q9PN94	METE_CAMJE		cytoplasm
Q9PN97	LUXS_CAMJE		extracellular
Q9PN98	GATB_CAMJE		cytoplasm
Q9PN99	GPDA_CAMJE	cytoplasm (probable).	cytoplasm
Q9PNA7	GIDA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PNB3	RS2_CAMJE		cytoplasm
Q9PNB8	KGUA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PNB9	TATA_CAMJE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane

Q9PNC3	Y1172_CAMJE		inner membrane
Q9PNE6	GMHA1_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PNH7	PRMA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PNI5	LFTR_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PNI7	CLPS_CAMJE		cytoplasm
Q9PNI9	SSRP_CAMJE	cytoplasm (potential).	cytoplasm
Q9PNJ0	ISPE_CAMJE		cytoplasm
Q9PNJ2	TRUB_CAMJE		cytoplasm
Q9PNJ6	PYRB_CAMJE		cytoplasm
Q9PNJ9	LNT_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PNL3	THIE_CAMJE		cytoplasm
Q9PNM6	Y1068_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9PNN2	GATA_CAMJE		cytoplasm
Q9PNN7	MURC_CAMJE	cytoplasm (probable).	cytoplasm
Q9PNP6	THIG_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PNQ2	MURG_CAMJE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9PNQ6	ATE_CAMJE	cytoplasm (potential).	cytoplasm
Q9PNR1	LEPA_CAMJE		cytoplasm
Q9PNT0	TGT_CAMJE		cytoplasm
Q9PNT2	AROB_CAMJE	cytoplasm (probable).	cytoplasm
Q9PNU1	Y999_CAMJE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9PNU3	GIDB_CAMJE		cytoplasm
Q9PNU6	OTC_CAMJE	cytoplasm (probable).	cytoplasm
Q9PNX5	RNPA_CAMJE		extracellular
Q9PNX6	Y959_CAMJE		inner membrane
Q9PNX7	OXAA_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PNX9	TRME_CAMJE		cytoplasm
Q9PNY0	PURL_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PNY2	PUR9_CAMJE		cytoplasm
Q9PNZ0	LOLA_CAMJE	periplasm (by similarity).	periplasm
Q9PP01	PPCK_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PP04	AMPA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PP15	KPRS_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PP26	ALR_CAMJE		cytoplasm
Q9PP65	MURA_CAMJE	cytoplasm (probable).	inner membrane
Q9PP70	GSA_CAMJE	cytoplasm (potential).	cytoplasm
Q9PP76	PSD_CAMJE		cytoplasm
Q9PP77	Y846_CAMJE		extracellular
Q9PP92	TRMA_CAMJE		cytoplasm
Q9PP96	TRUA_CAMJE		cytoplasm
Q9PP99	UPPS_CAMJE		inner membrane
Q9PPA2	GLMU_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PPA9	LPXK_CAMJE		cytoplasm
Q9PPB0	NADE_CAMJE		cytoplasm
Q9PPB4	DAPA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PPC1	RUVA_CAMJE		cytoplasm

Q9PPC2	DDL_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PPD9	NAPA_CAMJE	periplasm (by similarity).	periplasm
Q9PPE0	TPX_CAMJE		cytoplasm
Q9PPE4	SYV_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PPF1	PAAD_CAMJE		cytoplasm
Q9PPF2	COAD_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PPF3	KTHY_CAMJE		cytoplasm
Q9PPG2	HRCA_CAMJE		cytoplasm
Q9PPI1	CORA_CAMJE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9PPI9	DPO3A_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PPJ3	RL19_CAMJE		cytoplasm
Q9PPJ4	TRMD_CAMJE	cytoplasm (potential).	cytoplasm
Q9PPJ5	RIMM_CAMJE	cytoplasm (potential).	cytoplasm
Q9PPJ7	RS16_CAMJE		cytoplasm
Q9PPK2	Y705_CAMJE		cytoplasm
Q9PPL4	MRAW_CAMJE		cytoplasm
Q9PPL8	ACKA_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9PPM0	FLGH_CAMJE	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q9PPM1	ISPG_CAMJE		cytoplasm
Q9PPM7	UVRB_CAMJE	cytoplasm (by similarity).	cytoplasm
Q9R3W3	SIRC_SALTY		cytoplasm
Q9R4E4	AROA_AGRSC	cytoplasm (probable).	cytoplasm
Q9R6Z3	ACP_PHOPR	cytoplasm (by similarity).	cytoplasm
Q9R9L6	GPDA_RHIME	cytoplasm (probable).	cytoplasm
Q9R9M8	AAU3_RHIME		cytoplasm
Q9R9M9	LPSC_RHIME		cytoplasm
Q9R9N0	LPSD_RHIME		cytoplasm
Q9R9N1	LPSE_RHIME		cytoplasm
Q9R9N2	LPSB_RHIME		cytoplasm
Q9R9N3	ODP2_RHIME		inner membrane
Q9R9N4	ODPB_RHIME		cytoplasm
Q9R9N5	ODPA_RHIME		cytoplasm
Q9R9S0	YADS_AERPU	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9R9T9	SRPR_PSEPU		cytoplasm
Q9R9U0	SRPS_PSEPU		cytoplasm
Q9R9U2	RECA_PSEOL	cytoplasm (by similarity).	cytoplasm
Q9R9W0	FADA_PSEPU	cytoplasm (by similarity).	cytoplasm
Q9RA11	YU61_VIBMA		inner membrane
Q9RA13	Y267_VIBMA		inner membrane
Q9RA32	ACP_VIBMA	cytoplasm (by similarity).	cytoplasm
Q9RA35	PLSX_VIBMA		cytoplasm
Q9RA36	RL32_VIBMA		cytoplasm
Q9RA96	SLYA_SERMA		cytoplasm
Q9RAA9	DPO1_RICFE		cytoplasm
Q9RAE4	GLND_RHILV		cytoplasm
Q9RAE6	DADA_RHILV		cytoplasm
Q9RAE7	ALR2_RHILV		cytoplasm
Q9RAM6	KHSE_METFL		cytoplasm
Q9RAM7	TYSY_METFL	cytoplasm (by similarity).	cytoplasm
Q9RB09	SLYA_PECCC		cytoplasm
Q9RBJ0	ISPH_ACIAD		cytoplasm

Q9RBJ2	BCSA4_ACEXY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9RBJ3	UPP_XANCP		cytoplasm
Q9RCE7	TRPA_VIBME		cytoplasm
Q9RCE8	TRPB_VIBME		cytoplasm
Q9RCG7	PAXB_PASAE	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9RCT3	HLYEL_SHIFL		cytoplasm
Q9RDT9	RECX_PSETO	cytoplasm (potential).	cytoplasm
Q9RDU0	RECA_PSETO	cytoplasm (by similarity).	cytoplasm
Q9RDX2	HIS6_LEGPN	cytoplasm (by similarity).	cytoplasm
Q9RDX3	HIS5_LEGPN	cytoplasm (by similarity).	cytoplasm
Q9RDY2	G6PI_LEGPN	cytoplasm (by similarity).	cytoplasm
Q9RED7	RLMB_BURSP		cytoplasm
Q9RED9	KSGA_BURSP		cytoplasm
Q9REF2	HRCA_BRAJA		cytoplasm
Q9REF4	RISB_BARHE		cytoplasm
Q9REM0	HYP_A_RHOSH		cytoplasm
Q9REQ0	TRUA_ZYMMO		cytoplasm
Q9REQ3	RMUC_ZYMMO		cytoplasm
Q9REQ6	PURL_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9REQ9	MRAW_ZYMMO		cytoplasm
Q9RF98	OADB_HAEDU	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9RFA0	OADG_HAEDU	bacterial cell membrane; single-pass membrane protein (by similarity).	inner membrane
Q9RFA7	PHOU_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9RFD3	BCHE_RHOS4		cytoplasm
Q9RFD4	BCHN_RHOS4		cytoplasm
Q9RFD5	BCHH_RHOS4		cytoplasm
Q9RFD6	BCHL_RHOS4		cytoplasm
Q9RFF5	URE3_RHOS4	cytoplasm (by similarity).	cytoplasm
Q9RFV6	NQRF_VIBHA	bacterial cell inner membrane (potential).	inner membrane
Q9RFV7	NQRE_VIBHA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9RFV8	NQRD_VIBHA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9RFV9	NQRC_VIBHA	bacterial cell inner membrane (potential).	inner membrane
Q9RFW0	NQRB_VIBHA	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9RFW1	NQRA_VIBHA		inner membrane
Q9RFY6	CDTA_HELHP	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q9RG22	ACP_RHILE	cytoplasm (by similarity).	cytoplasm
Q9RGM3	MDCB_ACICA		cytoplasm
Q9RGQ1	IPYR_RHORT	cytoplasm (by similarity).	cytoplasm
Q9RGR0	LCBA_NEIME		cytoplasm

Q9RGS8	PHLN_BURPS		extracellular
Q9RGX2	RS21_HAEDU		cytoplasm
Q9RH04	HIS1_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9RH05	HISX_ZYMMO		cytoplasm
Q9RH11	NUDH_ZYMMO		cytoplasm
Q9RH13	Y1566_ZYMMO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9RH29	GVPN_ANCAQ		cytoplasm
Q9RH40	RPOC_RICCN		cytoplasm
Q9RH41	RPOB_RICCN		cytoplasm
Q9RH43	RPOB_RICMA		cytoplasm
Q9RH67	RECO_BRAJA		cytoplasm
Q9RH70	RPOZ_BRAJA		cytoplasm
Q9RH74	SSRP_BRAJA	cytoplasm (potential).	cytoplasm
Q9RH76	DAPA_BRAJA	cytoplasm (by similarity).	cytoplasm
Q9RHF4	SSB2_SALTI		cytoplasm
Q9RHM8	4OT_COMTE		cytoplasm
Q9RHS9	TRMA_PSEFL		cytoplasm
Q9RHV8	DSBA_BURCE	periplasm (by similarity).	periplasm
Q9RI00	HIS8_PSEST		cytoplasm
Q9RI02	SERC_PSEST	cytoplasm (by similarity).	cytoplasm
Q9RI18	LCRT_YERPE		cytoplasm
Q9RLB6	DPO1_RICHE		cytoplasm
Q9RLM3	T2D1_NEIMC		extracellular
Q9RLM4	MTD1_NEIMC		cytoplasm
Q9RLM5	VSN1_NEIMC		extracellular
Q9RM66	USPC_SALTY	cytoplasm (by similarity).	cytoplasm
Q9RMC1	G6PI_ACILW	cytoplasm (by similarity).	cytoplasm
Q9RME4	COAD_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9RME6	EX7S_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9RML6	PDXJ_NEIMC	cytoplasm (probable).	cytoplasm
Q9RMM3	MURI_NEIMC		cytoplasm
Q9RN37	USHA_SALPU	periplasm (by similarity).	periplasm
Q9RN77	AROD_SALEN		cytoplasm
Q9RND6	KCY_BORBR	cytoplasm (by similarity).	cytoplasm
Q9RND7	AROA_BORBR	cytoplasm (probable).	cytoplasm
Q9RNH6	GLGA_RHOS4		cytoplasm
Q9RNH7	GLGC_RHOS4		cytoplasm
Q9RNL1	UPPP_ZYMMO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9RNL5	OXAA_ZYMMO	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9RNL6	ENGB_ZYMMO		cytoplasm
Q9RNM2	MURE_ZYMMO	cytoplasm (probable).	cytoplasm
Q9RNM6	MURG_ZYMMO	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9RNM7	MURC_ZYMMO	cytoplasm (probable).	cytoplasm
Q9RNM8	MURB_ZYMMO	cytoplasm (probable).	inner membrane
Q9RNN5	DGTL1_ZYMMO		cytoplasm
Q9RNP3	PDXH_ZYMMO		cytoplasm

Q9RNX4	CCME_RHIET	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q9RNY0	RECA_PASHA	cytoplasm (by similarity).	cytoplasm
Q9RNY7	LIPA_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9RNZ1	ISPDF_ZYMMO		cytoplasm
Q9RNZ5	IHFA_ZYMMO		cytoplasm
Q9RNZ9	TFDD_COMAC		cytoplasm
Q9RP66	MUTL_CAUCR		cytoplasm
Q9RP67	SKGA_CAUCR		cytoplasm
Q9RPT0	RCSF1_PSEAE		inner membrane
Q9RPT1	RHLG_PSEAE		cytoplasm
Q9RPU2	HPAC_SALDU		cytoplasm
Q9RPU5	HPAG_SALDU		cytoplasm
Q9RPV0	HPAI_SALDU		cytoplasm
Q9RPW7	MCH_METOR	cytoplasm (by similarity).	cytoplasm
Q9RPW8	MCH_METCA	cytoplasm (by similarity).	cytoplasm
Q9RPX1	UBID_HAEDU		cytoplasm
Q9RQ01	MCH_XANAU	cytoplasm (by similarity).	cytoplasm
Q9RQ04	MCH_METFL	cytoplasm (by similarity).	cytoplasm
Q9RQ16	ALGH_PSEAE		inner membrane
Q9RQ33	TRPA_BUCMH		cytoplasm
Q9RQ35	TRPD_BUCMH		cytoplasm
Q9RQ47	ILVC_BUCMH		cytoplasm
Q9RQ48	ILVD_BUCMH		cytoplasm
Q9RQ51	ILVC_BUCSC		cytoplasm
Q9RQ52	ILVD_BUCSC		cytoplasm
Q9RQ55	ILVC_BUCDN		cytoplasm
Q9RQ56	ILVD_BUCDN		cytoplasm
Q9RQ65	ILVI_BUCSC		cytoplasm
Q9RQ74	ATPG_BUCMH		inner membrane
Q9RQ77	ATPG_BUCSC		inner membrane
Q9RQ80	ATPG_BUCDN		inner membrane
Q9RQ83	HIS1_BUCMH	cytoplasm (by similarity).	cytoplasm
Q9RQ86	HIS1_BUCSC	cytoplasm (by similarity).	cytoplasm
Q9RQ89	HIS1_BUCDN	cytoplasm (by similarity).	cytoplasm
Q9RQB9	CYCA_METME		periplasm
Q9RQI5	GLGB_NEIDE		extracellular
Q9RQJ6	MRAW_CAUCR		cytoplasm
Q9RQJ9	KTHY_CAUCR		cytoplasm
Q9RQV9	PDXJ_NEIMA	cytoplasm (probable).	cytoplasm
Q9RQW2	ACPS_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9S0K8	PTSO_SHEVI	cytoplasm (probable).	cytoplasm
Q9S0K9	YPTO_SHEVI		cytoplasm
Q9S0L2	RP54_SHEVI		cytoplasm
Q9S0Q8	RPOA_SHEVI		cytoplasm
Q9S0Q9	RS4_SHEVI		cytoplasm
Q9S0R0	RS11_SHEVI		cytoplasm
Q9S0R1	RS13_SHEVI		cytoplasm
Q9S0X3	PHI_METAM		cytoplasm
Q9S0X4	TAL_METAM	cytoplasm (probable).	cytoplasm
Q9S169	BL24_ECOLI		extracellular
Q9S170	CSPG_SHEVI	cytoplasm (by similarity).	cytoplasm
Q9S1B7	CSPA_SHEVI	cytoplasm (by similarity).	cytoplasm

Q9S1E4	NRFI_WOLSU	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9S1E6	NRFH_WOLSU	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (potential).	inner membrane
Q9S1F9	DHE4_PSYT1		cytoplasm
Q9S1G2	DPO1_RHILE		cytoplasm
Q9S1G8	SERD_THASE	cytoplasm (probable).	cytoplasm
Q9S340	SKP_PHOLU	periplasm (by similarity).	periplasm
Q9S341	YAET_PHOLU	bacterial cell outer membrane (by similarity).	outer membrane
Q9S358	TRPE_NEIMC		cytoplasm
Q9S3J9	SOTB_ERWCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9S3K0	SOTA_ERWCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9S3P9	MOTY_VIBAN	bacterial cell membrane; peripheral membrane protein (potential).	inner membrane
Q9S3Q5	MQO_PSEFL		cytoplasm
Q9S3S0	DINI_SERMA		inner membrane
Q9S3U4	TRPF_ZYMMP		cytoplasm
Q9S3U8	VIOD_CHRVO		cytoplasm
Q9S3U9	VIOC_CHRVO		cytoplasm
Q9S3V0	VIOB_CHRVO		cytoplasm
Q9S3V1	VIOA_CHRVO		cytoplasm
Q9S3W0	SLYA_YEREN		cytoplasm
Q9S3Z2	LUXD_VIBFI		cytoplasm
Q9S424	BL13_KLEPN		extracellular
Q9S493	DNAA_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9S498	THIH_SALTY		cytoplasm
Q9S4T0	HGD_LEGPH		cytoplasm
Q9S4T3	SURE_LEGPH	cytoplasm (potential).	periplasm
Q9S4U0	RRAA_PSEFL		cytoplasm
Q9S4Y6	ALLA_SALEN		extracellular
Q9S4Y8	SELU_SALEN		cytoplasm
Q9S586	GPH1_PSEAE		cytoplasm
Q9S5B5	ATPE_DESVM	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9S5G4	HIS4_ECO57	cytoplasm (by similarity).	cytoplasm
Q9S5G5	HIS7_ECO57	cytoplasm (by similarity).	cytoplasm
Q9S5G6	HIS8_ECO57		cytoplasm
Q9S642	RFBB_NEIMA		cytoplasm
Q9S646	PPK_PSEAE		cytoplasm
Q9S6P8	MUTS_ECO57		cytoplasm
Q9WW00	TRPE_NEIGO		cytoplasm
Q9WW25	RS21A_BURMA		cytoplasm
Q9WWG9	DNAK_PSESG		cytoplasm
Q9WWL3	CH10_METSS	cytoplasm (by similarity).	cytoplasm
Q9WWL4	CH60_METSS	cytoplasm (by similarity).	cytoplasm

Q9WWR1	CYOA_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9WWR2	CYOB_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9WWR3	CYOC_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9WWR4	CYOD_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9WWR5	CYOE_PSEPU	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9WWW0	MOBA_PSEPU	cytoplasm (by similarity).	cytoplasm
Q9WWW2	ALKJ_PSEPU	bacterial cell inner membrane (by similarity).	inner membrane
Q9WWX6	HLDD_BURPS		cytoplasm
Q9WWZ8	TTGC_PSEPU	bacterial cell outer membrane; lipid-anchor (potential).	inner membrane
Q9WWZ9	TTGA_PSEPU	bacterial cell inner membrane; lipid-anchor (potential).	inner membrane
Q9WX61	BCSA3_ACEXY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9WX62	BCSB3_ACEXY	bacterial cell inner membrane; single-pass type i membrane protein (by similarity).	inner membrane
Q9WX63	BCSC3_ACEXY	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
Q9WX64	BCSD3_ACEXY		inner membrane
Q9WX69	BCSX_ACEXY		cytoplasm
Q9WX70	BCSY_ACEXY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9WX71	BCSC4_ACEXY	bacterial cell outer membrane; peripheral membrane protein (potential).	outer membrane
Q9WX75	BCSA5_ACEXY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9WXA9	BCHI_ACIRU		cytoplasm
Q9WXB4	BCHN_ACIRU		cytoplasm
Q9WXB5	BCHB_ACIRU		cytoplasm
Q9WXB9	CATA_LEGPN		periplasm
Q9WXD8	PUFQ_RHOSU		cytoplasm
Q9WXE6	LINE_PSEPA		cytoplasm
Q9WXI5	PTHP_BUCAI	cytoplasm (by similarity).	cytoplasm
Q9WXI7	PTGA_BUCAI	cytoplasm (by similarity).	cytoplasm
Q9WXL5	Y451_HELPY		nucleoid
Q9X2N2	6PGL_PSEAE		extracellular
Q9X2N9	QACF_ENTAE	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q9X2S4	LIFO_ACILW	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
Q9X2T1	THIO_PSEAE		cytoplasm
Q9X2V8	MCJB_ECOLI	cytoplasm (potential).	cytoplasm
Q9X2V9	MCJC_ECOLI	cytoplasm (potential).	cytoplasm
Q9X2W0	MCJD_ECOLI	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9X3W1	HIS3_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9X3W2	Y1179_ZYMMO		cytoplasm
Q9X3W5	ISPE_ZYMMO		cytoplasm
Q9X3X1	FPG_ZYMMO		cytoplasm
Q9X3X2	UBIE_ZYMMO		cytoplasm
Q9X3X5	DUT_ZYMMO		extracellular
Q9X3X7	TOP1_ZYMMO		cytoplasm
Q9X447	YACK_RHIME		inner membrane
Q9X448	PTA_RHIME	cytoplasm (potential).	inner membrane
Q9X449	ACKA2_RHIME	cytoplasm (by similarity).	cytoplasm
Q9X4A0	Y684_HAEDU		inner membrane
Q9X4E0	KCY_RHOS4	cytoplasm (by similarity).	cytoplasm
Q9X4E2	IHFB_RHOS4		cytoplasm
Q9X4E3	TRPF_RHOS4		cytoplasm
Q9X4E5	TRPB_RHOS4		cytoplasm
Q9X4E8	TRPA_RHOS4		cytoplasm
Q9X4F5	HGD_RHIME		cytoplasm
Q9X4F7	MAAI_RHIME		cytoplasm
Q9X4G0	HGD_PSEAE		cytoplasm
Q9X4G9	PILC_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9X4H2	AROA_EDWIC	cytoplasm (probable).	cytoplasm
Q9X4I6	SPED_XANCP		cytoplasm
Q9X4M8	PPK_ACIBA		cytoplasm
Q9X4P2	NUDH_PSEAE		cytoplasm
Q9X4Q5	NQRC_VIBCH	bacterial cell inner membrane (potential).	inner membrane
Q9X4Q6	NQRD_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9X4Q7	NQRE_VIBCH	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9X4Q8	NQRF_VIBCH	bacterial cell inner membrane (potential).	inner membrane
Q9X4Q9	FLIG_VIBCH	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9X4W3	RTXC_VIBCH	cytoplasm (potential).	cytoplasm
Q9X4Y0	AGAL_RHIME		cytoplasm
Q9X528	RNPH_LEGPN		cytoplasm
Q9X576	CATE_RHIME		periplasm
Q9X5D8	RECA_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9X5E1	PHOU_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9X5E3	PSD_ZYMMO		cytoplasm
Q9X5E7	RS2_ZYMMO		cytoplasm

Q9X5E9	PYRH_ZYMMO		cytoplasm
Q9X5F0	RRF_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9X5F1	UPPS_ZYMMO		inner membrane
Q9X5F2	DXR_ZYMMO		cytoplasm
Q9X5F5	FABZ_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9X5F8	ILVC_ZYMMO		cytoplasm
Q9X5G1	MIAA_ZYMMO		cytoplasm
Q9X5H9	FTSQ_BARBA	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q9X5N0	CLPP2_MYXXA	cytoplasm (by similarity).	cytoplasm
Q9X5N1	CLPX_MYXXA		cytoplasm
Q9X5P5	RECA_XENNE	cytoplasm (by similarity).	cytoplasm
Q9X5U8	RS2_COXBU		cytoplasm
Q9X5V3	ATCU_RHILV	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9X5V4	HMMR_RHILV	cytoplasm (probable).	cytoplasm
Q9X5W3	MOAA_RHOCA		cytoplasm
Q9X5X3	ATCU_RHIME	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9X5X4	HMRR_RHIME	cytoplasm (probable).	cytoplasm
Q9X5Y0	FLGE_RHIME		flagellar
Q9X600	FLHD_ERWCA	cytoplasm (by similarity).	cytoplasm
Q9X601	FLHC_ERWCA		cytoplasm
Q9X6B0	CATA_YERPE		periplasm
Q9X6B1	C5621_YERPE	periplasm (by similarity).	periplasm
Q9X6B2	C561_YERPE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9X6E8	RL20_PSEFL		cytoplasm
Q9X6G2	YHIQ_SALTY		inner membrane
Q9X6M5	KGUA_SALTY	cytoplasm (by similarity).	cytoplasm
Q9X6N4	MURE_VIBCH	cytoplasm (probable).	cytoplasm
Q9X6P4	LPXA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9X6R0	SPEE1_PSEAE		cytoplasm
Q9X6U2	BDHA_RALEU		cytoplasm
Q9X6V6	RLPA_PSEAE	bacterial cell membrane; lipid-anchor (potential).	inner membrane
Q9X6V8	Y3998_PSEAE		inner membrane
Q9X6V9	LIPB_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9X6W6	PYRR_PSEAE		cytoplasm
Q9X6W7	TIG_AZOBR		cytoplasm
Q9X6W8	CLPP_AZOBR	cytoplasm (by similarity).	cytoplasm
Q9X6X4	LIPB_MYXXA	cytoplasm (by similarity).	cytoplasm
Q9X6Y8	MURB_BORPE	cytoplasm (probable).	inner membrane
Q9X6Y9	DAPB_BORPE	cytoplasm (by similarity).	cytoplasm
Q9X725	OXYR_ERWCH		cytoplasm
Q9X755	KEFF_KLEAE		cytoplasm
Q9X756	KEFC_KLEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9X772	DJLA_LEGPH	bacterial cell inner membrane; single-pass type iii membrane protein (by similarity).	inner membrane

Q9X7F4	COBT_METCH		cytoplasm
Q9X7G7	CBID_METCH		cytoplasm
Q9X7K0	MOBA_RHOCA	cytoplasm (by similarity).	cytoplasm
Q9X7K6	DCTA_RHIGA	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9X7L1	CLPP3_RHIME	cytoplasm (by similarity).	cytoplasm
Q9X7L2	LEU1_RHIME		cytoplasm
Q9X7L3	Y2378_RHIME		cytoplasm
Q9X7L4	Y2377_RHIME		cytoplasm
Q9X980	COAD_SERMA	cytoplasm (by similarity).	cytoplasm
Q9X981	FPG_SERMA		cytoplasm
Q9X9E0	EXUR_ERWCH		cytoplasm
Q9X9F2	FLHD_XENNE	cytoplasm (by similarity).	cytoplasm
Q9X9F3	FLHC_XENNE		cytoplasm
Q9X9F6	IHFA_YERPS		cytoplasm
Q9X9J3	FLGJ_VIBPA	periplasm (by similarity).	extracellular
Q9X9J4	FLGI2_VIBPA	periplasm (by similarity).	periplasm
Q9X9J5	FLGH1_VIBPA	bacterial cell outer membrane; lipid-anchor (probable).	inner membrane
Q9X9K2	CHER_VIBPA		cytoplasm
Q9XAT2	PUR5_RHILE	cytoplasm (by similarity).	cytoplasm
Q9XAU7	CH60_PSEHT	cytoplasm (by similarity).	cytoplasm
Q9XAY7	GLYA_NEIMA	cytoplasm (by similarity).	cytoplasm
Q9XAZ0	TRPE_NEIMA		cytoplasm
Q9XAZ1	GLYA_NEIMC	cytoplasm (by similarity).	cytoplasm
Q9XB01	GLYA_NEIGO	cytoplasm (by similarity).	cytoplasm
Q9XB38	FLAA_HELFE		flagellar
Q9XB42	YKFH_ECOLI		cytoplasm
Q9XB52	KDGT_PECCC	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9XB53	KDUI_PECCC		cytoplasm
Q9XB58	CARD_PECCC		cytoplasm
Q9XBC1	RECA_RHOPA	cytoplasm (by similarity).	cytoplasm
Q9XBF7	GIDB_CAUCR		cytoplasm
Q9XBF8	GIDA_CAUCR	cytoplasm (by similarity).	cytoplasm
Q9XBF9	TRME_CAUCR		cytoplasm
Q9XBG0	PHOU_BURSP	cytoplasm (by similarity).	cytoplasm
Q9XBG1	PSTB_BURSP	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9XBG6	TILS_BRAJA	cytoplasm (probable).	cytoplasm
Q9XBM3	TRPC_ZYMMO		cytoplasm
Q9XBM6	MTLD_KLEPN		cytoplasm
Q9XBM7	PTM3C_KLEPN	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9XBQ0	PKN3_MYXXA		cytoplasm
Q9XBR5	WRBA_ZYMMO		cytoplasm
Q9XBR7	Y1337_ZYMMO		cytoplasm
Q9XBS9	Y1353_ZYMMO		inner membrane
Q9XC07	GLND_VIBF1		cytoplasm
Q9XC89	COAD_KLEPN	cytoplasm (by similarity).	cytoplasm
Q9XC90	WAAE_KLEPN		cytoplasm
Q9XCA1	HLDD_KLEPN		cytoplasm

Q9XCI8	ENGA_SALTY		cytoplasm
Q9XCX7	RECO_PSEAE		cytoplasm
Q9XCX8	ERA_PSEAE		cytoplasm
Q9XCX9	RNC_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9XD50	ISPZ_VITS1	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9XD74	SODF_RHIME		periplasm
Q9XD77	RBL_NITWI		extracellular
Q9XD78	QUIA_XANCJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9XDB2	DSBD_PANCI	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9XDH6	DPO3A_PSEFL	cytoplasm (by similarity).	cytoplasm
Q9XDM6	PDUV_SALTY		cytoplasm
Q9XDP0	DSBI_ENTAM	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9Z377	EFE_PSEYS		cytoplasm
Q9Z3E6	RL17_XANCP		cytoplasm
Q9Z3F0	RS13_XANCP		cytoplasm
Q9Z3I3	NODI_BRASS	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9Z3Q0	CYA3_RHIME		cytoplasm
Q9Z3Q1	SYRB2_RHIME		cytoplasm
Q9Z3Q3	PHAG_RHIME	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9Z3Q6	RHRA_RHIME		cytoplasm
Q9Z3Q7	RHBF_RHIME		outer membrane
Q9Z3Q8	RHBE_RHIME		cytoplasm
Q9Z3Q9	RHBD_RHIME		cytoplasm
Q9Z3R0	RHBC_RHIME		outer membrane
Q9Z3R1	RHBB_RHIME		cytoplasm
Q9Z3R2	RHBA_RHIME		cytoplasm
Q9Z3R3	ACSA1_RHIME		cytoplasm
Q9Z3R4	AGLR_RHIME		cytoplasm
Q9Z3R5	AGLE_RHIME	periplasm (probable).	periplasm
Q9Z3R6	AGLF_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9Z3R7	AGLG_RHIME	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9Z3R8	AGLA_RHIME		extracellular
Q9Z3R9	AGLK_RHIME	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9Z3S0	EDD_RHIME		cytoplasm
Q9Z3S1	6PGL_RHIME		extracellular
Q9Z3S2	G6PD_RHIME		cytoplasm
Q9Z3S3	ORDL_RHIME		cytoplasm
Q9Z3S4	RS21A_RHIME		cytoplasm

Q9Z3S6	CSPA_RHIME	cytoplasm (by similarity).	cytoplasm
Q9Z3T0	EFE_PSESJ		cytoplasm
Q9Z3U0	LOLA_PSEU2	periplasm (by similarity).	periplasm
Q9Z3Z6	MURA_PSEPU	cytoplasm (probable).	inner membrane
Q9Z408	CSD_PSEPK		cytoplasm
Q9Z410	PHOU_PSEPU	cytoplasm (by similarity).	cytoplasm
Q9Z411	PSTB_PSEPU	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9Z426	PTHP_PSEPU	cytoplasm (by similarity).	cytoplasm
Q9Z427	YPTO_PSEPU		cytoplasm
Q9Z431	4OT3_PSEPU		cytoplasm
Q9Z439	UBIE_PSEPU		cytoplasm
Q9Z441	PYRR_PSEPU		cytoplasm
Q9Z448	MDCG_PSEPU		cytoplasm
Q9Z451	MDCC_PSEPU	cytoplasm (by similarity).	cytoplasm
Q9Z452	MDCB_PSEPU		cytoplasm
Q9Z462	CH60_PARDE	cytoplasm (by similarity).	cytoplasm
Q9Z463	CH10_PARDE	cytoplasm (by similarity).	cytoplasm
Q9Z4N4	MCEA_KLEPN		extracellular
Q9Z5D3	IDI_RHOS4	cytoplasm (by similarity).	cytoplasm
Q9Z5D6	BCHG_RHOS4	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9Z5D7	BCHJ_RHOS4		cytoplasm
Q9Z5D9	BCHB_RHOS4		cytoplasm
Q9Z5T8	FLGH_ZYMMO	bacterial cell outer membrane (probable).	inner membrane
Q9Z5T9	FLGI_ZYMMO	periplasm (by similarity).	periplasm
Q9Z5U2	RNC_ZYMMO	cytoplasm (by similarity).	cytoplasm
Q9Z5U7	RS4_ZYMMO		cytoplasm
Q9Z5V0	RS20_ZYMMO		cytoplasm
Q9Z5X1	LUXS_VIBHA		extracellular
Q9Z5Z2	NFSA_SALTY		cytoplasm
Q9Z613	COAD_BUCAP	cytoplasm (by similarity).	cytoplasm
Q9Z614	SYK3_BUCAP		cytoplasm
Q9Z615	FENR_BUCAP		cytoplasm
Q9Z616	IBP_BUCAP		cytoplasm
Q9Z617	HSLU_BUCAP	cytoplasm (by similarity).	cytoplasm
Q9Z645	DSBE_PANCI	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	periplasm
Q9Z647	CCME_PANCI	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q9Z651	CCMA_PANCI	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9Z660	ARLY_ZYMMO	cytoplasm (probable).	cytoplasm
Q9Z661	DCDA_ZYMMO		cytoplasm
Q9Z670	Gntp_ZYMMO	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9Z671	TYSY_ZYMMO	cytoplasm (by similarity).	cytoplasm

Q9Z6B9	TPIS_ENTCL		cytoplasm
Q9Z6C1	CDH_ENTCL	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9Z6C3	K6PF_ENTCL	cytoplasm (by similarity).	cytoplasm
Q9Z6D1	RS7_HAEDU		cytoplasm
Q9Z6D2	RS12_HAEDU		cytoplasm
Q9Z9H0	GLND_PSEAE		cytoplasm
Q9ZA11	ALDH_RHORT		cytoplasm
Q9ZA86	PTSO_PROMI	cytoplasm (probable).	cytoplasm
Q9ZA87	YPTO_PROMI		cytoplasm
Q9ZAB8	NPD_ACTAC	cytoplasm (probable).	cytoplasm
Q9ZAH3	RS6_CAMJE		cytoplasm
Q9ZAQ8	SSB_RHOS4		cytoplasm
Q9ZB35	RBL_HYDPS		extracellular
Q9ZBB6	LUXU_VIBHA		inner membrane
Q9ZC45	NQRE_YERPE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZC49	GLSA2_YERPE		extracellular
Q9ZC70	ASTE_YERPE		periplasm
Q9ZC74	HUTI_YERPE	cytoplasm (potential).	cytoplasm
Q9ZC83	DCUP_RICPR	cytoplasm (probable).	cytoplasm
Q9ZC84	HEMH_RICPR	cytoplasm (by similarity).	extracellular
Q9ZC85	Y883_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZC86	HEM6_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZC88	MUTL_RICPR		cytoplasm
Q9ZC89	RL33_RICPR		cytoplasm
Q9ZC90	RS16_RICPR		cytoplasm
Q9ZC91	LIPB_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZC92	Y875_RICPR		inner membrane
Q9ZC94	Y872_RICPR		cytoplasm
Q9ZC95	Y871_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZC99	Y867_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZCA0	Y866_RICPR		inner membrane
Q9ZCA2	Y864_RICPR		periplasm
Q9ZCA3	TRUA_RICPR		cytoplasm
Q9ZCA5	Y855_RICPR		inner membrane
Q9ZCA6	Y854_RICPR		inner membrane
Q9ZCA7	Y853_RICPR		inner membrane
Q9ZCB3	HEMK_RICPR		cytoplasm
Q9ZCB7	TIG_RICPR		cytoplasm
Q9ZCB8	HEM1_RICPR		inner membrane
Q9ZCB9	HTPG_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCC0	Y839_RICPR		inner membrane
Q9ZCC2	SSB_RICPR		cytoplasm
Q9ZCC3	UVRA_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCC5	P34_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q9ZCC6	Y831_RICPR		cytoplasm
Q9ZCC8	FER_RICPR		periplasm
Q9ZCD1	Y826_RICPR		cytoplasm
Q9ZCD3	AMPM_RICPR		cytoplasm
Q9ZCD6	Y821_RICPR		cytoplasm
Q9ZCE0	XERC_RICPR	cytoplasm (by similarity).	nucleoid
Q9ZCE2	Y815_RICPR	cytoplasm (potential).	inner membrane
Q9ZCE4	Y812_RICPR		cytoplasm
Q9ZCE8	Y808_RICPR		cytoplasm
Q9ZCE9	PBPA_RICPR	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q9ZCF2	ATPD_RICPR		inner membrane
Q9ZCF3	ATPE_RICPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZCF4	ACON_RICPR		cytoplasm
Q9ZCF6	NUOG_RICPR		cytoplasm
Q9ZCF8	NUOI_RICPR		cytoplasm
Q9ZCF9	CCMA_RICPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZCG0	NUOM_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZCG1	NUOL_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZCG2	NUOK_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZCG6	TATC_RICPR	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9ZCH0	RL32_RICPR		cytoplasm
Q9ZCH1	FABH_RICPR	cytoplasm (probable).	cytoplasm
Q9ZCH6	Y766_RICPR		cytoplasm
Q9ZCH7	KGUA_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCH9	ACP_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCI0	Y760_RICPR		inner membrane
Q9ZCI1	TRME_RICPR		cytoplasm
Q9ZCI4	Y756_RICPR		cytoplasm
Q9ZCI7	AK_RICPR		cytoplasm
Q9ZCI8	RL27_RICPR		cytoplasm
Q9ZCI9	RL21_RICPR		cytoplasm
Q9ZCJ1	TATA_RICPR	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9ZCJ2	Y748_RICPR		cytoplasm
Q9ZCJ3	Y747_RICPR		cytoplasm
Q9ZCJ8	SUR1_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZCJ9	DPO3E_RICPR		cytoplasm
Q9ZCK0	COAE_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCK3	RNH_RICPR	cytoplasm (potential).	extracellular

Q9ZCK4	Y725_RICPR		inner membrane
Q9ZCK6	Y723_RICPR		cytoplasm
Q9ZCK7	Y722_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZCK8	TGT_RICPR		cytoplasm
Q9ZCK9	DNLJ_RICPR		cytoplasm
Q9ZCL0	LPXK_RICPR		cytoplasm
Q9ZCL3	Y714_RICPR		cytoplasm
Q9ZCL4	Y713_RICPR		inner membrane
Q9ZCL5	Y712_RICPR		cytoplasm
Q9ZCL6	Y709_RICPR		cytoplasm
Q9ZCM4	LOLD_RICPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZCM7	Y697_RICPR		inner membrane
Q9ZCN1	CLPX_RICPR		cytoplasm
Q9ZCN2	Y691_RICPR		cytoplasm
Q9ZCN3	RMUC_RICPR		cytoplasm
Q9ZCN4	Y689_RICPR		inner membrane
Q9ZCN5	Y688_RICPR		inner membrane
Q9ZCN6	SYV_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCN9	KTHY_RICPR		cytoplasm
Q9ZCP3	UBIE_RICPR		cytoplasm
Q9ZCP6	ENGA_RICPR		cytoplasm
Q9ZCP8	EX7L_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCP9	CSPA_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCQ4	FUMC_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCQ6	RL4_RICPR		cytoplasm
Q9ZCQ7	RL23_RICPR		cytoplasm
Q9ZCQ8	RL2_RICPR		cytoplasm
Q9ZCQ9	RS19_RICPR		cytoplasm
Q9ZCR0	RL22_RICPR		cytoplasm
Q9ZCR1	RS3_RICPR		cytoplasm
Q9ZCR2	RL16_RICPR		cytoplasm
Q9ZCR3	RL29_RICPR		cytoplasm
Q9ZCR4	RS17_RICPR		cytoplasm
Q9ZCR5	RL14_RICPR		cytoplasm
Q9ZCR6	RL24_RICPR		cytoplasm
Q9ZCR7	RL5_RICPR		cytoplasm
Q9ZCR8	RS14_RICPR		cytoplasm
Q9ZCR9	RS8_RICPR		cytoplasm
Q9ZCS0	RL6_RICPR		cytoplasm
Q9ZCS1	RL18_RICPR		cytoplasm
Q9ZCS2	RS5_RICPR		cytoplasm
Q9ZCS3	RL30_RICPR		cytoplasm
Q9ZCS4	RL15_RICPR		cytoplasm
Q9ZCS7	RS13_RICPR		cytoplasm
Q9ZCS8	RS11_RICPR		cytoplasm
Q9ZCS9	RPOA_RICPR		cytoplasm
Q9ZCT0	RL17_RICPR		cytoplasm
Q9ZCT1	RS20_RICPR		cytoplasm
Q9ZCT3	Y630_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZCT4	GRPE_RICPR	cytoplasm (probable).	cytoplasm

Q9ZCT5	RNPH_RICPR		cytoplasm
Q9ZCT6	CH10_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCT7	CH60_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCT9	UBIG_RICPR		cytoplasm
Q9ZCU4	SYI_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCU6	RS21_RICPR		cytoplasm
Q9ZCU8	RNPA_RICPR		extracellular
Q9ZCV0	RL20_RICPR		cytoplasm
Q9ZCV1	RL35_RICPR		cytoplasm
Q9ZCV3	RL25_RICPR		cytoplasm
Q9ZCV4	PTH_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCV8	Y600_RICPR		cytoplasm
Q9ZCW0	MRAY_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZCW2	Y592_RICPR		cytoplasm
Q9ZCW3	Y591_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZCW4	MVIN_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZCW5	IPYR_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCW6	CCME_RICPR	bacterial cell inner membrane; single-pass type ii membrane protein; periplasmic side (potential).	inner membrane
Q9ZCW7	SCO22_RICPR		cytoplasm
Q9ZCW8	SECD_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZCW9	Y585_RICPR		inner membrane
Q9ZCX2	GYRB_RICPR		cytoplasm
Q9ZCX3	MURA_RICPR	cytoplasm (probable).	inner membrane
Q9ZCX4	RPOZ_RICPR		cytoplasm
Q9ZCX5	ACPS_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCX7	SECA_RICPR	bacterial cell membrane; peripheral membrane protein; cytoplasmic side (by similarity).	inner membrane
Q9ZCX8	Y573_RICPR		extracellular
Q9ZCX9	UVRC_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCY1	MRAZ_RICPR		inner membrane
Q9ZCY2	MRAW_RICPR		cytoplasm
Q9ZCY3	Y568_RICPR		inner membrane
Q9ZCY5	Y566_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZCZ4	SYI_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZCZ6	Y554_RICPR		inner membrane
Q9ZCZ7	NUSA_RICPR		cytoplasm
Q9ZCZ9	Y551_RICPR		cytoplasm
Q9ZD00	Y550_RICPR		inner membrane
Q9ZD01	Y549_RICPR		cytoplasm
Q9ZD02	Y548_RICPR		cytoplasm
Q9ZD03	Y547_RICPR		cytoplasm

Q9ZD04	RADA_RICPR		cytoplasm
Q9ZD05	Y545_RICPR		cytoplasm
Q9ZD06	Y544_RICPR		cytoplasm
Q9ZD07	Y543_RICPR		cytoplasm
Q9ZD08	DNAB_RICPR		nucleoid
Q9ZD09	PAAD_RICPR		cytoplasm
Q9ZD10	PRIA_RICPR		cytoplasm
Q9ZD12	Y538_RICPR		cytoplasm
Q9ZD15	SODF_RICPR		periplasm
Q9ZD18	Y532_RICPR		inner membrane
Q9ZD20	ODP2_RICPR		inner membrane
Q9ZD24	RHO_RICPR		inner membrane
Q9ZD26	DBHL_RICPR		cytoplasm
Q9ZD27	KCY_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZD28	RS1_RICPR		cytoplasm
Q9ZD29	CLPP_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZD32	FOLD_RICPR		cytoplasm
Q9ZD36	Y511_RICPR		cytoplasm
Q9ZD37	MIAA_RICPR		cytoplasm
Q9ZD40	Y507_RICPR		cytoplasm
Q9ZD41	Y506_RICPR		inner membrane
Q9ZD42	Y505_RICPR		cytoplasm
Q9ZD43	PNP_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZD44	RS15_RICPR		cytoplasm
Q9ZD45	Y502_RICPR		cytoplasm
Q9ZD46	TRUB_RICPR		cytoplasm
Q9ZD49	SCA4_RICPR	cytoplasm (probable).	cytoplasm
Q9ZD50	Y497_RICPR		cytoplasm
Q9ZD53	Y494_RICPR		cytoplasm
Q9ZD54	Y493_RICPR		cytoplasm
Q9ZD55	PPDK_RICPR		cytoplasm
Q9ZD57	Y489_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZD60	ISCS_RICPR		cytoplasm
Q9ZD61	NIFU_RICPR		cytoplasm
Q9ZD62	Y484_RICPR		inner membrane
Q9ZD66	Y478_RICPR		cytoplasm
Q9ZD70	Y474_RICPR		cytoplasm
Q9ZD71	Y473_RICPR		inner membrane
Q9ZD72	Y472_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZD73	Y471_RICPR		cytoplasm
Q9ZD74	Y470_RICPR		cytoplasm
Q9ZD77	HEM3_RICPR		cytoplasm
Q9ZD84	Y459_RICPR		cytoplasm
Q9ZD85	Y458_RICPR		cytoplasm
Q9ZD88	Y455_RICPR		inner membrane
Q9ZD90	IDI2_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZD92	LON_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZD94	Y448_RICPR		cytoplasm
Q9ZD95	UVRD_RICPR		cytoplasm
Q9ZD97	TRXB_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDA0	GPDA_RICPR	cytoplasm (probable).	cytoplasm
Q9ZDA2	PPNK_RICPR	cytoplasm (by similarity).	cytoplasm

Q9ZDA3	Y439_RICPR		cytoplasm
Q9ZDA4	RECR_RICPR		cytoplasm
Q9ZDA5	Y437_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDA7	UPPS_RICPR		inner membrane
Q9ZDA8	CDSA_RICPR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZDA9	Y423_RICPR		inner membrane
Q9ZDB2	Y420_RICPR		inner membrane
Q9ZDB3	DPO3B_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDB6	Y416_RICPR		cytoplasm
Q9ZDB7	DAPF_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDB9	Y413_RICPR		cytoplasm
Q9ZDC0	MURG_RICPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZDC2	MURD_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDC3	Y409_RICPR		cytoplasm
Q9ZDC4	LSPA_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9ZDC6	COX2_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDC7	Y404_RICPR		cytoplasm
Q9ZDC9	Y402_RICPR		extracellular
Q9ZDD0	Y401_RICPR		cytoplasm
Q9ZDD2	DUT_RICPR		extracellular
Q9ZDD5	Y396_RICPR		cytoplasm
Q9ZDD6	Y395_RICPR		cytoplasm
Q9ZDD7	Y394_RICPR		inner membrane
Q9ZDD9	ISPZ_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZDE1	RLPA_RICPR		cytoplasm
Q9ZDE5	RUVB_RICPR		cytoplasm
Q9ZDE6	RUVA_RICPR		cytoplasm
Q9ZDE8	GCH1_RICPR		cytoplasm
Q9ZDE9	Y382_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDF0	KDSB_RICPR	cytoplasm (potential).	cytoplasm
Q9ZDF1	PYRG_RICPR		cytoplasm
Q9ZDF3	MDH_RICPR		cytoplasm
Q9ZDF6	MAO2_RICPR		cytoplasm
Q9ZDF9	Y370_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDG1	Y368_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDG2	Y367_RICPR		cytoplasm

Q9ZDG3	LNT_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZDG4	FABI_RICPR	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZDG5	Y364_RICPR		cytoplasm
Q9ZDG6	Y363_RICPR		cytoplasm
Q9ZDG7	Y362_RICPR		cytoplasm
Q9ZDG8	XERD_RICPR	cytoplasm (by similarity).	nucleoid
Q9ZDG9	Y359_RICPR		cytoplasm
Q9ZDH2	NUOB_RICPR		cytoplasm
Q9ZDH3	NUOC_RICPR		inner membrane
Q9ZDH4	NUOD_RICPR		cytoplasm
Q9ZDH5	NUOE_RICPR		cytoplasm
Q9ZDH7	3MGH_RICPR		cytoplasm
Q9ZDH8	EX7S_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDI0	RIMM_RICPR	cytoplasm (potential).	cytoplasm
Q9ZDI3	RS4_RICPR		cytoplasm
Q9ZDJ7	Y331_RICPR		cytoplasm
Q9ZDJ8	RUVX_RICPR	cytoplasm (potential).	cytoplasm
Q9ZDK2	TOP1_RICPR		cytoplasm
Q9ZDK4	Y324_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDK5	CYAY_RICPR		inner membrane
Q9ZDK7	LPXB_RICPR		cytoplasm
Q9ZDK8	HSLU_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDK9	HSLV_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDL0	Y318_RICPR		inner membrane
Q9ZDL1	YHIT_RICPR		cytoplasm
Q9ZDL2	DHAS_RICPR		cytoplasm
Q9ZDL6	Y311_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDM1	TRMU_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDM3	COXZ_RICPR		inner membrane
Q9ZDM5	TOLB_RICPR	periplasm (potential).	periplasm
Q9ZDM6	THYX_RICPR		cytoplasm
Q9ZDM9	MUTS_RICPR		cytoplasm
Q9ZDN1	Y296_RICPR		cytoplasm
Q9ZDN2	Y295_RICPR		cytoplasm
Q9ZDN8	Y289_RICPR		inner membrane
Q9ZDN9	Y288_RICPR		inner membrane
Q9ZDP8	Y279_RICPR		cytoplasm
Q9ZDQ0	Y277_RICPR		cytoplasm
Q9ZDQ1	LEPA_RICPR		cytoplasm
Q9ZDQ3	HSP19_RICPR		cytoplasm
Q9ZDQ5	UCRI_RICPR	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane
Q9ZDQ6	Y269_RICPR		cytoplasm
Q9ZDQ7	Y268_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDQ9	Y266_RICPR		inner membrane

Q9ZDR0	IDH_RICPR		cytoplasm
Q9ZDR3	ODPB_RICPR		cytoplasm
Q9ZDR4	ODPA_RICPR		cytoplasm
Q9ZDR7	RLUC_RICPR		cytoplasm
Q9ZDS0	Y255_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDS1	LPXC_RICPR		cytoplasm
Q9ZDS3	Y252_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDS5	FTSQ_RICPR	bacterial cell inner membrane; single-pass type ii membrane protein (by similarity).	inner membrane
Q9ZDS6	DDL_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDS7	MURB_RICPR	cytoplasm (probable).	inner membrane
Q9ZDS8	MURC_RICPR	cytoplasm (probable).	cytoplasm
Q9ZDT1	Y244_RICPR		cytoplasm
Q9ZDT4	PSD_RICPR		cytoplasm
Q9ZDT5	Y240_RICPR		cytoplasm
Q9ZDT9	NUDH_RICPR		cytoplasm
Q9ZDU0	RS9_RICPR		cytoplasm
Q9ZDU1	RL13_RICPR		cytoplasm
Q9ZDU2	Y232_RICPR		cytoplasm
Q9ZDU3	Y231_RICPR		inner membrane
Q9ZDU9	Y225_RICPR		cytoplasm
Q9ZDV2	Y222_RICPR		cytoplasm
Q9ZDV4	Y215_RICPR		cytoplasm
Q9ZDV6	QUEA_RICPR	cytoplasm (potential).	cytoplasm
Q9ZDV8	DEF_RICPR		cytoplasm
Q9ZDV9	Y207_RICPR		inner membrane
Q9ZDW1	GLRX_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDW2	UVRB_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDW3	RNH2_RICPR	cytoplasm (potential).	extracellular
Q9ZDW4	HSCB_RICPR		inner membrane
Q9ZDW5	HSCA_RICPR		cytoplasm
Q9ZDW6	FER2_RICPR		cytoplasm
Q9ZDW7	Y198_RICPR		cytoplasm
Q9ZDX0	DAPD_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZDX1	Y193_RICPR		inner membrane
Q9ZDX2	Y192_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDX5	Y189_RICPR		cytoplasm
Q9ZDX6	Y188_RICPR		cytoplasm
Q9ZDX7	Y187_RICPR		cytoplasm
Q9ZDX9	DNAK_RICPR		cytoplasm
Q9ZDY0	DNAJ_RICPR	cytoplasm (probable).	cytoplasm
Q9ZDY1	Y183_RICPR	bacterial cell membrane; lipid-anchor (potential).	outer membrane
Q9ZDY2	RECN_RICPR		cytoplasm
Q9ZDY3	ODO1_RICPR		cytoplasm
Q9ZDY4	ODO2_RICPR		inner membrane
Q9ZDY5	PTPS_RICPR		extracellular
Q9ZDZ0	SRP54_RICPR		cytoplasm
Q9ZDZ2	DBH_RICPR		cytoplasm

Q9ZDZ4	Y169_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZDZ5	Y168_RICPR		cytoplasm
Q9ZDZ6	Y167_RICPR		cytoplasm
Q9ZDZ7	Y166_RICPR		cytoplasm
Q9ZDZ8	Y165_RICPR		cytoplasm
Q9ZE00	RRMJ_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZE01	NUSB_RICPR		cytoplasm
Q9ZE02	Y161_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9ZE04	Y158_RICPR		cytoplasm
Q9ZE07	PYRH_RICPR		cytoplasm
Q9ZE08	RRF_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZE09	GATC_RICPR		cytoplasm
Q9ZE10	GATA_RICPR		cytoplasm
Q9ZE11	GATB_RICPR		cytoplasm
Q9ZE14	DAPB_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZE15	Y147_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZE16	Y146_RICPR		cytoplasm
Q9ZE18	Y144_RICPR		cytoplasm
Q9ZE20	RPOC_RICPR		cytoplasm
Q9ZE21	RL7_RICPR		cytoplasm
Q9ZE22	RL10_RICPR		cytoplasm
Q9ZE23	RL1_RICPR		cytoplasm
Q9ZE24	RL11_RICPR		cytoplasm
Q9ZE27	MRP_RICPR		cytoplasm
Q9ZE28	Y120_RICPR		cytoplasm
Q9ZE29	RUVC_RICPR		cytoplasm
Q9ZE30	ERA_RICPR		cytoplasm
Q9ZE31	RNC_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZE32	LEP_RICPR	bacterial cell membrane; single-pass type ii membrane protein (potential).	inner membrane
Q9ZE33	NUOF_RICPR		periplasm
Q9ZE34	SECF_RICPR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZE35	Y113_RICPR		inner membrane
Q9ZE36	RL19_RICPR		cytoplasm
Q9ZE37	TRMD_RICPR	cytoplasm (potential).	cytoplasm
Q9ZE38	ACKA_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZE39	PTA_RICPR	cytoplasm (potential).	inner membrane
Q9ZE46	ENGB_RICPR		cytoplasm
Q9ZE47	RL31_RICPR		cytoplasm
Q9ZE48	RL28_RICPR		cytoplasm
Q9ZE49	Y098_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZE52	ALR_RICPR		cytoplasm
Q9ZE55	Y092_RICPR		extracellular
Q9ZE56	AAT_RICPR	cytoplasm (by similarity).	cytoplasm

Q9ZE57	Y090_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZE58	KDTA_RICPR	bacterial cell inner membrane; single-pass membrane protein (potential).	inner membrane
Q9ZE59	Y088_RICPR		inner membrane
Q9ZE61	RS2_RICPR		cytoplasm
Q9ZE63	Y084_RICPR		extracellular
Q9ZE65	Y082_RICPR		cytoplasm
Q9ZE67	Y080_RICPR		cytoplasm
Q9ZE68	SECG_RICPR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZE74	QUEF_RICPR	cytoplasm (probable).	cytoplasm
Q9ZE76	SECB_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZE77	DCD_RICPR		extracellular
Q9ZE79	PARC_RICPR	bacterial cell membrane; peripheral membrane protein (by similarity).	cytoplasm
Q9ZE80	Y066_RICPR		inner membrane
Q9ZE82	DGTL1_RICPR		cytoplasm
Q9ZE83	Y063_RICPR		inner membrane
Q9ZE84	KDSA_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZE85	Y061_RICPR		cytoplasm
Q9ZE87	PARB_RICPR		cytoplasm
Q9ZE89	GIDB_RICPR		cytoplasm
Q9ZE90	GIDA_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZE91	NDK_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZE93	Y052_RICPR		cytoplasm
Q9ZE96	PGSA_RICPR	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZE97	OXAA_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZE99	LGT_RICPR	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZEA1	DHSB_RICPR		cytoplasm
Q9ZEA3	TILS_RICPR	cytoplasm (probable).	cytoplasm
Q9ZEA4	RL9_RICPR		cytoplasm
Q9ZEA5	RS18_RICPR		cytoplasm
Q9ZEA6	RS6_RICPR		cytoplasm
Q9ZEA8	GCP_RICPR		extracellular
Q9ZEA9	CLPB_RICPR	cytoplasm (probable).	cytoplasm
Q9ZEB1	Y034_RICPR		cytoplasm
Q9ZEB3	Y032_RICPR		inner membrane
Q9ZEB4	SCO21_RICPR		cytoplasm
Q9ZEB5	Y030_RICPR		cytoplasm
Q9ZEB6	RECF_RICPR	cytoplasm (potential).	cytoplasm
Q9ZEB7	Y027_RICPR		cytoplasm
Q9ZEC0	Y024_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane

Q9ZEC2	ATPL_RICPR	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZEC4	ATPF_RICPR		inner membrane
Q9ZEC6	Y017_RICPR		extracellular
Q9ZEC7	Y016_RICPR		extracellular
Q9ZEC9	Y014_RICPR		inner membrane
Q9ZED0	Y013_RICPR		cytoplasm
Q9ZED1	Y012_RICPR		cytoplasm
Q9ZED2	DUS_RICPR		cytoplasm
Q9ZED3	LPXD_RICPR		cytoplasm
Q9ZED4	FABZ_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZED5	LPXA_RICPR	cytoplasm (by similarity).	cytoplasm
Q9ZED6	Y006_RICPR		cytoplasm
Q9ZED7	Y005_RICPR		cytoplasm
Q9ZEE0	THIO_RICPR		cytoplasm
Q9ZEE1	Y001_RICPR		cytoplasm
Q9ZEL6	LEP4_PSEST	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9ZEL7	COAE_PSEST	cytoplasm (by similarity).	cytoplasm
Q9ZEM5	LIFO2_BURCE	bacterial cell inner membrane; single-pass membrane protein (by similarity).	periplasm
Q9ZEP8	PANB_PSEFL		cytoplasm
Q9ZEQ8	MSRA_DICD3		cytoplasm
Q9ZER9	TRPE_BUCPS		cytoplasm
Q9ZES2	TRPE_BUCTC		cytoplasm
Q9ZES3	APAG_RHIET		inner membrane
Q9ZEU6	ECTC_CHRSL		cytoplasm
Q9ZEU7	ECTB_CHRSL		cytoplasm
Q9ZEU8	ECTA_CHRSL		cytoplasm
Q9ZEY8	LEU1_BUCAI		cytoplasm
Q9ZEY9	REPA2_BUCAI		nucleoid
Q9ZEZ0	YREP_BUCAI	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZEZ4	YREP_BUCPP	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZEZ5	LEUD_BUCPP		cytoplasm
Q9ZF60	GLTI_SALTY	periplasm (probable).	periplasm
Q9ZF69	COAE_BURPS	cytoplasm (by similarity).	cytoplasm
Q9ZF70	LEP4_BURPS	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9ZF89	DBHA_BURPS		cytoplasm
Q9ZF99	MDH_AQUAR		cytoplasm
Q9ZFA4	LEXA_RHOS4		cytoplasm
Q9ZFA6	MOAC_RHOS4		cytoplasm
Q9ZFA7	TRPC_RHOS4		cytoplasm
Q9ZFA8	TRPD_RHOS4		cytoplasm
Q9ZFC5	DNAJ_METSS	cytoplasm (probable).	cytoplasm
Q9ZFC6	DNAK_METSS		cytoplasm
Q9ZFC7	GRPE_METSS	cytoplasm (probable).	cytoplasm
Q9ZFD8	CH60_BURVI	cytoplasm (by similarity).	cytoplasm

Q9ZFD9	CH10_BURVI	cytoplasm (by similarity).	cytoplasm
Q9ZFE0	CH60_BURCE	cytoplasm (by similarity).	cytoplasm
Q9ZFE1	CH10_BURCE	cytoplasm (by similarity).	cytoplasm
Q9ZFE4	FABI_PSEAE	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZFF6	RECA_SHISO	cytoplasm (by similarity).	cytoplasm
Q9ZFF7	AROA_SHISO	cytoplasm (probable).	cytoplasm
Q9ZFK4	KDSA_PSEAE	cytoplasm (by similarity).	cytoplasm
Q9ZFL0	ISPH2_BURPS		cytoplasm
Q9ZFN4	GLGC_RHORU		cytoplasm
Q9ZFR5	URED_YERPE		cytoplasm
Q9ZFR7	UREF_YERPE		cytoplasm
Q9ZFR8	UREE_YERPE	cytoplasm (by similarity).	cytoplasm
Q9ZFR9	URE1_YERPE	cytoplasm (by similarity).	cytoplasm
Q9ZFU5	YPFK_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZFU6	YPFL_SALTY		cytoplasm
Q9ZFU7	EUTR_SALTY		cytoplasm
Q9ZFU8	EUTK_SALTY		cytoplasm
Q9ZFV2	EUTA_SALTY		cytoplasm
Q9ZFV4	EUTT_SALTY		cytoplasm
Q9ZFV5	EUTQ_SALTY		cytoplasm
Q9ZFV7	EUTS_SALTY		cytoplasm
Q9ZFV8	MAO2_SALTY		cytoplasm
Q9ZG89	ENGB_CAUCR		cytoplasm
Q9ZGW3	Y4209_YERPE		inner membrane
Q9ZGW6	SYCT_YERPE		cytoplasm
Q9ZH79	TONB_PASMU	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
Q9ZHC7	SILP_SALTY	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZHC9	SILA_SALTY	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZHD0	SILB_SALTY		inner membrane
Q9ZHD2	SILC_SALTY	bacterial cell outer membrane; lipid-anchor (potential).	outer membrane
Q9ZHD3	SILR_SALTY	cytoplasm (probable).	cytoplasm
Q9ZHD4	SILS_SALTY	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9ZHD6	HEM1_NEIG1		cytoplasm
Q9ZHD8	DCD_BUCAP		extracellular
Q9ZHD9	6PGD_BUCAP		cytoplasm
Q9ZHE0	HIS2_BUCAP	cytoplasm (by similarity).	cytoplasm
Q9ZHE1	HIS6_BUCAP	cytoplasm (by similarity).	cytoplasm
Q9ZHE2	HIS4_BUCAP	cytoplasm (by similarity).	cytoplasm
Q9ZHE3	HIS5_BUCAP	cytoplasm (by similarity).	cytoplasm
Q9ZHE4	HIS7_BUCAP	cytoplasm (by similarity).	cytoplasm
Q9ZHE5	HIS8_BUCAP		cytoplasm
Q9ZHE6	HISX_BUCAP		cytoplasm

Q9ZHE7	HIS1_BUCAP	cytoplasm (by similarity).	cytoplasm
Q9ZHE8	Y091_BUCAP		inner membrane
Q9ZHE9	AROC_BUCAP		cytoplasm
Q9ZHF0	BCP_BUCAP		cytoplasm
Q9ZHF3	PILQ_NEIME	bacterial cell outer membrane; peripheral membrane protein (by similarity).	outer membrane
Q9ZHI1	THIL_CHRVO	cytoplasm (potential).	cytoplasm
Q9ZHI2	PHAC_CHRVO	cytoplasm (potential).	cytoplasm
Q9ZHI4	SOXR_CHRVO		cytoplasm
Q9ZHI5	PAND_RALEU		cytoplasm
Q9ZHR1	HYBF_MORMO		cytoplasm
Q9ZHV9	EXBD2_VIBCH	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
Q9ZHW3	1A1D_ENTCL		cytoplasm
Q9ZHY2	RECO_NEIG1		cytoplasm
Q9ZHZ1	RBL1_THIIN		extracellular
Q9ZHZ4	RBL2_THINE		extracellular
Q9ZI33	RBS_BRAJA		cytoplasm
Q9ZI34	RBL_BRAJA		extracellular
Q9ZI54	4OT_PSEST		cytoplasm
Q9ZI76	HFQ_PECCC		cytoplasm
Q9ZI86	LNT_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZIB7	TYRR_ENTAG		cytoplasm
Q9ZIE4	NIFH_ACEDI		cytoplasm
Q9ZIJ1	GNUT_PSEAE	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZIQ0	RECA_RHOVI	cytoplasm (by similarity).	cytoplasm
Q9ZIX8	OTCC_HAEGA	cytoplasm (probable).	cytoplasm
Q9ZIY0	SYK3_HAEGA		cytoplasm
Q9ZJ12	SYK3_SALTY		cytoplasm
Q9ZJ24	YF88_HELPJ		inner membrane
Q9ZJ25	YF87_HELPJ		inner membrane
Q9ZJ27	GCP_HELPJ		extracellular
Q9ZJ28	PDXA_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZJ29	PDXJ_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZJ38	RLPA_HELPJ	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q9ZJ42	ENGB_HELPJ		cytoplasm
Q9ZJ48	FTSW_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZJ57	MFD_HELPJ		cytoplasm
Q9ZJ65	SECF_HELPJ	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZJ66	SECD_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZJ67	YF51_HELPJ		inner membrane
Q9ZJ70	RS2_HELPJ		cytoplasm
Q9ZJ74	F16P_HELPJ	cytoplasm (potential).	cytoplasm

Q9ZJ75	RPE_HELPJ		cytoplasm
Q9ZJ80	RECN_HELPJ		cytoplasm
Q9ZJ81	PPNK_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJ93	THYX_HELPJ		cytoplasm
Q9ZJ94	GLMS_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJ96	DNAA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJA1	RECG_HELPJ		cytoplasm
Q9ZJA6	NUSA_HELPJ		cytoplasm
Q9ZJA7	SELA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJB1	Y1402_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZJC3	PTH_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJC4	RL25_HELPJ		cytoplasm
Q9ZJC5	TAL_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZJC6	MURE_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZJC8	YE91_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZJD1	AN36_HELPJ		inner membrane
Q9ZJD5	YE84_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZJD7	EX7S_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJE3	PAAD_HELPJ		cytoplasm
Q9ZJE4	COAD_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJE5	KTHY_HELPJ		cytoplasm
Q9ZJE9	DPO1_HELPJ		cytoplasm
Q9ZJF1	ILVE_HELPJ		cytoplasm
Q9ZJF9	DPO3A_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJG0	YE59_HELPJ		cytoplasm
Q9ZJG6	TRME_HELPJ		cytoplasm
Q9ZJG8	OXAA_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZJG9	YE49_HELPJ		inner membrane
Q9ZJH0	RNPA_HELPJ		extracellular
Q9ZJH1	EXBL2_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZJH2	SSRP_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZJH3	ISPE_HELPJ		cytoplasm
Q9ZJH4	CSRA_HELPJ		cytoplasm
Q9ZJH5	PPIA_HELPJ		cytoplasm
Q9ZJI4	YE28_HELPJ		cytoplasm
Q9ZJI5	YE29_HELPJ		cytoplasm
Q9ZJI6	YE30_HELPJ		inner membrane
Q9ZJI7	KSGA_HELPJ		cytoplasm
Q9ZJJ0	YE23_HELPJ		cytoplasm
Q9ZJJ1	SYI_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJJ3	FLII_HELPJ	cytoplasm (potential).	inner membrane
Q9ZJJ4	MURB_HELPJ	cytoplasm (probable).	inner membrane
Q9ZJJ7	MIAA_HELPJ		cytoplasm
Q9ZJJ9	QUEF_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZJK8	BIOB_HELPJ		cytoplasm
Q9ZJL6	FABZ_HELPJ	cytoplasm (by similarity).	cytoplasm

Q9ZJL7	LPXA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJL8	CLPX_HELPJ		cytoplasm
Q9ZJM4	YD63_HELPJ		inner membrane
Q9ZJM5	DNAB_HELPJ		nucleoid
Q9ZJN0	PSD_HELPJ		cytoplasm
Q9ZJN1	NADA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJN2	NADC_HELPJ		cytoplasm
Q9ZJN8	PLSC_HELPJ	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q9ZJN9	UNG_HELPJ	cytoplasm (by similarity).	extracellular
Q9ZJP2	CORA_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZJP4	TONB_HELPJ	bacterial cell inner membrane; single-pass membrane protein; periplasmic side (by similarity).	inner membrane
Q9ZJP5	EXBD_HELPJ	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
Q9ZJP7	NIKR_HELPJ		cytoplasm
Q9ZJP8	NADD_HELPJ		cytoplasm
Q9ZJQ0	TRMU_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJQ2	DNAJ_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZJQ3	YD31_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZJQ4	YD30_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZJQ9	FUMC_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJR1	RNH2_HELPJ	cytoplasm (potential).	extracellular
Q9ZJR4	RL3_HELPJ		cytoplasm
Q9ZJR5	RL4_HELPJ		cytoplasm
Q9ZJR6	RL2_HELPJ		cytoplasm
Q9ZJR7	RS19_HELPJ		cytoplasm
Q9ZJR8	RL22_HELPJ		cytoplasm
Q9ZJR9	RS3_HELPJ		cytoplasm
Q9ZJS0	RL16_HELPJ		cytoplasm
Q9ZJS1	RL29_HELPJ		cytoplasm
Q9ZJS2	RL14_HELPJ		cytoplasm
Q9ZJS3	RL24_HELPJ		cytoplasm
Q9ZJS4	RL5_HELPJ		cytoplasm
Q9ZJS5	RS14_HELPJ		cytoplasm
Q9ZJS6	RL6_HELPJ		cytoplasm
Q9ZJS7	RL18_HELPJ		cytoplasm
Q9ZJS8	RL15_HELPJ		cytoplasm
Q9ZJT0	AMPM_HELPJ		cytoplasm
Q9ZJT2	RS13_HELPJ		cytoplasm
Q9ZJT3	RS11_HELPJ		cytoplasm
Q9ZJT4	RS4_HELPJ		cytoplasm
Q9ZJT5	RPOA_HELPJ		cytoplasm
Q9ZJT6	RL17_HELPJ		cytoplasm
Q9ZJU5	TRPE_HELPJ		cytoplasm
Q9ZJU6	TRPG_HELPJ		cytoplasm

Q9ZJU7	TRPD_HELPJ		cytoplasm
Q9ZJU8	TRPC_HELPJ		cytoplasm
Q9ZJU9	TRPB_HELPJ		cytoplasm
Q9ZJV0	TRPA_HELPJ		cytoplasm
Q9ZJW8	NPD_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZJX0	PYRE_HELPJ		cytoplasm
Q9ZJX1	RRF_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZJX2	SECG_HELPJ	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZJX8	AROE_HELPJ		cytoplasm
Q9ZJX9	RNR_HELPJ		cytoplasm
Q9ZJY0	Y1168_HELPJ		cytoplasm
Q9ZJY1	RS6_HELPJ		cytoplasm
Q9ZJY2	SSB_HELPJ		cytoplasm
Q9ZJY6	Y1161_HELPJ	cytoplasm (potential).	inner membrane
Q9ZJY8	AMIF_HELPJ		cytoplasm
Q9ZJY9	CARA_HELPJ		cytoplasm
Q9ZJZ2	YC34_HELPJ	bacterial cell membrane; multi-pass membrane protein (probable).	inner membrane
Q9ZJZ7	AK_HELPJ		cytoplasm
Q9ZJZ8	NUDH_HELPJ		cytoplasm
Q9ZK01	CRCB_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZK05	UPPS_HELPJ		inner membrane
Q9ZK07	PUR2_HELPJ		cytoplasm
Q9ZK12	ATPL_HELPJ		inner membrane
Q9ZK14	CYSE_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZK20	NUSG_HELPJ		cytoplasm
Q9ZK21	RL1_HELPJ		cytoplasm
Q9ZK22	RL10_HELPJ		cytoplasm
Q9ZK23	RPOBC_HELPJ		cytoplasm
Q9ZK28	DHAS_HELPJ		cytoplasm
Q9ZK31	SOTB_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZK37	DEOB_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZK38	DEOD_HELPJ		cytoplasm
Q9ZK41	GLUP_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9ZK47	CSTA_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZK49	G6PI_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZK53	FLAV_HELPJ		cytoplasm
Q9ZK54	YB60_HELPJ		inner membrane
Q9ZK56	PROC_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZK59	MURG_HELPJ	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZK61	SYV_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZK62	SRP54_HELPJ		cytoplasm
Q9ZK63	RS16_HELPJ		cytoplasm

Q9ZK65	RIMM_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZK66	TRMD_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZK67	RL19_HELPJ		cytoplasm
Q9ZK72	FMT_HELPJ		cytoplasm
Q9ZK75	PARB_HELPJ		cytoplasm
Q9ZK77	ATPF_HELPJ		inner membrane
Q9ZK79	ATPA_HELPJ	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZK80	ATPG_HELPJ		inner membrane
Q9ZK81	ATPB_HELPJ		inner membrane
Q9ZK82	ATPE_HELPJ	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZK83	EXBL1_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZK84	EXDL1_HELPJ	bacterial cell inner membrane; single-pass type ii membrane protein (probable).	inner membrane
Q9ZK86	TOLB_HELPJ	periplasm (potential).	periplasm
Q9ZK89	SLYD_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZKA0	UVRB_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZKA2	PUR8_HELPJ		cytoplasm
Q9ZKB0	GLK_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZKB1	6PGL_HELPJ		extracellular
Q9ZKB2	G6PD_HELPJ		cytoplasm
Q9ZKB3	EDD_HELPJ		cytoplasm
Q9ZKB5	HCPC_HELPJ	secreted protein (potential).	extracellular
Q9ZKC0	TRUA_HELPJ		cytoplasm
Q9ZKC2	PIMT_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZKC3	RIR2_HELPJ		cytoplasm
Q9ZKD0	DCD_HELPJ		extracellular
Q9ZKD2	Y374_HELPJ		cytoplasm
Q9ZKD4	HEMH_HELPJ	cytoplasm (by similarity).	extracellular
Q9ZKD8	DHE4_HELPJ		cytoplasm
Q9ZKE4	PRIA_HELPJ		cytoplasm
Q9ZKE6	SODF_HELPJ		periplasm
Q9ZKE7	TPX_HELPJ		cytoplasm
Q9ZKF2	Y395_HELPJ		cytoplasm
Q9ZKF3	Y396_HELPJ		cytoplasm
Q9ZKF6	RS1_HELPJ		cytoplasm
Q9ZKF7	AROA_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZKG4	GUAA_HELPJ		cytoplasm
Q9ZKG7	Y415_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZKH6	Y960_HELPJ		inner membrane
Q9ZKM3	FTSA_HELPJ		inner membrane
Q9ZKM5	BIOA_HELPJ		cytoplasm
Q9ZKM6	GATC_HELPJ		cytoplasm
Q9ZKM7	GPMI_HELPJ		cytoplasm
Q9ZKP0	GPDA_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZKP2	Y959_HELPJ		cytoplasm
Q9ZKP5	Y956_HELPJ		cytoplasm

Q9ZKP6	LGT_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZKQ2	Y883_HELPJ		inner membrane
Q9ZKQ6	Y944_HELPJ		cytoplasm
Q9ZKQ7	Y943_HELPJ		cytoplasm
Q9ZKQ9	ALR_HELPJ		cytoplasm
Q9ZKR8	PTPS_HELPJ		extracellular
Q9ZKS0	SURE_HELPJ	cytoplasm (potential).	periplasm
Q9ZKS2	GCH1_HELPJ		cytoplasm
Q9ZKS4	HTPX_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9ZKS5	TRUD_HELPJ		cytoplasm
Q9ZKS6	RECR_HELPJ		cytoplasm
Q9ZKS7	Y858_HELPJ		cytoplasm
Q9ZKT1	Y920_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZKT2	CARB_HELPJ		cytoplasm
Q9ZKU4	PTA_HELPJ	cytoplasm (potential).	inner membrane
Q9ZKU5	ACKA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZKU8	HYPB_HELPJ		cytoplasm
Q9ZKW1	VDLC_HELPJ		cytoplasm
Q9ZKW3	Y888_HELPJ	bacterial cell inner membrane; peripheral membrane protein (potential).	inner membrane
Q9ZKW7	MVIN_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZKW9	RUVA_HELPJ		cytoplasm
Q9ZKX3	RUVC_HELPJ		cytoplasm
Q9ZKX5	CATA_HELPJ	cytoplasm (probable).	periplasm
Q9ZKX9	CDH_HELPJ	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9ZKY0	FLGE_HELPJ		flagellar
Q9ZKY2	LPXB_HELPJ		cytoplasm
Q9ZKY3	DUT_HELPJ		extracellular
Q9ZKY5	PGBB_HELPJ	the plasminogen-binding region is localized on the surface the bacterium (probable).	cytoplasm
Q9ZKY8	GMHB_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZKZ0	HLDE_HELPJ		cytoplasm
Q9ZKZ1	GMHA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZKZ2	GUAC_HELPJ		cytoplasm
Q9ZKZ9	THIM_HELPJ		cytoplasm
Q9ZL00	THID_HELPJ		cytoplasm
Q9ZL01	THIE_HELPJ		cytoplasm
Q9ZL08	DBH_HELPJ		cytoplasm
Q9ZL09	ENGA_HELPJ		cytoplasm
Q9ZL11	SPEE_HELPJ		cytoplasm
Q9ZL12	COAE_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZL13	GATA_HELPJ		cytoplasm
Q9ZL14	IMDH_HELPJ		cytoplasm

Q9ZL18	TRXB_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZL19	Y823_HELPJ		inner membrane
Q9ZL20	DHOM_HELPJ		cytoplasm
Q9ZL21	UVRC_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZL36	ACPS_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZL40	RIBB_HELPJ		cytoplasm
Q9ZL42	GCH2_HELPJ		extracellular
Q9ZL44	MOAE_HELPJ		cytoplasm
Q9ZL45	MOG_HELPJ		cytoplasm
Q9ZL46	MOAC_HELPJ		cytoplasm
Q9ZL47	HAAA_HELPJ	bacterial cell outer membrane; lipid-anchor (probable).	outer membrane
Q9ZL49	TIG_HELPJ		cytoplasm
Q9ZL50	CLPP_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZL51	DEF_HELPJ		cytoplasm
Q9ZL57	SECA_HELPJ	bacterial cell inner membrane; peripheral membrane protein; cytoplasmic side (by similarity).	inner membrane
Q9ZL58	LOLA_HELPJ	periplasm (by similarity).	periplasm
Q9ZL64	ACON2_HELPJ		cytoplasm
Q9ZL66	PYRH_HELPJ		cytoplasm
Q9ZL69	SYU_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZL73	FLHB_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZL74	MOBA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZL75	MOAA_HELPJ		cytoplasm
Q9ZL80	FTSY_HELPJ	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZL83	Y760_HELPJ		inner membrane
Q9ZL96	TRMB_HELPJ		cytoplasm
Q9ZL98	Y745_HELPJ		cytoplasm
Q9ZLA0	RODA_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZLA1	KPRS_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLA5	DDL_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLA7	Y736_HELPJ		cytoplasm
Q9ZLB5	TILS_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZLB6	DUS_HELPJ		cytoplasm
Q9ZLB9	ASPG_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZLC8	RP54_HELPJ		cytoplasm
Q9ZLD4	MRAW_HELPJ		cytoplasm
Q9ZLD6	UVRA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLD9	GYRA_HELPJ		cytoplasm
Q9ZLE0	KDGL_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZLE3	SCOA_HELPJ		cytoplasm
Q9ZLE4	SCOB_HELPJ		cytoplasm
Q9ZLF3	FEOB_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZLF6	GLMU_HELPJ	cytoplasm (by similarity).	cytoplasm

Q9ZLF9	RIR1_HELPJ		cytoplasm
Q9ZLH0	HEMN_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLH1	AROC_HELPJ		cytoplasm
Q9ZLH2	RNC_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLH3	RNH_HELPJ	cytoplasm (potential).	extracellular
Q9ZLH6	GATB_HELPJ		cytoplasm
Q9ZLI1	FTN_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLI5	ASPA_HELPJ		cytoplasm
Q9ZLI6	MURA_HELPJ	cytoplasm (probable).	inner membrane
Q9ZLL0	ISPG_HELPJ		cytoplasm
Q9ZLL2	MURC_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZLL4	MUTS2_HELPJ		cytoplasm
Q9ZLL5	IPYR_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLM1	DNLJ_HELPJ		cytoplasm
Q9ZLM8	DCUP_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZLN3	BIOF_HELPJ		cytoplasm
Q9ZLP5	Y58A_HELPJ		cytoplasm
Q9ZLQ0	PYRC_HELPJ		cytoplasm
Q9ZLQ4	FOLD_HELPJ		cytoplasm
Q9ZLQ5	LEP_HELPJ	bacterial cell membrane; single-pass type ii membrane protein (potential).	inner membrane
Q9ZLR1	AMPA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLR4	Y567_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZLR5	DAPF_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLR9	RS21_HELPJ		cytoplasm
Q9ZLS1	ACP_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLS3	ACCA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLS8	Y552_HELPJ		cytoplasm
Q9ZLS9	RHO_HELPJ		inner membrane
Q9ZLT0	MURI_HELPJ		cytoplasm
Q9ZLT1	CAGA_HELPJ		cytoplasm
Q9ZLT4	CAGE_HELPJ		cytoplasm
Q9ZLU4	CAGS_HELPJ		cytoplasm
Q9ZLU5	CAGT_HELPJ	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q9ZLW0	ERA_HELPJ		cytoplasm
Q9ZLW1	HSLU_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLW2	HSLV_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLW3	RL9_HELPJ		cytoplasm
Q9ZLW6	DAPB_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLW8	PGBA_HELPJ	the plasminogen-binding region is localized on the surface the bacterium (probable).	cytoplasm
Q9ZLX3	GYRB_HELPJ		cytoplasm
Q9ZLX4	DPO3B_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLX8	Y496_HELPJ		cytoplasm
Q9ZLX9	Y495_HELPJ		cytoplasm
Q9ZLY0	MURD_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZLY1	MRAY_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZLZ3	TYPA_HELPJ		cytoplasm

Q9ZM10	PPK_HELPJ		cytoplasm
Q9ZM11	PYRD_HELPJ	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZM13	DAPA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZM19	ISPDF_HELPJ		cytoplasm
Q9ZM26	FUR_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZM34	FLHF_HELPJ		cytoplasm
Q9ZM35	HPPK_HELPJ		cytoplasm
Q9ZM37	AROQ_HELPJ		cytoplasm
Q9ZM39	RS15_HELPJ		cytoplasm
Q9ZM40	FLHA_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZM43	YA44_HELPJ		extracellular
Q9ZM44	YA46_HELPJ		inner membrane
Q9ZM45	RBFA_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZM48	KHSE_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZM50	LPXC_HELPJ		cytoplasm
Q9ZM51	MINC_HELPJ		inner membrane
Q9ZM56	PANB_HELPJ		cytoplasm
Q9ZM57	RUVB_HELPJ		cytoplasm
Q9ZM58	TATB_HELPJ	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane
Q9ZM59	TATC_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (probable).	inner membrane
Q9ZM60	QUEA_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZM61	GIDB_HELPJ		cytoplasm
Q9ZM64	CHEY_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZM65	PRMA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZM67	YA70_HELPJ		cytoplasm
Q9ZM70	COPP_HELPJ		periplasm
Q9ZM74	NIXA_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZM81	PYRB_HELPJ		cytoplasm
Q9ZM93	LEPA_HELPJ		cytoplasm
Q9ZM94	DXS_HELPJ		cytoplasm
Q9ZM99	PYRG_HELPJ		cytoplasm
Q9ZMA1	Y347_HELPJ		cytoplasm
Q9ZMA5	RUVX_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZMA7	MINE_HELPJ		inner membrane
Q9ZMA8	MIND_HELPJ	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZMA9	ILVC_HELPJ		cytoplasm
Q9ZMB0	NADE_HELPJ		cytoplasm
Q9ZMB1	LPXK_HELPJ		cytoplasm
Q9ZMB3	FLGH_HELPJ	bacterial cell outer membrane (probable).	inner membrane
Q9ZMB7	KGUA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZMB8	TATA_HELPJ	bacterial cell inner membrane; single-pass membrane protein (by similarity).	inner membrane

Q9ZMD0	GSA_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZMD3	Y303_HELPJ		cytoplasm
Q9ZMD8	RL27_HELPJ		cytoplasm
Q9ZMD9	RL21_HELPJ		cytoplasm
Q9ZME1	AMIE_HELPJ		cytoplasm
Q9ZME5	DCDA_HELPJ		cytoplasm
Q9ZMF0	Y285_HELPJ		cytoplasm
Q9ZMF2	AROB_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZMF4	TGT_HELPJ		cytoplasm
Q9ZMG1	Y274_HELPJ		inner membrane
Q9ZMG6	Y269_HELPJ		cytoplasm
Q9ZMG9	PYRX_HELPJ		extracellular
Q9ZMH1	CLPB_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZMH7	EX7L_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZMH8	Y242_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	extracellular
Q9ZMI1	PURA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZMI9	FLGI_HELPJ	periplasm (by similarity).	periplasm
Q9ZMJ1	NAPA_HELPJ		cytoplasm
Q9ZMJ5	HEM1_HELPJ		cytoplasm
Q9ZMJ7	HEM3_HELPJ		cytoplasm
Q9ZMJ9	HCPE_HELPJ	secreted protein (potential).	extracellular
Q9ZMK2	Y217_HELPJ	bacterial cell membrane; lipid-anchor (probable).	inner membrane
Q9ZMK4	KDSB_HELPJ	cytoplasm (potential).	cytoplasm
Q9ZMK8	MSRAB_HELPJ		cytoplasm
Q9ZMK9	RADA_HELPJ		cytoplasm
Q9ZML2	ISCS_HELPJ		cytoplasm
Q9ZML6	DXR_HELPJ		cytoplasm
Q9ZML7	CDSA_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZML9	GIDA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZMM1	HCPA_HELPJ	secreted protein (by similarity).	extracellular
Q9ZMM2	HTPG_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZMM5	MRP_HELPJ		cytoplasm
Q9ZMN0	FABH_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZMN1	PLSX_HELPJ		cytoplasm
Q9ZMN4	NDK_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZMN6	LPXD_HELPJ		cytoplasm
Q9ZMN7	FABI_HELPJ	bacterial cell inner membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZMN8	TPIS_HELPJ		cytoplasm
Q9ZMN9	FRDC_HELPJ	bacterial cell membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZMP0	FRDA_HELPJ		periplasm
Q9ZMP1	FRDB_HELPJ		cytoplasm
Q9ZMP2	Y190_HELPJ	bacterial cell membrane; single-pass membrane protein (potential).	inner membrane

Q9ZMP3	Y189_HELPJ	bacterial cell membrane; multi-pass membrane protein (potential).	inner membrane
Q9ZMP7	GLYA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZMQ0	LNT_HELPJ	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZMQ6	ALF_HELPJ		cytoplasm
Q9ZMQ7	Y175_HELPJ		cytoplasm
Q9ZMR3	Y155_HELPJ		cytoplasm
Q9ZMR8	HEM2_HELPJ		cytoplasm
Q9ZMR9	Y162_HELPJ		inner membrane
Q9ZMS0	HCPD_HELPJ	secreted protein (by similarity).	extracellular
Q9ZMS3	AROK_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZMS6	ENO_HELPJ	cytoplasm (by similarity).	extracellular
Q9ZMS7	RECA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZMU4	BCP_HELPJ		cytoplasm
Q9ZMU7	SDHL_HELPJ		cytoplasm
Q9ZMV1	RL20_HELPJ		cytoplasm
Q9ZMV4	PPSA_HELPJ		cytoplasm
Q9ZMV6	Y117_HELPJ		cytoplasm
Q9ZMV7	TOP1_HELPJ		cytoplasm
Q9ZMV8	FLAB_HELPJ		flagellar
Q9ZMW2	HRCA_HELPJ		cytoplasm
Q9ZMW3	GRPE_HELPJ	cytoplasm (probable).	cytoplasm
Q9ZMW4	DNAK_HELPJ		cytoplasm
Q9ZMW6	CYSM_HELPJ		cytoplasm
Q9ZMW8	LUXS_HELPJ		extracellular
Q9ZMX5	THRC_HELPJ		cytoplasm
Q9ZMY2	MTNN_HELPJ		cytoplasm
Q9ZMY3	RPOD_HELPJ		cytoplasm
Q9ZMY5	MQO_HELPJ	bacterial cell membrane; peripheral membrane protein (by similarity).	inner membrane
Q9ZMY6	RL13_HELPJ		cytoplasm
Q9ZMZ1	RS20_HELPJ		cytoplasm
Q9ZMZ2	GLMM_HELPJ		cytoplasm
Q9ZMZ3	LSPA_HELPJ	bacterial cell membrane; multi-pass membrane protein (by similarity).	extracellular
Q9ZMZ4	URE23_HELPJ		cytoplasm
Q9ZMZ6	UREF_HELPJ		cytoplasm
Q9ZMZ7	UREG_HELPJ		cytoplasm
Q9ZMZ8	UREH_HELPJ		cytoplasm
Q9ZN29	Y035_HELPJ		inner membrane
Q9ZN30	PAND_HELPJ		cytoplasm
Q9ZN32	CLPS_HELPJ		cytoplasm
Q9ZN34	BIOD_HELPJ		cytoplasm
Q9ZN36	IDH_HELPJ		cytoplasm
Q9ZN37	CISY_HELPJ		cytoplasm
Q9ZN49	PRIM_HELPJ		cytoplasm
Q9ZN50	CH60_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZN52	PANC_HELPJ		cytoplasm
Q9ZN53	PYRF_HELPJ		cytoplasm
Q9ZN54	CYNT_HELPJ		cytoplasm

Q9ZN55	KDSA_HELPJ	cytoplasm (by similarity).	cytoplasm
Q9ZN56	RISB_HELPJ		cytoplasm
Q9ZN57	NUSB_HELPJ		cytoplasm
Q9ZN70	PPX_PSEAE		cytoplasm
Q9ZN79	LINR_PSEPA		cytoplasm
Q9ZN99	CATA_DESVM		periplasm
Q9ZNA5	SAHH_ROSDE	cytoplasm (by similarity).	cytoplasm
Q9ZNA8	TNAA_PROIN		cytoplasm
Q9ZNB7	ALGL_HALMR	periplasm (by similarity).	periplasm
Q9ZNE7	SECE_VIBAL	bacterial cell inner membrane; multi-pass membrane protein (by similarity).	inner membrane
Q9ZNH4	PPCK_RHOPA	cytoplasm (by similarity).	cytoplasm