

Online Supporting Information B

Predicted subcellular locations for the 7,962 human proteins

| Accession number | Swiss-Prot code | Uncertain annotation in Swiss-Prot database | Predicted location by Hum-mPLoc |
|------------------|-----------------|--|-----------------------------------|
| O00124 | UBXD6_HUMAN | | secreted protein. |
| O00139 | KIF2_HUMAN | | cytoplasm. nucleus. |
| O00144 | FZD9_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | membrane. |
| O00148 | DDX39_HUMAN | nucleus (by similarity). | nucleus. |
| O00151 | PDLI1_HUMAN | cytoplasm (by similarity). cytoskeleton (by similarity). associates with the actin stress fibers (by similarity). | cytoplasm. |
| O00159 | MYO1C_HUMAN | | cytoskeleton. |
| O00160 | MYO1F_HUMAN | | cytoskeleton. |
| O00167 | EYA2_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| O00178 | GTPB1_HUMAN | | cytoplasm. |
| O00180 | KCNK1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O00182 | LEG9_HUMAN | | cytoplasm. |
| O00192 | ARVC_HUMAN | | cytoplasm. cytoskeleton. nucleus. |
| O00198 | HRK_HUMAN | | cytoplasm. |
| O00203 | AP3B1_HUMAN | Golgi apparatus. cytoplasmic vesicle; cytoplasmic vesicle membrane; peripheral membrane protein; cytoplasmic side (by similarity). component of the coat surrounding the cytoplasmic face of coated vesicles located at the Golgi complex (by similarity). | Golgi. |
| O00204 | ST2B1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O00212 | RHOD_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| O00213 | APBB1_HUMAN | | cytoplasm. nucleus. |
| O00214 | LEG8_HUMAN | cytoplasm (probable). | secreted protein. |
| O00217 | NUIM_HUMAN | | mitochondrion. |
| O00219 | HAS3_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O00221 | IKBE_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O00231 | PSD11_HUMAN | | centriole. cytoplasm. endosome. |
| O00232 | PSD12_HUMAN | | cytoplasm. |
| O00233 | PSD9_HUMAN | | cytoplasm. |
| O00237 | RN103_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. |
| O00244 | ATOX1_HUMAN | | cytoplasm. secreted protein. |
| O00253 | AGRP_HUMAN | secreted protein (by similarity). | secreted protein. |
| O00264 | PGRC1_HUMAN | microsome; microsomal membrane; single-pass membrane protein (by | membrane. |
| O00294 | TULP1_HUMAN | | centriole. |
| O00295 | TULP2_HUMAN | | centriole. cytoplasm. |
| O00303 | IF35_HUMAN | | cytoplasm. |
| O00305 | CACB4_HUMAN | | membrane. |

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| O00308 | WWP2_HUMAN | | cytoplasm. |
| O00327 | BMAL1_HUMAN | nucleus (by similarity). | nucleus. |
| O00329 | PK3CD_HUMAN | | cytoplasm. |
| O00358 | FOXE1_HUMAN | nucleus (potential). | nucleus. |
| O00391 | QSCN6_HUMAN | membrane; single-pass membrane protein (potential). isoform 1 may be found in the membrane. extracellular. isoform 2 is found in the extracellular medium of quiescent cells and is not found in | endosome; membrane. |
| O00399 | DCTN6_HUMAN | | mitochondrion. |
| O00400 | ACATN_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). | endoplasmic reticulum; membrane. |
| O00401 | WASL_HUMAN | cytoplasm (by similarity). nucleus (by similarity). preferentially localized in the cytoplasm when phosphorylated and in the nucleus when unphosphorylated (by similarity). | cytoplasm. |
| O00408 | PDE2A_HUMAN | membrane; peripheral membrane protein (potential). | cytoplasm. |
| O00409 | FOXN3_HUMAN | nucleus (probable). | nucleus. |
| O00418 | EF2K_HUMAN | | cytoplasm. |
| O00421 | CCRL2_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| O00422 | SAP18_HUMAN | | nucleus. |
| O00423 | EMAL1_HUMAN | | cytoplasm. |
| O00444 | PLK4_HUMAN | | cytoplasm. |
| O00445 | SYT5_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein (by similarity). endosome; recycling endosome; recycling endosomal membrane; multi-pass membrane protein (by similarity). in mast cells, localizes to the endocytic recycling compartment (by | membrane. |
| O00451 | GFRA2_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | secreted protein; membrane. |
| O00458 | IFRD1_HUMAN | | cytoplasm. secreted protein. |
| O00459 | P85B_HUMAN | | cytoplasm. |
| O00463 | TRAF5_HUMAN | cytoplasm (probable). | cytoplasm. |
| O00468 | AGRIN_HUMAN | secreted protein; extracellular space; extracellular matrix. synaptic basal lamina at the neuromuscular junction (by | cytoplasm. |
| O00470 | MEIS1_HUMAN | nucleus (by similarity). | nucleus. |
| O00478 | BT3A3_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| O00479 | HMGN4_HUMAN | nucleus (by similarity). | nucleus. |
| O00481 | BT3A1_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| O00482 | NR5A2_HUMAN | nucleus (probable). | nucleus. |
| O00487 | PSDE_HUMAN | | cytoplasm. |
| O00505 | IMA3_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| O00507 | USP9Y_HUMAN | | nucleus. |
| O00512 | BCL9_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| O00522 | KRIT1_HUMAN | membrane; peripheral membrane protein (probable). | cytoplasm. |

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| O00526 | UPK2_HUMAN | membrane; single-pass type i membrane protein (potential). heterodimer formation with uroplakin ia is a prerequisite to exit out of the endoplasmic reticulum (er) (by similarity). | endoplasmic reticulum. |
| O00533 | CHL1_HUMAN | membrane; single-pass type i membrane protein. soluble forms produced by cleavage/shedding also exist (by | membrane. |
| O00534 | LHR2A_HUMAN | | cytoplasm. |
| O00570 | SOX1_HUMAN | nucleus (probable). | nucleus. |
| O00584 | RNT2_HUMAN | secreted protein (potential). | secreted protein. |
| O00587 | MFNG_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. |
| O00592 | PODXL_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O00629 | IMA4_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| O00634 | NET2_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| O00635 | TRI38_HUMAN | | cytoplasm. |
| O00743 | PPP6_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O00748 | EST2_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| O00757 | F16P2_HUMAN | | lysosome. |
| O00762 | UBE2C_HUMAN | | cytoplasm. nucleus. |
| O00767 | ACOD_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). | endoplasmic reticulum. |
| O14490 | DLGP1_HUMAN | cell membrane; peripheral membrane protein (by similarity). found in postsynaptic density of neuronal cells (by | cytoplasm. |
| O14493 | CLD4_HUMAN | cell membrane; cell-cell junction; tight junction; multi-pass membrane protein (by similarity). | membrane. |
| O14508 | SOCS2_HUMAN | | cytoplasm. |
| O14511 | NRG2_HUMAN | pro-neuregulin-2, membrane-bound isoform: cell membrane; single-pass type i membrane protein (by similarity). neuregulin-2: secreted protein (by similarity). exists also as a proteolytically released soluble growth factor form. the membrane-bound form does not seem to be active (by similarity). | secreted protein. |
| O14519 | CDKA1_HUMAN | | nucleus. |
| O14523 | TMM24_HUMAN | | membrane. |
| O14524 | K0286_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O14525 | ASTN_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O14529 | CUTL2_HUMAN | nucleus (by similarity). | nucleus. |
| O14530 | TXND9_HUMAN | | centriole. cytoplasm. |
| O14531 | DPYL4_HUMAN | cytoplasm (potential). | secreted protein. |
| O14543 | SOCS3_HUMAN | | cytoplasm. |
| O14544 | SOCS6_HUMAN | | cytoplasm. |
| O14548 | COX7R_HUMAN | mitochondrion (by similarity). | mitochondrion. |

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|--------|-------------|--|-----------------------------------|
| O14556 | G3PT_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O14558 | HSPB6_HUMAN | | cytoplasm. |
| O14559 | TCGAP_HUMAN | | cytoplasm. |
| O14569 | C56D2_HUMAN | membrane; multi-pass membrane protein (probable). | mitochondrion. |
| O14576 | DC111_HUMAN | | cytoplasm. |
| O14578 | CTRO_HUMAN | cytoplasm (by similarity). | cytoplasm. nucleus. |
| O14579 | COPE_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). the coatomer is cytoplasmic or polymerized on the cytoplasmic side of the Golgi, as well as on the vesicles/buds originating from it (by similarity). | cytoplasm. |
| O14594 | CSPG3_HUMAN | | secreted protein. |
| O14595 | CTDS2_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| O14598 | VCY1_HUMAN | | cytoplasm. |
| O14599 | VCY2_HUMAN | | cytoplasm. |
| O14602 | IF1AY_HUMAN | | cytoplasm. |
| O14603 | PRY_HUMAN | | cytoplasm. |
| O14607 | UTY_HUMAN | nucleus (potential). | centriole. nucleus. |
| O14609 | XKRY_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| O14610 | GBGT2_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| O14617 | AP3D1_HUMAN | | Golgi. |
| O14627 | CDX4_HUMAN | nucleus (by similarity). | nucleus. |
| O14628 | ZN195_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O14633 | LCE2B_HUMAN | | cytoplasm. secreted protein. |
| O14639 | ABLM1_HUMAN | cytoplasm (by similarity). cytoskeleton (by similarity). associated with the cytoskeleton (by similarity). | cytoplasm. |
| O14640 | DVL1_HUMAN | | cytoplasm. |
| O14645 | IDLC_HUMAN | | cytoplasm. |
| O14647 | CHD2_HUMAN | nucleus (by similarity). | nucleus. |
| O14649 | KCNK3_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O14653 | GOSR2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type iv membrane protein (potential). | endoplasmic reticulum. Golgi. |
| O14657 | TOR1B_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | cytoplasm. endoplasmic reticulum. |
| O14668 | TMG1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O14681 | EI24_HUMAN | | cytoplasm. nucleus. |
| O14683 | P5I11_HUMAN | | cytoplasm. |
| O14684 | PTGES_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O14686 | MLL2_HUMAN | nucleus (probable). | nucleus. |
| O14709 | ZN197_HUMAN | nucleus (probable). | nucleus. |
| O14717 | DNMT2_HUMAN | nucleus (probable). | nucleus. |
| O14718 | OPSX_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| O14730 | RIOK3_HUMAN | | cytoplasm. |
| O14732 | IMPA2_HUMAN | | cytoplasm. |
| O14733 | MP2K7_HUMAN | | cytoplasm. |

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|--------|-------------|--|-----------------------------------|
| O14735 | CDIPT_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). Golgi apparatus; Golgi membrane; multi-pass membrane protein (by similarity). cell membrane; multi-pass membrane protein | endoplasmic reticulum. |
| O14737 | PDCD5_HUMAN | | cytoplasm. |
| O14753 | OVOL1_HUMAN | nucleus (by similarity). | nucleus. |
| O14770 | MEIS2_HUMAN | nucleus (probable). | nucleus. |
| O14771 | ZN213_HUMAN | nucleus (potential). | nucleus. |
| O14775 | GBB5_HUMAN | | cytoplasm. |
| O14782 | KIF3C_HUMAN | | cytoplasm. nucleus. |
| O14787 | TNPO2_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| O14788 | TNF11_HUMAN | isoform 1, isoform 3: cell membrane; single- pass type ii membrane protein. isoform 2, processed isoform 1: secreted protein (by similarity). | membrane. |
| O14792 | OST1_HUMAN | Golgi apparatus; Golgi lumen (probable). | membrane. |
| O14795 | UN13B_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). cell membrane (by similarity). localized to synapses. translocated to the plasma membrane in response to phorbol ester binding (by similarity). | cytoplasm. |
| O14796 | SH21B_HUMAN | | centriole. cytoplasm. |
| O14807 | RASM_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| O14810 | CPLX1_HUMAN | cytoplasm; cytosol. enriched at synaptic-releasing sites in mature neurons (by similarity). | cytoplasm; membrane. |
| O14813 | PHX2A_HUMAN | nucleus (by similarity). | nucleus. |
| O14815 | CAN9_HUMAN | | cytoplasm. |
| O14829 | PPE1_HUMAN | | cytoplasm. |
| O14841 | OPLA_HUMAN | | cytoplasm. |
| O14862 | AIM2_HUMAN | | secreted protein. |
| O14863 | ZNT4_HUMAN | endosome; endosomal membrane; multi-pass membrane protein (probable). | membrane. |
| O14867 | BACH1_HUMAN | nucleus (by similarity). | nucleus. |
| O14879 | IFIT3_HUMAN | | centriole. cytoplasm. |
| O14880 | MGST3_HUMAN | microsome; microsomal membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. microsome. |
| O14896 | IRF6_HUMAN | nucleus (potential). | nucleus. |
| O14910 | LIN7A_HUMAN | cell membrane; peripheral membrane protein. cell membrane; cell-cell junction; peripheral membrane protein (by similarity). localized to postsynaptic densities of neurons and epithelial cell-cell junctions (by similarity). mainly basolateral in renal epithelial cells. | Golgi. |
| O14917 | PCD17_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| O14921 | RGS13_HUMAN | | cytoplasm. |

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| O14925 | TIM23_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (by similarity). mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| O14926 | FSCN2_HUMAN | | cytoplasm. |
| O14931 | NCTR3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O14933 | UB2L6_HUMAN | | cytoplasm. |
| O14936 | CSKP_HUMAN | cytoplasm (potential). | cytoplasm. |
| O14939 | PLD2_HUMAN | membrane; peripheral membrane protein (by similarity). | membrane. |
| O14940 | MOS1A_HUMAN | | cytoplasm. |
| O14966 | RAB7L_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| O14972 | DSCR3_HUMAN | | centriole. cytoplasm. |
| O14976 | GAK_HUMAN | cytoplasm; perinuclear region. Golgi apparatus; trans-Golgi network. cell membrane; cell-matrix junction; focal adhesion (probable). localizes to the perinuclear area and to the trans-Golgi network. also seen on the plasma membrane, probably at focal adhesions. | nucleus. |
| O14977 | AZIN1_HUMAN | | cytoplasm. |
| O14978 | ZN263_HUMAN | nucleus (by similarity). | nucleus. |
| O14986 | PI51B_HUMAN | intracytoplasmic membrane (by similarity). associated with membranes (by similarity). | membrane. |
| O14990 | IPP4_HUMAN | | cytoplasm. |
| O15013 | ARHGA_HUMAN | | cytoplasm. |
| O15015 | ZN646_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O15016 | TRI66_HUMAN | | cytoplasm. |
| O15018 | PDZD2_HUMAN | nucleus (by similarity). cytoplasm. endoplasmic reticulum. at cell-cell contacts in lung epithelial cells. the proteolytic processed protein is secreted (by | cytoplasm. |
| O15020 | SPTN2_HUMAN | | cytoplasm. |
| O15027 | K0310_HUMAN | | cytoplasm. |
| O15033 | K0317_HUMAN | | cytoplasm. |
| O15034 | RIMB2_HUMAN | cell membrane (by similarity). synapse (by similarity). synaptic plasma membrane (by similarity). | cytoplasm. Golgi. |
| O15037 | K0323_HUMAN | | nucleus. |
| O15040 | K0329_HUMAN | | cytoplasm. |
| O15041 | SEM3E_HUMAN | secreted protein (by similarity). | secreted protein; membrane. |
| O15047 | SET1A_HUMAN | nucleus (probable). | nucleus. |
| O15055 | PER2_HUMAN | nucleus (potential). | nucleus. |
| O15056 | SYNJ2_HUMAN | membrane (by similarity). predominantly associated with the particulate fractions (by similarity). | nucleus. |
| O15060 | ZBT39_HUMAN | nucleus (potential). | nucleus. |
| O15061 | DMN_HUMAN | cytoplasm (probable). there are at least two distinct dmN subpopulations, one in which dmN interacts with desmin within the z-lines, and another in which it interacts with both alpha-dystrobrevin and desmin at the costamere. | cytoplasm. |

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| O15062 | ZBTB5_HUMAN | nucleus (potential). | nucleus. |
| O15063 | K0355_HUMAN | | nucleus. |
| O15066 | KIF3B_HUMAN | | cytoplasm. nucleus. |
| O15067 | PUR4_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O15072 | ATS3_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| O15075 | DCAK1_HUMAN | | cytoplasm. |
| O15083 | ERC2_HUMAN | cytoplasm. in neurons, localized to synapses, and colocalizes with pclo. localized to the active zone of presynaptic density (by similarity). | cytoplasm. |
| O15084 | ANR28_HUMAN | | cytoplasm. |
| O15091 | K0391_HUMAN | | cytoplasm. |
| O15105 | SMAD7_HUMAN | nucleus. interaction with nedd4l induces translocation from the nucleus to the cytoplasm (by similarity). | nucleus. |
| O15116 | LSM1_HUMAN | nucleus (potential). | nucleus. |
| O15117 | FYB_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. nucleus. |
| O15119 | TBX3_HUMAN | nucleus (potential). | nucleus. |
| O15120 | PLCB_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| O15143 | ARC1B_HUMAN | | cytoplasm. |
| O15144 | ARPC2_HUMAN | | cytoplasm. |
| O15145 | ARPC3_HUMAN | | cytoplasm. |
| O15146 | MUSK_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O15155 | BET1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type iv membrane protein (by similarity). | endoplasmic reticulum. |
| O15164 | TIF1A_HUMAN | nucleus (potential). | nucleus. |
| O15165 | CR001_HUMAN | cell membrane; single-pass membrane protein (potential). | membrane. |
| O15173 | PGRC2_HUMAN | | membrane. |
| O15194 | CTDSL_HUMAN | nucleus (potential). | nucleus. |
| O15195 | VILL_HUMAN | | cytoplasm. |
| O15198 | SMAD9_HUMAN | cytoplasm (by similarity). in the cytoplasm in the absence of ligand. nucleus (by similarity). migration to the nucleus when complexed with smad4 (by similarity). | cytoplasm. nucleus. |
| O15204 | ADEC1_HUMAN | secreted protein (by similarity). | membrane. |
| O15205 | UBD_HUMAN | | cytoplasm. |
| O15209 | ZN297_HUMAN | nucleus (probable). | nucleus. |
| O15211 | RGL2_HUMAN | | cytoplasm. |
| O15212 | PFD6_HUMAN | | cytoplasm. |
| O15213 | WDR46_HUMAN | | centriole. cytoplasm. |
| O15225 | INE1_HUMAN | | centriole. cytoplasm. |
| O15228 | GNPAT_HUMAN | peroxisome; peroxisomal membrane; matrix side (by similarity). exclusively localized to the luminal side of the peroxisomal membrane (by similarity). | cytoplasm. endoplasmic reticulum. mitochondrion. peroxisome. |
| O15231 | ZN185_HUMAN | nucleus (potential). | cytoplasm. nucleus. |

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| O15234 | CASC3_HUMAN | cytoplasm; perinuclear region (by similarity). predominantly found in the perinuclear region (by similarity). nucleus (by similarity). shuttles between the nucleus and the cytoplasm in a crm1-dependent manner (by similarity). | nucleus. |
| O15240 | VGF_HUMAN | stored in secretory vesicles and then secreted (by similarity). | cytoplasm. |
| O15243 | OBRG_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O15247 | CLIC2_HUMAN | | nucleus; membrane. |
| O15254 | ACOX3_HUMAN | peroxisome (potential). | mitochondrion. |
| O15255 | CXX1_HUMAN | cell membrane; lipid-anchor (potential). | secreted protein. |
| O15259 | NPHP1_HUMAN | cell membrane; cell-cell junction; adherens junction (by similarity). localizes at or near the cell-cell adherens junctions (by similarity). | cytoplasm; membrane. |
| O15260 | SURF4_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| O15264 | MK13_HUMAN | | cytoplasm. |
| O15266 | SHOX_HUMAN | nucleus (by similarity). | nucleus. |
| O15269 | LCB1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| O15270 | LCB2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| O15294 | OGT1_HUMAN | cytoplasm (potential). nucleus (potential). | cytoplasm. |
| O15296 | LX15B_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O15297 | PP2CD_HUMAN | | centriole. cytoplasm. nucleus. |
| O15315 | RA51B_HUMAN | nucleus (probable). | nucleus. |
| O15318 | RPO3G_HUMAN | nucleus (potential). | nucleus. |
| O15320 | CTGE5_HUMAN | | nucleus. |
| O15321 | TM9S1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O15327 | INP4B_HUMAN | | cytoplasm. |
| O15335 | CHAD_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| O15342 | VA0H_HUMAN | membrane; multi-pass membrane protein (potential). | mitochondrion. |
| O15347 | HMGB3_HUMAN | nucleus (by similarity). | nucleus. |
| O15355 | PP2CG_HUMAN | cytoplasm (potential). | centriole. cytoplasm. |
| O15371 | IF37_HUMAN | | cytoplasm. |
| O15372 | IF33_HUMAN | | cytoplasm. |
| O15397 | IPO8_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| O15400 | STX7_HUMAN | endosome; early endosome; early endosomal membrane; single-pass type iv membrane protein (by similarity). | membrane. |
| O15409 | FOXP2_HUMAN | nucleus (probable). | nucleus. |
| O15428 | PINL_HUMAN | | centriole. cytoplasm. |
| O15431 | COPT1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O15432 | COPT2_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |

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| O15442 | MPPD1_HUMAN | | centriole. cytoplasm. |
| O15453 | NBR2_HUMAN | | centriole. cytoplasm. |
| O15455 | TLR3_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| O15457 | MSH4_HUMAN | | nucleus. |
| O15466 | SIA8E_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |
| O15479 | MAGB2_HUMAN | | nucleus. |
| O15480 | MAGB3_HUMAN | | nucleus. |
| O15481 | MAGB4_HUMAN | | nucleus. |
| O15482 | TEX28_HUMAN | | membrane. |
| O15484 | CAN5_HUMAN | | cytoplasm. lysosome. |
| O15488 | GLYG2_HUMAN | | cytoplasm. |
| O15492 | RGS16_HUMAN | | cytoplasm. |
| O15499 | GSCL_HUMAN | nucleus (by similarity). | nucleus. |
| O15503 | INSI1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| O15511 | ARPC5_HUMAN | | cytoplasm. |
| O15516 | CLOCK_HUMAN | nucleus (potential). | nucleus. |
| O15519 | CFLAR_HUMAN | | cytoplasm. |
| O15520 | FGF10_HUMAN | secreted protein (potential). | secreted protein. |
| O15522 | NKX28_HUMAN | nucleus (probable). | nucleus. |
| O15524 | SOCS1_HUMAN | | cytoplasm. |
| O15533 | TPSN_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (probable). | endoplasmic reticulum. |
| O15534 | PER1_HUMAN | nucleus (by similarity). | nucleus. |
| O15535 | ZN193_HUMAN | nucleus (potential). | nucleus. |
| O15539 | RGS5_HUMAN | | cytoplasm. |
| O15541 | R113A_HUMAN | | nucleus. |
| O15544 | GR6_HUMAN | | membrane. |
| O15550 | UTX_HUMAN | nucleus (potential). | centriole. nucleus. |
| O15551 | CLD3_HUMAN | cell membrane; cell-cell junction; tight junction; multi-pass membrane protein (by similarity). | membrane. |
| O42043 | ENK7_HUMAN | | endosome; membrane. |
| O43155 | FLRT2_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| O43156 | K0406_HUMAN | | cytoplasm. |
| O43166 | SI1L1_HUMAN | | cytoplasm. |
| O43167 | ZBT24_HUMAN | nucleus (potential). | nucleus. |
| O43169 | CYB5B_HUMAN | mitochondrion; mitochondrial outer membrane (by similarity). | mitochondrion. |
| O43172 | PRP4_HUMAN | nucleus; nucleoplasm; nuclear speckle (by similarity). colocalizes with spliceosomal snrnps (by similarity). | nucleus. |
| O43173 | SIA8C_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |
| O43175 | SERA_HUMAN | | cytoplasm. |
| O43186 | CRX_HUMAN | nucleus (by similarity). | nucleus. |
| O43187 | IRAK2_HUMAN | | cytoplasm. |
| O43189 | PHF1_HUMAN | nucleus (potential). | nucleus. |
| O43196 | MSH5_HUMAN | | nucleus. |
| O43236 | SEPT4_HUMAN | | cytoplasm. |
| O43237 | DC1L2_HUMAN | | cytoplasm. |
| O43240 | KLK10_HUMAN | secreted protein (probable). | secreted protein. |

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|--------|-------------|---|---------------------------------|
| O43242 | PSD3_HUMAN | | centriole. cytoplasm. endosome. |
| O43246 | CTR4_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O43247 | EAN57_HUMAN | | cytoplasm. |
| O43251 | RBM9_HUMAN | | nucleus. |
| O43252 | PAPS1_HUMAN | | cytoplasm. mitochondrion. |
| O43255 | SIAH2_HUMAN | cytoplasm. predominantly cytoplasmic (probable). nucleus (probable). partially nuclear (probable). | cytoplasm. nucleus. |
| O43257 | ZNHI1_HUMAN | | cytoplasm. |
| O43261 | LEU1_HUMAN | | cytoplasm. nucleus. |
| O43262 | LEU2_HUMAN | | cytoplasm. |
| O43280 | TREA_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | lysosome. |
| O43281 | EFS_HUMAN | | cytoplasm. |
| O43291 | SPIT2_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome. secreted protein. |
| O43293 | DAPK3_HUMAN | nucleus. cytoplasm (by similarity). relocates to the cytoplasm on binding pawr where the complex appears to interact with actin filaments (by similarity). associates to centromeres from prophase to anaphase. | nucleus. |
| O43295 | SRGP2_HUMAN | | cytoplasm. |
| O43296 | ZN264_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O43298 | Z297B_HUMAN | nucleus (probable). | nucleus. |
| O43300 | LRTM2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O43301 | HS12A_HUMAN | | cytoplasm. |
| O43309 | ZNF96_HUMAN | nucleus (potential). | nucleus. |
| O43310 | K0427_HUMAN | | cytoplasm. |
| O43312 | MTSS1_HUMAN | | cytoplasm. |
| O43313 | K0431_HUMAN | | nucleus. |
| O43318 | M3K7_HUMAN | | cytoplasm. |
| O43320 | FGF16_HUMAN | secreted protein (by similarity). | secreted protein. |
| O43323 | DHH_HUMAN | the c-terminal peptide diffuses from the cell, while the n-terminal peptide remains associated with the cell surface. is also secreted in either cleaved or uncleaved form to mediate signaling to other cells (by similarity). | secreted protein. |
| O43325 | YG01_HUMAN | | nucleus. |
| O43345 | ZN208_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O43347 | MS11H_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| O43353 | RIPK2_HUMAN | cytoplasm (probable). | cytoplasm. |
| O43399 | TPD54_HUMAN | | cytoplasm. |
| O43402 | CX4NB_HUMAN | | cytoplasm. |
| O43422 | P52K_HUMAN | | nucleus. |
| O43423 | AN32C_HUMAN | | cytoplasm. |
| O43424 | GRID2_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| O43426 | SYNJ1_HUMAN | cytoplasm (by similarity). | nucleus. |
| O43432 | IF4G3_HUMAN | | cytoplasm. |
| O43435 | TBX1_HUMAN | nucleus (potential). | nucleus. |
| O43439 | MTG8R_HUMAN | nucleus (probable). | nucleus. |

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|--------|-------------|--|--|
| O43448 | KCAB3_HUMAN | cytoplasm (potential). | membrane. |
| O43462 | MBTP2_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O43482 | OIP5_HUMAN | | cytoplasm. |
| O43488 | ARK72_HUMAN | Golgi apparatus (by similarity). | cytoplasm. |
| O43490 | PROM1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O43491 | E41L2_HUMAN | | cytoplasm. |
| O43502 | RA51C_HUMAN | nucleus (probable). | nucleus. |
| O43511 | S26A4_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O43516 | WASIP_HUMAN | cytoplasmic vesicle (by similarity). cytoskeleton (by similarity). vesicle surfaces and along actin tails. co-localized with actin stress fibers. when co-expressed with wasl, no longer associated with actin filaments but accumulated in perinuclear and cortical areas like wasl (by similarity). | cytoplasm. |
| O43521 | BIM_HUMAN | membrane; peripheral membrane protein. associated with intracytoplasmic membranes (by similarity). | cytoplasm. |
| O43529 | CHSTA_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | Golgi. |
| O43531 | RG141_HUMAN | | cytoplasm. |
| O43541 | SMAD6_HUMAN | | nucleus. |
| O43542 | XRCC3_HUMAN | nucleus (probable). | nucleus. |
| O43556 | SGCE_HUMAN | sarcolemma; sarcolemmal membrane; single-pass type i membrane protein (potential). | membrane. |
| O43566 | RGS14_HUMAN | | cytoplasm. |
| O43567 | RNF13_HUMAN | nucleus (by similarity). | cytoplasm. |
| O43581 | SYT7_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein (by similarity). | membrane. |
| O43583 | DENR_HUMAN | | centriole. cytoplasm. |
| O43586 | PPIP1_HUMAN | cytoplasm (by similarity). cytoskeleton (by similarity). colocalized with the cortical actin cytoskeleton during interphase, lamellipodia and actin-rich cytokinetic cleavage furrow. colocalized with was to filamentous structures within the cytoplasm. colocalized with ptpn12 in the cytoplasm and the perinuclear region. colocalized with cd2ap and was in the actin cytoskeleton. colocalized with cd2, cd2ap and was at the site of t-cell:apc contact (by similarity). | cytoplasm. |
| O43598 | RCL_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| O43612 | OREX_HUMAN | endoplasmic reticulum; rough endoplasmic reticulum (by similarity). associated with perikaryal rough endoplasmic reticulum as well as cytoplasmic large granular vesicles at synapses (by similarity). | endoplasmic reticulum. secreted protein. |
| O43617 | TPPC3_HUMAN | Golgi apparatus; cis-Golgi network (by similarity). | endoplasmic reticulum. |
| O43623 | SLUG_HUMAN | nucleus (probable). | nucleus. |

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| O43633 | CHM2A_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O43638 | FREA_HUMAN | nucleus (potential). | nucleus. |
| O43665 | RGS10_HUMAN | | cytoplasm. |
| O43670 | ZN207_HUMAN | nucleus (probable). | nucleus. |
| O43681 | ARSA1_HUMAN | | membrane. |
| O43683 | BUB1_HUMAN | nucleus (by similarity). nuclear in interphase cells. kinetochore localization is required for normal mitotic timing and checkpoint response to spindle damage | nucleus. |
| O43704 | ST1B1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O43708 | MAAI_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O43709 | WBS22_HUMAN | nucleus (potential). | nucleus. |
| O43711 | TLX3_HUMAN | nucleus (probable). | nucleus. |
| O43716 | 15E2_HUMAN | | centriole. cytoplasm. |
| O43734 | CIKS_HUMAN | | cytoplasm. |
| O43736 | ITM2A_HUMAN | membrane; single-pass type ii membrane protein (potential). | endosome; membrane. |
| O43739 | CYH3_HUMAN | | cytoplasm. |
| O43741 | AAKB2_HUMAN | | cytoplasm. mitochondrion. |
| O43745 | CHP2_HUMAN | | cytoplasm. |
| O43752 | STX6_HUMAN | Golgi apparatus; Golgi membrane; single-pass type iv membrane protein (potential). | Golgi. |
| O43763 | TLX2_HUMAN | nucleus (probable). | nucleus. |
| O43765 | SGTA_HUMAN | | cytoplasm. |
| O43766 | LIAS_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| O43768 | ENSA_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O43781 | DYRK3_HUMAN | | cytoplasm. |
| O43790 | KRHB6_HUMAN | | cytoplasm. |
| O43795 | MYO1B_HUMAN | | cytoskeleton. |
| O43815 | STRN_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| O43820 | HYAL3_HUMAN | secreted protein (by similarity). lysosome (by similarity). | lysosome. |
| O43822 | CU002_HUMAN | | cytoplasm. |
| O43825 | B3GT2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | Golgi. |
| O43826 | G6PT1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | membrane. |
| O43827 | ANGL7_HUMAN | secreted protein (probable). | cytoplasm. |
| O43830 | ZNF73_HUMAN | nucleus (probable). | nucleus. |
| O43847 | NRDC_HUMAN | | membrane. |
| O43865 | SAHH2_HUMAN | | cytoplasm. |
| O43896 | KIF1C_HUMAN | | cytoplasm. nucleus. |
| O43897 | TLL1_HUMAN | secreted protein (probable). | secreted protein. |
| O43900 | LMO6_HUMAN | | cytoplasm. |
| O43903 | GAS2_HUMAN | cytoskeleton (by similarity). membrane; peripheral membrane protein (by similarity). component of the microfilament system. colocalizes with actin fibers at the cell border and along the stress fibers in growth-arrested fibroblasts. mainly membrane-associated. when hyperphosphorylated, accumulates at membrane ruffles (by similarity). | cytoplasm. |

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|--------|-------------|---|------------------------------|
| O43916 | CHST1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| O43921 | EFNA2_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | secreted protein. |
| O43924 | PDE6D_HUMAN | | centriole. |
| O43930 | PRKY_HUMAN | | cytoplasm. |
| O60218 | AK1BA_HUMAN | cytoplasm (probable). | cytoplasm. |
| O60224 | SSX4_HUMAN | | nucleus. |
| O60225 | SSX5_HUMAN | | nucleus. |
| O60229 | HAPIP_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O60232 | SSA27_HUMAN | | cytoplasm. |
| O60234 | GMFG_HUMAN | | cytoplasm. |
| O60240 | PLIN_HUMAN | lipid droplet surface-associated (by similarity). | cytoplasm. |
| O60243 | H6ST1_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| O60248 | SOX15_HUMAN | nucleus (potential). | nucleus. |
| O60256 | KPRB_HUMAN | | cytoplasm. secreted protein. |
| O60262 | GBG7_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| O60268 | K0513_HUMAN | | nucleus. |
| O60269 | GRIN2_HUMAN | | nucleus. |
| O60281 | ZN292_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O60282 | KIF5C_HUMAN | | cytoplasm. nucleus. |
| O60285 | NUAK1_HUMAN | | cytoplasm. |
| O60291 | MGRN1_HUMAN | | cytoplasm. |
| O60292 | SI1L3_HUMAN | | cytoplasm. |
| O60294 | LCMT2_HUMAN | | cytoplasm. |
| O60296 | TRAK2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O60299 | K0552_HUMAN | | cytoplasm. |
| O60304 | ZN500_HUMAN | nucleus (potential). | nucleus. |
| O60307 | MAST3_HUMAN | | cytoplasm. |
| O60308 | K0562_HUMAN | | cytoplasm. |
| O60309 | LR37A_HUMAN | | cytoplasm. |
| O60312 | AT10A_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| O60318 | MCM3A_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| O60330 | PCDGC_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| O60333 | KIF1B_HUMAN | cytoplasmic vesicle (by similarity). predominantly found on the vesicle- and tubule-like structures within the cytoplasm (by similarity). | cytoplasm. |
| O60343 | TBCD4_HUMAN | | cytoplasm. |
| O60346 | PHLPP_HUMAN | cytoplasm (probable). membrane; peripheral membrane protein (by similarity). may be membrane-associated | Golgi. |
| O60347 | TBC12_HUMAN | | cytoplasm. |
| O60361 | NDK8_HUMAN | | cytoplasm. |
| O60383 | GDF9_HUMAN | secreted protein (by similarity). | secreted protein. |
| O60391 | NMD3B_HUMAN | cell membrane; multi-pass membrane protein (by similarity). requires the presence of grin1 to be targeted at the plasma membrane (by similarity). | membrane. |

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| O60397 | COX7S_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| O60443 | DFNA5_HUMAN | | centriole. cytoplasm. secreted protein. membrane. |
| O60449 | LY75_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| O60469 | DSCAM_HUMAN | isoform long: cell membrane; single-pass type i membrane protein (potential). isoform short: secreted protein (probable). the short isoform may be secreted. | membrane. |
| O60477 | FAM5A_HUMAN | secreted protein (potential). | cytoplasm. |
| O60478 | G137B_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O60479 | DLX3_HUMAN | nucleus (potential). | nucleus. |
| O60486 | PLXC1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O60487 | EVA1_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| O60488 | ACSL4_HUMAN | mitochondrion; mitochondrial outer membrane; single-pass type iii membrane protein (by similarity). peroxisome; peroxisomal membrane; single-pass type iii membrane protein (by similarity). microsome; microsomal membrane; single-pass type iii membrane protein (by similarity). | endoplasmic reticulum. peroxisome. |
| O60493 | SNX3_HUMAN | | cytoplasm. |
| O60494 | CUBN_HUMAN | endosome; endosomal membrane; peripheral membrane protein (by similarity). lysosome; lysosomal membrane; peripheral membrane protein (by similarity). colocalizes with amn and lrp2 in the endocytotic apparatus of epithelial cells (by similarity). | membrane. |
| O60496 | DOK2_HUMAN | | cytoplasm. |
| O60499 | STX10_HUMAN | Golgi apparatus; Golgi membrane; single-pass type iv membrane protein (potential). | endosome. Golgi. |
| O60500 | NPHN_HUMAN | membrane; single-pass type i membrane protein (potential). predominantly located at podocyte slit diaphragm between podocyte foot processes. also associated with podocyte apical plasma membrane. | membrane. |
| O60504 | VINEX_HUMAN | isoform alpha, isoform beta: cell membrane; cell-matrix junction (by similarity). isoform beta: nucleus. localized at cell-extracellular matrix junctions (by similarity). both isoforms were localized at focal adhesion and cell-cell adhesion sites, vinexin beta was also found in the nucleus. | cytoskeleton. |
| O60506 | HNRPQ_HUMAN | nucleus (by similarity). isoform 1, isoform 2, isoform 3: nucleus; nucleoplasm (by similarity). isoform 1, isoform 2 and isoform 3 are expressed predominantly in the nucleoplasm. cytoplasm (by similarity). microsome (by similarity). the tyrosine phosphorylated form bound to rna is found in microsomes (by similarity). | nucleus. |

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| O60507 | TPS1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | Golgi; membrane. |
| O60516 | 4EBP3_HUMAN | | cytoplasm. |
| O60518 | RNBP6_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. nucleus. |
| O60524 | SDCG1_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| O60543 | CIDEA_HUMAN | | cytoplasm. |
| O60547 | GMDS_HUMAN | | centriole. cytoplasm. |
| O60548 | FOXD2_HUMAN | nucleus (by similarity). | nucleus. |
| O60565 | GREM1_HUMAN | secreted protein (probable). | secreted protein. |
| O60573 | IF4E3_HUMAN | | cytoplasm. |
| O60575 | ISK4_HUMAN | secreted protein (by similarity). | secreted protein. |
| O60584 | EDDF1_HUMAN | | secreted protein. |
| O60602 | TLR5_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| O60603 | TLR2_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| O60610 | DIAP1_HUMAN | | cytoplasm. |
| O60635 | TSN1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O60636 | TSN2_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O60637 | TSN3_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| O60641 | AP180_HUMAN | cell membrane; coated pit; cytoplasmic side (by similarity). component of the coat surrounding the cytoplasmic face of coated vesicles in the plasma membrane (by similarity). | Golgi. |
| O60645 | EXOC3_HUMAN | | cytoplasm. |
| O60658 | PDE8A_HUMAN | | cytoplasm. |
| O60663 | LMX1B_HUMAN | nucleus (potential). | nucleus. |
| O60673 | DPOLZ_HUMAN | nucleus (potential). | nucleus. |
| O60674 | JAK2_HUMAN | intracytoplasmic membrane; peripheral membrane protein. wholly intracellular, possibly membrane associated (by similarity). | cytoplasm. |
| O60683 | PEX10_HUMAN | peroxisome; peroxisomal membrane; peripheral membrane protein (potential). | peroxisome. |
| O60684 | IMA7_HUMAN | | cytoplasm. |
| O60688 | YPEL1_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| O60701 | UGDH_HUMAN | | mitochondrion. |
| O60704 | TPS2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. |
| O60706 | ABCC9_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O60729 | CC14B_HUMAN | nucleus; nucleolus (by similarity). nucleolar during interphase. | cytoplasm. |
| O60732 | MAGC1_HUMAN | | nucleus. |
| O60739 | EIF1B_HUMAN | | cytoplasm. |
| O60749 | SNX2_HUMAN | | cytoplasm. Golgi. |
| O60756 | BCE1_HUMAN | | membrane. |
| O60765 | TCF17_HUMAN | nucleus (by similarity). | nucleus. |
| O60774 | FMO6_HUMAN | microsome (by similarity). | microsome. |

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|--------|-------------|--|------------------------------|
| O60784 | TOM1_HUMAN | cytoplasm (probable). | cytoplasm. endosome. |
| O60806 | TBX19_HUMAN | nucleus (potential). | nucleus. |
| O60810 | PRAM4_HUMAN | | cytoplasm. |
| O60811 | PRAM2_HUMAN | | cytoplasm. |
| O60812 | HNRCL_HUMAN | nucleus (by similarity). component of ribonucleosomes (by similarity). | nucleus. |
| O60825 | F262_HUMAN | | cytoplasm. |
| O60826 | CCD22_HUMAN | | cytoplasm. |
| O60829 | GAGC1_HUMAN | | centriole. cytoplasm. |
| O60831 | PRAF2_HUMAN | membrane; multi-pass membrane protein (potential). | Golgi. |
| O60841 | IF2P_HUMAN | | cytoplasm. |
| O60844 | ZG16_HUMAN | secreted protein. stored in zymogen granules (by similarity). | secreted protein. |
| O60858 | RFP2_HUMAN | | cytoplasm. |
| O60861 | GAS7_HUMAN | cytoplasm (by similarity). | membrane. |
| O60880 | SH21A_HUMAN | cytoplasm (potential). | cytoplasm. |
| O60882 | MMP20_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| O60884 | DNJA2_HUMAN | or: membrane; lipid-anchor (potential). or: nucleus. | cytoplasm. |
| O60885 | BRD4_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| O60888 | CUTA_HUMAN | | cytoplasm. Golgi. |
| O60890 | OPHN1_HUMAN | | cytoplasm. |
| O60907 | TBL1X_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O60908 | YG02_HUMAN | | centriole. cytoplasm. |
| O60911 | CATL2_HUMAN | lysosome (potential). | lysosome. |
| O60925 | PFD1_HUMAN | | cytoplasm. |
| O60927 | PP1RB_HUMAN | | cytoplasm. secreted protein. |
| O60930 | RNH1_HUMAN | cytoplasm (potential). | secreted protein. |
| O60934 | NBN_HUMAN | nucleus (by similarity). localizes to discrete nuclear foci after treatment with genotoxic agents (by similarity). | nucleus. |
| O60938 | KERA_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | centriole. secreted protein. |
| O60942 | MCE1_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| O71037 | ENK3_HUMAN | | endosome; membrane. |
| O75027 | ABCB7_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (potential). | mitochondrion; membrane. |
| O75031 | HSF2B_HUMAN | | cytoplasm. |
| O75037 | KI21B_HUMAN | cytoplasm (by similarity). | cytoplasm. nucleus. |
| O75044 | FNBP2_HUMAN | | cytoplasm. |
| O75051 | PLXA2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O75052 | CAPON_HUMAN | | cytoplasm. |
| O75061 | AUXI_HUMAN | | cytoplasm. |
| O75063 | FA20B_HUMAN | secreted protein (potential). | membrane. |
| O75069 | TMCC2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| O75071 | K0494_HUMAN | | cytoplasm. |
| O75072 | FCMD_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| O75074 | LRP3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |

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|--------|-------------|---|---------------------------|
| O75077 | ADA23_HUMAN | cell membrane; single-pass type i membrane protein (potential). isoform gamma: secreted protein. | membrane. |
| O75083 | WDR1_HUMAN | | cytoplasm. |
| O75093 | SLIT1_HUMAN | secreted protein (by similarity). | secreted protein. |
| O75094 | SLIT3_HUMAN | secreted protein (by similarity). | cytoplasm. |
| O75095 | MEGF6_HUMAN | | cytoplasm. |
| O75096 | LRP4_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O75106 | AOC2_HUMAN | | secreted protein. |
| O75110 | ATP9A_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| O75116 | ROCK2_HUMAN | cytoplasm (by similarity). cell membrane; peripheral membrane protein (by similarity). cytoplasmic, and associated with actin microfilaments and the plasma membrane (by similarity). | cytoplasm. |
| O75121 | MFA3L_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| O75123 | ZN623_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| O75127 | PTCD1_HUMAN | | cytoplasm. |
| O75131 | CPNE3_HUMAN | | cytoplasm. |
| O75132 | ZBED4_HUMAN | | nucleus. |
| O75140 | DEPD5_HUMAN | | cytoplasm. |
| O75143 | K0652_HUMAN | | membrane. |
| O75144 | ICOSL_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| O75145 | LIPA3_HUMAN | cytoplasm (by similarity). cell surface (by similarity). colocalizes with ptpfr at the cell surface (by similarity). | cytoplasm. |
| O75146 | HIP1R_HUMAN | cytoplasm; perinuclear region (by similarity). intracytoplasmic membrane (by similarity). membrane- associated protein, mainly localized at the endocytic compartments and in the perinuclear | cytoplasm. |
| O75147 | OBSL1_HUMAN | | nucleus. |
| O75151 | PHF2_HUMAN | nucleus (potential). | nucleus. |
| O75152 | ZC11A_HUMAN | | cytoplasm. nucleus. |
| O75153 | IF3X_HUMAN | | cytoplasm. |
| O75155 | CAND2_HUMAN | nucleus (by similarity). | nucleus. |
| O75157 | T22D2_HUMAN | | nucleus. |
| O75159 | SOCS5_HUMAN | | cytoplasm. |
| O75161 | NPHP4_HUMAN | | cytoplasm. |
| O75165 | DNJCD_HUMAN | | cytoplasm. |
| O75167 | PHAR2_HUMAN | | cytoplasm. |
| O75170 | SAPS2_HUMAN | | cytoplasm. |
| O75173 | ATS4_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| O75175 | CNOT3_HUMAN | cytoplasm (probable). nucleus (probable). | nucleus. |
| O75177 | S18L1_HUMAN | | cytoplasm. |
| O75182 | SIN3B_HUMAN | nucleus (by similarity). | nucleus. |
| O75185 | AT2C2_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| O75190 | DNJB6_HUMAN | | cytoplasm. |
| O75191 | XYLB_HUMAN | | cytoplasm. mitochondrion. |

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| O75197 | LRP5_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O75208 | COQ9_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| O75223 | CG024_HUMAN | | cytoplasm. |
| O75251 | NUKM_HUMAN | | mitochondrion. |
| O75293 | GA45B_HUMAN | | secreted protein. |
| O75298 | RTN2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| O75309 | CAD16_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| O75317 | UBP12_HUMAN | | nucleus. |
| O75323 | NIPS2_HUMAN | | mitochondrion. |
| O75324 | SNN_HUMAN | | endosome; membrane. |
| O75325 | LRRN5_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O75330 | HMMR_HUMAN | cell surface. cytoplasm (by similarity). | cytoplasm. |
| O75333 | TBX10_HUMAN | nucleus (potential). | nucleus. |
| O75335 | LIPA4_HUMAN | cytoplasm (by similarity). cell surface (by similarity). colocalizes with ptpfr at the cell surface (by similarity). | cytoplasm. |
| O75340 | PDCD6_HUMAN | | cytoplasm. |
| O75342 | LX12B_HUMAN | | cytoplasm. |
| O75344 | FKBP6_HUMAN | | cytoplasm. |
| O75346 | ZN253_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O75347 | TBCA_HUMAN | | cytoplasm. |
| O75348 | VATG1_HUMAN | | mitochondrion. |
| O75352 | MPU1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| O75355 | ENP3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. |
| O75356 | ENP5_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| O75360 | PROP1_HUMAN | nucleus (by similarity). | nucleus. |
| O75362 | ZN217_HUMAN | nucleus (probable). | nucleus. |
| O75364 | PITX3_HUMAN | nucleus (by similarity). | nucleus. |
| O75366 | AVIL_HUMAN | | cytoplasm. |
| O75368 | SH3L1_HUMAN | | centriole. cytoplasm. nucleus. |
| O75376 | NCOR1_HUMAN | nucleus (by similarity). | nucleus. |
| O75379 | VAMP4_HUMAN | Golgi apparatus; trans-Golgi network; trans- Golgi network membrane; single-pass type iv membrane protein (probable). associated with trans Golgi network (tgn) and newly formed immature secretory granules (isg). not found on the mature secretory organelles. | endosome. lysosome. |
| O75382 | TRIM3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O75385 | ULK1_HUMAN | | cytoplasm. |
| O75386 | TULP3_HUMAN | | cytoplasm. nucleus. |
| O75387 | LAT3_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O75391 | SPAG7_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| O75396 | SC22B_HUMAN | membrane; single-pass type iv membrane protein (probable). | endoplasmic reticulum. |
| O75400 | PRP40_HUMAN | nucleus; nucleoplasm; nuclear speckle (by similarity). | nucleus. |

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|--------|-------------|--|--|
| O75414 | NDK6_HUMAN | | cytoplasm. |
| O75417 | DPOLQ_HUMAN | | nucleus. |
| O75420 | PERQ1_HUMAN | | cytoplasm. |
| O75425 | MSPD3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| O75426 | FBX24_HUMAN | | cytoplasm. nucleus. |
| O75427 | LRCH4_HUMAN | | cytoplasm. |
| O75437 | ZN254_HUMAN | nucleus (probable). | nucleus. |
| O75439 | MPPB_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | cytoplasm. mitochondrion; membrane. |
| O75443 | TECTA_HUMAN | cell membrane; lipid-anchor; gpi-anchor; extracellular side (probable). found in the non-collagenous matrix of the tectorial membrane (by similarity). | secreted protein. |
| O75444 | MAF_HUMAN | nucleus (by similarity). | nucleus. |
| O75459 | GAGB1_HUMAN | | secreted protein. |
| O75467 | ZN324_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O75469 | PXR_HUMAN | nucleus (by similarity). | nucleus. |
| O75474 | FRAT2_HUMAN | | secreted protein. |
| O75477 | SPFH1_HUMAN | membrane; peripheral membrane protein (probable). | cytoplasm. Golgi. |
| O75487 | GPC4_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| O75493 | CAH11_HUMAN | secreted protein (potential). | cytoplasm. |
| O75496 | GEMI_HUMAN | | cytoplasm. |
| O75509 | TNR21_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| O75553 | DAB1_HUMAN | | membrane. |
| O75556 | SG2A1_HUMAN | | cytoplasm. |
| O75558 | STX11_HUMAN | membrane; peripheral membrane protein (potential). | Golgi. |
| O75562 | HFS1_HUMAN | | membrane. |
| O75564 | JERKY_HUMAN | nucleus (potential). | nucleus. |
| O75569 | PRKRA_HUMAN | | cytoplasm. microsome. nucleus. |
| O75575 | CRCP_HUMAN | | cytoplasm. |
| O75581 | LRP6_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O75594 | PGRP_HUMAN | secreted protein; cytoplasmic granule (by similarity). | secreted protein. |
| O75596 | CLC3A_HUMAN | | secreted protein. |
| O75603 | GCM2_HUMAN | nucleus (by similarity). | nucleus. |
| O75604 | UBP2_HUMAN | cytoplasm (by similarity). | nucleus. |
| O75608 | LYPA1_HUMAN | cytoplasm (by similarity). | cytoplasm. cytoskeleton. mitochondrion. membrane. |
| O75616 | ERAL_HUMAN | | cytoplasm. |
| O75618 | DEDD_HUMAN | cytoplasm. translocated to the nucleus during cd95-mediated apoptosis where it is localized in the nucleoli (by similarity). following apoptosis induction, the mono and/or diubiquitination form increases and forms filamentous structures that colocalize with krt8 and krt18 intermediate filament network in simple epithelial cells. | nucleus. |
| O75626 | PRDM1_HUMAN | nucleus (by similarity). | nucleus. |
| O75628 | REM1_HUMAN | | cytoplasm. |

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|--------|-------------|--|---|
| O75631 | UPK3A_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). heterodimer formation with uroplakin ib is a prerequisite to exit out of the endoplasmic reticulum (er) (by similarity). | endoplasmic reticulum. |
| O75635 | SPB7_HUMAN | cytoplasm (by similarity). | secreted protein. |
| O75638 | CTAG2_HUMAN | | membrane. |
| O75642 | IF1AH_HUMAN | | cytoplasm. |
| O75674 | TM1L1_HUMAN | Golgi apparatus; Golgi stack. endosome; endosomal membrane (potential). | cytoplasm. |
| O75677 | RFPL1_HUMAN | | cytoplasm. |
| O75678 | RFPL2_HUMAN | | cytoplasm. |
| O75679 | RFPL3_HUMAN | | cytoplasm. |
| O75683 | SURF6_HUMAN | nucleus; nucleoplasm (by similarity). granular component of the nucleolus (by similarity). | nucleus. |
| O75688 | PP2CB_HUMAN | | cytoplasm. |
| O75690 | KRA58_HUMAN | | secreted protein. |
| O75694 | NU155_HUMAN | nucleus; nuclear envelope; nuclear pore complex; peripheral membrane protein; cytoplasmic side (by similarity). nucleus; nuclear envelope; nuclear pore complex; peripheral membrane protein; nucleoplasmic side (by similarity). in mitosis, assumes a diffuse cytoplasmic distribution probably as a monomer, before reversing back into a punctate nuclear surface localization at the end of mitosis | nucleus. |
| O75698 | HUG1_HUMAN | | nucleus. |
| O75711 | SCRG1_HUMAN | secreted protein (probable). | secreted protein. |
| O75716 | STK16_HUMAN | membrane; lipid-anchor (probable). | cytoplasm. |
| O75717 | WDHD1_HUMAN | nucleus; nucleoplasm (by similarity). | cytoplasm. nucleus. |
| O75718 | CRTAP_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | centriole. cytoplasm. secreted protein. |
| O75747 | P3C2G_HUMAN | membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| O75762 | TRPA1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O75764 | TCEA3_HUMAN | nucleus (by similarity). | nucleus. |
| O75771 | RA51D_HUMAN | nucleus (probable). | nucleus. |
| O75781 | PALM_HUMAN | membrane; peripheral membrane protein (by similarity). | membrane. |
| O75787 | RENH_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O75791 | GRAP2_HUMAN | | cytoplasm. |
| O75792 | RNH2A_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O75794 | CD123_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O75800 | ZMY10_HUMAN | | cytoplasm. |
| O75815 | BCAR3_HUMAN | | cytoplasm. |
| O75817 | POP7_HUMAN | nucleus; nucleolus (potential). | nucleus. |
| O75818 | RPP40_HUMAN | nucleus; nucleolus (potential). | nucleus. |
| O75820 | ZN189_HUMAN | nucleus (probable). | nucleus. |
| O75821 | IF34_HUMAN | | cytoplasm. |
| O75822 | IF31_HUMAN | | cytoplasm. |
| O75828 | DHC3_HUMAN | cytoplasm (potential). | cytoplasm. |

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|--------|-------------|--|------------------------------|
| O75830 | SPI2_HUMAN | secreted protein (probable). | secreted protein. |
| O75832 | PSD10_HUMAN | | cytoplasm. |
| O75838 | KIP2_HUMAN | | cytoplasm. |
| O75840 | KLF7_HUMAN | nucleus (probable). | nucleus. |
| O75844 | FACE1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein. Golgi apparatus; Golgi membrane; multi-pass membrane protein | membrane. |
| O75845 | SC5D_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). | endoplasmic reticulum. |
| O75865 | TPC6A_HUMAN | Golgi apparatus; cis-Golgi network (by similarity). | endoplasmic reticulum. |
| O75886 | STAM2_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | Golgi. |
| O75896 | TUSC2_HUMAN | | cytoplasm. |
| O75907 | DGAT1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| O75909 | CCNK_HUMAN | | nucleus. |
| O75911 | DHRS3_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| O75914 | PAK3_HUMAN | | cytoplasm. |
| O75915 | PRAF3_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). cytoplasm (by similarity). also exists as a soluble form in the cytoplasm. associated with microtubules (by similarity). | endoplasmic reticulum. |
| O75916 | RGS9_HUMAN | | membrane. |
| O75920 | SERF1_HUMAN | | cytoplasm. secreted protein. |
| O75923 | DYSF_HUMAN | sarcolemma; sarcolemmal membrane; single- pass type ii membrane protein (probable). | membrane. |
| O75937 | DNJC8_HUMAN | | cytoplasm. |
| O75947 | ATP5H_HUMAN | | mitochondrion. |
| O75949 | TMM28_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| O75951 | LYZL6_HUMAN | secreted protein (probable). | lysosome. |
| O75953 | DNJB5_HUMAN | | endoplasmic reticulum. |
| O75954 | TSN9_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O75956 | CDKA2_HUMAN | | nucleus. |
| O75962 | TRIO_HUMAN | | cytoplasm. |
| O75964 | ATP5L_HUMAN | | mitochondrion. |

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|--------|-------------|--|------------------------------|
| O75970 | MPDZ_HUMAN | cell membrane; cytoplasmic side. associated with membranes. colocalizes with htr2c on the apical membrane of epithelial choroid plexus cells. highly enriched in postsynaptic densities (psd). localized to punctae on dendrites of hippocampal neurons and colocalizes with the synaptic marker dlg4. localized mainly in the schmidt-lanterman incisures of myelinating schwann cells (by similarity). in the retina, localizes to the sub-apical region adjacent to the adherens junction complex at the outer limiting membrane. enriched at the tight junctions of epithelial cells. association to the tight junctions depends on cxadr. | cytoplasm. |
| O75976 | CBPD_HUMAN | membrane; single-pass type i membrane protein (potential). | secreted protein; membrane. |
| O75995 | SLY_HUMAN | | nucleus. |
| O76009 | K1HA_HUMAN | | cytoplasm. |
| O76011 | K1H4_HUMAN | | cytoplasm. |
| O76013 | K1H6_HUMAN | | cytoplasm. |
| O76014 | K1H7_HUMAN | | cytoplasm. |
| O76015 | K1H8_HUMAN | | cytoplasm. |
| O76027 | ANXA9_HUMAN | | cytoplasm. |
| O76036 | NCTR1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O76039 | CDKL5_HUMAN | | cytoplasm. |
| O76041 | NEBL_HUMAN | | cytoplasm. |
| O76042 | CC051_HUMAN | | cytoplasm. secreted protein. |
| O76061 | STC2_HUMAN | secreted protein (potential). | secreted protein. |
| O76074 | PDE5A_HUMAN | | cytoplasm. |
| O76076 | WISP2_HUMAN | secreted protein (probable). | secreted protein. |
| O76080 | ZFAN5_HUMAN | | cytoplasm. |
| O76083 | PDE9A_HUMAN | | cytoplasm. |
| O76087 | GAGE7_HUMAN | | centriole. cytoplasm. |
| O76093 | FGF18_HUMAN | secreted protein (by similarity). | secreted protein. |
| O76095 | JTB_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O77932 | DOM3Z_HUMAN | | cytoplasm. |
| O94760 | DDAH1_HUMAN | | cytoplasm. |
| O94772 | LY6H_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| O94776 | MTA2_HUMAN | nucleus (by similarity). | nucleus. |
| O94779 | CNTN5_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| O94804 | STK10_HUMAN | | cytoplasm. |
| O94806 | KPCD3_HUMAN | | cytoplasm. |
| O94808 | GFPT2_HUMAN | | cytoplasm. |
| O94810 | RGS11_HUMAN | | cytoplasm. |
| O94811 | P25A_HUMAN | | nucleus. |
| O94812 | BAIP3_HUMAN | | synapse. |
| O94817 | ATG12_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O94819 | KBTBB_HUMAN | | cytoplasm. |
| O94822 | ZN294_HUMAN | | cytoplasm. |

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|--------|-------------|--|---------------------|
| O94826 | TOM70_HUMAN | mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| O94832 | MYO1D_HUMAN | | cytoskeleton. |
| O94833 | BPAEA_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O94842 | CN092_HUMAN | nucleus (probable). | nucleus. |
| O94844 | RHBT1_HUMAN | | cytoplasm. |
| O94850 | DEND_HUMAN | | nucleus. |
| O94851 | MICA2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O94855 | SC24D_HUMAN | cytoplasm (by similarity). cytoplasm; perinuclear region (by similarity). | cytoplasm. |
| O94874 | K0776_HUMAN | | nucleus. |
| O94876 | TMCC1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| O94880 | PHF14_HUMAN | | nucleus. |
| O94885 | SASH1_HUMAN | | cytoplasm. |
| O94887 | FARP2_HUMAN | | cytoplasm. |
| O94888 | UBXD7_HUMAN | | nucleus. |
| O94889 | KLH18_HUMAN | | cytoplasm. |
| O94892 | ZN432_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| O94900 | TOX_HUMAN | nucleus (by similarity). | nucleus. |
| O94901 | UN84A_HUMAN | nucleus; nuclear membrane; multi-pass membrane protein (potential). membrane; perinuclear region; multi-pass membrane protein (potential). | endosome. |
| O94903 | PROSC_HUMAN | | cytoplasm. |
| O94905 | SPFH2_HUMAN | membrane; peripheral membrane protein (potential). | secreted protein. |
| O94919 | ENDD1_HUMAN | secreted protein (potential). | secreted protein. |
| O94923 | GLCE_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endosome. |
| O94925 | GLSK_HUMAN | | mitochondrion. |
| O94927 | K0841_HUMAN | | cytoplasm. |
| O94929 | ABLM3_HUMAN | cytoplasm (probable). | cytoplasm. |
| O94933 | SLIK3_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| O94941 | RNF37_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| O94952 | FBX21_HUMAN | | cytoplasm. |
| O94955 | RHBT3_HUMAN | | cytoplasm. |
| O94964 | K0889_HUMAN | | nucleus. |
| O94967 | WDR47_HUMAN | | cytoplasm. |
| O94973 | AP2A2_HUMAN | cell membrane; coated pit; cytoplasmic side (by similarity). component of the coat surrounding the cytoplasmic face of coated vesicles in the plasma membrane (by similarity). | Golgi. |
| O94983 | CMTA2_HUMAN | nucleus (probable). | nucleus. |
| O94985 | CSTN1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O94988 | FA13A_HUMAN | | nucleus. |
| O94989 | ARHGF_HUMAN | | cytoplasm. |
| O94991 | SLIK5_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| O95025 | SEM3D_HUMAN | secreted protein (by similarity). | membrane. |
| O95045 | UPP2_HUMAN | | cytoplasm. |

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|--------|-------------|--|------------------------------|
| O95049 | ZO3_HUMAN | | cytoplasm. |
| O95050 | INMT_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O95057 | DIRA1_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| O95059 | RPP14_HUMAN | nucleus (potential). | nucleus. |
| O95067 | CCNB2_HUMAN | | nucleus. |
| O95069 | KCNK2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O95072 | REC8L_HUMAN | nucleus (by similarity). localized along chromosome axis, but not chromosome ends (by similarity). | centriole. |
| O95073 | FSBP_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| O95076 | ALX3_HUMAN | nucleus (by similarity). | nucleus. |
| O95081 | HRBL_HUMAN | | cytoplasm. |
| O95084 | PRS23_HUMAN | secreted protein (potential). | secreted protein. |
| O95096 | NKX22_HUMAN | nucleus (by similarity). | nucleus. |
| O95104 | SFR15_HUMAN | nucleus (by similarity). | nucleus. |
| O95147 | DUS14_HUMAN | | cytoplasm. |
| O95150 | TNF15_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| O95155 | UBE4B_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| O95156 | NXPH2_HUMAN | secreted protein (potential). | secreted protein. |
| O95157 | NXPH3_HUMAN | secreted protein (potential). | secreted protein. |
| O95158 | NXPH4_HUMAN | secreted protein (potential). | secreted protein. |
| O95159 | ZFPL1_HUMAN | | cytoplasm. nucleus. |
| O95164 | UBL3_HUMAN | cell membrane; lipid-anchor (by similarity). | cytoplasm. |
| O95166 | GBRAP_HUMAN | intracytoplasmic membrane (by similarity). cytoskeleton (by similarity). largely associated with intracellular membrane structures including the Golgi apparatus and post-synaptic cisternae. colocalizes with microtubules (by similarity). | cytoplasm. Golgi. |
| O95177 | CP003_HUMAN | | centriole. cytoplasm. |
| O95183 | VAMP5_HUMAN | cell membrane; single-pass type iv membrane protein (probable). intracytoplasmic membrane; single-pass type iv membrane protein (probable). Golgi apparatus; trans-Golgi network; trans-Golgi network membrane; single-pass type iv membrane protein (probable). associated with the plasma membrane as well as intracellular perinuclear and peripheral vesicular structures of myotubes. associated with the trans-Golgi, but not with the cis- Golgi apparatus (by similarity). | membrane. |
| O95185 | UNC5C_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| O95190 | OAZ2_HUMAN | | cytoplasm. secreted protein. |

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|--------|-------------|--|------------------------|
| O95196 | CSPG5_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). Golgi apparatus; Golgi membrane; single-pass type i membrane protein (by similarity). in neurons, localizes to synaptic junctions (by similarity). also detected in the endoplasmic reticulum and the Golgi (by similarity). partially enriched in lipid rafts | membrane. |
| O95197 | RTN3_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| O95198 | KLHL2_HUMAN | | cytoplasm. |
| O95199 | RCBT2_HUMAN | | cytoplasm. |
| O95201 | ZN205_HUMAN | nucleus (probable). | nucleus. |
| O95206 | PCDH8_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| O95210 | GET1_HUMAN | membrane; single-pass type iii membrane protein (probable). distributed in the transverse tubules and/or near the junctional sarcoplasmic reticulum. | membrane. |
| O95214 | LERL1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| O95218 | ZRAB2_HUMAN | nucleus (potential). | cytoplasm. |
| O95219 | SNX4_HUMAN | | cytoplasm. |
| O95235 | KI20A_HUMAN | Golgi apparatus (by similarity). | cytoplasm. nucleus. |
| O95236 | APOL3_HUMAN | cytoplasm (probable). | secreted protein. |
| O95237 | LRAT_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| O95238 | SPDEF_HUMAN | nucleus (probable). | nucleus. |
| O95249 | GOSR1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type iv membrane protein (by similarity). enriched on vesicular components at the terminal rims of the Golgi (by similarity). | endoplasmic reticulum. |
| O95255 | MRP6_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| O95256 | I18RA_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O95257 | GA45G_HUMAN | | cytoplasm. |
| O95258 | UCP5_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| O95260 | ATE1_HUMAN | isoform ate1-1: nucleus (by similarity). isoform ate1-1, isoform ate1-2: cytoplasm (by similarity). | cytoplasm. |
| O95263 | PDE8B_HUMAN | | cytoplasm. |
| O95279 | KCNK5_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |

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|--------|-------------|--|-------------------------------------|
| O95292 | VAPB_HUMAN | cell membrane; single-pass type iv membrane protein (by similarity). intracytoplasmic membrane; single-pass type iv membrane protein (by similarity). present in the plasma membrane and in intracellular vesicles (by similarity). | membrane. |
| O95294 | RASL1_HUMAN | | cytoplasm. |
| O95295 | S25BP_HUMAN | or: cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). or: cytoplasmic vesicle; cytoplasmic vesicle membrane (by similarity). may be cytoplasmic and peripheral membrane bound or anchored to the vesicular membrane through an n-terminal signal anchor (by similarity). | synapse. |
| O95297 | MPZL1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O95302 | FKBP9_HUMAN | endoplasmic reticulum (by similarity). | endoplasmic reticulum. |
| O95336 | 6PGL_HUMAN | | endoplasmic reticulum. lysosome. |
| O95337 | YZX3_HUMAN | | nucleus. |
| O95340 | PAPS2_HUMAN | | cytoplasm. mitochondrion. |
| O95352 | ATG7_HUMAN | cytoplasm (probable). | cytoplasm. |
| O95363 | SYFM_HUMAN | mitochondrion; mitochondrial matrix (potential). | cytoplasm. |
| O95365 | ZBT7A_HUMAN | nucleus (by similarity). | nucleus. |
| O95372 | LYPA2_HUMAN | cytoplasm (probable). | cytoplasm. |
| O95382 | M3K6_HUMAN | | cytoplasm. |
| O95389 | WISP3_HUMAN | secreted protein (probable). | secreted protein. |
| O95390 | GDF11_HUMAN | secreted protein (probable). | secreted protein. |
| O95393 | BMP10_HUMAN | secreted protein (by similarity). | secreted protein. |
| O95394 | AGM1_HUMAN | | cytoplasm. |
| O95396 | MOCS3_HUMAN | | cytoplasm. |
| O95406 | CNIH_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O95415 | BI3_HUMAN | | endosome; membrane. |
| O95416 | SOX14_HUMAN | nucleus (potential). | nucleus. |
| O95424 | MYLE_HUMAN | | membrane. |
| O95427 | PIGN_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| O95447 | CU013_HUMAN | | nucleus. |
| O95450 | ATS2_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| O95455 | TGDS_HUMAN | | cytoplasm. mitochondrion. |
| O95456 | DSCR2_HUMAN | | membrane. |
| O95466 | FMNL_HUMAN | cytoplasm (by similarity). | centriole. cytoplasm. |
| O95475 | SIX6_HUMAN | nucleus (by similarity). | nucleus. |
| O95477 | ABCA1_HUMAN | | membrane. |
| O95497 | VNN1_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | endosome. |
| O95498 | VNN2_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | endosome. |
| O95502 | NPTXR_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| O95503 | CBX6_HUMAN | nucleus (by similarity). | nucleus. |

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|--------|-------------|---|---------------------------------------|
| O95521 | PRAM1_HUMAN | | cytoplasm. |
| O95522 | PRA12_HUMAN | | cytoplasm. |
| O95544 | NADK_HUMAN | | cytoplasm. |
| O95562 | SFT2B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. Golgi. |
| O95563 | BR44_HUMAN | | mitochondrion. |
| O95568 | CA156_HUMAN | | cytoplasm. |
| O95573 | ACSL3_HUMAN | mitochondrion; mitochondrial outer membrane; single-pass type iii membrane protein (by similarity). peroxisome; peroxisomal membrane; single-pass type iii membrane protein (by similarity). microsome; microsomal membrane; single-pass type iii membrane protein (by similarity). | endoplasmic reticulum. peroxisome. |
| O95620 | DUS4L_HUMAN | | mitochondrion. |
| O95625 | ZBT11_HUMAN | nucleus (potential). | nucleus. |
| O95626 | AN32D_HUMAN | | cytoplasm. |
| O95628 | CNOT4_HUMAN | cytoplasm (probable). nucleus (probable). | nucleus. |
| O95631 | NET1_HUMAN | secreted protein; extracellular space (by similarity). | secreted protein. |
| O95633 | FSTL3_HUMAN | secreted protein (potential). ref.3: nucleus. | secreted protein. |
| O95650 | KI16_HUMAN | | membrane. |
| O95661 | DIRA3_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| O95670 | VATG2_HUMAN | | mitochondrion. |
| O95671 | ASML_HUMAN | | cytoplasm. |
| O95672 | ECEL1_HUMAN | membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| O95674 | CDS2_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein; matrix side (potential). | membrane. |
| O95677 | EYA4_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| O95685 | PPR3D_HUMAN | | cytoplasm. |
| O95704 | APBB3_HUMAN | | cytoplasm. |
| O95714 | HERC2_HUMAN | | cytoplasm. |
| O95716 | RAB3D_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| O95718 | ERR2_HUMAN | nucleus (potential). | nucleus. |
| O95741 | CPNE6_HUMAN | | cytoplasm. |
| O95747 | OXS1_HUMAN | | cytoplasm. |
| O95757 | HS74L_HUMAN | cytoplasm (by similarity). nucleus (by similarity). may translocate to the nucleus after heat shock (by similarity). | cytoplasm. |
| O95758 | ROD1_HUMAN | | nucleus. |
| O95759 | TBCD8_HUMAN | | Golgi. |
| O95766 | CG028_HUMAN | | cytoplasm. |
| O95777 | LSM8_HUMAN | nucleus (potential). | nucleus. |
| O95780 | ZN682_HUMAN | nucleus (probable). | cytoplasm. nucleus. |

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|--------|-------------|--|--|
| O95782 | AP2A1_HUMAN | Golgi apparatus. cytoplasmic vesicle; cytoplasmic vesicle membrane; peripheral membrane protein; cytoplasmic side (by similarity). component of the coat surrounding the cytoplasmic face of coated vesicles located at the Golgi complex (by similarity). | Golgi. |
| O95789 | ZMYM6_HUMAN | nucleus (potential). | nucleus. |
| O95801 | TTC4_HUMAN | | cytoplasm. |
| O95803 | NDST3_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endosome. |
| O95807 | TM50A_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| O95813 | CER1_HUMAN | secreted protein (probable). | secreted protein. |
| O95816 | BAG2_HUMAN | | cytoplasm. |
| O95817 | BAG3_HUMAN | | cytoplasm. |
| O95819 | M4K4_HUMAN | | cytoplasm. |
| O95825 | QORL_HUMAN | | cytoplasm. |
| O95832 | CLD1_HUMAN | cell membrane; cell-cell junction; tight junction; multi-pass membrane protein (by similarity). | membrane. |
| O95834 | EMAL2_HUMAN | | cytoplasm. |
| O95837 | GNA14_HUMAN | | cytoplasm. |
| O95843 | GUC1C_HUMAN | | cytoplasm. |
| O95857 | TSN13_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O95858 | TSN15_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| O95859 | TSN12_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| O95861 | BPNT1_HUMAN | | cytoplasm. |
| O95863 | SNAI_HUMAN | nucleus (potential). | nucleus. |
| O95865 | DDAH2_HUMAN | | cytoplasm. |
| O95867 | LYG6C_HUMAN | secreted protein (potential). | membrane. |
| O95870 | BAT5_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O95872 | BAT4_HUMAN | | centriole. nucleus. |
| O95873 | CF047_HUMAN | | cytoplasm. |
| O95886 | DLGP3_HUMAN | cell membrane; peripheral membrane protein (by similarity). postsynaptic density of neuronal cells (by similarity). | cytoplasm. |
| O95897 | NOE2_HUMAN | secreted protein (probable). | cytoplasm. Golgi. |
| O95905 | SGT1_HUMAN | | nucleus. |
| O95922 | TTLL1_HUMAN | | centriole. cytoplasm. secreted protein. mitochondrion. |
| O95925 | EPPI_HUMAN | secreted protein (potential). | secreted protein. |
| O95932 | TGM3L_HUMAN | | cytoplasm. |
| O95935 | TBX18_HUMAN | nucleus (potential). | nucleus. |
| O95936 | EOMES_HUMAN | nucleus (potential). | nucleus. |
| O95944 | NCTR2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| O95947 | TBX6_HUMAN | nucleus (potential). | nucleus. |
| O95968 | SG1D1_HUMAN | secreted protein (potential). | secreted protein. |
| O95969 | SG1D2_HUMAN | secreted protein (potential). | secreted protein. |
| O95970 | LGI1_HUMAN | secreted protein (potential). | cytoplasm. secreted protein. |

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|--------|-------------|--|------------------------|
| O95985 | TOP3B_HUMAN | | nucleus. |
| O95988 | TCL1B_HUMAN | | cytoplasm. |
| O95989 | NUDT3_HUMAN | cytoplasm (probable). | cytoplasm. |
| O95992 | CH25H_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| O95994 | AGR2_HUMAN | secreted protein (probable). | cytoplasm. |
| O95995 | GAS8_HUMAN | cytoplasm (by similarity). associates with microtubules. flagellum (by similarity). in spermatozoa, it localizes to the flagellum (by similarity). | cytoplasm. |
| O95998 | I18BP_HUMAN | secreted protein (potential). | secreted protein. |
| O96001 | GSUB_HUMAN | | secreted protein. |
| O96002 | CX001_HUMAN | | centriole. cytoplasm. |
| O96004 | HAND1_HUMAN | | nucleus. |
| O96005 | CLPT1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| O96007 | MOS2L_HUMAN | | cytoplasm. |
| O96008 | TOM40_HUMAN | mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| O96009 | NAPSA_HUMAN | | secreted protein. |
| O96013 | PAK4_HUMAN | | cytoplasm. |
| O96015 | DNAL4_HUMAN | | cytoplasm. |
| O96018 | APBA3_HUMAN | | Golgi. |
| O96033 | MOS2S_HUMAN | | cytoplasm. |
| P00156 | CYB_HUMAN | | mitochondrion. |
| P00374 | DYR_HUMAN | | cytoplasm. |
| P00414 | COX3_HUMAN | | mitochondrion. |
| P00439 | PH4H_HUMAN | | endoplasmic reticulum. |
| P00491 | PNPH_HUMAN | | cytoplasm. |
| P00540 | MOS_HUMAN | | cytoplasm. |
| P00736 | C1R_HUMAN | | secreted protein. |
| P00739 | HPTR_HUMAN | secreted protein (potential). | secreted protein. |
| P00742 | FA10_HUMAN | | secreted protein. |
| P00749 | UROK_HUMAN | | secreted protein. |
| P00813 | ADA_HUMAN | | cytoplasm. |
| P00966 | ASSY_HUMAN | | cytoplasm. |
| P01023 | A2MG_HUMAN | | secreted protein. |
| P01031 | CO5_HUMAN | | secreted protein. |
| P01034 | CYTC_HUMAN | | secreted protein. |
| P01036 | CYTS_HUMAN | | secreted protein. |
| P01037 | CYTN_HUMAN | | secreted protein. |
| P01124 | BLYM_HUMAN | | secreted protein. |
| P01127 | PDGFB_HUMAN | | secreted protein. |
| P01189 | COLI_HUMAN | | secreted protein. |
| P01591 | IGJ_HUMAN | | secreted protein. |
| P01593 | KV1A_HUMAN | | secreted protein. |
| P01594 | KV1B_HUMAN | | secreted protein. |
| P01595 | KV1C_HUMAN | | secreted protein. |
| P01596 | KV1D_HUMAN | | secreted protein. |
| P01597 | KV1E_HUMAN | | secreted protein. |
| P01598 | KV1F_HUMAN | | secreted protein. |
| P01599 | KV1G_HUMAN | | secreted protein. |
| P01600 | KV1H_HUMAN | | secreted protein. |
| P01603 | KV1K_HUMAN | | secreted protein. |

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|--------|------------|--|-------------------|
| P01604 | KV1L_HUMAN | | secreted protein. |
| P01605 | KV1M_HUMAN | | secreted protein. |
| P01606 | KV1N_HUMAN | | secreted protein. |
| P01607 | KV1O_HUMAN | | secreted protein. |
| P01608 | KV1P_HUMAN | | secreted protein. |
| P01609 | KV1Q_HUMAN | | secreted protein. |
| P01610 | KV1R_HUMAN | | secreted protein. |
| P01611 | KV1S_HUMAN | | secreted protein. |
| P01612 | KV1T_HUMAN | | secreted protein. |
| P01613 | KV1U_HUMAN | | secreted protein. |
| P01614 | KV2A_HUMAN | | secreted protein. |
| P01615 | KV2B_HUMAN | | secreted protein. |
| P01616 | KV2C_HUMAN | | membrane. |
| P01617 | KV2D_HUMAN | | secreted protein. |
| P01619 | KV3A_HUMAN | | membrane. |
| P01620 | KV3B_HUMAN | | secreted protein. |
| P01622 | KV3D_HUMAN | | secreted protein. |
| P01623 | KV3E_HUMAN | | secreted protein. |
| P01624 | KV3F_HUMAN | | secreted protein. |
| P01625 | KV4A_HUMAN | | secreted protein. |
| P01699 | LV1A_HUMAN | | secreted protein. |
| P01700 | LV1B_HUMAN | | secreted protein. |
| P01701 | LV1C_HUMAN | | secreted protein. |
| P01702 | LV1D_HUMAN | | secreted protein. |
| P01703 | LV1E_HUMAN | | secreted protein. |
| P01704 | LV2A_HUMAN | | secreted protein. |
| P01705 | LV2B_HUMAN | | secreted protein. |
| P01706 | LV2C_HUMAN | | secreted protein. |
| P01707 | LV2D_HUMAN | | secreted protein. |
| P01708 | LV2E_HUMAN | | secreted protein. |
| P01709 | LV2F_HUMAN | | secreted protein. |
| P01710 | LV2G_HUMAN | | secreted protein. |
| P01711 | LV2H_HUMAN | | secreted protein. |
| P01712 | LV2I_HUMAN | | secreted protein. |
| P01713 | LV2J_HUMAN | | secreted protein. |
| P01714 | LV3A_HUMAN | | secreted protein. |
| P01715 | LV4A_HUMAN | | secreted protein. |
| P01716 | LV4B_HUMAN | | secreted protein. |
| P01717 | LV4C_HUMAN | | secreted protein. |
| P01718 | LV4D_HUMAN | | secreted protein. |
| P01719 | LV5A_HUMAN | | secreted protein. |
| P01720 | LV7A_HUMAN | | secreted protein. |
| P01721 | LV6A_HUMAN | | secreted protein. |
| P01722 | LV6B_HUMAN | | secreted protein. |
| P01733 | TVB1_HUMAN | | membrane. |
| P01737 | TVA3_HUMAN | | membrane. |
| P01742 | HV1A_HUMAN | | secreted protein. |
| P01743 | HV1B_HUMAN | | secreted protein. |
| P01760 | HV1D_HUMAN | | secreted protein. |
| P01761 | HV1E_HUMAN | | secreted protein. |
| P01762 | HV3A_HUMAN | | secreted protein. |
| P01763 | HV3B_HUMAN | | secreted protein. |
| P01764 | HV3C_HUMAN | | secreted protein. |
| P01765 | HV3D_HUMAN | | secreted protein. |
| P01766 | HV3E_HUMAN | | secreted protein. |
| P01767 | HV3F_HUMAN | | secreted protein. |

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|--------|-------------|--|---|
| P01768 | HV3G_HUMAN | | secreted protein. |
| P01769 | HV3H_HUMAN | | secreted protein. |
| P01770 | HV3I_HUMAN | | secreted protein. |
| P01771 | HV3J_HUMAN | | secreted protein. |
| P01772 | HV3K_HUMAN | | secreted protein. |
| P01773 | HV3L_HUMAN | | secreted protein. |
| P01774 | HV3M_HUMAN | | secreted protein. |
| P01775 | HV3N_HUMAN | | secreted protein. |
| P01776 | HV3O_HUMAN | | secreted protein. |
| P01777 | HV3P_HUMAN | | secreted protein. |
| P01778 | HV3Q_HUMAN | | secreted protein. |
| P01779 | HV3R_HUMAN | | secreted protein. |
| P01780 | HV3S_HUMAN | | secreted protein. |
| P01781 | HV3T_HUMAN | | secreted protein. |
| P01782 | HV3U_HUMAN | | secreted protein. |
| P01814 | HV2A_HUMAN | | secreted protein. |
| P01815 | HV2B_HUMAN | | secreted protein. |
| P01816 | HV2C_HUMAN | | secreted protein. |
| P01817 | HV2D_HUMAN | | secreted protein. |
| P01818 | HV2E_HUMAN | | secreted protein. |
| P01824 | HV2F_HUMAN | | secreted protein. |
| P01825 | HV2G_HUMAN | | secreted protein. |
| P01834 | KAC_HUMAN | | secreted protein. |
| P01842 | LAC_HUMAN | | secreted protein. |
| P01848 | TCA_HUMAN | | membrane. |
| P01850 | TCB_HUMAN | | membrane. |
| P01854 | IGHE_HUMAN | | secreted protein. |
| P01857 | IGHG1_HUMAN | | cytoplasm. |
| P01859 | IGHG2_HUMAN | | cytoplasm. secreted protein; membrane. |
| P01860 | IGHG3_HUMAN | | cytoplasm. secreted protein; membrane. |
| P01861 | IGHG4_HUMAN | | cytoplasm. secreted protein; membrane. |
| P01871 | MUC_HUMAN | | cytoplasm. secreted protein; membrane. |
| P01876 | IGHA1_HUMAN | | secreted protein. |
| P01877 | IGHA2_HUMAN | | cytoplasm. secreted protein; membrane. |
| P01880 | IGHD_HUMAN | | secreted protein. |
| P01893 | HLAH_HUMAN | | membrane. |
| P01906 | HA26_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P01907 | HA25_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P01908 | HA21_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| P01909 | HA23_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P01912 | HB2B_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| P01914 | HB2C_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| P01918 | HB21_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |

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| P01919 | HB22_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P01920 | HB24_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P02008 | HBAZ_HUMAN | | endoplasmic reticulum. |
| P02042 | HBD_HUMAN | | endoplasmic reticulum. |
| P02100 | HBE_HUMAN | | endoplasmic reticulum. |
| P02144 | MYG_HUMAN | | endoplasmic reticulum. |
| P02452 | CO1A1_HUMAN | | secreted protein. |
| P02458 | CO2A1_HUMAN | | secreted protein. |
| P02461 | CO3A1_HUMAN | | secreted protein. |
| P02462 | CO4A1_HUMAN | | secreted protein. |
| P02489 | CRYAA_HUMAN | | cytoplasm. |
| P02511 | CRYAB_HUMAN | | cytoplasm. |
| P02533 | K1C14_HUMAN | | cytoplasm. |
| P02538 | K2C6A_HUMAN | | cytoplasm. |
| P02549 | SPTA1_HUMAN | | cytoplasm. |
| P02585 | TNNC2_HUMAN | | cytoplasm. |
| P02689 | MYP2_HUMAN | | cytoplasm. |
| P02735 | SAA_HUMAN | | secreted protein. |
| P02748 | CO9_HUMAN | | secreted protein. |
| P02792 | FRIL_HUMAN | | secreted protein. |
| P02794 | FRIH_HUMAN | | endoplasmic reticulum. secreted protein. |
| P02795 | MT2_HUMAN | | cytoplasm. nucleus. |
| P02811 | PRPE_HUMAN | | secreted protein. |
| P03886 | NU1M_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (probable). | mitochondrion. |
| P03897 | NU3M_HUMAN | | mitochondrion. |
| P03901 | NU4LM_HUMAN | | mitochondrion. |
| P03905 | NU4M_HUMAN | | mitochondrion. |
| P03915 | NU5M_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (probable). | mitochondrion. |
| P03923 | NU6M_HUMAN | | mitochondrion. |
| P03952 | KLKB1_HUMAN | | secreted protein. |
| P03956 | MMP1_HUMAN | | secreted protein. |
| P03979 | TVC_HUMAN | | membrane. |
| P03986 | TCC_HUMAN | | membrane. |
| P03992 | HB25_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P04003 | C4BP_HUMAN | | secreted protein. |
| P04049 | RAF1_HUMAN | | cytoplasm. |
| P04070 | PROC_HUMAN | | secreted protein. |
| P04075 | ALDOA_HUMAN | | cytoplasm. |
| P04083 | ANXA1_HUMAN | | cytoplasm. |
| P04085 | PDGFA_HUMAN | | secreted protein. |
| P04180 | LCAT_HUMAN | | cytoplasm. |
| P04206 | KV3G_HUMAN | | secreted protein. |
| P04207 | KV3H_HUMAN | | secreted protein. |
| P04208 | LV1F_HUMAN | | secreted protein. |
| P04209 | LV2K_HUMAN | | secreted protein. |
| P04211 | LV0A_HUMAN | | secreted protein. |
| P04220 | MUCB_HUMAN | | cytoplasm. secreted protein; membrane. |

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| P04226 | HA22_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P04233 | HG2A_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| P04259 | K2C6B_HUMAN | | cytoplasm. |
| P04264 | K2C1_HUMAN | | cytoplasm. cytoskeleton. |
| P04278 | SHBG_HUMAN | secreted protein (by similarity). in testis, it is synthesized by the sertoli cells, secreted into the lumen of the seminiferous tubule and transported to the epididymis (by similarity). | secreted protein. |
| P04280 | PRP1_HUMAN | | secreted protein. |
| P04281 | PRP5_HUMAN | | centriole. cytoplasm. |
| P04350 | TBB4_HUMAN | | centriole. |
| P04424 | ARLY_HUMAN | | cytoplasm. |
| P04430 | KV1V_HUMAN | | secreted protein. |
| P04431 | KV1W_HUMAN | | secreted protein. |
| P04432 | KV1X_HUMAN | | secreted protein. |
| P04435 | TVB2_HUMAN | | membrane. |
| P04437 | TVA2_HUMAN | | membrane. |
| P04438 | HV2H_HUMAN | | secreted protein. |
| P04440 | HB2P_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P04629 | NTRK1_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). endocytosed to the endosomes upon treatment of cells with ngf (by similarity). | membrane. |
| P04632 | CPNS1_HUMAN | cytoplasm (by similarity). cell membrane (by similarity). translocates to the plasma membrane upon calcium binding (by similarity). | centriole. cytoplasm. |
| P04731 | MT1A_HUMAN | | cytoplasm. |
| P04732 | MT1E_HUMAN | | cytoplasm. |
| P04733 | MT1F_HUMAN | | cytoplasm. |
| P04818 | TYSY_HUMAN | | cytoplasm. |
| P04899 | GNAI2_HUMAN | | cytoplasm. Golgi. |
| P05062 | ALDOB_HUMAN | | cytoplasm. |
| P05093 | CP17A_HUMAN | | endoplasmic reticulum. |
| P05109 | S10A8_HUMAN | | secreted protein. |
| P05129 | KPCG_HUMAN | | cytoplasm. |
| P05160 | F13B_HUMAN | | secreted protein. |
| P05162 | LEG2_HUMAN | | secreted protein. |
| P05198 | IF2A_HUMAN | | cytoplasm. |
| P05230 | FGF1_HUMAN | | secreted protein. |
| P05386 | RLA1_HUMAN | | cytoplasm. |
| P05387 | RLA2_HUMAN | | cytoplasm. |
| P05388 | RLA0_HUMAN | | cytoplasm. |
| P05451 | REG1A_HUMAN | | secreted protein. |
| P05455 | LA_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P05536 | HA27_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P05537 | HB23_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P05538 | HB2X_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P05546 | HEP2_HUMAN | | secreted protein. |

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| P05771 | KPCB_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | cytoplasm. |
| P05783 | K1C18_HUMAN | | cytoplasm. |
| P05787 | K2C8_HUMAN | | cytoplasm. |
| P05813 | CRBA1_HUMAN | | cytoplasm. |
| P05937 | CALB1_HUMAN | | cytoplasm. |
| P05976 | MLE1_HUMAN | | cytoskeleton. |
| P05997 | CO5A2_HUMAN | | secreted protein. |
| P06276 | CHLE_HUMAN | | secreted protein. |
| P06307 | CCKN_HUMAN | | secreted protein. |
| P06310 | KV2F_HUMAN | | secreted protein. |
| P06311 | KV3K_HUMAN | | secreted protein. |
| P06313 | KV4B_HUMAN | | secreted protein. |
| P06314 | KV4C_HUMAN | | secreted protein. |
| P06315 | KV5A_HUMAN | | secreted protein. |
| P06316 | LV1G_HUMAN | | secreted protein. |
| P06317 | LV6C_HUMAN | | membrane. |
| P06318 | LV6D_HUMAN | | secreted protein. |
| P06319 | LV6E_HUMAN | | secreted protein. |
| P06326 | HV1F_HUMAN | | secreted protein. |
| P06331 | HV2I_HUMAN | | secreted protein. |
| P06493 | CDC2_HUMAN | nucleus (by similarity). | nucleus. |
| P06702 | S10A9_HUMAN | | secreted protein. |
| P06703 | S10A6_HUMAN | | endosome. nucleus. |
| P06730 | IF4E_HUMAN | | cytoplasm. |
| P06737 | PYGL_HUMAN | | cytoplasm. |
| P06741 | MLE3_HUMAN | | cytoskeleton. |
| P06753 | TPM3_HUMAN | | cytoplasm. |
| P06870 | KLK1_HUMAN | | secreted protein. |
| P06881 | CALCA_HUMAN | | secreted protein. |
| P06887 | LV1H_HUMAN | | secreted protein. |
| P06888 | LV1I_HUMAN | | secreted protein. |
| P06889 | LV4E_HUMAN | | secreted protein. |
| P07101 | TY3H_HUMAN | | endoplasmic reticulum. |
| P07108 | ACBP_HUMAN | | cytoplasm. |
| P07196 | NFL_HUMAN | | cytoplasm. |
| P07197 | NFM_HUMAN | | cytoplasm. |
| P07205 | PGK2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P07237 | PDIA1_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen. highly abundant. in some cell types, seems to be also secreted or associated with the plasma membrane, where it undergoes constant shedding and replacement from intracellular sources (probable). localizes near cd4-enriched regions on lymphoid cell surfaces. | endoplasmic reticulum. |
| P07311 | ACYP1_HUMAN | | cytoplasm. |
| P07315 | CRGC_HUMAN | | cytoplasm. |
| P07316 | CRGB_HUMAN | | cytoplasm. |
| P07320 | CRGD_HUMAN | | centriole. |
| P07332 | FES_HUMAN | | cytoplasm. |
| P07384 | CAN1_HUMAN | cytoplasm (by similarity). cell membrane (by similarity). translocates to the plasma membrane upon ca(2+) binding (by similarity). | cytoplasm. lysosome. |
| P07437 | TBB2_HUMAN | | centriole. |

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| P07438 | MT1B_HUMAN | | cytoplasm. |
| P07737 | PROF1_HUMAN | | cytoplasm. |
| P07738 | PMGE_HUMAN | | lysosome. |
| P07814 | SYEP_HUMAN | | cytoplasm. |
| P07902 | GALT_HUMAN | | nucleus. |
| P07948 | LYN_HUMAN | | cytoplasm. |
| P07951 | TPM2_HUMAN | | cytoplasm. |
| P07996 | TSP1_HUMAN | | secreted protein. |
| P08107 | HSP71_HUMAN | | cytoplasm. |
| P08123 | CO1A2_HUMAN | | secreted protein. |
| P08133 | ANXA6_HUMAN | stress fibers (by similarity). | cytoplasm. |
| P08134 | RHOC_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| P08237 | K6PF_HUMAN | | cytoplasm. |
| P08243 | ASNS_HUMAN | | cytoplasm. mitochondrion. |
| P08246 | ELNE_HUMAN | | secreted protein. |
| P08253 | MMP2_HUMAN | | secreted protein. |
| P08254 | MMP3_HUMAN | | secreted protein. |
| P08311 | CATG_HUMAN | | secreted protein. |
| P08397 | HEM3_HUMAN | cytoplasm (probable). | cytoplasm. |
| P08519 | APOA_HUMAN | | secreted protein. |
| P08547 | LIN1_HUMAN | | cytoplasm. |
| P08567 | PLEK_HUMAN | | cytoplasm. |
| P08572 | CO4A2_HUMAN | | secreted protein. |
| P08590 | MYL3_HUMAN | | cytoskeleton. |
| P08620 | FGF4_HUMAN | | secreted protein. |
| P08621 | RU17_HUMAN | | nucleus. |
| P08637 | FCG3A_HUMAN | cell membrane; single-pass type i membrane protein (potential). secreted protein. exists also as a soluble receptor. | membrane. |
| P08670 | VIME_HUMAN | | cytoplasm. cytoskeleton. |
| P08708 | RS17_HUMAN | | cytoplasm. |
| P08709 | FA7_HUMAN | | secreted protein. |
| P08727 | K1C19_HUMAN | | cytoplasm. |
| P08729 | K2C7_HUMAN | | cytoplasm. |
| P08754 | GNAI3_HUMAN | | Golgi. |
| P08758 | ANXA5_HUMAN | | cytoplasm. |
| P08779 | K1C16_HUMAN | | cytoplasm. |
| P08861 | ELA3B_HUMAN | | secreted protein. |
| P08910 | LBH2_HUMAN | | membrane. |
| P09038 | FGF2_HUMAN | | secreted protein. |
| P09093 | ELA3A_HUMAN | | secreted protein. |
| P09104 | ENOG_HUMAN | cytoplasm (by similarity). cell membrane (by similarity). can translocate to the plasma membrane in either the homodimeric (alpha/alpha) or heterodimeric (alpha/gamma) form (by | membrane. |
| P09105 | HBAT_HUMAN | | endoplasmic reticulum. |
| P09131 | P3_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| P09211 | GSTP1_HUMAN | | cytoplasm. |
| P09237 | MMP7_HUMAN | | secreted protein. |
| P09238 | MMP10_HUMAN | | secreted protein. |
| P09327 | VILI_HUMAN | | cytoplasm. |
| P09382 | LEG1_HUMAN | | cytoplasm. |
| P09417 | DHPR_HUMAN | | cytoplasm. |

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|--------|-------------|---|--|
| P09466 | PAEP_HUMAN | | secreted protein. |
| P09467 | F16P1_HUMAN | | cytoplasm. |
| P09471 | GNAO1_HUMAN | | cytoplasm. Golgi. |
| P09486 | SPRC_HUMAN | | secreted protein. |
| P09493 | TPM1_HUMAN | | cytoplasm. |
| P09525 | ANXA4_HUMAN | | cytoplasm. |
| P09565 | IG2R_HUMAN | | cytoplasm. secreted protein. |
| P09603 | CSF1_HUMAN | cell membrane; single-pass membrane protein (by similarity). processed form: secreted protein; extracellular space (by similarity). | membrane. |
| P09769 | FGR_HUMAN | | cytoplasm. |
| P09871 | C1S_HUMAN | | secreted protein. |
| P09912 | INI2_HUMAN | | mitochondrion. |
| P09913 | IFIT2_HUMAN | | centriole. cytoplasm. |
| P09914 | IFIT1_HUMAN | | cytoplasm. |
| P09972 | ALDOC_HUMAN | | cytoplasm. |
| P0C024 | NUDT7_HUMAN | peroxisome (by similarity). | cytoplasm. |
| P0C025 | NUD17_HUMAN | mitochondrion (potential). | mitochondrion. |
| P0C091 | FREM3_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | membrane. |
| P0C0E4 | RB40L_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| P0C0L4 | CO4A_HUMAN | | secreted protein. |
| P0C0L5 | CO4B_HUMAN | | secreted protein. |
| P0C0P6 | NPS_HUMAN | secreted protein (potential). | secreted protein. |
| P10070 | GLI2_HUMAN | nucleus (potential). | nucleus. |
| P10072 | HKR1_HUMAN | nucleus (potential). | nucleus. |
| P10075 | GLI4_HUMAN | nucleus (potential). | nucleus. |
| P10114 | RAP2A_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| P10124 | PGSG_HUMAN | | cytoplasm. |
| P10153 | RNAS2_HUMAN | lysosome (probable). matrix of eosinophils large specific granule. | secreted protein. |
| P10163 | PRB4S_HUMAN | | secreted protein. |
| P10265 | VPK10_HUMAN | | secreted protein. |
| P10266 | POK10_HUMAN | | cytoplasm. secreted protein. mitochondrion. nucleus. |
| P10267 | ENK10_HUMAN | | membrane. |
| P10301 | RRAS_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (by similarity). inner surface of plasma membrane possibly with attachment requiring acylation of the c-terminal cysteine (by similarity with ras). | cytoplasm. Golgi. |
| P10323 | ACRO_HUMAN | | secreted protein. |
| P10398 | ARAF_HUMAN | | cytoplasm. |
| P10523 | ARRS_HUMAN | | cytoplasm. |
| P10588 | EAR2_HUMAN | nucleus (potential). | nucleus. |
| P10589 | COT1_HUMAN | nucleus (potential). | nucleus. |
| P10620 | MGST1_HUMAN | microsome (by similarity). mitochondrion; mitochondrial outer membrane; peripheral membrane protein (by similarity). | cytoplasm. microsome. |
| P10643 | CO7_HUMAN | | secreted protein. |
| P10644 | KAPO_HUMAN | | cytoplasm. |
| P10746 | HEM4_HUMAN | | cytoplasm. |

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|--------|-------------|--|------------------------------|
| P10912 | GHR_HUMAN | cell membrane; single-pass type i membrane protein. on growth hormone binding, ghr is ubiquitinated, internalized, down-regulated and transported into a degradative or non-degradative pathway (by similarity). isoform 2 remains fixed to the cell membrane and is not internalized. ghbp: secreted protein. the proteolytic processed protein, ghbp, is secreted. | membrane. |
| P10915 | HPLN1_HUMAN | | secreted protein. |
| P10916 | MLRV_HUMAN | | cytoplasm. |
| P10966 | CD8B_HUMAN | cell membrane; single-pass type i membrane protein. isoform 3: secreted protein (probable). | membrane. |
| P11086 | PNMT_HUMAN | | cytoplasm. |
| P11137 | MAP2_HUMAN | | cytoplasm. |
| P11166 | GTR1_HUMAN | cell membrane; multi-pass membrane protein (by similarity). localizes primarily at the cell surface (by similarity). | membrane. |
| P11172 | PYR5_HUMAN | | centriole. cytoplasm. |
| P11216 | PYGB_HUMAN | | centriole. cytoplasm. |
| P11217 | PYGM_HUMAN | | cytoplasm. |
| P11226 | MBL2_HUMAN | | secreted protein. |
| P11233 | RALA_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| P11234 | RALB_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| P11274 | BCR_HUMAN | | cytoplasm. |
| P11277 | SPTB1_HUMAN | | cytoplasm. |
| P11387 | TOP1_HUMAN | | cytoplasm. |
| P11413 | G6PD_HUMAN | | cytoplasm. |
| P11441 | UBL4A_HUMAN | | cytoplasm. |
| P11464 | PSG1_HUMAN | secreted protein (potential). | secreted protein. |
| P11465 | PSG2_HUMAN | secreted protein (potential). | cytoplasm. secreted protein. |
| P11487 | FGF3_HUMAN | | secreted protein. |
| P11488 | GNAT1_HUMAN | | membrane. |
| P11532 | DMD_HUMAN | | cytoplasm. |
| P11802 | CDK4_HUMAN | | nucleus. |
| P11844 | CRGA_HUMAN | | cytoplasm. |
| P11908 | PRPS2_HUMAN | | cytoplasm. |
| P11926 | DCOR_HUMAN | | cytoplasm. |
| P12018 | VPREB_HUMAN | | secreted protein. |
| P12034 | FGF5_HUMAN | | secreted protein. |
| P12035 | K2C3_HUMAN | | cytoplasm. |
| P12036 | NFH_HUMAN | | cytoplasm. |
| P12107 | COBA1_HUMAN | | secreted protein. |
| P12109 | CO6A1_HUMAN | | secreted protein. |
| P12111 | CO6A3_HUMAN | | secreted protein. |
| P12259 | FA5_HUMAN | | secreted protein. |
| P12268 | IMDH2_HUMAN | | cytoplasm. mitochondrion. |
| P12273 | PIP_HUMAN | secreted protein (probable). | secreted protein. |
| P12429 | ANXA3_HUMAN | | cytoplasm. |
| P12525 | MYCL2_HUMAN | nucleus (potential). | nucleus. |
| P12757 | SKIL_HUMAN | | nucleus. |
| P12829 | MYL4_HUMAN | | cytoskeleton. |
| P12931 | SRC_HUMAN | | cytoplasm. |

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|--------|-------------|---|-----------------------------------|
| P12955 | PEPD_HUMAN | | secreted protein. |
| P12980 | LYL1_HUMAN | nucleus (potential). | nucleus. |
| P13056 | TR2_HUMAN | nucleus (potential). | nucleus. |
| P13164 | IFM1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| P13385 | TDGF1_HUMAN | | secreted protein. |
| P13497 | BMP1_HUMAN | | secreted protein. |
| P13498 | CY24A_HUMAN | | mitochondrion. |
| P13640 | MT1G_HUMAN | | cytoplasm. nucleus. |
| P13645 | K1C10_HUMAN | | cytoplasm. |
| P13646 | K1C13_HUMAN | | cytoplasm. |
| P13647 | K2C5_HUMAN | | cytoplasm. |
| P13682 | ZNF35_HUMAN | nucleus (potential). | nucleus. |
| P13716 | HEM2_HUMAN | | cytoplasm. |
| P13762 | HB2K_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P13763 | HB2Q_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P13805 | TNNT1_HUMAN | | cytoplasm. |
| P13807 | GYS1_HUMAN | | centriole. |
| P13861 | KAP2_HUMAN | | cytoplasm. |
| P13928 | ANXA8_HUMAN | | cytoplasm. secreted protein. |
| P13929 | ENOB_HUMAN | cytoplasm. localized to the z line. some colocalization with ckm at m-band (by similarity). | membrane. |
| P13942 | COBA2_HUMAN | | secreted protein. |
| P13994 | 9KD_HUMAN | | cytoplasm. secreted protein. |
| P14174 | MIF_HUMAN | | secreted protein. |
| P14207 | FOLR2_HUMAN | cell membrane; lipid-anchor; gpi-anchor. secreted protein (probable). | secreted protein; membrane. |
| P14209 | CD99_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P14210 | HGF_HUMAN | | secreted protein. |
| P14314 | GLU2B_HUMAN | | endoplasmic reticulum. |
| P14317 | HCLS1_HUMAN | membrane; peripheral membrane protein (probable). cytoplasm (probable). mitochondrion (probable). | nucleus. |
| P14373 | TRI27_HUMAN | or: nucleus (by similarity). or: cytoplasm (by similarity). nuclear or cytoplasmic depending on the cell type (by similarity). colocalized with pml and eif3s6 in nuclear | nucleus. |
| P14550 | AK1A1_HUMAN | | cytoplasm. |
| P14618 | KPYM_HUMAN | | cytoplasm. |
| P14619 | KGP1B_HUMAN | | cytoplasm. |
| P14621 | ACYP2_HUMAN | | cytoplasm. |
| P14635 | CCNB1_HUMAN | | nucleus. |
| P14649 | MYL6B_HUMAN | | cytoskeleton. |
| P14780 | MMP9_HUMAN | | secreted protein. |
| P14854 | CX6B1_HUMAN | mitochondrion; mitochondrial intermembrane space (by similarity). | mitochondrion. |
| P14902 | I23O_HUMAN | | cytoplasm. endoplasmic reticulum. |
| P15090 | FABPA_HUMAN | cytoplasm (potential). | cytoplasm. |

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| P15144 | AMPN_HUMAN | cell membrane; single-pass type ii membrane protein. cytoplasm; cytosol (potential). a soluble form has also been detected. | membrane. |
| P15170 | GSPT1_HUMAN | | cytoplasm. |
| P15259 | PGAM2_HUMAN | | lysosome. |
| P15309 | PPAP_HUMAN | | cytoplasm. |
| P15311 | EZRI_HUMAN | membrane. localization to the apical membrane of parietal cells depends on the interaction with mpp5. localizes to cell extensions and peripheral processes of astrocytes (by similarity). microvillar peripheral membrane protein (cytoplasmic side). | cytoplasm. |
| P15328 | FOLR1_HUMAN | cell membrane; lipid-anchor; gpi-anchor. secreted protein (probable). | membrane. |
| P15498 | VAV_HUMAN | | cytoplasm. |
| P15509 | CSF2R_HUMAN | cell membrane; single-pass type i membrane protein. isoform 3, isoform 4, isoform 6: secreted protein (probable). | membrane. |
| P15514 | AREG_HUMAN | | membrane. |
| P15515 | HIS1_HUMAN | | secreted protein. |
| P15621 | ZNF44_HUMAN | nucleus (probable). | nucleus. |
| P15622 | ZN250_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P15735 | PHKG2_HUMAN | | cytoplasm. |
| P15814 | IGLL1_HUMAN | secreted protein (potential). | membrane. |
| P15880 | RS2_HUMAN | | cytoplasm. |
| P15882 | CHIN_HUMAN | | cytoplasm. |
| P15884 | ITF2_HUMAN | nucleus (probable). | nucleus. |
| P15954 | COX7C_HUMAN | | mitochondrion. |
| P16112 | PGCA_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein; membrane. |
| P16118 | F261_HUMAN | | cytoplasm. |
| P16157 | ANK1_HUMAN | isoform er1: cytoplasm. probably the other erythrocyte (er) isoforms, are located near the surface of erythrocytic plasma membrane. isoform mu17: membrane. isoform mu18, isoform mu19, isoform mu20: sarcoplasmic reticulum (probable). isoform mu17 colocalizes with obscn isoform 3/obscurin at the m line in differentiated skeletal muscle cells. | cytoplasm. |
| P16298 | PP2BB_HUMAN | | cytoplasm. |
| P16499 | PDE6A_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| P16520 | GBB3_HUMAN | | membrane. |
| P16562 | CRIS2_HUMAN | secreted protein (probable). | secreted protein. |
| P16885 | PLCG2_HUMAN | | cytoplasm. secreted protein. |
| P16930 | FAAA_HUMAN | | membrane. |
| P17014 | ZNF12_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P17017 | ZNF14_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P17019 | ZN708_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P17020 | ZNF16_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P17022 | ZNF18_HUMAN | nucleus (potential). | nucleus. |
| P17023 | ZNF19_HUMAN | nucleus (potential). | nucleus. |
| P17024 | ZNF20_HUMAN | nucleus (potential). | cytoplasm. nucleus. |

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| P17025 | ZN182_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P17026 | ZNF22_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| P17027 | ZNF23_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P17028 | ZNF24_HUMAN | nucleus (potential). | nucleus. |
| P17029 | ZKSC1_HUMAN | nucleus (potential). | nucleus. |
| P17030 | ZNF25_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P17031 | ZNF26_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P17032 | ZN37A_HUMAN | nucleus (potential). | nucleus. |
| P17035 | ZNF28_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P17036 | ZNF38_HUMAN | nucleus (potential). | nucleus. |
| P17038 | ZNF43_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P17039 | ZNF30_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P17040 | ZNF31_HUMAN | nucleus (potential). | nucleus. |
| P17041 | ZNF32_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P17066 | HSP76_HUMAN | | cytoplasm. |
| P17097 | ZNF7_HUMAN | nucleus (probable). | nucleus. |
| P17098 | ZNF8_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P17152 | TMM11_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| P17252 | KPCA_HUMAN | | cytoplasm. |
| P17257 | SEC_HUMAN | | cytoplasm. secreted protein. |
| P17542 | TAL1_HUMAN | | nucleus. |
| P17612 | KAPCA_HUMAN | inactive holoenzyme, monomeric catalytic subunit: cytoplasm (by similarity). monomeric catalytic subunit: nucleus (by similarity). translocates into the nucleus (monomeric catalytic subunit). | cytoplasm. nucleus. |
| P17643 | TYRP1_HUMAN | melanosome; melanosomal membrane; single- pass type i membrane protein (potential). | membrane. |
| P17707 | DCAM_HUMAN | | cytoplasm. |
| P17735 | ATTY_HUMAN | | cytoplasm. |
| P17752 | TPH1_HUMAN | | endoplasmic reticulum. |
| P17812 | PYRG1_HUMAN | | cytoplasm. |
| P17858 | K6PL_HUMAN | | cytoplasm. |
| P17980 | PRS6A_HUMAN | cytoplasm (potential). nucleus (potential). | cytoplasm. |
| P18077 | RL35A_HUMAN | | cytoplasm. |
| P18085 | ARF4_HUMAN | | Golgi. |
| P18124 | RL7_HUMAN | | cytoplasm. |
| P18135 | KV3L_HUMAN | | secreted protein. |
| P18136 | KV3M_HUMAN | | secreted protein. |
| P18545 | CNRG_HUMAN | | centriole. |
| P18621 | RL17_HUMAN | | cytoplasm. |
| P18669 | PGAM1_HUMAN | | cytoplasm. |
| P18859 | ATP5J_HUMAN | | mitochondrion. |
| P18887 | XRCC1_HUMAN | nucleus (probable). | nucleus. |
| P19012 | K1C15_HUMAN | | cytoplasm. |
| P19013 | K2C4_HUMAN | | cytoplasm. |
| P19087 | GNAT2_HUMAN | | membrane. |
| P19105 | MLRM_HUMAN | | cytoplasm. |
| P19113 | DCHS_HUMAN | | cytoplasm. |
| P19174 | PLCG1_HUMAN | | cytoplasm. |
| P19237 | TNNI1_HUMAN | | cytoplasm. |
| P19429 | TNNI3_HUMAN | | cytoplasm. |
| P19525 | E2AK2_HUMAN | | cytoplasm. |
| P19623 | SPEE_HUMAN | | centriole. cytoplasm. |

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|--------|-------------|---|-----------------------|
| P19784 | CSK22_HUMAN | | cytoplasm. |
| P19823 | ITIH2_HUMAN | | secreted protein. |
| P19827 | ITIH1_HUMAN | | secreted protein. |
| P19835 | CEL_HUMAN | | secreted protein. |
| P19961 | AMYC_HUMAN | secreted protein (by similarity). | lysosome. |
| P19971 | TYPH_HUMAN | | secreted protein. |
| P20036 | HA2Q_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| P20042 | IF2B_HUMAN | | cytoplasm. |
| P20073 | ANXA7_HUMAN | | cytoplasm. |
| P20151 | KLK2_HUMAN | | secreted protein. |
| P20248 | CCNA2_HUMAN | | nucleus. |
| P20292 | FLAP_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| P20336 | RAB3A_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| P20337 | RAB3B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| P20339 | RAB5A_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (by similarity). endosome; early endosome; early endosomal membrane; lipid-anchor (by similarity). | endosome. |
| P20393 | NR1D1_HUMAN | nucleus (potential). | nucleus. |
| P20472 | PRVA_HUMAN | | cytoplasm. |
| P20585 | MSH3_HUMAN | | nucleus. |
| P20711 | DDC_HUMAN | | cytoplasm. |
| P20774 | MIME_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| P20794 | MAK_HUMAN | | cytoplasm. |
| P20810 | ICAL_HUMAN | | secreted protein. |
| P20839 | IMDH1_HUMAN | | centriole. cytoplasm. |
| P20848 | A1ATR_HUMAN | | secreted protein. |
| P20849 | CO9A1_HUMAN | | secreted protein. |
| P20851 | C4BB_HUMAN | | secreted protein. |
| P20908 | CO5A1_HUMAN | | secreted protein. |
| P20929 | NEBU_HUMAN | | cytoplasm. |
| P20930 | FILA_HUMAN | | cytoplasm. |
| P20931 | YYY3_HUMAN | | secreted protein. |
| P20962 | PTMS_HUMAN | | nucleus. |
| P21108 | PRPS3_HUMAN | | cytoplasm. |
| P21128 | PP11_HUMAN | | secreted protein. |
| P21283 | VATC_HUMAN | | mitochondrion. |
| P21359 | NF1_HUMAN | | cytoplasm. |
| P21506 | ZNF10_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P21583 | SCF_HUMAN | isoform 1, isoform 2: cell membrane; single- pass type i membrane protein (by similarity). processed isoform 1: secreted protein. also exists as a secreted soluble form (isoform 1 only) (by similarity). | membrane. |
| P21728 | DRD1_HUMAN | cell membrane; multi-pass membrane protein. endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein. transport from the endoplasmic reticulum to the cell surface is regulated by interaction with dnajc14 (by | membrane. |

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|--------|-------------|--|------------------------|
| P21810 | PGS1_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| P21817 | RYR1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| P21941 | MATN1_HUMAN | | secreted protein. |
| P21980 | TGM2_HUMAN | | cytoplasm. |
| P22004 | BMP6_HUMAN | secreted protein (by similarity). | secreted protein. |
| P22090 | RS4Y1_HUMAN | | cytoplasm. |
| P22102 | PUR2_HUMAN | | cytoplasm. |
| P22234 | PUR6_HUMAN | | cytoplasm. |
| P22303 | ACES_HUMAN | | secreted protein. |
| P22314 | UBE1_HUMAN | | cytoplasm. |
| P22612 | KAPCG_HUMAN | | cytoplasm. |
| P22614 | SAA3_HUMAN | | secreted protein. |
| P22676 | CALB2_HUMAN | | cytoplasm. |
| P22694 | KAPCB_HUMAN | inactive holoenzyme, monomeric catalytic subunit: cytoplasm (by similarity). monomeric catalytic subunit: nucleus (by similarity). translocates into the nucleus (monomeric) (by similarity). | cytoplasm. |
| P22760 | AAAD_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (by similarity). microsome; microsomal membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. |
| P22914 | CRBS_HUMAN | | centriole. cytoplasm. |
| P23083 | HV1G_HUMAN | | secreted protein. |
| P23109 | AMPD1_HUMAN | | cytoplasm. |
| P23381 | SYW_HUMAN | | cytoplasm. |
| P23396 | RS3_HUMAN | | cytoplasm. |
| P23443 | KS6B1_HUMAN | cytoplasm. synaptosome (by similarity). also found in the soluble synaptosomal fractions (by similarity). | cytoplasm. |
| P23470 | PTPRG_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| P23508 | CRCM_HUMAN | | cytoplasm. |
| P23588 | IF4B_HUMAN | | nucleus. |
| P23610 | F8I2_HUMAN | | nucleus. |
| P23677 | IP3KA_HUMAN | | cytoplasm. |
| P23743 | DGKA_HUMAN | | cytoplasm. |
| P23919 | DTYMK_HUMAN | | cytoplasm. |
| P24158 | PRTN3_HUMAN | | secreted protein. |
| P24311 | COX7B_HUMAN | | mitochondrion. |
| P24347 | MMP11_HUMAN | | secreted protein. |
| P24385 | CCND1_HUMAN | | cytoplasm. nucleus. |
| P24386 | RAE1_HUMAN | | cytoplasm. |
| P24387 | CRHBP_HUMAN | | cytoplasm. |
| P24522 | GA45A_HUMAN | | nucleus. |
| P24534 | EF1B_HUMAN | | cytoplasm. |
| P24723 | KPCL_HUMAN | | cytoplasm. |
| P24844 | MLRN_HUMAN | | cytoskeleton. |
| P24941 | CDK2_HUMAN | | nucleus. |
| P25054 | APC_HUMAN | | cytoplasm. nucleus. |
| P25067 | CO8A2_HUMAN | | secreted protein. |
| P25098 | ARBK1_HUMAN | | cytoplasm. |

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|--------|-------------|---|---------------------------|
| P25440 | BRD2_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| P25686 | DNJB2_HUMAN | | endoplasmic reticulum. |
| P25713 | MT3_HUMAN | | cytoplasm. |
| P25791 | RBTN2_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P25800 | RBTN1_HUMAN | nucleus (by similarity). | nucleus. |
| P25815 | S100P_HUMAN | | cytoplasm. |
| P25940 | CO5A3_HUMAN | | secreted protein. |
| P26038 | MOES_HUMAN | cytoplasm (by similarity). phosphorylated form is enriched in microvilli-like structures at apical membrane (by similarity). | cytoplasm. |
| P26232 | CTNA2_HUMAN | | cytoplasm. |
| P26371 | KRA59_HUMAN | | cytoplasm. |
| P26373 | RL13_HUMAN | | cytoplasm. |
| P26374 | RAE2_HUMAN | | cytoplasm. |
| P26378 | ELAV4_HUMAN | | cytoplasm. nucleus. |
| P26447 | S10A4_HUMAN | | cytoplasm. |
| P26640 | SYV_HUMAN | | cytoplasm. |
| P26641 | EF1G_HUMAN | | cytoplasm. |
| P26885 | FKBP2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; peripheral membrane protein (probable). | cytoplasm. |
| P26927 | HGFL_HUMAN | | secreted protein. |
| P26998 | CRBB3_HUMAN | | cytoplasm. |
| P27361 | MK03_HUMAN | | cytoplasm. nucleus. |
| P27448 | MARK3_HUMAN | | cytoplasm. |
| P27469 | GOS2_HUMAN | | cytoplasm. |
| P27482 | CALL3_HUMAN | | cytoplasm. |
| P27544 | LASS1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). isoform 1 may recycle from the Golgi to the | endoplasmic reticulum. |
| P27635 | RL10_HUMAN | | cytoplasm. mitochondrion. |
| P27658 | CO8A1_HUMAN | | secreted protein. |
| P27816 | MAP4_HUMAN | | cytoplasm. |
| P27986 | P85A_HUMAN | | cytoplasm. |
| P27987 | IP3KB_HUMAN | | cytoplasm. |
| P28039 | AOAH_HUMAN | | secreted protein. |
| P28062 | PSB8_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| P28065 | PSB9_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| P28070 | PSB4_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| P28223 | 5HT2A_HUMAN | cell membrane; multi-pass membrane protein. localizes to the post-synaptic thickening of axo-dendritic synapses (by similarity). | membrane. |
| P28289 | TMOD1_HUMAN | | cytoplasm. |
| P28325 | CYTD_HUMAN | | secreted protein. |
| P28329 | CLAT_HUMAN | | cytoplasm. |
| P28482 | MK01_HUMAN | | cytoplasm. |
| P28562 | DUS1_HUMAN | | cytoplasm. |
| P28749 | RBL1_HUMAN | | nucleus. |
| P29034 | S10A2_HUMAN | | cytoplasm. |
| P29074 | PTN4_HUMAN | | cytoplasm. |

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|--------|-------------|---|------------------------|
| P29279 | CTGF_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). secreted protein (by similarity). | secreted protein. |
| P29372 | 3MG_HUMAN | nucleus (potential). | nucleus. |
| P29375 | JAD1A_HUMAN | nucleus (potential). | nucleus. |
| P29377 | S100G_HUMAN | | cytoplasm. |
| P29400 | CO4A5_HUMAN | | secreted protein. |
| P29401 | TKT_HUMAN | | cytoplasm. |
| P29474 | NOS3_HUMAN | | cytoplasm. |
| P29597 | TYK2_HUMAN | | cytoplasm. |
| P29692 | EF1D_HUMAN | | cytoplasm. |
| P29777 | GNAO2_HUMAN | | cytoplasm. Golgi. |
| P29966 | MARCS_HUMAN | | cytoplasm; membrane. |
| P29992 | GNA11_HUMAN | | cytoplasm. |
| P30039 | MAWBP_HUMAN | | centriole. cytoplasm. |
| P30041 | PRDX6_HUMAN | cytoplasm (by similarity). lysosome (by similarity). cytoplasmic vesicle (by similarity). and also found in lung secretory organelles (by similarity). | cytoplasm. |
| P30042 | ES1_HUMAN | mitochondrion (potential). | mitochondrion. |
| P30043 | BLVRB_HUMAN | cytoplasm (potential). | cytoplasm. |
| P30046 | DOPD_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P30047 | GFRP_HUMAN | | cytoplasm. |
| P30050 | RL12_HUMAN | | cytoplasm. |
| P30086 | PEBP1_HUMAN | cytoplasm (by similarity). | secreted protein. |
| P30101 | PDIA3_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| P30153 | 2AAA_HUMAN | | cytoplasm. nucleus. |
| P30154 | 2AAB_HUMAN | | cytoplasm. nucleus. |
| P30279 | CCND2_HUMAN | | cytoplasm. nucleus. |
| P30304 | MPIP1_HUMAN | | nucleus. |
| P30531 | SC6A1_HUMAN | cell membrane; multi-pass membrane protein. membrane; multi-pass membrane protein. localized at the plasma membrane and in a subset of intracellular vesicles. localized at the presynaptic terminals of interneurons (by similarity). | membrane. |
| P30566 | PUR8_HUMAN | | cytoplasm. |
| P30613 | KPYR_HUMAN | | cytoplasm. |
| P30679 | GNA15_HUMAN | | membrane. |
| P30740 | ILEU_HUMAN | cytoplasm (by similarity). | secreted protein. |
| P30793 | GCH1_HUMAN | | cytoplasm. |
| P31146 | COR1A_HUMAN | in non-infected macrophages, associated with the cortical microtubule network. in mycobacteria-infected macrophages, becomes progressively relocalized and retained around the mycobacterial phagosomes. retention on the phagosomal membrane is strictly dependent on mycobacterial viability and not due to impaired acidification (by similarity). | cytoplasm. |
| P31150 | GDIA_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P31152 | MK04_HUMAN | | cytoplasm. |
| P31153 | METK2_HUMAN | | cytoplasm. |
| P31314 | TLX1_HUMAN | nucleus (probable). | nucleus. |
| P31321 | KAP1_HUMAN | | cytoplasm. |

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|--------|-------------|---|------------------------------------|
| P31323 | KAP3_HUMAN | | cytoplasm. |
| P31689 | DNJA1_HUMAN | membrane; lipid-anchor (potential). | cytoplasm. |
| P31751 | AKT2_HUMAN | | cytoplasm. |
| P31939 | PUR9_HUMAN | | cytoplasm. |
| P31941 | ABC3A_HUMAN | | cytoplasm. |
| P31944 | CASPE_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P31948 | STIP1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | centriole. nucleus. |
| P31949 | S10AB_HUMAN | | cytoplasm. |
| P32019 | I5P2_HUMAN | | cytoplasm. endosome. |
| P32121 | ARRB2_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. nucleus. |
| P32242 | OTX1_HUMAN | nucleus (probable). | nucleus. |
| P32243 | OTX2_HUMAN | nucleus (probable). | nucleus. |
| P32298 | GRK4_HUMAN | | cytoplasm. |
| P32320 | CDD_HUMAN | | cytoplasm. |
| P32321 | DCTD_HUMAN | | cytoplasm. |
| P32322 | P5CR1_HUMAN | | mitochondrion. |
| P32455 | GBP1_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| P32456 | GBP2_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| P32745 | SSR3_HUMAN | membrane; multi-pass membrane protein. internalized into endoplasmic vesicles upon somatostatin- stimulation (by | membrane. |
| P32754 | HPPD_HUMAN | | endoplasmic reticulum. |
| P32926 | DSG3_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| P32930 | ONCO_HUMAN | | cytoplasm. |
| P32969 | RL9_HUMAN | | cytoplasm. |
| P33121 | ACSL1_HUMAN | mitochondrion; mitochondrial outer membrane; single-pass type iii membrane protein (by similarity). peroxisome; peroxisomal membrane; single-pass type iii membrane protein (by similarity). microsome; microsomal membrane; single-pass type iii membrane protein (by similarity). | endoplasmic reticulum. peroxisome. |
| P33151 | CADH5_HUMAN | cell membrane; cell-cell junction; single-pass type i membrane protein. cell membrane; cell-matrix junction; single-pass type i membrane protein (probable). found at cell-cell boundaries and probably at cell-matrix boundaries. | membrane. |
| P33176 | KINH_HUMAN | uniformly distributed between soma and neurites in hippocampal neurons (by similarity). | cytoplasm. nucleus. |
| P33552 | CKS2_HUMAN | | cytoplasm. nucleus. |
| P33763 | S10A5_HUMAN | | cytoplasm. |
| P33764 | S10A3_HUMAN | | cytoplasm. nucleus. |
| P33981 | TTK_HUMAN | | cytoplasm. |
| P33991 | MCM4_HUMAN | nucleus (by similarity). | nucleus. |
| P33993 | MCM7_HUMAN | nucleus (by similarity). | nucleus. |
| P34925 | RYK_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P34931 | HS70L_HUMAN | | cytoplasm. |

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|--------|-------------|---|-------------------------------|
| P34932 | HSP74_HUMAN | cytoplasm (probable). | cytoplasm. |
| P34947 | GRK5_HUMAN | | cytoplasm. |
| P34949 | MANA_HUMAN | cytoplasm (probable). | cytoplasm. |
| P35080 | PROF2_HUMAN | | cytoplasm. |
| P35125 | UBP6_HUMAN | | cytoplasm. nucleus. |
| P35219 | CAH8_HUMAN | | cytoplasm. |
| P35228 | NOS2A_HUMAN | | cytoplasm. |
| P35232 | PHB_HUMAN | mitochondrion; mitochondrial inner membrane (by similarity). | nucleus. |
| P35243 | RECO_HUMAN | | membrane. |
| P35249 | RFC4_HUMAN | nucleus (probable). | nucleus. |
| P35250 | RFC2_HUMAN | nucleus (probable). | nucleus. |
| P35268 | RL22_HUMAN | | cytoplasm. |
| P35398 | RORA_HUMAN | nucleus (probable). | nucleus. |
| P35442 | TSP2_HUMAN | | secreted protein. |
| P35443 | TSP4_HUMAN | | secreted protein. |
| P35527 | K1C9_HUMAN | | cytoplasm. |
| P35542 | SAA4_HUMAN | | secreted protein. |
| P35544 | UBIM_HUMAN | | centriole. cytoplasm. |
| P35555 | FBN1_HUMAN | | secreted protein. |
| P35556 | FBN2_HUMAN | | secreted protein. |
| P35557 | HXK4_HUMAN | | cytoplasm. |
| P35568 | IRS1_HUMAN | | nucleus. |
| P35573 | GDE_HUMAN | | lysosome. |
| P35579 | MYH9_HUMAN | | cytoskeleton. |
| P35580 | MYH10_HUMAN | | cytoskeleton. |
| P35606 | COPB2_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). the coatomer is cytoplasmic or polymerized on the cytoplasmic side of the Golgi, as well as on the vesicles/buds originating from it (by similarity). | endoplasmic reticulum. Golgi. |
| P35611 | ADDA_HUMAN | | cytoplasm. |
| P35612 | ADDB_HUMAN | | cytoplasm. |
| P35613 | BASI_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). colocalizes with slc16a1 and slc16a8 (by similarity). | membrane. |
| P35626 | ARBK2_HUMAN | | cytoplasm. |
| P35659 | DEK_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P35670 | ATP7B_HUMAN | Golgi apparatus; trans-Golgi network; trans- Golgi network membrane; multi-pass membrane protein (by similarity). predominantly found in the trans-Golgi network (tgn). not redistributed to the plasma membrane in response to elevated copper levels. plasma membrane in response to elevated copper levels. isoform 2: cytoplasm. wnd/140 kda: mitochondrion. the cleaved form wnd/140 kda is located in mitochondria. | membrane. |
| P35711 | SOX5_HUMAN | nucleus (by similarity). | nucleus. |
| P35712 | SOX6_HUMAN | nucleus (potential). | nucleus. |
| P35713 | SOX18_HUMAN | nucleus (potential). | nucleus. |
| P35716 | SOX11_HUMAN | nucleus (potential). | cytoplasm. nucleus. |

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|--------|-------------|---|-----------------------------------|
| P35789 | ZNF93_HUMAN | nucleus (probable). | nucleus. |
| P35813 | PP2CA_HUMAN | | cytoplasm. |
| P35908 | K22E_HUMAN | | cytoplasm. |
| P36021 | MOT8_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | membrane. |
| P36268 | GGTL2_HUMAN | | cytoplasm. |
| P36269 | GGT5_HUMAN | membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| P36404 | ARL2_HUMAN | | Golgi. |
| P36405 | ARL3_HUMAN | | Golgi. |
| P36507 | MP2K2_HUMAN | | cytoplasm. |
| P36508 | ZNF76_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P36543 | VATE_HUMAN | | mitochondrion. |
| P36575 | ARRC_HUMAN | | centriole. cytoplasm. |
| P36578 | RL4_HUMAN | | cytoplasm. |
| P36639 | 8ODP_HUMAN | | nucleus. |
| P36915 | GNL1_HUMAN | | cytoplasm. |
| P36959 | GMPR1_HUMAN | | cytoplasm. mitochondrion. |
| P37058 | DHB3_HUMAN | | cytoplasm. endoplasmic reticulum. |
| P37059 | DHB2_HUMAN | membrane; single-pass type ii membrane protein (potential). | endoplasmic reticulum. |
| P37235 | HPCL1_HUMAN | | cytoplasm. |
| P37802 | TAGL2_HUMAN | | cytoplasm; membrane. |
| P37837 | TALDO_HUMAN | cytoplasm (probable). | cytoplasm. |
| P38405 | GNAL_HUMAN | | cytoplasm. Golgi. |
| P38606 | VATA1_HUMAN | | mitochondrion. |
| P38607 | VATA2_HUMAN | | mitochondrion. |
| P38935 | SMBP2_HUMAN | nucleus (potential). | nucleus. |
| P39019 | RS19_HUMAN | | cytoplasm. |
| P39059 | COFA1_HUMAN | | secreted protein. |
| P39060 | COIA1_HUMAN | | secreted protein. |
| P39900 | MMP12_HUMAN | | secreted protein. |
| P40123 | CAP2_HUMAN | cell membrane (by similarity). | Golgi. |
| P40205 | NCYM_HUMAN | | secreted protein. |
| P40222 | TXLNA_HUMAN | | secreted protein. |
| P40305 | INI7_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| P40306 | PSB10_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| P40313 | CTRL_HUMAN | | secreted protein. |
| P40424 | PBX1_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P40425 | PBX2_HUMAN | nucleus (probable). | nucleus. |
| P40426 | PBX3_HUMAN | nucleus (probable). | nucleus. |
| P40429 | RL13A_HUMAN | | mitochondrion. |
| P40616 | ARL1_HUMAN | | Golgi. |
| P40818 | UBP8_HUMAN | | nucleus. |
| P40879 | S26A3_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| P40937 | RFC5_HUMAN | nucleus (probable). | nucleus. |
| P40938 | RFC3_HUMAN | nucleus (probable). | nucleus. |
| P40967 | PME17_HUMAN | membrane; single-pass type i membrane protein (potential). me20-s: secreted protein. there is also a secreted soluble form, me20-s, probably product of | membrane. |
| P41091 | IF2G_HUMAN | | cytoplasm. |

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|--------|-------------|---|---------------------------|
| P41162 | ETV3_HUMAN | nucleus (potential). | nucleus. |
| P41182 | BCL6_HUMAN | nucleus (by similarity). | nucleus. |
| P41219 | PERI_HUMAN | | cytoplasm. |
| P41220 | RGS2_HUMAN | | cytoplasm. |
| P41223 | BUD31_HUMAN | nucleus (potential). | nucleus. |
| P41226 | UBE1L_HUMAN | | cytoplasm. |
| P41229 | JAD1C_HUMAN | | nucleus. |
| P41236 | IPP2_HUMAN | | cytoplasm. |
| P41238 | ABEC1_HUMAN | | nucleus. |
| P41240 | CSK_HUMAN | cytoplasm (by similarity). mainly cytoplasmic. cell membrane (by similarity). also present in lipid rafts (by similarity). | cytoplasm. |
| P41247 | PLPL4_HUMAN | | centriole. cytoplasm. |
| P41250 | SYG_HUMAN | | cytoplasm. |
| P41567 | EIF1_HUMAN | | cytoplasm. |
| P42025 | ACTY_HUMAN | | cytoplasm. |
| P42331 | RHG25_HUMAN | | cytoplasm. |
| P42336 | PK3CA_HUMAN | | cytoplasm. |
| P42338 | PK3CB_HUMAN | | cytoplasm. |
| P42345 | FRAP_HUMAN | | cytoplasm. |
| P42356 | PI4KA_HUMAN | | cytoplasm. |
| P42357 | HUTH_HUMAN | | cytoplasm. |
| P42566 | EP15_HUMAN | cytoplasm (by similarity). cell membrane (by similarity). recruited to the plasma membrane upon egfr activation and localizes to coated pits (by similarity). | cytoplasm. |
| P42568 | AF9_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| P42575 | CASP2_HUMAN | | cytoplasm. |
| P42658 | DPP6_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| P42677 | RS27_HUMAN | | cytoplasm. |
| P42680 | TEC_HUMAN | cytoplasm (probable). | cytoplasm. |
| P42681 | TXK_HUMAN | cytoplasm (probable). | cytoplasm. |
| P42685 | FRK_HUMAN | cytoplasm (probable). | cytoplasm. |
| P42694 | HELZ_HUMAN | nucleus (probable). | nucleus. |
| P42695 | K0056_HUMAN | | cytoplasm. |
| P42704 | LPPRC_HUMAN | | cytoplasm. |
| P42766 | RL35_HUMAN | | cytoplasm. mitochondrion. |
| P42768 | WASP_HUMAN | | cytoplasm. |
| P42771 | CD2A1_HUMAN | | nucleus. |
| P42772 | CDN2B_HUMAN | | cytoplasm. |
| P42773 | CDN2C_HUMAN | | nucleus. |
| P42857 | NSG1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | cytoplasm. endosome. |
| P42898 | MTHR_HUMAN | | cytoplasm. |
| P43034 | LIS1_HUMAN | redistributes to axons during neuronal development. also localizes to the microtubules of the manchette in elongating spermatids and to the meiotic spindle in spermatocytes (by similarity). cytoplasm. localizes to the plus end of microtubules and to the centrosome. may localize to the nuclear membrane. | cytoplasm. |

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|--------|-------------|--|---------------------------|
| P43155 | CACP_HUMAN | endoplasmic reticulum (potential). peroxisome (potential). mitochondrion; mitochondrial inner membrane; matrix side (potential). | mitochondrion. |
| P43166 | CAH7_HUMAN | cytoplasm (probable). | cytoplasm. |
| P43234 | CATO_HUMAN | | lysosome. |
| P43246 | MSH2_HUMAN | nucleus (potential). | nucleus. |
| P43320 | CRBB2_HUMAN | | centriole. |
| P43351 | RAD52_HUMAN | nucleus (potential). | nucleus. |
| P43353 | AL3B1_HUMAN | | cytoplasm. mitochondrion. |
| P43354 | NR4A2_HUMAN | nucleus (probable). | nucleus. |
| P43356 | MAGA2_HUMAN | | cytoplasm. |
| P43357 | MAGA3_HUMAN | | cytoplasm. |
| P43358 | MAGA4_HUMAN | | centriole. cytoplasm. |
| P43359 | MAGA5_HUMAN | | centriole. cytoplasm. |
| P43360 | MAGA6_HUMAN | | centriole. cytoplasm. |
| P43361 | MAGA8_HUMAN | | centriole. cytoplasm. |
| P43362 | MAGA9_HUMAN | | centriole. cytoplasm. |
| P43363 | MAGAA_HUMAN | | nucleus. |
| P43365 | MAGAC_HUMAN | | centriole. cytoplasm. |
| P43366 | MAGB1_HUMAN | | nucleus. |
| P43378 | PTN9_HUMAN | cytoplasm (probable). | cytoplasm. |
| P43403 | ZAP70_HUMAN | | cytoplasm. |
| P43405 | KSYK_HUMAN | | cytoplasm. |
| P43487 | RANG_HUMAN | | cytoplasm. nucleus. |
| P43490 | NAMPT_HUMAN | cytoplasm (by similarity). | secreted protein. |
| P45378 | TNNT3_HUMAN | | cytoplasm. |
| P45379 | TNNT2_HUMAN | | cytoplasm. |
| P45381 | ACY2_HUMAN | | lysosome. |
| P45452 | MMP13_HUMAN | | secreted protein. |
| P45974 | UBP5_HUMAN | | cytoplasm. nucleus. |
| P45983 | MK08_HUMAN | | cytoplasm. |
| P45984 | MK09_HUMAN | | cytoplasm. |
| P45985 | MP2K4_HUMAN | | cytoplasm. |
| P46019 | KPB2_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| P46020 | KPB1_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| P46060 | RGP1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P46108 | CRK_HUMAN | cytoplasm (by similarity). cell membrane (by similarity). translocated to the plasma membrane upon cell adhesion (by similarity). | cytoplasm. |
| P46109 | CRKL_HUMAN | | cytoplasm. |
| P46379 | BAT3_HUMAN | | cytoplasm. |
| P46527 | CDN1B_HUMAN | nucleus (by similarity). cytoplasm (by similarity). nuclear and cytoplasmic in quiescent cells. upon cell cycle progression, mostly cytoplasmic (by similarity). | cytoplasm. nucleus. |
| P46531 | NOTC1_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). nicd: nucleus (by similarity). following proteolytical processing nicd is translocated to the nucleus (by similarity). | nucleus. |
| P46597 | HIOM_HUMAN | | cytoplasm. |

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| P46695 | IEX1_HUMAN | membrane; single-pass type ii membrane protein (potential). | endosome; membrane. |
| P46734 | MP2K3_HUMAN | | cytoplasm. |
| P46736 | BRCC3_HUMAN | | cytoplasm. |
| P46776 | RL27A_HUMAN | | cytoplasm. |
| P46778 | RL21_HUMAN | | cytoplasm. |
| P46779 | RL28_HUMAN | | cytoplasm. |
| P46781 | RS9_HUMAN | | cytoplasm. |
| P46782 | RS5_HUMAN | | cytoplasm. |
| P46821 | MAP1B_HUMAN | | cytoplasm. |
| P46926 | GNPI_HUMAN | cytoplasm (by similarity). | lysosome. |
| P46934 | NEDD4_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P46937 | YAP1_HUMAN | | cytoplasm. |
| P46976 | GLYG_HUMAN | | centriole. cytoplasm. |
| P47224 | MSS4_HUMAN | | cytoplasm. |
| P47755 | CAZA2_HUMAN | | cytoplasm. |
| P47756 | CAPZB_HUMAN | | cytoplasm. |
| P47813 | IF1AX_HUMAN | | cytoplasm. |
| P47895 | AL1A3_HUMAN | cytoplasm (by similarity). | mitochondrion. |
| P47914 | RL29_HUMAN | | cytoplasm. |
| P47929 | LEG7_HUMAN | cytoplasm. nucleus. secreted protein (potential). may be secreted by a non-classical secretory pathway. | nucleus. |
| P47944 | MT4_HUMAN | | cytoplasm. |
| P47972 | NPTX2_HUMAN | secreted protein (by similarity). | endoplasmic reticulum. secreted protein. |
| P47974 | TISD_HUMAN | nucleus (potential). | nucleus. |
| P48059 | PINC_HUMAN | cytoplasm (potential). | cytoplasm. nucleus. |
| P48060 | GLIP1_HUMAN | | endosome; membrane. |
| P48304 | REG1B_HUMAN | | secreted protein. |
| P48357 | LEPR_HUMAN | cell membrane; single-pass type i membrane protein. isoform e: secreted protein (probable). | membrane. |
| P48378 | RFX2_HUMAN | nucleus (probable). | nucleus. |
| P48380 | RFX3_HUMAN | nucleus (probable). | nucleus. |
| P48426 | PI52A_HUMAN | | cytoplasm. |
| P48436 | SOX9_HUMAN | nucleus (potential). | nucleus. |
| P48444 | COPD_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). the coatomer is cytoplasmic or polymerized on the cytoplasmic side of the Golgi, as well as on the vesicles/buds originating from it (by similarity). | endosome. |
| P48448 | AL3B2_HUMAN | | cytoplasm. mitochondrion. |
| P48449 | ERG7_HUMAN | | cytoplasm. |
| P48454 | PP2BC_HUMAN | | cytoplasm. |
| P48506 | GSH1_HUMAN | | cytoplasm. |
| P48507 | GSH0_HUMAN | | mitochondrion. |
| P48539 | PEP19_HUMAN | | cytoplasm. secreted protein. nucleus. |
| P48553 | TMEM1_HUMAN | Golgi apparatus; cis-Golgi network (by similarity). | endoplasmic reticulum. |
| P48556 | PSD8_HUMAN | | cytoplasm. |
| P48595 | SPB10_HUMAN | cytoplasm (by similarity). | secreted protein. |
| P48637 | GSHB_HUMAN | | cytoplasm. |

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|--------|-------------|--|------------------------|
| P48666 | K2C6C_HUMAN | | cytoplasm. |
| P48668 | K2C6E_HUMAN | | cytoplasm. |
| P48681 | NEST_HUMAN | | cytoplasm. |
| P48736 | PK3CG_HUMAN | | cytoplasm. |
| P48739 | PIPNB_HUMAN | cytoplasm (by similarity). Golgi apparatus (by similarity). | cytoplasm. |
| P48740 | MASP1_HUMAN | | secreted protein. |
| P48742 | LHX1_HUMAN | nucleus (probable). | nucleus. |
| P48775 | T23O_HUMAN | | endoplasmic reticulum. |
| P48788 | TNNI2_HUMAN | | cytoplasm. |
| P49006 | MRP_HUMAN | | membrane. |
| P49023 | PAXI_HUMAN | | cytoplasm. |
| P49116 | TR4_HUMAN | nucleus (probable). | nucleus. |
| P49137 | MAPK2_HUMAN | | cytoplasm. |
| P49189 | AL9A1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P49207 | RL34_HUMAN | | cytoplasm. |
| P49221 | TGM4_HUMAN | | cytoplasm. |
| P49223 | SPIT3_HUMAN | secreted protein (probable). | secreted protein. |
| P49247 | RPIA_HUMAN | | cytoplasm. |
| P49279 | NRAM1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| P49281 | NRAM2_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| P49336 | CDK8_HUMAN | | nucleus. |
| P49354 | PFTA_HUMAN | | cytoplasm. |
| P49356 | PFTB_HUMAN | | cytoplasm. |
| P49366 | DHYS_HUMAN | | centriole. cytoplasm. |
| P49407 | ARRB1_HUMAN | | cytoplasm. |
| P49419 | AL7A1_HUMAN | | mitochondrion. |
| P49427 | UB2R1_HUMAN | | cytoplasm. |
| P49441 | INPP_HUMAN | | cytoplasm. |
| P49459 | UBE2A_HUMAN | | nucleus. |
| P49585 | PCY1A_HUMAN | cytoplasm; cytosol (by similarity). membrane; peripheral membrane protein (by similarity). it can interconvert between an inactive cytosolic form and an active membrane-bound form (by similarity). | membrane. |
| P49588 | SYA_HUMAN | cytoplasm (potential). | cytoplasm. |
| P49590 | SYHH_HUMAN | | cytoplasm. |
| P49591 | SYS_HUMAN | | cytoplasm. |
| P49593 | PPM1F_HUMAN | | cytoplasm. |
| P49641 | MA2A2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endosome. Golgi. |
| P49642 | PRI1_HUMAN | | nucleus. |
| P49643 | PRI2_HUMAN | | nucleus. |
| P49646 | YYY1_HUMAN | | nucleus. |
| P49662 | CASP4_HUMAN | | cytoplasm. |
| P49674 | KC1E_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P49703 | ARL4D_HUMAN | nucleus; nucleolus (by similarity). | Golgi. nucleus. |
| P49736 | MCM2_HUMAN | nucleus (probable). | nucleus. |
| P49746 | TSP3_HUMAN | | secreted protein. |
| P49750 | YLPM1_HUMAN | nucleus; nucleoplasm; nuclear speckle (by similarity). | nucleus. |
| P49754 | VPS41_HUMAN | | cytoplasm. |

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|--------|-------------|---|------------------------|
| P49755 | TMEDA_HUMAN | Golgi apparatus; Golgi membrane; single-pass type i membrane protein; luminal side (by similarity). | endoplasmic reticulum. |
| P49756 | RBM25_HUMAN | | nucleus. |
| P49760 | CLK2_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| P49761 | CLK3_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| P49770 | EI2BB_HUMAN | | cytoplasm. |
| P49788 | TIG1_HUMAN | membrane; single-pass type ii membrane protein (potential). | endosome; membrane. |
| P49798 | RGS4_HUMAN | | cytoplasm. |
| P49802 | RGS7_HUMAN | | cytoplasm. |
| P49840 | GSK3A_HUMAN | | cytoplasm. |
| P49841 | GSK3B_HUMAN | | cytoplasm. |
| P49895 | IOD1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| P49901 | MCSP_HUMAN | cytoplasm. becomes associated with the spermatid mitochondrion capsule at step 16 of spermatogenesis (by similarity). | cytoplasm. |
| P49903 | SPS1_HUMAN | | cytoplasm. |
| P49910 | ZN165_HUMAN | nucleus (potential). | nucleus. |
| P49918 | CDN1C_HUMAN | nucleus (by similarity). | nucleus. |
| P49959 | MRE11_HUMAN | nucleus (by similarity). localizes to discrete nuclear foci after treatment with genotoxic agents (by similarity). | nucleus. |
| P49961 | ENP1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| P50053 | KHK_HUMAN | | cytoplasm. |
| P50148 | GNAQ_HUMAN | | cytoplasm. |
| P50150 | GBG4_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| P50151 | GBG10_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| P50221 | MEOX1_HUMAN | nucleus (potential). | nucleus. |
| P50222 | MEOX2_HUMAN | nucleus (potential). | nucleus. |
| P50238 | CRIP1_HUMAN | | cytoplasm. |
| P50281 | MMP14_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P50290 | BB1_HUMAN | | cytoplasm. nucleus. |
| P50336 | PPOX_HUMAN | mitochondrion; mitochondrial inner membrane; peripheral membrane protein; intermembrane side (by similarity). | mitochondrion. |
| P50395 | GDIB_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | cytoplasm. |
| P50458 | LHX2_HUMAN | nucleus (probable). | nucleus. |
| P50461 | CSRP3_HUMAN | nucleus (potential). cytoplasm. cytoskeleton (potential). nuclear, associates with the actin cytoskeleton | nucleus. |
| P50479 | PDLI4_HUMAN | | cytoplasm. |
| P50502 | F10A1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P50549 | ETV1_HUMAN | nucleus (probable). | nucleus. |
| P50553 | ASCL1_HUMAN | nucleus (probable). | nucleus. |
| P50579 | AMPM2_HUMAN | | cytoplasm. |
| P50583 | AP4A_HUMAN | | cytoplasm. |
| P50591 | TNF10_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |

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|--------|-------------|--|-----------------------------|
| P50607 | TUB_HUMAN | | centriole. |
| P50616 | TOB1_HUMAN | | cytoplasm. |
| P50851 | LRBA_HUMAN | | centriole. cytoplasm. |
| P50876 | UB7I4_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| P50914 | RL14_HUMAN | | cytoplasm. |
| P50995 | ANX11_HUMAN | cytoplasm (by similarity). nucleus (by similarity). found throughout the nucleoplasm at interphase and at the mitotic phase is concentrated at the loop-like structure around the mitotic apparatus (by similarity). | cytoplasm. nucleus. |
| P51148 | RAB5C_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (by similarity). endosome; early endosome; early endosomal membrane; lipid-anchor (by similarity). | Golgi. |
| P51151 | RAB9A_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| P51157 | RAB28_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| P51160 | PDE6C_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| P51178 | PLCD1_HUMAN | | secreted protein. |
| P51397 | DAP1_HUMAN | | cytoplasm. |
| P51449 | RORG_HUMAN | nucleus (probable). | nucleus. |
| P51451 | BLK_HUMAN | | cytoplasm. |
| P51452 | DUS3_HUMAN | | cytoplasm. |
| P51504 | ZNF80_HUMAN | nucleus (probable). | nucleus. |
| P51508 | ZNF81_HUMAN | nucleus (probable). | nucleus. |
| P51511 | MMP15_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P51512 | MMP16_HUMAN | isoform long: cell membrane; single-pass type i membrane protein (potential). isoform short: secreted protein. cell surface. localized at the cell surface of melanoma cells. | secreted protein; membrane. |
| P51522 | ZNF83_HUMAN | nucleus (probable). | nucleus. |
| P51523 | ZNF84_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P51570 | GALK1_HUMAN | | cytoplasm. |
| P51587 | BRCA2_HUMAN | | nucleus. |
| P51606 | RENBP_HUMAN | | centriole. cytoplasm. |
| P51617 | IRAK1_HUMAN | | cytoplasm. |
| P51636 | CAV2_HUMAN | Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). membrane protein of caveolae. potential hairpin-like structure in the membrane (by similarity). | membrane. |
| P51649 | SSDH_HUMAN | | mitochondrion. |
| P51654 | GPC3_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| P51665 | PSD7_HUMAN | | cytoplasm. |
| P51668 | UB2D1_HUMAN | | cytoplasm. |
| P51689 | ARSD_HUMAN | lysosome (potential). | lysosome. |
| P51692 | STA5B_HUMAN | cytoplasm (by similarity). nucleus (by similarity). translocated into the nucleus in response to phosphorylation (by similarity). | cytoplasm. |
| P51786 | ZN157_HUMAN | nucleus (potential). | nucleus. |

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|--------|-------------|--|------------------------|
| P51805 | PLXA3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P51808 | DYLT3_HUMAN | | centriole. cytoplasm. |
| P51809 | SYBL1_HUMAN | membrane; single-pass type iv membrane protein (probable). | endoplasmic reticulum. |
| P51811 | XK_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| P51812 | KS6A3_HUMAN | | cytoplasm. |
| P51813 | BMX_HUMAN | cytoplasm (probable). | cytoplasm. |
| P51814 | ZNF41_HUMAN | nucleus (potential). | nucleus. |
| P51815 | ZNF75_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P51816 | AFF2_HUMAN | | nucleus. |
| P51817 | PRKX_HUMAN | | cytoplasm. |
| P51825 | AFF1_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| P51854 | TKT2_HUMAN | | cytoplasm. |
| P51861 | CDR1_HUMAN | | endoplasmic reticulum. |
| P51864 | TDGF2_HUMAN | | secreted protein. |
| P51878 | CASP5_HUMAN | | cytoplasm. |
| P51884 | LUM_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| P51911 | CNN1_HUMAN | | cytoskeleton. |
| P51956 | NEK3_HUMAN | | nucleus. |
| P51957 | NEK4_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| P51965 | UB2E1_HUMAN | | cytoplasm. |
| P51991 | ROA3_HUMAN | nucleus (by similarity). component of ribonucleosomes (by similarity). | nucleus. |
| P52179 | MYOM1_HUMAN | | cytoskeleton. |
| P52198 | RND2_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). colocalizes with racgap1 in Golgi-derived proacrosomal vesicles and the acrosome. | cytoplasm. Golgi. |
| P52209 | 6PGD_HUMAN | | cytoplasm. |
| P52306 | GDS1_HUMAN | | cytoplasm. |
| P52333 | JAK3_HUMAN | intracytoplasmic membrane; peripheral membrane protein. wholly intracellular, possibly membrane associated (by similarity). | cytoplasm. |
| P52429 | DGKE_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. |
| P52564 | MP2K6_HUMAN | | cytoplasm. |
| P52732 | KIF11_HUMAN | | cytoplasm. nucleus. |
| P52735 | VAV2_HUMAN | | cytoplasm. |
| P52736 | ZN133_HUMAN | nucleus (potential). | nucleus. |
| P52737 | ZN136_HUMAN | nucleus (potential). | nucleus. |
| P52738 | ZN140_HUMAN | nucleus (potential). | nucleus. |
| P52739 | ZN131_HUMAN | nucleus (potential). | nucleus. |
| P52740 | ZN132_HUMAN | nucleus (potential). | nucleus. |
| P52741 | ZN134_HUMAN | nucleus (potential). | nucleus. |
| P52742 | ZN135_HUMAN | nucleus (potential). | nucleus. |
| P52743 | ZN137_HUMAN | nucleus (potential). | nucleus. |
| P52746 | ZN142_HUMAN | nucleus (potential). | nucleus. |
| P52747 | ZN143_HUMAN | nucleus (potential). | nucleus. |
| P52756 | RBM5_HUMAN | nucleus (potential). | nucleus. |
| P52757 | CHIO_HUMAN | membrane; peripheral membrane protein (potential). | cytoplasm. |
| P52788 | SPSY_HUMAN | | centriole. cytoplasm. |

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|--------|-------------|---|-----------------------------------|
| P52789 | HXK2_HUMAN | mitochondrion; mitochondrial outer membrane. its hydrophobic n-terminal sequence may be involved in membrane binding (by similarity). | cytoplasm. |
| P52790 | HXK3_HUMAN | | cytoplasm. |
| P52798 | EFNA4_HUMAN | isoform 1: cell membrane; lipid-anchor; gpi- anchor. isoform 2: secreted protein (probable). | membrane. |
| P52824 | DGKQ_HUMAN | | cytoplasm. |
| P52849 | NDST2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endosome. |
| P52895 | AK1C2_HUMAN | cytoplasm (potential). | cytoplasm. |
| P52907 | CAZA1_HUMAN | | cytoplasm. |
| P52943 | CRIP2_HUMAN | | cytoplasm. nucleus. |
| P52951 | GBX2_HUMAN | nucleus (probable). | nucleus. |
| P52952 | NKX25_HUMAN | nucleus (probable). | nucleus. |
| P52954 | LBX1_HUMAN | nucleus (probable). | nucleus. |
| P53365 | ARFP2_HUMAN | | cytoplasm. |
| P53367 | ARFP1_HUMAN | | cytoplasm. secreted protein. |
| P53370 | NUDT6_HUMAN | | cytoplasm. |
| P53384 | NUBP1_HUMAN | | cytoplasm. nucleus. |
| P53420 | CO4A4_HUMAN | cell surface (potential). | secreted protein. |
| P53582 | AMPM1_HUMAN | | cytoplasm. |
| P53602 | ERG19_HUMAN | | cytoplasm. |
| P53609 | PGT1_HUMAN | | cytoplasm. nucleus. |
| P53611 | PGTB_HUMAN | | cytoplasm. |
| P53618 | COPB_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). the coatomer is cytoplasmic or polymerized on the cytoplasmic side of the Golgi, as well as on the vesicles/buds originating from it (by similarity). | Golgi. |
| P53621 | COPA_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). the coatomer is cytoplasmic or polymerized on the cytoplasmic side of the Golgi, as well as on the vesicles/buds originating from it (by similarity). xenin: | cytoplasm. endoplasmic reticulum. |
| P53672 | CRBA2_HUMAN | | cytoplasm. |
| P53673 | CRBA4_HUMAN | | cytoplasm. |
| P53674 | CRBB1_HUMAN | | cytoplasm. |
| P53675 | CLH2_HUMAN | cytoplasmic face of coated pits and vesicles (by similarity). | cytoplasm. |
| P53677 | AP3M2_HUMAN | | Golgi. |
| P53701 | CCHL_HUMAN | mitochondrion; mitochondrial inner membrane (potential). | mitochondrion. |
| P53801 | PTTG_HUMAN | membrane; single-pass type i membrane protein (potential). ref.2: cytoplasm. | cytoplasm. |
| P53804 | TTC3_HUMAN | | cytoplasm. nucleus. |
| P53805 | CCP1_HUMAN | | nucleus. |
| P53816 | HRSL3_HUMAN | | membrane. |
| P53990 | K0174_HUMAN | | cytoplasm. |

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| P54253 | ATX1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| P54257 | HAP1_HUMAN | | cytoplasm. |
| P54259 | ATN1_HUMAN | | cytoplasm. nucleus. |
| P54277 | PMS1_HUMAN | nucleus (potential). | nucleus. |
| P54284 | CACB3_HUMAN | | membrane. |
| P54296 | MYOM2_HUMAN | | cytoskeleton. |
| P54315 | LIPR1_HUMAN | secreted protein (potential). | secreted protein. |
| P54368 | OAZ1_HUMAN | | cytoplasm. secreted protein. |
| P54578 | UBP14_HUMAN | | nucleus. |
| P54619 | AAKG1_HUMAN | | cytoplasm. |
| P54646 | AAPK2_HUMAN | | cytoplasm. |
| P54652 | HSP72_HUMAN | | cytoplasm. |
| P54710 | ATNG_HUMAN | membrane; single-pass type iii membrane protein (potential). | membrane. |
| P54725 | RD23A_HUMAN | nucleus (probable). | nucleus. |
| P54727 | RD23B_HUMAN | nucleus. cytoplasm (by similarity). | nucleus. |
| P54750 | PDE1A_HUMAN | | cytoplasm. |
| P54792 | DVL1L_HUMAN | cytoplasm (potential). | cytoplasm. |
| P54793 | ARSF_HUMAN | secreted protein (potential). | lysosome. |
| P54802 | ANAG_HUMAN | | lysosome. |
| P54826 | GAS1_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | cytoplasm. |
| P54829 | PTN5_HUMAN | | cytoplasm. |
| P54840 | GYS2_HUMAN | | cytoplasm. |
| P54922 | ADPRH_HUMAN | | cytoplasm. |
| P55008 | AIF1_HUMAN | | centriole. nucleus. |
| P55010 | IF5_HUMAN | | cytoplasm. |
| P55036 | PSD4_HUMAN | | cytoplasm. |
| P55042 | RAD_HUMAN | | cytoplasm. Golgi. |
| P55055 | NR1H2_HUMAN | nucleus (potential). | nucleus. |
| P55061 | BI1_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| P55082 | MFAP3_HUMAN | cell membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| P55103 | INHBC_HUMAN | secreted protein (by similarity). | secreted protein. |
| P55145 | ARMET_HUMAN | secreted protein (potential). | centriole. cytoplasm. |
| P55160 | NCKPL_HUMAN | | membrane. |
| P55197 | AF10_HUMAN | nucleus (potential). | nucleus. |
| P55198 | AF17_HUMAN | nucleus (potential). | nucleus. |
| P55201 | BRPF1_HUMAN | nucleus (probable). | nucleus. |
| P55211 | CASP9_HUMAN | | cytoplasm. |
| P55259 | GP2_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). secreted protein (by similarity). secreted after cleavage in the pancreatic juice. | secreted protein. |
| P55263 | ADK_HUMAN | | cytoplasm. |
| P55269 | BAXD_HUMAN | cytoplasm (potential). | cytoplasm. |
| P55273 | CDN2D_HUMAN | | cytoplasm. nucleus. |
| P55316 | FOXGA_HUMAN | nucleus (potential). | nucleus. |
| P55327 | TPD52_HUMAN | | cytoplasm. |
| P55345 | ANM2_HUMAN | | nucleus. |
| P55347 | PKNX1_HUMAN | nucleus (probable). | nucleus. |
| P55735 | SEC13_HUMAN | | cytoplasm. |
| P55786 | PSA_HUMAN | cytoplasm (potential). nucleus (potential). | cytoplasm. |
| P55789 | ALR_HUMAN | | cytoplasm. |

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| P55808 | XG_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P55822 | SH3BG_HUMAN | | cytoplasm. |
| P55884 | IF39_HUMAN | | cytoplasm. nucleus. |
| P55916 | UCP3_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| P55957 | BID_HUMAN | cytoplasm (by similarity). when uncleaved, it is predominantly cytoplasmic. mitochondrion; mitochondrial membrane (by similarity). p15 bid translocates to mitochondria as an integral membrane protein. p13 and p22 bid are associated with the mitochondrial membrane (by similarity). isoform 1, isoform 3: cytoplasm. isoform 2: mitochondrion; mitochondrial membrane. a significant proportion of isoform 2 localizes to mitochondria, it may be cleaved constitutively. | cytoplasm. mitochondrion. |
| P56134 | ATPK_HUMAN | | mitochondrion. |
| P56159 | GFRA1_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | cytoplasm. secreted protein; membrane. |
| P56177 | DLX1_HUMAN | nucleus (potential). | nucleus. |
| P56178 | DLX5_HUMAN | nucleus (potential). | nucleus. |
| P56179 | DLX6_HUMAN | nucleus (potential). | nucleus. |
| P56192 | SYM_HUMAN | cytoplasm (probable). | cytoplasm. |
| P56202 | CATW_HUMAN | | lysosome. |
| P56211 | ARP19_HUMAN | cytoplasm (by similarity). | secreted protein. |
| P56270 | MAZ_HUMAN | nucleus (probable). in brains of alzheimer disease patients, present in a plaque-like structures. | nucleus. |
| P56278 | MTCPB_HUMAN | | secreted protein. |
| P56378 | 68MP_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| P56385 | ATP5I_HUMAN | | mitochondrion. |
| P56470 | LEG4_HUMAN | | membrane. |
| P56539 | CAV3_HUMAN | Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). membrane protein of caveolae. potential hairpin-like structure in the membrane (by similarity). | endosome; membrane. |
| P56545 | CTBP2_HUMAN | nucleus (potential). | nucleus. |
| P56555 | DSCR4_HUMAN | | centriole. cytoplasm. |
| P56557 | TM50B_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| P56559 | ARL4C_HUMAN | nucleus (by similarity). | Golgi. nucleus. |
| P56597 | NDK5_HUMAN | | cytoplasm. |
| P56645 | PER3_HUMAN | nucleus (potential). | nucleus. |
| P56696 | KCNQ4_HUMAN | cell membrane; basal cell membrane; multi- pass membrane protein. situated at the basal membrane of cochlear outer hair cells (by similarity). | membrane. |
| P56715 | RP1_HUMAN | | centriole. cytoplasm. |
| P56747 | CLD6_HUMAN | cell membrane; cell-cell junction; tight junction; multi-pass membrane protein (by similarity). | membrane. |

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| P56748 | CLD8_HUMAN | cell membrane; cell-cell junction; tight junction; multi-pass membrane protein (by similarity). | membrane. |
| P56846 | TCL6_HUMAN | | centriole. cytoplasm. |
| P56847 | TNG2_HUMAN | | centriole. cytoplasm. |
| P56851 | EP3B_HUMAN | secreted protein (potential). | secreted protein. |
| P56945 | BCAR1_HUMAN | cell membrane; cell-matrix junction; focal adhesion (by similarity). cytoplasm (by similarity). unphosphorylated form localizes in the cytoplasm and can move to the membrane upon tyrosine phosphorylation (by similarity). | cytoplasm. |
| P56962 | STX17_HUMAN | membrane; single-pass type iv membrane protein. appears to be associated with a membrane compartment, perhaps a subset of the er such as exit or entrance sites (by similarity). | Golgi. |
| P56975 | NRG3_HUMAN | pro-neuregulin-3, membrane-bound isoform: cell membrane; single-pass type i membrane protein (by similarity). neuregulin-3: secreted protein (by similarity). exists also as a proteolytically released soluble growth factor form. the membrane-bound form does not seem to be active (by similarity). | membrane. |
| P57043 | ILK2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P57052 | RBM11_HUMAN | | nucleus. |
| P57054 | PIGP_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| P57055 | DSCR6_HUMAN | | centriole. cytoplasm. |
| P57056 | CU032_HUMAN | | nucleus. |
| P57057 | GLPT_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| P57058 | HUNK_HUMAN | | cytoplasm. |
| P57059 | SN1L1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). may translocate to the nucleus on activation (by similarity). | cytoplasm. |
| P57060 | CU006_HUMAN | | centriole. cytoplasm. |
| P57071 | PRD15_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P57073 | SOX8_HUMAN | nucleus (potential). | nucleus. |
| P57076 | CU059_HUMAN | | nucleus. |
| P57077 | TAK1L_HUMAN | | cytoplasm. |
| P57078 | RIPK4_HUMAN | | cytoplasm. |
| P57081 | WDR4_HUMAN | | centriole. cytoplasm. |
| P57082 | TBX4_HUMAN | nucleus (potential). | nucleus. |
| P57086 | SCND1_HUMAN | nucleus (potential). | nucleus. |
| P57087 | JAM2_HUMAN | cell membrane; cell-cell junction; tight junction; single-pass type i membrane protein (by similarity). localized at tight junctions of both epithelial and endothelial cells (by similarity). | membrane. |
| P57088 | TMM33_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| P57105 | SYJ2B_HUMAN | mitochondrion; mitochondrial outer membrane (by similarity). | mitochondrion. |
| P57679 | EVC_HUMAN | | endosome; membrane. |

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| P57682 | KLF3_HUMAN | nucleus (probable). | nucleus. |
| P57729 | RAB38_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| P57730 | ICBR_HUMAN | | cytoplasm. |
| P57735 | RAB25_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| P57737 | CORO7_HUMAN | cytoplasm; cytosol (by similarity). predominantly cytosolic. membrane (by similarity). also found in membrane fractions. detected on vesicle-like cytoplasmic structures and on the cis-Golgi. not associated with actin filaments (by similarity). | Golgi. |
| P57738 | TCTA_HUMAN | | membrane. |
| P57739 | CLD2_HUMAN | cell membrane; cell-cell junction; tight junction; multi-pass membrane protein (by similarity). | membrane. |
| P57764 | GSDC1_HUMAN | | cytoplasm. |
| P57768 | SNX16_HUMAN | | cytoplasm. |
| P57771 | RGS8_HUMAN | | cytoplasm. |
| P57772 | SELB_HUMAN | cytoplasm (potential). nucleus (potential). | cytoplasm. |
| P57775 | FBXW4_HUMAN | | cytoplasm. |
| P57789 | KCNKA_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| P57796 | CABP4_HUMAN | presynapses from both types of photoreceptors: rod spherules and cone pedicles (by similarity). | cytoplasm. |
| P58004 | SESN2_HUMAN | | centriole. |
| P58005 | SESN3_HUMAN | | centriole. cytoplasm. |
| P58062 | ISK7_HUMAN | secreted protein (potential). | secreted protein. |
| P58107 | EPIPL_HUMAN | | cytoplasm. |
| P58166 | INHBE_HUMAN | secreted protein (by similarity). | secreted protein. |
| P58215 | LOXL3_HUMAN | secreted protein; extracellular space (potential). | secreted protein. |
| P58397 | ATS12_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| P58400 | NRX1B_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P58401 | NRX2B_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P58417 | NXPH1_HUMAN | secreted protein (potential). | nucleus. |
| P58418 | USH3A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| P58505 | CU058_HUMAN | | cytoplasm. |
| P58511 | CU051_HUMAN | | centriole. cytoplasm. |
| P58512 | CU067_HUMAN | | nucleus. |
| P58513 | CU042_HUMAN | | cytoplasm. |
| P58546 | MTPN_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P58549 | FXVD7_HUMAN | | membrane. |
| P58550 | FXVD8_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P58556 | CU069_HUMAN | | nucleus. |
| P58557 | CU057_HUMAN | | cytoplasm. |
| P58658 | CU063_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |

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| P58743 | S26A5_HUMAN | cell membrane; multi-pass membrane protein (by similarity). lateral wall of outer hair cells (by similarity). | membrane. |
| P58753 | TIRAP_HUMAN | cytoplasm (by similarity). | membrane. |
| P58872 | RHBD4_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| P59020 | DSCR9_HUMAN | | centriole. cytoplasm. |
| P59022 | DSC10_HUMAN | | centriole. cytoplasm. |
| P59025 | RTP1_HUMAN | cell membrane; single-pass type iii membrane protein (by similarity). effective cell surface expression depends upon interaction with olfactory receptors (by similarity). | endosome; membrane. |
| P59036 | CU082_HUMAN | | cytoplasm. |
| P59037 | CU084_HUMAN | | secreted protein. |
| P59043 | CU100_HUMAN | | membrane. |
| P59045 | NAL11_HUMAN | | cytoplasm. |
| P59047 | NALP5_HUMAN | | cytoplasm. |
| P59051 | CU087_HUMAN | | nucleus. |
| P59052 | CU088_HUMAN | | membrane. |
| P59074 | CG301_HUMAN | | cytoplasm. |
| P59088 | CU085_HUMAN | | endosome; membrane. |
| P59089 | CU086_HUMAN | | nucleus. |
| P59090 | CU090_HUMAN | | secreted protein. |
| P59091 | CU093_HUMAN | | secreted protein. |
| P59092 | CU094_HUMAN | | membrane. |
| P59093 | CU099_HUMAN | | nucleus. |
| P59095 | STAR6_HUMAN | | secreted protein. |
| P59190 | RAB15_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| P59282 | P25B_HUMAN | | cytoplasm. |
| P59510 | ATS20_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| P59646 | FXYD4_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P59768 | GBG2_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| P59773 | U258_HUMAN | | endosome; membrane. |
| P59780 | AP3S2_HUMAN | Golgi apparatus. cytoplasmic vesicle; cytoplasmic vesicle membrane; peripheral membrane protein; cytoplasmic side (by similarity). component of the coat surrounding the cytoplasmic face of coated vesicles located at the Golgi complex (by similarity). | cytoplasm. |
| P59796 | GPX6_HUMAN | secreted protein (by similarity). | cytoplasm. |
| P59797 | SELV_HUMAN | | cytoplasm. |
| P59817 | SUHW1_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| P59826 | LPLC3_HUMAN | secreted protein (by similarity). ref.1: cytoplasm. | cytoplasm. |
| P59827 | LPLC4_HUMAN | secreted protein (by similarity). | cytoplasm. |
| P59861 | DB131_HUMAN | secreted protein (potential). | secreted protein. |
| P59894 | DCDC1_HUMAN | | cytoplasm. |
| P59910 | DNJBD_HUMAN | | cytoplasm. |
| P59923 | ZN445_HUMAN | nucleus (potential). | nucleus. |
| P59990 | KR121_HUMAN | | cytoplasm. |

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|--------|-------------|--|---------------------------|
| P59991 | KR122_HUMAN | | cytoplasm. |
| P59998 | ARPC4_HUMAN | | cytoplasm. |
| P60002 | ELOF1_HUMAN | nucleus (by similarity). | nucleus. |
| P60006 | CK051_HUMAN | | cytoplasm. |
| P60014 | KR10A_HUMAN | | cytoplasm. |
| P60059 | SC61G_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| P60153 | RNAS9_HUMAN | secreted protein (probable). | secreted protein. |
| P60174 | TPIS_HUMAN | | cytoplasm. |
| P60321 | NANO2_HUMAN | | cytoplasm. |
| P60323 | NANO3_HUMAN | | cytoplasm. |
| P60328 | KR123_HUMAN | | cytoplasm. |
| P60329 | KR124_HUMAN | | cytoplasm. |
| P60331 | KR101_HUMAN | | cytoplasm. |
| P60368 | KR102_HUMAN | | cytoplasm. |
| P60369 | KR103_HUMAN | | cytoplasm. |
| P60370 | KR105_HUMAN | | cytoplasm. |
| P60371 | KR106_HUMAN | | cytoplasm. |
| P60372 | KR104_HUMAN | | cytoplasm. |
| P60409 | KR107_HUMAN | | cytoplasm. |
| P60410 | KR108_HUMAN | | cytoplasm. |
| P60411 | KR109_HUMAN | | cytoplasm. |
| P60412 | KR10B_HUMAN | | cytoplasm. |
| P60413 | KR10C_HUMAN | | cytoplasm. |
| P60507 | EFC1_HUMAN | | endosome; membrane. |
| P60508 | EFR1_HUMAN | | endosome; membrane. |
| P60509 | ERB1_HUMAN | | endosome; membrane. |
| P60520 | GBRL2_HUMAN | Golgi apparatus (by similarity). | cytoplasm. |
| P60602 | CT052_HUMAN | | endosome; membrane. |
| P60604 | UB2G2_HUMAN | | cytoplasm. |
| P60606 | CTXN1_HUMAN | | endosome; membrane. |
| P60608 | EFC2_HUMAN | | membrane. |
| P60660 | MYL6_HUMAN | | cytoplasm. |
| P60673 | PROF3_HUMAN | | cytoplasm. |
| P60827 | C1QT8_HUMAN | secreted protein (potential). | secreted protein. |
| P60842 | IF4A1_HUMAN | | nucleus. |
| P60866 | RS20_HUMAN | | cytoplasm. mitochondrion. |
| P60880 | SNP25_HUMAN | | cytoplasm. |
| P60891 | PRPS1_HUMAN | | cytoplasm. |
| P60893 | GPR85_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| P60896 | DSS1_HUMAN | | cytoplasm. |
| P60903 | S10AA_HUMAN | | cytoplasm. |
| P60953 | CDC42_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| P60981 | DEST_HUMAN | | cytoplasm. |
| P60983 | GMFB_HUMAN | | cytoplasm. |
| P60985 | U467_HUMAN | secreted protein (potential). | nucleus. |
| P61006 | RAB8A_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| P61009 | SPCS3_HUMAN | microsome; microsomal membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. |
| P61018 | RAB4B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |

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| P61024 | CKS1_HUMAN | | cytoplasm. |
| P61026 | RAB10_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| P61077 | UB2D3_HUMAN | | cytoplasm. |
| P61081 | UBC12_HUMAN | | cytoplasm. |
| P61086 | UBC1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P61088 | UBE2N_HUMAN | | nucleus. |
| P61106 | RAB14_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. endosome. Golgi. lysosome. |
| P61109 | ANRE_HUMAN | secreted protein (potential). | cytoplasm. |
| P61129 | ZC3H6_HUMAN | | cytoplasm. nucleus. |
| P61158 | ARP3_HUMAN | | cytoplasm. |
| P61160 | ARP2_HUMAN | | cytoplasm. |
| P61163 | ACTZ_HUMAN | | cytoplasm. |
| P61165 | CK010_HUMAN | | membrane. |
| P61204 | ARF3_HUMAN | | Golgi. |
| P61225 | RAP2B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| P61236 | YPEL3_HUMAN | | nucleus. |
| P61247 | RS3A_HUMAN | | cytoplasm. mitochondrion. |
| P61254 | RL26_HUMAN | | mitochondrion. |
| P61266 | STX1C_HUMAN | membrane; single-pass type iv membrane protein (potential). | Golgi; membrane. |
| P61289 | PSME3_HUMAN | | cytoplasm. |
| P61296 | HAND2_HUMAN | | nucleus. |
| P61313 | RL15_HUMAN | | mitochondrion. |
| P61326 | MGN_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| P61328 | FGF12_HUMAN | nucleus (probable). | centriole. cytoplasm. |
| P61353 | RL27_HUMAN | | mitochondrion. |
| P61421 | VA0D_HUMAN | | mitochondrion. |
| P61513 | RL37A_HUMAN | | mitochondrion. |
| P61549 | ENH5_HUMAN | | cytoplasm. |
| P61550 | ENT1_HUMAN | | endosome; membrane. |
| P61565 | ENK1_HUMAN | | endosome; membrane. |
| P61566 | ENK11_HUMAN | | endosome; membrane. |
| P61567 | ENK12_HUMAN | | endosome; membrane. |
| P61568 | ENK13_HUMAN | | endosome; membrane. |
| P61569 | ENK16_HUMAN | | centriole. |
| P61570 | ENK17_HUMAN | | endosome; membrane. |
| P61571 | REC1_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and the cytoplasm. when in the nucleus, resides in the nucleolus (by | cytoplasm. |
| P61572 | REC3_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and the cytoplasm. when in the nucleus, resides in the nucleolus (by | cytoplasm. |
| P61573 | REC4_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and the cytoplasm. when in the nucleus, resides in the nucleolus (by | cytoplasm. |
| P61574 | REC5_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and the cytoplasm. when in the nucleus, resides in the nucleolus (by | cytoplasm. |

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| P61575 | REC6_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and the cytoplasm. when in the nucleus, resides in the nucleolus (by | cytoplasm. |
| P61576 | REC9_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and the cytoplasm. when in the nucleus, resides in the nucleolus (by | cytoplasm. |
| P61577 | REC15_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and the cytoplasm. when in the nucleus, resides in the nucleolus (by | cytoplasm. |
| P61578 | REC16_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and the cytoplasm. when in the nucleus, resides in the nucleolus (by | cytoplasm. |
| P61579 | REC17_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and cytoplasm. when in the nucleus, resides in the nucleolus (by | cytoplasm. |
| P61580 | NP10_HUMAN | nucleus (by similarity). when overexpressed (by similarity). | centriole. |
| P61582 | NP12_HUMAN | nucleus (by similarity). when overexpressed (by similarity). | centriole. |
| P61583 | NP8_HUMAN | nucleus (by similarity). when overexpressed (by similarity). | centriole. |
| P61599 | NAT5_HUMAN | | cytoplasm. |
| P61601 | NCALD_HUMAN | | cytoplasm. |
| P61626 | LYSC_HUMAN | | secreted protein. |
| P61647 | SIA8F_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |
| P61803 | DAD1_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| P61916 | NPC2_HUMAN | secreted protein (potential). | cytoplasm. |
| P61923 | COPZ1_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). the coatomer is cytoplasmic or polymerized on the cytoplasmic side of the Golgi, as well as on the vesicles/buds originating from it (by similarity). | endosome. |
| P61925 | IPKA_HUMAN | | cytoplasm. |
| P61927 | RL37_HUMAN | | cytoplasm. |
| P61952 | GBG11_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| P61962 | WDR68_HUMAN | cytoplasm (probable). | cytoplasm. |
| P61964 | WDR5_HUMAN | | cytoplasm. |
| P61968 | LMO4_HUMAN | | nucleus. |
| P61981 | 1433G_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P62068 | UBP46_HUMAN | | nucleus. |
| P62070 | RRAS2_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (by similarity). inner surface of plasma membrane possibly with attachment requiring acylation of the c-terminal cysteine (by similarity with ras). | cytoplasm. |

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| P62079 | TSN5_HUMAN | membrane; multi-pass membrane protein (probable). | mitochondrion. |
| P62081 | RS7_HUMAN | | cytoplasm. mitochondrion. |
| P62136 | PP1A_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P62140 | PP1B_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P62158 | CALM_HUMAN | | cytoplasm. |
| P62195 | PRS8_HUMAN | cytoplasm (potential). nucleus (potential). | cytoplasm. |
| P62241 | RS8_HUMAN | | cytoplasm. |
| P62244 | RS15A_HUMAN | | mitochondrion. |
| P62249 | RS16_HUMAN | | cytoplasm. mitochondrion. |
| P62253 | UB2G1_HUMAN | | cytoplasm. |
| P62256 | UBE2H_HUMAN | | cytoplasm. |
| P62258 | 1433E_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P62263 | RS14_HUMAN | | cytoplasm. |
| P62266 | RS23_HUMAN | | cytoplasm. mitochondrion. |
| P62273 | RS29_HUMAN | | cytoplasm. mitochondrion. |
| P62277 | RS13_HUMAN | | mitochondrion. |
| P62280 | RS11_HUMAN | | cytoplasm. |
| P62310 | LSM3_HUMAN | nucleus (potential). | nucleus. |
| P62312 | LSM6_HUMAN | nucleus (potential). | nucleus. |
| P62324 | BTG1_HUMAN | | cytoplasm. nucleus. |
| P62330 | ARF6_HUMAN | | Golgi. |
| P62333 | PRS10_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| P62341 | SELT_HUMAN | | cytoplasm. |
| P62380 | TBPL1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| P62424 | RL7A_HUMAN | | mitochondrion. |
| P62508 | ERR3_HUMAN | nucleus (potential). | nucleus. |
| P62633 | CNBP_HUMAN | cytoplasm (by similarity). endoplasmic reticulum (by similarity). | cytoplasm. nucleus. |
| P62683 | GAK1_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |
| P62684 | GAK5_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |
| P62685 | GAK6_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |
| P62686 | GAK13_HUMAN | | centriole. |
| P62687 | GAK14_HUMAN | | centriole. |
| P62688 | GAK15_HUMAN | | cytoplasm. Golgi. |
| P62689 | GAK17_HUMAN | | cytoplasm. Golgi. |
| P62690 | GAK18_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |
| P62691 | GAK19_HUMAN | | cytoplasm. Golgi. |
| P62699 | YPEL5_HUMAN | | cytoplasm. |
| P62701 | RS4X_HUMAN | | cytoplasm. |
| P62706 | FBS1_HUMAN | secreted protein (by similarity). | secreted protein. |
| P62750 | RL23A_HUMAN | | cytoplasm. |
| P62753 | RS6_HUMAN | | cytoplasm. |
| P62760 | VISL1_HUMAN | | cytoplasm. |
| P62829 | RL23_HUMAN | | mitochondrion. |
| P62837 | UB2D2_HUMAN | | cytoplasm. |
| P62841 | RS15_HUMAN | | cytoplasm. mitochondrion. |
| P62847 | RS24_HUMAN | | cytoplasm. mitochondrion. |
| P62851 | RS25_HUMAN | | cytoplasm. mitochondrion. |
| P62854 | RS26_HUMAN | | cytoplasm. |

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| P62857 | RS28_HUMAN | | cytoplasm. mitochondrion. |
| P62861 | RS30_HUMAN | | cytoplasm. mitochondrion. |
| P62873 | GBB1_HUMAN | | nucleus. |
| P62879 | GBB2_HUMAN | | nucleus. |
| P62888 | RL30_HUMAN | | mitochondrion. |
| P62891 | RL39_HUMAN | | mitochondrion. |
| P62899 | RL31_HUMAN | | mitochondrion. |
| P62906 | RL10A_HUMAN | | mitochondrion. |
| P62910 | RL32_HUMAN | | mitochondrion. |
| P62913 | RL11_HUMAN | | cytoplasm. |
| P62952 | BC10_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| P62955 | CCG7_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| P62979 | RS27A_HUMAN | | cytoplasm. mitochondrion. |
| P62987 | RL40_HUMAN | | mitochondrion. |
| P62993 | GRB2_HUMAN | | cytoplasm. |
| P63000 | RAC1_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (by similarity). inner surface of plasma membrane possibly with attachment requiring prenylation of the c-terminal cysteine (by similarity). | Golgi. |
| P63092 | GNAS_HUMAN | | cytoplasm. |
| P63096 | GNAI1_HUMAN | | cytoplasm. |
| P63098 | CANB1_HUMAN | | cytoplasm. |
| P63117 | GFC1_HUMAN | cell membrane (potential). | cytoplasm. |
| P63118 | GAH2_HUMAN | cell membrane (potential). | membrane. |
| P63119 | VPK1_HUMAN | | secreted protein. |
| P63120 | VPK3_HUMAN | | secreted protein. |
| P63121 | VPK5_HUMAN | | secreted protein. |
| P63122 | VPK6_HUMAN | | secreted protein. |
| P63123 | VPK7_HUMAN | | secreted protein. |
| P63124 | VPK9_HUMAN | | secreted protein. |
| P63125 | VPK17_HUMAN | | secreted protein. |
| P63126 | GAK4_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |
| P63127 | VPK4_HUMAN | | secreted protein. |
| P63128 | POK4_HUMAN | | nucleus. |
| P63129 | VPK11_HUMAN | | secreted protein. |
| P63130 | GAK12_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |
| P63131 | VPK12_HUMAN | | secreted protein. |
| P63132 | POK5_HUMAN | | nucleus. |
| P63133 | POK6_HUMAN | | nucleus. |
| P63134 | POK9_HUMAN | | microsome. nucleus. |
| P63135 | POK12_HUMAN | | nucleus. |
| P63136 | POK17_HUMAN | | nucleus. |
| P63145 | GAK11_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |
| P63146 | UBE2B_HUMAN | cell membrane (by similarity). nucleus (by similarity). in peripheral neurons, expressed both at the plasma membrane and in nuclei (by similarity). | cytoplasm. |
| P63151 | 2ABA_HUMAN | | cytoplasm. |
| P63172 | DYLT1_HUMAN | | cytoplasm. |
| P63173 | RL38_HUMAN | | mitochondrion. |

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| P63208 | SKP1_HUMAN | | cytoplasm. |
| P63211 | GBG1_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| P63215 | GBG3_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| P63218 | GBG5_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| P63220 | RS21_HUMAN | | cytoplasm. mitochondrion. |
| P63241 | IF5A1_HUMAN | | cytoplasm. |
| P63244 | GBLP_HUMAN | | cytoplasm. |
| P63279 | UBC9_HUMAN | | cytoplasm. |
| P63316 | TNNC1_HUMAN | | cytoplasm. |
| P67870 | CSK2B_HUMAN | | cytoplasm. |
| P67936 | TPM4_HUMAN | | cytoplasm. |
| P68036 | UB2L3_HUMAN | | cytoplasm. |
| P68106 | FKB1B_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P68363 | TBAK_HUMAN | | centriole. |
| P68366 | TBA1_HUMAN | | centriole. |
| P68371 | TBB2C_HUMAN | | centriole. |
| P68400 | CSK21_HUMAN | | cytoplasm. |
| P68543 | UBXD4_HUMAN | | nucleus. |
| P68871 | HBB_HUMAN | | endosome. endoplasmic reticulum. |
| P69849 | NOMO3_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome. |
| P69891 | HBG1_HUMAN | | endoplasmic reticulum. |
| P69892 | HBG2_HUMAN | | endoplasmic reticulum. |
| P69905 | HBA_HUMAN | | endoplasmic reticulum. |
| P78312 | CD008_HUMAN | | nucleus. |
| P78314 | 3BP2_HUMAN | | cytoplasm. |
| P78316 | NOP14_HUMAN | nucleus; nucleolus (by similarity). | nucleus. |
| P78318 | IGBP1_HUMAN | cytoplasm (potential). | cytoplasm. |
| P78330 | SERB_HUMAN | | cytoplasm. |
| P78332 | RBM6_HUMAN | nucleus (probable). | nucleus. |
| P78333 | GPC5_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| P78344 | IF4G2_HUMAN | | cytoplasm. |
| P78345 | RPP38_HUMAN | nucleus; nucleolus (potential). | nucleus. |
| P78346 | RPP30_HUMAN | nucleus; nucleolus (potential). | nucleus. |
| P78347 | GTF2I_HUMAN | cytoplasm (potential). nucleus (potential). | nucleus. |
| P78348 | ACCN2_HUMAN | cell membrane; multi-pass membrane protein (by similarity). localizes in synaptosomes at dendritic synapses of neurons. colocalizes with dlg4 (by similarity). | membrane. |
| P78356 | PI52B_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; peripheral membrane protein (by similarity). cell membrane; peripheral membrane protein (by similarity). associated with the plasma membrane and the endoplasmic reticulum | cytoplasm. endoplasmic reticulum. |
| P78357 | CNTP1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| P78358 | CTG1B_HUMAN | | cytoplasm. |
| P78367 | NKX32_HUMAN | nucleus (probable). | nucleus. |

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| P78383 | S35B1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| P78385 | KRHB3_HUMAN | | cytoplasm. |
| P78386 | KRHB5_HUMAN | | cytoplasm. |
| P78395 | MAPE_HUMAN | | cytoplasm. |
| P78396 | CCNA1_HUMAN | | nucleus. |
| P78410 | BT3A2_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| P78411 | IRX5_HUMAN | nucleus (probable). | nucleus. |
| P78412 | IRX6_HUMAN | nucleus (probable). | nucleus. |
| P78413 | IRX4_HUMAN | nucleus (probable). | nucleus. |
| P78414 | IRX1_HUMAN | nucleus (probable). | nucleus. |
| P78415 | IRX3_HUMAN | nucleus (probable). | nucleus. |
| P78424 | PO6F2_HUMAN | nucleus (probable). | nucleus. |
| P78426 | NKX61_HUMAN | nucleus (probable). | nucleus. |
| P78509 | RELN_HUMAN | secreted protein (by similarity). | secreted protein. |
| P78524 | ST5_HUMAN | | cytoplasm. |
| P78537 | BL1S1_HUMAN | | cytoplasm. |
| P78543 | BTG2_HUMAN | | nucleus. |
| P78559 | MAP1A_HUMAN | | cytoplasm. |
| P78560 | CRADD_HUMAN | | cytoplasm. |
| P78562 | PHEX_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| P78563 | RED1_HUMAN | | nucleus. |
| P79522 | PRR3_HUMAN | | cytoplasm. nucleus. |
| P80192 | M3K9_HUMAN | | cytoplasm. |
| P80294 | MT1H_HUMAN | | cytoplasm. |
| P80295 | MT1I_HUMAN | | cytoplasm. nucleus. |
| P80297 | MT1X_HUMAN | | cytoplasm. nucleus. |
| P80362 | KV1Y_HUMAN | | secreted protein. |
| P80419 | HV3V_HUMAN | | secreted protein. |
| P80421 | HV1H_HUMAN | | secreted protein. |
| P80422 | LV2L_HUMAN | | secreted protein. |
| P80511 | S10AC_HUMAN | | secreted protein. |
| P80748 | LV3B_HUMAN | | secreted protein. |
| P81133 | SIM1_HUMAN | nucleus (potential). | nucleus. |
| P81408 | COTE1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| P81877 | SSBP2_HUMAN | nucleus (by similarity). | nucleus. |
| P82094 | TMF1_HUMAN | | nucleus. |
| P82251 | BAT1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| P82673 | RT35_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| P82980 | RET5_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| P83105 | HTRA4_HUMAN | | secreted protein. |
| P83110 | HTRA3_HUMAN | secreted protein (by similarity). | secreted protein. |
| P83111 | LACTB_HUMAN | | endosome. |
| P83731 | RL24_HUMAN | | mitochondrion. |
| P84022 | SMAD3_HUMAN | cytoplasm (by similarity). in the cytoplasm in the absence of ligand. nucleus (by similarity). migration to the nucleus when complexed with smad4 (by similarity). | nucleus. |
| P84074 | HPCA_HUMAN | | cytoplasm. |
| P84077 | ARF1_HUMAN | | Golgi. |
| P84085 | ARF5_HUMAN | | Golgi. |

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| P84090 | ERH_HUMAN | | cytoplasm. |
| P84095 | RHOG_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| P84098 | RL19_HUMAN | | mitochondrion. |
| P84101 | SERF2_HUMAN | | nucleus. |
| P84157 | TMAP1_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| P84550 | LBXCO_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| P87893 | POK11_HUMAN | | microsome. nucleus. |
| P98066 | TSG6_HUMAN | | secreted protein. |
| P98073 | ENTK_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| P98077 | SHC2_HUMAN | | cytoplasm. |
| P98082 | DAB2_HUMAN | | secreted protein. |
| P98161 | PKD1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| P98168 | ZXDA_HUMAN | nucleus (potential). | nucleus. |
| P98169 | ZXDB_HUMAN | nucleus (potential). | nucleus. |
| P98173 | FAM3A_HUMAN | secreted protein (potential). | centriole. cytoplasm. |
| P98174 | FGD1_HUMAN | cytoplasm (by similarity). associated with membrane ruffles and lamellipodia (by similarity). | cytoplasm. |
| P98175 | RBM10_HUMAN | nucleus (by similarity). | nucleus. |
| P98179 | RBM3_HUMAN | | cytoplasm. nucleus. |
| P98182 | ZN200_HUMAN | nucleus (probable). | nucleus. |
| P98187 | CP4F8_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q00005 | 2ABB_HUMAN | | cytoplasm. |
| Q00266 | METK1_HUMAN | | cytoplasm. |
| Q00526 | CDK3_HUMAN | | nucleus. |
| Q00534 | CDK6_HUMAN | | nucleus. |
| Q00535 | CDK5_HUMAN | cytoplasm. in axonal growth cone with extension to the peripheral lamellipodia (by similarity). | nucleus. |
| Q00536 | PCK1_HUMAN | | cytoplasm. |
| Q00537 | PCK2_HUMAN | | cytoplasm. |
| Q00604 | NDP_HUMAN | secreted protein (potential). | secreted protein. |
| Q00722 | PLCB2_HUMAN | | secreted protein. |
| Q00765 | REEP5_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q00796 | DHSO_HUMAN | | cytoplasm. |
| Q00872 | MYPC1_HUMAN | | cytoplasm. cytoskeleton. |
| Q00887 | PSG9_HUMAN | secreted protein (potential). | secreted protein. |
| Q00888 | PSG4_HUMAN | secreted protein (potential). | secreted protein. |
| Q00889 | PSG6_HUMAN | secreted protein (potential). | secreted protein. |
| Q00994 | BEX3_HUMAN | nucleus. cytoplasm. shuttles between the cytoplasm and the nucleus. associates with replicating mitochondria (by similarity). | nucleus. |
| Q01082 | SPTB2_HUMAN | | cytoplasm. |
| Q01101 | INSM1_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q01415 | GALK2_HUMAN | | cytoplasm. |
| Q01432 | AMPD3_HUMAN | | cytoplasm. |
| Q01433 | AMPD2_HUMAN | | cytoplasm. |
| Q01449 | MLRA_HUMAN | | cytoplasm. cytoskeleton. |
| Q01484 | ANK2_HUMAN | | cytoplasm. |

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|--------|-------------|---|-----------------------|
| Q01518 | CAP1_HUMAN | cell membrane (by similarity). | cytoplasm. Golgi. |
| Q01546 | K22O_HUMAN | | cytoplasm. |
| Q01628 | IFM3_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q01629 | IFM2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q01650 | LAT1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q01804 | OTUD4_HUMAN | | centriole. cytoplasm. |
| Q01813 | K6PP_HUMAN | | cytoplasm. |
| Q01831 | XPC_HUMAN | nucleus (probable). | nucleus. |
| Q01850 | CDR2_HUMAN | | nucleus. |
| Q01954 | BNC1_HUMAN | nucleus (by similarity). cytoplasm (by similarity). relocates to the midpiece of the flagellum during late spermiogenesis in spermatids (by similarity). | nucleus. |
| Q01955 | CO4A3_HUMAN | cell surface (potential). | secreted protein. |
| Q01968 | OCRL_HUMAN | | cytoplasm. |
| Q01995 | TAGL_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q02040 | XE7_HUMAN | membrane; peripheral membrane protein (probable). | cytoplasm. Golgi. |
| Q02045 | MLR5_HUMAN | | cytoskeleton. |
| Q02083 | ASAHL_HUMAN | lysosome (probable). | cytoplasm. |
| Q02156 | KPCE_HUMAN | | cytoplasm. |
| Q02388 | CO7A1_HUMAN | | secreted protein. |
| Q02410 | APBA1_HUMAN | | endosome. Golgi. |
| Q02413 | DSG1_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q02509 | OC90_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q02543 | RL18A_HUMAN | | cytoplasm. |
| Q02575 | HEN1_HUMAN | | nucleus. |
| Q02577 | HEN2_HUMAN | | nucleus. |
| Q02750 | MP2K1_HUMAN | | cytoplasm. |
| Q02779 | M3K10_HUMAN | | cytoplasm. |
| Q02818 | NUCB1_HUMAN | Golgi apparatus; cis-Golgi network; cis-Golgi network membrane; peripheral membrane protein; luminal side (by similarity). cytoplasm (by similarity). a small fraction of the protein may be cytoplasmic (by similarity). | cytoplasm. |
| Q02833 | RASF7_HUMAN | | nucleus. |
| Q02878 | RL6_HUMAN | | cytoplasm. |
| Q02930 | CREB5_HUMAN | nucleus (probable). | nucleus. |
| Q02952 | AKA12_HUMAN | cytoplasm; cell cortex (probable). cytoskeleton (probable). may be part of the cortical cytoskeleton. | cytoplasm. |
| Q03014 | HHEX_HUMAN | nucleus (probable). | nucleus. |
| Q03167 | TGBR3_HUMAN | secreted protein. secreted protein; extracellular space. cell membrane; single-pass type i membrane protein (potential). exists both as a membrane-bound form and as soluble form in serum and in the extracellular matrix. | membrane. |
| Q03169 | TNAP2_HUMAN | | secreted protein. |
| Q03393 | PTPS_HUMAN | | cytoplasm. |

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|--------|-------------|---|----------------------------------|
| Q03405 | UPAR_HUMAN | isoform 1: cell membrane; lipid-anchor; gpi- anchor. isoform 2: secreted protein (probable). | membrane. |
| Q03468 | ERCC6_HUMAN | nucleus (probable). | nucleus. |
| Q03692 | COAA1_HUMAN | | secreted protein. |
| Q03936 | ZNF92_HUMAN | nucleus (probable). | nucleus. |
| Q04118 | PRB3_HUMAN | | centriole. cytoplasm. |
| Q04323 | SAKS1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q04446 | GLGB_HUMAN | | lysosome. |
| Q04637 | IF4G1_HUMAN | | cytoplasm. |
| Q04656 | ATP7A_HUMAN | Golgi apparatus; trans-Golgi network; trans- Golgi network membrane; multi-pass membrane protein. cell membrane; multi-pass membrane protein. cycles constitutively between the trans-Golgi network (tgn) and the plasma membrane. predominantly found in the tgn and relocalized to the plasma membrane in response to elevated copper levels. isoform 3: cytoplasm; cytosol (probable). isoform 5: endoplasmic reticulum. | endoplasmic reticulum; membrane. |
| Q04695 | K1C17_HUMAN | | cytoplasm. |
| Q04741 | EMX1_HUMAN | nucleus (probable). | nucleus. |
| Q04743 | EMX2_HUMAN | nucleus (probable). | nucleus. |
| Q04759 | KPCT_HUMAN | | cytoplasm. |
| Q04760 | LGUL_HUMAN | | cytoplasm. |
| Q04762 | YYY5_HUMAN | | secreted protein. |
| Q04864 | REL_HUMAN | nucleus (by similarity). | nucleus. |
| Q04917 | 1433F_HUMAN | | cytoplasm. |
| Q05086 | UBE3A_HUMAN | nucleus (probable). | nucleus. |
| Q05215 | EGR4_HUMAN | nucleus (probable). | nucleus. |
| Q05469 | LIPS_HUMAN | | mitochondrion. |
| Q05481 | ZNF91_HUMAN | nucleus (probable). | nucleus. |
| Q05513 | KPCZ_HUMAN | cytoplasm. in the retina, localizes in the terminals of the rod bipolar cells (by similarity). endosome. associates with endosomes. | cytoplasm. |
| Q05524 | ENO1B_HUMAN | cytoplasm (potential). | membrane. |
| Q05586 | NMDZ1_HUMAN | cell membrane; multi-pass membrane protein (by similarity). enriched in post-synaptic plasma membrane and post-synaptic densities (by similarity). | membrane. |
| Q05639 | EF1A2_HUMAN | nucleus (by similarity). | cytoplasm. |
| Q05655 | KPCD_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | cytoplasm. |
| Q05682 | CALD1_HUMAN | on thin filaments in smooth muscle and on stress fibers in fibroblasts (nonmuscle) (by similarity). | cytoplasm. |
| Q05707 | COEA1_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q05877 | HHCM_HUMAN | | cytoplasm. |
| Q06033 | ITIH3_HUMAN | | secreted protein. |
| Q06187 | BTK_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | cytoplasm. |
| Q06190 | 2ACA_HUMAN | | cytoplasm. |
| Q06203 | PUR1_HUMAN | | cytoplasm. |

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| Q06210 | GFPT1_HUMAN | | cytoplasm. |
| Q06250 | WIT1_HUMAN | | cytoplasm. |
| Q06323 | PSME1_HUMAN | | cytoplasm. |
| Q06416 | PO5FL_HUMAN | nucleus (by similarity). | nucleus. |
| Q06455 | MTG8_HUMAN | nucleus (potential). | nucleus. |
| Q06481 | APLP2_HUMAN | cell membrane; single-pass type i membrane protein (potential). nucleus (potential). | nucleus. |
| Q06547 | GABP2_HUMAN | | nucleus. |
| Q06643 | TNFC_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q06730 | ZN33A_HUMAN | nucleus (potential). | nucleus. |
| Q06732 | ZN11B_HUMAN | nucleus (potential). | nucleus. |
| Q06889 | EGR3_HUMAN | nucleus (probable). | nucleus. |
| Q07002 | PCTK3_HUMAN | | cytoplasm. |
| Q07092 | COGA1_HUMAN | | secreted protein. |
| Q07283 | TRHY_HUMAN | | cytoplasm. |
| Q07326 | PIGF_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q07343 | PDE4B_HUMAN | | cytoplasm. |
| Q07627 | KRA11_HUMAN | | secreted protein. |
| Q07687 | DLX2_HUMAN | nucleus (potential). | nucleus. |
| Q07817 | BCLX_HUMAN | mitochondrion; mitochondrial membrane (by similarity). mitochondrial membranes and perinuclear envelope (by similarity). | mitochondrion. |
| Q07820 | MCL1_HUMAN | membrane; single-pass membrane protein (potential). cytoplasm. mitochondrion. cytoplasmic, associated with mitochondria. nucleus; nucleoplasm. | mitochondrion. |
| Q07837 | SLC31_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q07866 | KLC1_HUMAN | | cytoplasm. |
| Q07889 | SOS1_HUMAN | | cytoplasm. |
| Q07890 | SOS2_HUMAN | | cytoplasm. |
| Q07912 | ACK1_HUMAN | | cytoplasm. |
| Q07954 | LRP1_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). processed form: cytoplasm (by similarity). processed form: nucleus (by similarity). after cleavage, the intracellular domain (Irp1cd) is detected both in the cytoplasm and in the nucleus (by similarity). | membrane. |
| Q08043 | ACTN3_HUMAN | | cytoplasm. |
| Q08116 | RGS1_HUMAN | | secreted protein. nucleus; membrane. |
| Q08117 | AES_HUMAN | nucleus (by similarity). | nucleus. |
| Q08174 | PCDH1_HUMAN | cell membrane; cell-cell junction; single-pass type i membrane protein. cell membrane; cell-matrix junction; single-pass type i membrane protein (probable). found at cell-cell boundaries and probably at cell-matrix boundaries. | membrane. |
| Q08188 | TGM3_HUMAN | | cytoplasm. |

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| Q08209 | PP2BA_HUMAN | nucleus (by similarity). colocalizes with actn1 and myoz2 at the z line in heart and skeletal muscle (by similarity). | cytoplasm. nucleus. |
| Q08289 | CACB2_HUMAN | sarcolemma; sarcolemmal membrane; peripheral membrane protein; cytoplasmic side (by similarity). | membrane. |
| Q08397 | LOXL1_HUMAN | secreted protein; extracellular space (potential). | secreted protein. |
| Q08477 | CP4F3_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q08493 | PDE4C_HUMAN | | cytoplasm. |
| Q08495 | DEMA_HUMAN | | cytoplasm. |
| Q08499 | PDE4D_HUMAN | cytoplasm (by similarity). membrane (by similarity). cytoskeleton (by similarity). centrosome (by similarity). found in the soluble fraction, associated with membranes, and associated with the cytoskeleton and the centrosome (by similarity). | cytoplasm. |
| Q08623 | HDD1A_HUMAN | | centriole. cytoplasm. |
| Q08708 | CM35A_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q08881 | ITK_HUMAN | cell membrane (by similarity). localizes to cell surface receptors in the plasma membrane after stimulation with respective receptors (tcr, cd28, cd2) in t-cells (by similarity). | cytoplasm. |
| Q09013 | DMPK_HUMAN | | cytoplasm. |
| Q0ZLH3 | PJVK_HUMAN | | centriole. cytoplasm. secreted protein. membrane. |
| Q10571 | MN1_HUMAN | | cytoplasm. |
| Q10713 | MPPA_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | membrane. |
| Q12756 | KIF1A_HUMAN | | cytoplasm. nucleus. |
| Q12765 | SCRN1_HUMAN | cytoplasm (by similarity). | nucleus. |
| Q12767 | K0195_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q12768 | K0196_HUMAN | | cytoplasm. |
| Q12770 | SCAP_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). Golgi apparatus; Golgi membrane; multi-pass membrane protein (by similarity). moves from the endoplasmic reticulum to the Golgi in the absence of sterols (by | endoplasmic reticulum. |
| Q12774 | ARHG5_HUMAN | | cytoplasm. |
| Q12792 | TWF1_HUMAN | cytoplasm. diffuse cytoplasmic localization with perinuclear and g-actin-rich cortical actin structures sublocalization. also found at membrane ruffles and cell-cell contacts (by similarity). | cytoplasm. |
| Q12796 | PNRC1_HUMAN | nucleus (potential). | nucleus. |
| Q12799 | TCP10_HUMAN | | cytoplasm. |
| Q12802 | LBC_HUMAN | | cytoplasm. |
| Q12816 | TROP_HUMAN | | membrane. |

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| Q12829 | RB40B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| Q12834 | CDC20_HUMAN | | nucleus. |
| Q12840 | KIF5A_HUMAN | cytoplasm; perinuclear region. concentrated in the cell body of the neurons, particularly in the perinuclear | cytoplasm. nucleus. |
| Q12841 | FSTL1_HUMAN | secreted protein (potential). | cytoplasm. secreted protein. |
| Q12846 | STX4_HUMAN | membrane; single-pass type iv membrane protein (potential). | endoplasmic reticulum. Golgi. |
| Q12851 | M4K2_HUMAN | cytoplasm (by similarity). cell membrane; basolateral cell membrane; peripheral membrane protein (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q12852 | M3K12_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | cytoplasm. |
| Q12864 | CAD17_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q12866 | MERTK_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q12870 | TCF15_HUMAN | nucleus (by similarity). | nucleus. |
| Q12872 | SFRS8_HUMAN | nucleus (potential). | nucleus. |
| Q12893 | TM115_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q12894 | IFRD2_HUMAN | | cytoplasm. secreted protein. |
| Q12899 | TRI26_HUMAN | | cytoplasm. |
| Q12901 | ZN155_HUMAN | nucleus (potential). | nucleus. |
| Q12904 | MCA1_HUMAN | | secreted protein. |
| Q12912 | LRMP_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type iv membrane protein (by similarity). processed form: cytoplasm (by similarity). a soluble form has also been detected (by | endoplasmic reticulum. |
| Q12923 | PTN13_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q12926 | ELAV2_HUMAN | | cytoplasm. nucleus. |
| Q12929 | EPS8_HUMAN | | cytoplasm. |
| Q12955 | ANK3_HUMAN | | cytoplasm. |
| Q12959 | DLG1_HUMAN | membrane; peripheral membrane protein (by similarity). colocalizes with epb41 at regions of intercellular contacts. basolateral in epithelial cells. may also associate with endoplasmic reticulum membranes. mainly found in neurons soma, moderately found at postsynaptic | membrane. |
| Q12965 | MYO1E_HUMAN | | cytoskeleton. |
| Q12967 | GNDS_HUMAN | | cytoplasm. |
| Q12979 | ABR_HUMAN | | cytoplasm. |
| Q12980 | U171_HUMAN | | centriole. cytoplasm. |
| Q12988 | HSPB3_HUMAN | | cytoplasm. |
| Q13007 | IL24_HUMAN | secreted protein (potential). | secreted protein. |
| Q13009 | TIAM1_HUMAN | | cytoplasm. |
| Q13011 | ECH1_HUMAN | mitochondrion (by similarity). peroxisome (by similarity). | mitochondrion. |
| Q13014 | BAK2_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q13015 | AF1Q_HUMAN | | cytoplasm. secreted protein. |

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| Q13021 | MALL_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q13033 | STRN3_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | centriole. nucleus. |
| Q13045 | FLII_HUMAN | nucleus (by similarity). cytoplasm (by similarity). centrosome (by similarity). colocalizes to actin-rich structures in blastocysts and, together with hras1, rhoa and cdc42, in migrating fibroblasts. localizes to centrosomes (by similarity). | nucleus. |
| Q13046 | PSG7_HUMAN | secreted protein (potential). | secreted protein. |
| Q13057 | COASY_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q13064 | MKRN3_HUMAN | | nucleus. |
| Q13065 | GAGE1_HUMAN | | secreted protein. |
| Q13066 | GAGE2_HUMAN | | centriole. cytoplasm. |
| Q13067 | GAGE3_HUMAN | | cytoplasm. |
| Q13068 | GAGE4_HUMAN | | cytoplasm. |
| Q13069 | GAGE5_HUMAN | | cytoplasm. |
| Q13070 | GAGE6_HUMAN | | cytoplasm. |
| Q13075 | BIRC1_HUMAN | | cytoplasm. |
| Q13077 | TRAF1_HUMAN | | cytoplasm. |
| Q13087 | PDIA2_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| Q13094 | LCP2_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q13099 | IFT88_HUMAN | | mitochondrion. |
| Q13106 | ZN154_HUMAN | nucleus (potential). | nucleus. |
| Q13114 | TRAF3_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q13129 | RLF_HUMAN | nucleus (probable). | nucleus. |
| Q13131 | AAPK1_HUMAN | | cytoplasm. |
| Q13133 | NR1H3_HUMAN | nucleus (potential). | nucleus. |
| Q13144 | EI2BE_HUMAN | | cytoplasm. |
| Q13145 | BAMBI_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome. |
| Q13151 | ROA0_HUMAN | nucleus (by similarity). component of ribonucleosomes (by similarity). | nucleus. |
| Q13155 | MCA2_HUMAN | | cytoplasm. |
| Q13158 | FADD_HUMAN | | cytoplasm. |
| Q13163 | MP2K5_HUMAN | | cytoplasm. |
| Q13164 | MK07_HUMAN | | cytoplasm. |
| Q13166 | CATR1_HUMAN | | centriole. cytoplasm. |
| Q13177 | PAK2_HUMAN | | cytoplasm. |
| Q13183 | S13A2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q13185 | CBX3_HUMAN | nucleus (potential). associates with euchromatin and is largely excluded from constitutive heterochromatin. may be associated with microtubules and mitotic poles during mitosis (potential). | nucleus. |
| Q13188 | STK3_HUMAN | cytoplasm. the caspase-cleaved form cycles between nucleus and cytoplasm (by similarity). | cytoplasm. |
| Q13190 | STX5_HUMAN | endoplasmic reticulum-Golgi intermediate compartment; endoplasmic reticulum-Golgi intermediate compartment membrane; single-pass type iv membrane protein (by similarity). | Golgi; membrane. |

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| Q13200 | PSD2_HUMAN | | cytoplasm. |
| Q13201 | MMRN1_HUMAN | | membrane. |
| Q13202 | DUS8_HUMAN | | cytoplasm. |
| Q13203 | MYBPH_HUMAN | | cytoplasm. |
| Q13206 | DDX10_HUMAN | | nucleus. |
| Q13207 | TBX2_HUMAN | nucleus (potential). | nucleus. |
| Q13214 | SEM3B_HUMAN | secreted protein (by similarity). endoplasmic reticulum. accumulates in the endoplasmic reticulum. | membrane. |
| Q13216 | ERCC8_HUMAN | nucleus (probable). | nucleus. |
| Q13217 | DNJC3_HUMAN | cytoplasm (by similarity). endoplasmic reticulum (by similarity). associated with the endoplasmic reticulum (er) (by | cytoplasm. |
| Q13227 | GPS2_HUMAN | | cytoplasm. |
| Q13228 | SBP1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q13232 | NDK3_HUMAN | | cytoplasm. |
| Q13237 | KGP2_HUMAN | | cytoplasm. |
| Q13239 | SLAP1_HUMAN | cytoplasm (by similarity). endosome (by similarity). colocalizes with endosomes (by similarity). | cytoplasm. |
| Q13263 | TIF1B_HUMAN | nucleus. associated with centromeric heterochromatin during cell differentiation through cbx1 (by similarity). | nucleus. |
| Q13275 | SEM3F_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q13277 | STX3_HUMAN | membrane; single-pass type iv membrane protein (potential). | Golgi; membrane. |
| Q13283 | G3BP_HUMAN | cytoplasm. cytoplasm; cytosol. cell membrane. nucleus. cytoplasmic in proliferating cells, can be recruited to the plasma membrane in exponentially growing cells (by similarity). cytosolic and partially nuclear in resting cells. recruited to stress granules (sgs) upon either arsenite or high temperature treatment. recruitment to sgs is influenced by hras. | cytoplasm. nucleus. |
| Q13296 | SG2A2_HUMAN | | centriole. cytoplasm. |
| Q13303 | KCAB2_HUMAN | cytoplasm (potential). | membrane. |
| Q13309 | SKP2_HUMAN | | cytoplasm. |
| Q13316 | DMP1_HUMAN | | secreted protein. |
| Q13319 | CD5R2_HUMAN | | nucleus. |
| Q13322 | GRB10_HUMAN | | cytoplasm. |
| Q13325 | IFIT5_HUMAN | | centriole. cytoplasm. |
| Q13326 | SGCG_HUMAN | sarcolemma; sarcolemmal membrane; single- pass type ii membrane protein (by similarity). | membrane. |
| Q13347 | IF32_HUMAN | | cytoplasm. |
| Q13360 | ZN177_HUMAN | nucleus (probable). | nucleus. |
| Q13363 | CTBP1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. nucleus. |
| Q13367 | AP3B2_HUMAN | Golgi apparatus. cytoplasmic vesicle; cytoplasmic vesicle membrane; peripheral membrane protein; cytoplasmic side (by similarity). component of the coat surrounding the cytoplasmic face of coated vesicles located at the Golgi complex (by similarity). | Golgi. |

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| Q13368 | MPP3_HUMAN | | cytoplasm. |
| Q13370 | PDE3B_HUMAN | membrane; peripheral membrane protein (potential). | cytoplasm. |
| Q13371 | PHLP_HUMAN | | cytoplasm. |
| Q13393 | PLD1_HUMAN | cytoplasm; perinuclear region (by similarity). endoplasmic reticulum (by similarity). Golgi apparatus (by similarity). endosome; late endosome (by similarity). | endoplasmic reticulum. |
| Q13398 | ZN211_HUMAN | nucleus (probable). | nucleus. |
| Q13402 | MYO7A_HUMAN | cytoplasm (probable). in the photoreceptor cells, mainly localized in the inner and base of outer segments as well as in the synaptic ending region. | cytoplasm. cytoskeleton. |
| Q13404 | UB2V1_HUMAN | nucleus. isoform 3, isoform 5: membrane; multi-pass membrane protein (potential). | nucleus. |
| Q13409 | DC1I2_HUMAN | | cytoplasm. |
| Q13412 | 3CL_HUMAN | | cytoplasm. secreted protein. |
| Q13423 | NNTM_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein; matrix side (potential). | mitochondrion. |
| Q13424 | SNTA1_HUMAN | sarcolemma; sarcolemmal membrane; peripheral membrane protein; cytoplasmic side (by similarity). in skeletal muscle, it localizes at the cytoplasmic side of the sarcolemmal membrane and at neuromuscular junctions (by similarity). | cytoplasm. |
| Q13425 | SNTB2_HUMAN | membrane-associated. in muscle, it is exclusively localized at the neuromuscular junction (by similarity). in insulinoma cell line, it is enriched in secretory granules. | cytoplasm. |
| Q13428 | TCOF_HUMAN | nucleus; nucleolus (potential). | nucleus. |
| Q13432 | UN119_HUMAN | | centriole. cytoplasm. |
| Q13434 | MKRN4_HUMAN | | cytoplasm. nucleus. |
| Q13435 | SF3B2_HUMAN | nucleus (by similarity). | nucleus. |
| Q13438 | OS9_HUMAN | | secreted protein. |
| Q13442 | HAP28_HUMAN | | cytoplasm. |
| Q13445 | TMED1_HUMAN | cell membrane; single-pass type i membrane protein (potential). | membrane. |
| Q13454 | TUSC3_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). | endoplasmic reticulum. |
| Q13464 | ROCK1_HUMAN | cytoplasm. associated with the mother centriole and an intercentriolar linker (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein. a small proportion is associated with Golgi membranes. | cytoplasm. |
| Q13465 | MDS1_HUMAN | | nucleus. |
| Q13467 | FZD5_HUMAN | cell membrane; multi-pass membrane protein (by similarity). Golgi apparatus; Golgi membrane; multi-pass membrane protein (by similarity). localized at the plasma membrane and also found at the Golgi (by similarity). | membrane. |
| Q13472 | TOP3A_HUMAN | | nucleus. |
| Q13474 | DRP2_HUMAN | | cytoplasm. nucleus. |

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| Q13480 | GAB1_HUMAN | | cytoplasm. |
| Q13488 | VPP3_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q13489 | BIRC3_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q13490 | BIRC2_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q13491 | GPM6B_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q13495 | CG1_HUMAN | | centriole. cytoplasm. |
| Q13496 | MTM1_HUMAN | | cytoplasm. |
| Q13501 | SQSTM_HUMAN | cytoplasm. in cardiac muscles localizes to the sarcomeric band (by similarity). localizes to late endosomes. may also localize to the nucleus. accumulates in neurofibrillary tangles and in lewy bodies of neurons from individuals with alzheimer and parkinson disease respectively. enriched in rosenthal fibers of pilocytic astrocytoma. in liver cells, accumulates in mallory bodies associated with alcoholic hepatitis, wilson disease, indian childhood cirrhosis and in hyaline bodies associated with hepatocellular carcinoma. | cytoplasm. nucleus. |
| Q13503 | MED21_HUMAN | nucleus (potential). | nucleus. |
| Q13505 | MTX1_HUMAN | mitochondrion; mitochondrial outer membrane (by similarity). | endoplasmic reticulum. Golgi. mitochondrion. |
| Q13506 | NAB1_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q13509 | TBB3_HUMAN | | centriole. |
| Q13516 | OLIG2_HUMAN | nucleus (by similarity). cytoplasm (by similarity). the nls contained in the bhlh domain could be masked in the native form and translocation to the nucleus could be mediated by interaction either with class e bhlh partner protein or with nkx2-2 (by similarity). | nucleus. |
| Q13520 | AQP6_HUMAN | cytoplasmic vesicle; cytoplasmic vesicle membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q13522 | IPP1_HUMAN | | cytoplasm. |
| Q13530 | SERC3_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q13541 | 4EBP1_HUMAN | | cytoplasm. |
| Q13542 | 4EBP2_HUMAN | | cytoplasm. |
| Q13554 | KCC2B_HUMAN | | cytoplasm. |
| Q13555 | KCC2G_HUMAN | | cytoplasm. |
| Q13557 | KCC2D_HUMAN | | cytoplasm. |
| Q13562 | NDF1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| Q13563 | PKD2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q13572 | ITPK1_HUMAN | | cytoplasm. |
| Q13576 | IQGA2_HUMAN | | cytoplasm. |
| Q13588 | GRAP_HUMAN | | cytoplasm. |
| Q13596 | SNX1_HUMAN | | cytoplasm. Golgi. |
| Q13613 | MTMR1_HUMAN | | cytoplasm. |
| Q13616 | CUL1_HUMAN | | nucleus. |
| Q13617 | CUL2_HUMAN | | nucleus. |

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|--------|-------------|---|-----------------------|
| Q13619 | CUL4A_HUMAN | | nucleus. |
| Q13620 | CUL4B_HUMAN | | cytoplasm. nucleus. |
| Q13630 | FCL_HUMAN | | cytoplasm. |
| Q13636 | RAB31_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q13641 | TPBG_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q13643 | FHL3_HUMAN | | cytoplasm. nucleus. |
| Q13653 | CO031_HUMAN | | secreted protein. |
| Q13685 | AAMP_HUMAN | | secreted protein. |
| Q13686 | ALKB1_HUMAN | | nucleus. |
| Q13748 | TBA2_HUMAN | | centriole. |
| Q13769 | CV019_HUMAN | | nucleus. |
| Q13772 | NCOA4_HUMAN | | cytoplasm. nucleus. |
| Q13794 | APR_HUMAN | | cytoplasm. |
| Q13796 | APXL_HUMAN | | cytoplasm. |
| Q13813 | SPTA2_HUMAN | | cytoplasm. |
| Q13825 | AUMH_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| Q13829 | TNAP1_HUMAN | | nucleus. |
| Q13868 | EXOS2_HUMAN | | cytoplasm. nucleus. |
| Q13884 | SNTB1_HUMAN | sarcolemma; sarcolemmal membrane; peripheral membrane protein; cytoplasmic side (by similarity). in skeletal muscle, it localizes at the cytoplasmic side of the sarcolemmal membrane and at neuromuscular junctions (by similarity). | cytoplasm. |
| Q13890 | BT3L1_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q13891 | BT3L2_HUMAN | nucleus (potential). | nucleus. |
| Q13892 | BT3L3_HUMAN | nucleus (potential). | nucleus. |
| Q13905 | RPGF1_HUMAN | | cytoplasm. |
| Q13951 | PEBB_HUMAN | nucleus (potential). | nucleus. |
| Q13956 | CNCG_HUMAN | | centriole. |
| Q13972 | GMRP_HUMAN | | cytoplasm. |
| Q13976 | KGP1A_HUMAN | | cytoplasm. |
| Q14002 | CEAM7_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | membrane. |
| Q14004 | CD2L5_HUMAN | | cytoplasm. |
| Q14008 | CKAP5_HUMAN | | centriole. |
| Q14011 | CIRBP_HUMAN | nucleus; nucleoplasm (by similarity). | nucleus. |
| Q14012 | KCC1A_HUMAN | cytoplasm (by similarity). predominantly cytoplasmic. nucleus (by similarity). | cytoplasm. |
| Q14019 | COTL1_HUMAN | | cytoplasm. |
| Q14031 | CO4A6_HUMAN | cell surface (potential). | secreted protein. |
| Q14050 | CO9A3_HUMAN | | secreted protein. |
| Q14055 | CO9A2_HUMAN | | secreted protein. |
| Q14061 | COX17_HUMAN | mitochondrion; mitochondrial intermembrane space (by similarity). | mitochondrion. |
| Q14088 | RB33A_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q14094 | CCNI_HUMAN | | cytoplasm. |
| Q14114 | LRP8_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). secreted protein (by similarity). isoforms that contain the exon coding for a furin-type cleavage site are proteolytically processed, leading to a secreted receptor fragment (by | membrane. |

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|--------|-------------|--|-------------------------------------|
| Q14117 | DPYS_HUMAN | | cytoplasm. |
| Q14123 | PDE1C_HUMAN | | cytoplasm. |
| Q14135 | VGLL4_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q14138 | Y0125_HUMAN | | membrane. |
| Q14139 | UBE4A_HUMAN | | cytoplasm. |
| Q14140 | SRTD2_HUMAN | | cytoplasm. |
| Q14141 | SEPT6_HUMAN | | cytoplasm. |
| Q14142 | TRI14_HUMAN | | cytoplasm. |
| Q14146 | K0133_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q14147 | DHX34_HUMAN | | nucleus. |
| Q14149 | MORC3_HUMAN | | cytoplasm. nucleus. |
| Q14153 | FA53B_HUMAN | | centriole. cytoplasm. |
| Q14154 | K0141_HUMAN | | cytoplasm. mitochondrion. |
| Q14155 | ARHG7_HUMAN | | cytoplasm. |
| Q14157 | UBP2L_HUMAN | | cytoplasm. |
| Q14161 | GIT2_HUMAN | | cytoplasm. |
| Q14162 | SREC_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q14164 | IKKE_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q14165 | K0152_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q14166 | TTL12_HUMAN | | cytoplasm. |
| Q14168 | MPP2_HUMAN | | membrane. |
| Q14183 | DOC2A_HUMAN | synaptic vesicle; synaptic vesicle membrane; peripheral membrane protein (probable). colocalizes to synaptic | synapse. |
| Q14184 | DOC2B_HUMAN | membrane; peripheral membrane protein (by similarity). | membrane. |
| Q14185 | DOCK1_HUMAN | cytoplasm (probable). membrane (probable). recruited to membranes via its interaction with phosphatidylinositol 3,4,5-triphosphate (probable). | cytoplasm. |
| Q14194 | DPYL1_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q14195 | DPYL3_HUMAN | cytoplasm (by similarity). colocalizes with synaptic vesicle protein 2 in the central region of the growth cone (by similarity). | cytoplasm. secreted protein. |
| Q14197 | ICT1_HUMAN | | centriole. cytoplasm. |
| Q14201 | BTG3_HUMAN | | cytoplasm. |
| Q14202 | ZMYM3_HUMAN | nucleus (by similarity). | nucleus. |
| Q14206 | CCP2_HUMAN | | cytoplasm. |
| Q14232 | EI2BA_HUMAN | | cytoplasm. |
| Q14240 | IF4A2_HUMAN | | nucleus. |
| Q14241 | ELOA1_HUMAN | nucleus (probable). | nucleus. |
| Q14247 | SRC8_HUMAN | cytoplasm (by similarity). associated with membrane ruffles and lamellipodia (by similarity). | cytoplasm. |
| Q14258 | TRI25_HUMAN | | cytoplasm. |
| Q14264 | ENR1_HUMAN | | membrane. |
| Q14296 | FASTK_HUMAN | | cytoplasm. |
| Q14318 | FKBP8_HUMAN | | cytoplasm. |
| Q14320 | FA50A_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q14324 | MYPE2_HUMAN | | cytoplasm. cytoskeleton. |
| Q14353 | GAMT_HUMAN | | cytoplasm. nucleus. |
| Q14376 | GALE_HUMAN | | cytoplasm. lysosome. mitochondrion. |

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|--------|-------------|--|------------------------------|
| Q14390 | GGTL4_HUMAN | | cytoplasm. |
| Q14397 | GCKR_HUMAN | | cytoplasm. |
| Q14409 | GLPK3_HUMAN | mitochondrion; mitochondrial outer membrane; peripheral membrane protein; cytoplasmic side (by similarity). cytoplasm (by similarity). in sperm the majority of the enzyme is bound to mitochondria (by similarity). | cytoplasm. |
| Q14410 | GLPK2_HUMAN | mitochondrion; mitochondrial outer membrane; peripheral membrane protein; cytoplasmic side (by similarity). cytoplasm (by similarity). in sperm the majority of the enzyme is bound to mitochondria (by similarity). | cytoplasm. |
| Q14432 | PDE3A_HUMAN | membrane; peripheral membrane protein (potential). | cytoplasm. |
| Q14444 | GP137_HUMAN | cell membrane; lipid-anchor; gpi-anchor (probable). | membrane. |
| Q14451 | GRB7_HUMAN | | cytoplasm. |
| Q14457 | BCN1_HUMAN | cytoplasm (potential). Golgi apparatus; Golgi membrane; peripheral membrane protein (by similarity). expressed in dendrites and cell bodies of cerebellar purkinje cells (by similarity). | cytoplasm. |
| Q14494 | NF2L1_HUMAN | nucleus (by similarity). | nucleus. |
| Q14507 | EP3A_HUMAN | secreted protein (potential). | secreted protein. |
| Q14508 | WFDC2_HUMAN | secreted protein (potential). | secreted protein. |
| Q14515 | SPRL1_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q14517 | FATH_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q14525 | K1HB_HUMAN | | cytoplasm. |
| Q14532 | K1H2_HUMAN | | cytoplasm. |
| Q14533 | KRHB1_HUMAN | | cytoplasm. |
| Q14541 | HNF4G_HUMAN | nucleus (by similarity). | nucleus. |
| Q14554 | PDIA5_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| Q14558 | KPRA_HUMAN | | cytoplasm. secreted protein. |
| Q14563 | SEM3A_HUMAN | secreted protein (by similarity). | cytoplasm; membrane. |
| Q14565 | DMC1_HUMAN | nucleus (potential). | nucleus. |
| Q14566 | MCM6_HUMAN | nucleus (by similarity). | nucleus. |
| Q14576 | ELAV3_HUMAN | | cytoplasm. |
| Q14582 | MAD4_HUMAN | nucleus (by similarity). | nucleus. |
| Q14584 | ZN266_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q14585 | ZN345_HUMAN | nucleus (probable). | nucleus. |
| Q14586 | ZN267_HUMAN | nucleus (probable). | nucleus. |
| Q14587 | ZN268_HUMAN | nucleus (probable). | nucleus. |
| Q14588 | ZN234_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q14591 | ZN271_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q14592 | ZN460_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q14593 | ZN273_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q14596 | NBR1_HUMAN | | cytoplasm. nucleus. |

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| Q14623 | IHH_HUMAN | the c-terminal peptide diffuses from the cell, while the n-terminal peptide remains associated with the cell surface. is also secreted in either cleaved or uncleaved form to mediate signaling to other cells (by similarity). | secreted protein. |
| Q14624 | ITIH4_HUMAN | | secreted protein. |
| Q14641 | INSL4_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q14656 | ITBA1_HUMAN | | centriole. cytoplasm. |
| Q14657 | LAGE3_HUMAN | | centriole. cytoplasm. |
| Q14667 | K0100_HUMAN | secreted protein (potential). | mitochondrion. |
| Q14669 | TRIPC_HUMAN | | cytoplasm. |
| Q14671 | PUM1_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q14679 | TTLL4_HUMAN | | cytoplasm. mitochondrion. |
| Q14680 | MELK_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q14681 | KCTD2_HUMAN | | nucleus; membrane. |
| Q14689 | DIP2A_HUMAN | nucleus (potential). | nucleus. |
| Q14691 | PSF1_HUMAN | nucleus (by similarity). | centriole. |
| Q14692 | BMS1_HUMAN | nucleus; nucleolus (by similarity). | nucleus. |
| Q14693 | LPIN1_HUMAN | nucleus (by similarity). | nucleus. |
| Q14694 | UBP10_HUMAN | | nucleus. |
| Q14695 | K0087_HUMAN | | membrane. |
| Q14696 | MESD2_HUMAN | | cytoplasm. secreted protein. |
| Q14697 | GANAB_HUMAN | endoplasmic reticulum (by similarity). Golgi apparatus (by similarity). | cytoplasm. |
| Q14714 | SSPN_HUMAN | sarcolemma; sarcolemmal membrane; multi-pass membrane protein (by similarity). also found in myotendinous junctions and in the postsynaptic membrane of neuromuscular junctions (by similarity). | membrane. |
| Q14722 | KCAB1_HUMAN | cytoplasm (potential). | membrane. |
| Q14746 | COG2_HUMAN | Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (probable). | Golgi. |
| Q14761 | PTCA_HUMAN | | membrane. |
| Q14767 | LTBP2_HUMAN | secreted protein (by similarity). localized in nuchal ligament and aorta to the fibrillin-containing, microfibrillar component of elastic fibers (by similarity). | secreted protein. |
| Q14774 | HLX1_HUMAN | nucleus (potential). | nucleus. |
| Q14802 | FXVD3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q14847 | LASP1_HUMAN | cytoplasm; cell cortex (by similarity). cytoskeleton (by similarity). associated with the f-actin rich cortical cytoskeleton (by similarity). | cytoplasm. |
| Q14865 | ARI5B_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q14872 | MTF1_HUMAN | nucleus (potential). | nucleus. |
| Q14896 | MYPC3_HUMAN | | cytoskeleton. |
| Q14919 | DRAP1_HUMAN | | nucleus. |
| Q14929 | ZN169_HUMAN | nucleus (potential). | nucleus. |
| Q14956 | GPNMB_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q14964 | RB39A_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |

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|--------|-------------|--|------------------------|
| Q14982 | OPCM_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| Q14990 | ODFP_HUMAN | | cytoskeleton. |
| Q14993 | COJA1_HUMAN | | secreted protein. |
| Q14994 | NR1I3_HUMAN | nucleus (by similarity). | nucleus. |
| Q14995 | NR1D2_HUMAN | nucleus (by similarity). | nucleus. |
| Q14999 | CUL7_HUMAN | | cytoplasm. |
| Q15005 | SPCS2_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q15006 | K0103_HUMAN | nucleus (by similarity). | nucleus. |
| Q15008 | PSD6_HUMAN | | cytoplasm. |
| Q15012 | LAP4A_HUMAN | intracytoplasmic membrane; multi-pass membrane protein (potential). may reside in an intracellular membrane-bound compartment (potential). | Golgi. |
| Q15018 | K0157_HUMAN | | nucleus. |
| Q15019 | SEPT2_HUMAN | | cytoplasm. |
| Q15022 | SUZ12_HUMAN | nucleus (probable). | nucleus. |
| Q15027 | CENB1_HUMAN | | cytoplasm. |
| Q15031 | SYLM_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | cytoplasm. |
| Q15032 | R3HD1_HUMAN | | secreted protein. |
| Q15035 | TRAM2_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q15038 | DAZP2_HUMAN | | cytoplasm. |
| Q15040 | JOS1_HUMAN | | cytoplasm. |
| Q15048 | LRC14_HUMAN | | cytoplasm. |
| Q15049 | MLC1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q15051 | IQCB1_HUMAN | | cytoplasm. |
| Q15052 | ARHG6_HUMAN | | cytoplasm. |
| Q15053 | K0040_HUMAN | | nucleus. |
| Q15056 | IF4H_HUMAN | cytoplasm; perinuclear region (by | cytoplasm. nucleus. |
| Q15057 | CENB2_HUMAN | | cytoplasm. |
| Q15058 | KIF14_HUMAN | | cytoplasm. nucleus. |
| Q15059 | BRD3_HUMAN | nucleus (potential). | nucleus. |
| Q15078 | CD5R1_HUMAN | cyclin-dependent kinase 5 activator 1, p35: cell membrane; lipid-anchor; cytoplasmic side (probable). cyclin-dependent kinase 5 activator 1, p25: nucleus. cytoplasm; perinuclear region. the conversion of p35 to p25 relocalizes the protein from the cell periphery to the cytoplasm, in nuclear and perinuclear regions. in the primary cortical neurons, p25 is primarily concentrated in the cell soma and is largely absent from neurites, whereas p35 is present in the peripheries and nerve terminals. | cytoplasm. |
| Q15084 | PDIA6_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| Q15131 | CDK10_HUMAN | | cytoplasm. |
| Q15139 | KPCD1_HUMAN | | cytoplasm. |
| Q15147 | PLCB4_HUMAN | | cytoplasm. |
| Q15149 | PLEC1_HUMAN | | cytoplasm. |
| Q15155 | NOMO1_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |

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| Q15166 | PON3_HUMAN | secreted protein; extracellular space (by similarity). | secreted protein. |
| Q15181 | IPYR_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q15198 | PGFRL_HUMAN | secreted protein (probable). | cytoplasm. |
| Q15235 | PSG10_HUMAN | secreted protein (potential). | secreted protein. |
| Q15238 | PSG5_HUMAN | secreted protein (potential). | secreted protein. |
| Q15257 | PTPA_HUMAN | | cytoplasm. |
| Q15274 | NADC_HUMAN | | centriole. cytoplasm. |
| Q15286 | RAB35_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q15323 | K1H1_HUMAN | | cytoplasm. |
| Q15332 | C11X_HUMAN | | nucleus. |
| Q15345 | LRC41_HUMAN | | cytoplasm. |
| Q15369 | ELOC_HUMAN | nucleus (probable). | nucleus. |
| Q15370 | ELOB_HUMAN | nucleus (probable). | nucleus. |
| Q15382 | RHEB_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| Q15386 | UBE3C_HUMAN | nucleus (potential). | cytoplasm. |
| Q15388 | TOM20_HUMAN | mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (potential). | mitochondrion. |
| Q15390 | MTFR1_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| Q15399 | TLR1_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). phagosome; phagosomal membrane; single- pass type i membrane protein (by | membrane. |
| Q15404 | RSU1_HUMAN | | cytoplasm. |
| Q15406 | NR6A1_HUMAN | nucleus (probable). | nucleus. |
| Q15413 | RYR3_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q15414 | RBY1A_HUMAN | | nucleus. |
| Q15417 | CNN3_HUMAN | | cytoskeleton. |
| Q15418 | KS6A1_HUMAN | | cytoplasm. |
| Q15427 | SF3B4_HUMAN | nucleus (by similarity). | nucleus. |
| Q15428 | SF3A2_HUMAN | nucleus (by similarity). | nucleus. |
| Q15431 | SYCP1_HUMAN | nucleus (by similarity). in tripartite segments of synaptonemal complexes, between lateral elements in the nucleus. found only where the chromosome cores are synapsed. its n-terminus is found towards the centre of the synaptonemal complex while the c-terminus extends well into the lateral domain of the | cytoplasm. nucleus. |
| Q15434 | RBMS2_HUMAN | nucleus (potential). | nucleus. |
| Q15435 | PP1R7_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q15437 | SC23B_HUMAN | | cytoplasm. endoplasmic reticulum. |
| Q15438 | CYH1_HUMAN | | cytoplasm. |
| Q15459 | SF3A1_HUMAN | nucleus (by similarity). | nucleus. |

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|--------|-------------|---|-------------------------------------|
| Q15465 | SHH_HUMAN | sonic hedgehog protein c-product: secreted protein; extracellular space (by similarity). sonic hedgehog protein n-product: cell membrane; lipid-anchor (by similarity). the c-terminal peptide diffuses from the cell, while the n-product either remains associated with lipid rafts at the cell surface, or forms freely diffusible active multimers with its hydrophobic lipid-modified n- and c-termini buried inside (by | membrane. |
| Q15477 | SKIV2_HUMAN | nucleus (potential). | nucleus. |
| Q15486 | GUSL2_HUMAN | | secreted protein. |
| Q15487 | GUSL3_HUMAN | | centriole. cytoplasm. |
| Q15488 | GUSL4_HUMAN | | centriole. cytoplasm. |
| Q15493 | RGN_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q15506 | SP17_HUMAN | membrane; peripheral membrane protein (potential). | cytoplasm. secreted protein. Golgi. |
| Q15513 | SPHAR_HUMAN | | cytoplasm. |
| Q15526 | SURF1_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| Q15527 | SURF2_HUMAN | | centriole. cytoplasm. |
| Q15528 | SURF5_HUMAN | | cytoplasm. |
| Q15531 | STX1B_HUMAN | membrane; single-pass type iv membrane protein (potential). | membrane. |
| Q15532 | SSXT_HUMAN | | cytoplasm. nucleus. |
| Q15546 | PAQRB_HUMAN | endosome; late endosome; late endosomal membrane; multi-pass membrane protein (by similarity). lysosome; lysosomal membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q15569 | TESK1_HUMAN | | cytoplasm. |
| Q15628 | TRADD_HUMAN | | cytoplasm. |
| Q15642 | CIP4_HUMAN | | cytoplasm. |
| Q15645 | TRP13_HUMAN | | nucleus. |
| Q15646 | OASL_HUMAN | | microsome. nucleus. |
| Q15649 | ZNHI3_HUMAN | | cytoplasm. nucleus. |
| Q15652 | JHD2C_HUMAN | nucleus (by similarity). | nucleus. |
| Q15653 | IKBB_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| Q15654 | TRIP6_HUMAN | | cytoplasm. |
| Q15669 | RHOH_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| Q15678 | PTN14_HUMAN | | cytoplasm. |
| Q15695 | U2AFL_HUMAN | nucleus (potential). | nucleus. |
| Q15696 | U2AFM_HUMAN | nucleus (potential). | nucleus. |
| Q15699 | CART1_HUMAN | nucleus (probable). | nucleus. |
| Q15700 | DLG2_HUMAN | membrane; lipid-anchor (by similarity). concentrated in soma and postsynaptic density of a subset of neurons (by similarity). | membrane. |
| Q15714 | T22D1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| Q15717 | ELAV1_HUMAN | | cytoplasm. nucleus. |
| Q15735 | PI5PA_HUMAN | cytoplasm (by similarity). predominantly localized to membrane ruffles (by | cytoplasm. |

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| Q15738 | NSDHL_HUMAN | | endoplasmic reticulum. |
| Q15742 | NAB2_HUMAN | nucleus (by similarity). isoform 2 is not localized to the nucleus (by similarity). | cytoplasm. nucleus. |
| Q15746 | MYLK_HUMAN | | cytoplasm. |
| Q15750 | TAB1_HUMAN | | cytoplasm. |
| Q15759 | MK11_HUMAN | | cytoplasm. |
| Q15762 | CD226_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q15771 | RAB30_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q15773 | MLF2_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | centriole. cytoplasm. |
| Q15776 | ZN192_HUMAN | nucleus (potential). | nucleus. |
| Q15777 | MPPD2_HUMAN | | secreted protein. |
| Q15782 | CH3L2_HUMAN | | secreted protein. lysosome. |
| Q15784 | NDF2_HUMAN | nucleus (probable). | nucleus. |
| Q15788 | NCOA1_HUMAN | nucleus (by similarity). | nucleus. |
| Q15800 | ERG25_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). | endoplasmic reticulum. |
| Q15811 | ITSN1_HUMAN | intracytoplasmic membrane (by similarity). synaptosome (by similarity). enriched in synaptosomes (by similarity). | cytoplasm. |
| Q15813 | TBCE_HUMAN | | cytoplasm. |
| Q15814 | TBCC_HUMAN | | cytoplasm. |
| Q15818 | NPTX1_HUMAN | secretory vesicle (potential). | cytoplasm. |
| Q15819 | UB2V2_HUMAN | | nucleus. |
| Q15831 | STK11_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q15833 | STXB2_HUMAN | | cytoplasm. |
| Q15834 | DIPA_HUMAN | | centriole. cytoplasm. |
| Q15836 | VAMP3_HUMAN | membrane; single-pass type iv membrane protein (probable). | membrane. |
| Q15846 | CLUL1_HUMAN | secreted protein (probable). | membrane. |
| Q15847 | APM2_HUMAN | | centriole. cytoplasm. |
| Q15858 | SCN9A_HUMAN | membrane; multi-pass membrane protein. in neurite terminals (by similarity). | membrane. |
| Q15884 | CI061_HUMAN | | centriole. cytoplasm. |
| Q15904 | VAS1_HUMAN | vacuole; vacuolar membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| Q15907 | RB11B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q15910 | EZH2_HUMAN | nucleus (probable). | nucleus. |
| Q15928 | ZN141_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q15935 | ZNF77_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q15937 | ZNF79_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q15973 | ZN124_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q16048 | MCHL1_HUMAN | | secreted protein. |
| Q16082 | HSPB2_HUMAN | | cytoplasm. |
| Q16181 | SEPT7_HUMAN | | cytoplasm. |
| Q16186 | ADRM1_HUMAN | | membrane. |
| Q16204 | CCDC6_HUMAN | cytoplasm. cytoskeleton (probable). may be a cytoskeletal protein. | cytoplasm. |
| Q16342 | PDCD2_HUMAN | nucleus (potential). | nucleus. |
| Q16352 | AINX_HUMAN | | cytoplasm. cytoskeleton. |
| Q16384 | SSX1_HUMAN | | nucleus. |

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|--------|-------------|--|---------------------------------|
| Q16401 | PSD5_HUMAN | | centriole. cytoplasm. endosome. |
| Q16512 | PKN1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q16513 | PKN2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q16515 | ACCN1_HUMAN | cell membrane; multi-pass membrane protein (by similarity). localized at the plasma membrane of neurons, in the soma and punctated peripheral processes (by similarity). | membrane. |
| Q16517 | NNAT_HUMAN | | secreted protein. |
| Q16518 | RPE65_HUMAN | cytoplasm (by similarity). cell membrane; lipid-anchor (by similarity). attached to the membrane by a lipid anchor when palmitoylated (membrane form), soluble when unpalmitoylated (by similarity). | membrane. |
| Q16527 | CSRP2_HUMAN | nucleus (by similarity). | nucleus. |
| Q16534 | HLF_HUMAN | nucleus (probable). | nucleus. |
| Q16538 | GP162_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q16539 | MK14_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q16549 | PCSK7_HUMAN | Golgi apparatus; trans-Golgi network; trans- Golgi network membrane; single-pass type i membrane protein (by similarity). seems to be localized intracellularly to the trans Golgi network | secreted protein; membrane. |
| Q16553 | LY6E_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| Q16555 | DPYL2_HUMAN | cytoplasm (potential). | cytoplasm. secreted protein. |
| Q16557 | PSG3_HUMAN | secreted protein (potential). | secreted protein. |
| Q16559 | TAL2_HUMAN | | nucleus. |
| Q16563 | SYPL1_HUMAN | cytoplasmic vesicle; cytoplasmic vesicle membrane; multi-pass membrane protein (by similarity). cytoplasmic transport vesicles (by similarity). | membrane. |
| Q16566 | KCC4_HUMAN | cytoplasm (by similarity). nucleus (by similarity). substantial localization in certain neuronal nuclei. in spermatids associated with chromatin and nuclear matrix (by similarity). | cytoplasm. |
| Q16568 | CART_HUMAN | secreted protein (potential). | secreted protein. |
| Q16585 | SGCB_HUMAN | sarcolemma; sarcolemmal membrane; single- pass type ii membrane protein (by similarity). | membrane. |
| Q16586 | SGCA_HUMAN | sarcolemma; sarcolemmal membrane; single- pass type i membrane protein (by similarity). | membrane. |
| Q16589 | CCNG2_HUMAN | cytoplasm (by similarity). | nucleus. |
| Q16600 | ZN239_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q16611 | BAK_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q16612 | NP311_HUMAN | | nucleus. |
| Q16613 | SNAT_HUMAN | | cytoplasm. |
| Q16617 | NKG7_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |

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|--------|-------------|--|---|
| Q16623 | STX1A_HUMAN | cell membrane; single-pass type iv membrane protein. isoform 2: secreted protein (probable). | cytoplasm; membrane. |
| Q16626 | MEA1_HUMAN | | cytoplasm. cytoskeleton. nucleus. |
| Q16650 | TBR1_HUMAN | nucleus (potential). | nucleus. |
| Q16653 | MOG_HUMAN | isoform 1, isoform 5: cell membrane; multi-pass membrane protein (potential). isoform 2, isoform 3, isoform 4, isoform 6, isoform 7, isoform 8, isoform 9: cell membrane; single-pass type i membrane protein | membrane. |
| Q16655 | MAR1_HUMAN | | membrane. |
| Q16658 | FSCN1_HUMAN | | cytoplasm. |
| Q16659 | MK06_HUMAN | | cytoplasm. |
| Q16667 | CDKN3_HUMAN | | cytoplasm. |
| Q16690 | DUS5_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q16706 | MA2A1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endosome. endoplasmic reticulum. Golgi. lysosome. |
| Q16763 | UBE2S_HUMAN | | cytoplasm. |
| Q16769 | QPCT_HUMAN | | secreted protein. |
| Q16774 | KGUA_HUMAN | | cytoplasm. |
| Q16775 | GLO2_HUMAN | | cytoplasm. |
| Q16816 | PHKG1_HUMAN | | cytoplasm. |
| Q16821 | PPR3A_HUMAN | membrane; single-pass membrane protein (by similarity). | endosome; membrane. |
| Q16825 | PTN21_HUMAN | | cytoplasm. |
| Q16831 | UPP1_HUMAN | | cytoplasm. |
| Q16842 | SIA4B_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (by similarity). membrane-bound form in trans cisternae of Golgi. secreted protein (by similarity). secreted into the body fluid (by similarity). | secreted protein. |
| Q16850 | CP51A_HUMAN | microsome (potential). | endoplasmic reticulum. |
| Q16851 | UGPA2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q16864 | VATF_HUMAN | | mitochondrion. |
| Q16875 | F263_HUMAN | | cytoplasm. |
| Q16877 | F264_HUMAN | | cytoplasm. |
| Q16878 | CDO1_HUMAN | | cytoplasm. |
| Q16880 | CGT_HUMAN | | microsome. |
| Q16881 | TRXR1_HUMAN | cytoplasm (by similarity). | mitochondrion. |
| Q16890 | TPD53_HUMAN | | centriole. cytoplasm. |
| Q18PE1 | DOK7_HUMAN | membrane; peripheral membrane protein. accumulates at neuromuscular junctions (by similarity). | cytoplasm. Golgi. |
| Q2I0M5 | RSPO4_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q2M218 | ZN630_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q2M243 | CCD27_HUMAN | | cytoplasm. |
| Q2M329 | CCD96_HUMAN | | centriole. |
| Q2M3C6 | CO027_HUMAN | | nucleus. |
| Q2M3R5 | TMM20_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q2M3V2 | ANR43_HUMAN | | cytoplasm. |
| Q2M3W8 | ZN181_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q2M3X9 | ZN674_HUMAN | nucleus (probable). | cytoplasm. nucleus. |

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| Q2MJR0 | SPRE3_HUMAN | membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q2MKA7 | RSPO1_HUMAN | secreted protein (by similarity). | mitochondrion. |
| Q2MV58 | TECT1_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q2TAP0 | CJ132_HUMAN | | nucleus. |
| Q2TAY7 | SMU1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | centriole. cytoplasm. |
| Q2TBC4 | CF049_HUMAN | | cytoplasm. nucleus. |
| Q30KP8 | DB136_HUMAN | secreted protein (potential). | secreted protein. |
| Q30KP9 | DB135_HUMAN | secreted protein (potential). | secreted protein. |
| Q30KQ1 | DB133_HUMAN | secreted protein (potential). | secreted protein. |
| Q30KQ2 | DB130_HUMAN | secreted protein (potential). | secreted protein. |
| Q30KQ4 | DB116_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q30KQ5 | DB115_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q30KQ6 | DB114_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q30KQ7 | DB113_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q30KQ8 | DB112_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q30KQ9 | DB111_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q30KR0 | DB110_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q30KR1 | DB109_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q32M78 | ZN699_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q32M84 | BTBDG_HUMAN | | nucleus. |
| Q32M92 | CO032_HUMAN | | nucleus. |
| Q32MQ0 | ZN750_HUMAN | | nucleus. |
| Q32P28 | P3H1_HUMAN | endoplasmic reticulum (by similarity). secreted protein; extracellular space; extracellular matrix (by similarity). secreted into the extracellular matrix as a chondroitin sulfate proteoglycan (cspg). | endoplasmic reticulum. |
| Q3B7T1 | EDRF1_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q3LHN2 | KR192_HUMAN | | cytoplasm. |
| Q3LI58 | KR211_HUMAN | | cytoplasm. |
| Q3LI59 | KR212_HUMAN | | cytoplasm. |
| Q3LI61 | KR202_HUMAN | | cytoplasm. |
| Q3LI63 | KR201_HUMAN | | cytoplasm. |
| Q3LI64 | KRA61_HUMAN | | cytoplasm. |
| Q3LI66 | KRA62_HUMAN | | cytoplasm. |
| Q3LI67 | KRA63_HUMAN | | cytoplasm. |
| Q3LI70 | KR196_HUMAN | | cytoplasm. |
| Q3LI72 | KR195_HUMAN | | cytoplasm. |
| Q3LI73 | KR194_HUMAN | | cytoplasm. |
| Q3LI76 | KR151_HUMAN | | cytoplasm. |
| Q3LI77 | KR134_HUMAN | | cytoplasm. |
| Q3LI83 | KR241_HUMAN | | cytoplasm. |
| Q3LXA3 | DAK_HUMAN | | cytoplasm. |
| Q3MIR4 | CC50B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q3MJ13 | WDR72_HUMAN | | cytoplasm. |
| Q3MJ16 | PA24E_HUMAN | cytoplasm; cytosol. translocates to lysosomal membranes in a calcium-dependent fashion (by similarity). | secreted protein. |
| Q3SXZ3 | ZF718_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q3SY05 | CA157_HUMAN | | nucleus. |
| Q3SY46 | KR133_HUMAN | | cytoplasm. |
| Q3SY56 | SP6_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q3SYF9 | KR197_HUMAN | | cytoplasm. |

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| Q3SYG4 | PTHB1_HUMAN | | centriole. |
| Q3T8J9 | GON4L_HUMAN | nucleus (potential). | nucleus. |
| Q400G9 | AMZ1_HUMAN | | secreted protein. |
| Q460N5 | PRP14_HUMAN | nucleus. cytoplasm. in steady state cells the protein is mostly nuclear. a minor proportion is detected in the cytoplasm (by similarity). | nucleus. |
| Q494V2 | CCD37_HUMAN | | centriole. |
| Q494W8 | CRFM7_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q495M9 | USH1G_HUMAN | | centriole. secreted protein. |
| Q495T6 | MMEL1_HUMAN | membrane; single-pass type ii membrane protein. secreted protein. a secreted form produced by proteolytic cleavage also exists (by similarity). | membrane. |
| Q495X7 | TRI60_HUMAN | | cytoplasm. |
| Q496J9 | SV2C_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein. enriched in small synaptic vesicles and adrenal microsomes, not present in chromaffin granules. associated with both insulin granules and synaptic-like microvesicles in insulin-secreting cells of the pancreas (by similarity). | membrane. |
| Q496M8 | CI094_HUMAN | | secreted protein. |
| Q499Z4 | ZN672_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q4G0F5 | VP26B_HUMAN | cytoplasm. membrane; peripheral membrane protein. does not localize to endosomes (by similarity). | endosome. |
| Q4G163 | FBX43_HUMAN | | cytoplasm. |
| Q4KMG0 | CDON_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q4KMQ2 | TM16F_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| Q4KMX7 | F106A_HUMAN | | membrane. |
| Q4KN68 | CT062_HUMAN | | membrane. |
| Q4QY38 | DB134_HUMAN | secreted protein (potential). | secreted protein. |
| Q4UJ75 | A20A2_HUMAN | | nucleus. |
| Q4VC05 | BCL7A_HUMAN | | nucleus. |
| Q4VC31 | CCD58_HUMAN | | cytoplasm. |
| Q4ZG46 | RBP24_HUMAN | | cytoplasm. |
| Q4ZIN3 | MBRL_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q502W7 | CCD38_HUMAN | | nucleus. |
| Q502X0 | MORN2_HUMAN | | cytoplasm. |
| Q50LG9 | LRC24_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q52LD8 | CB011_HUMAN | | nucleus. |
| Q52LG2 | KR132_HUMAN | | cytoplasm. |
| Q52LJ0 | FA98B_HUMAN | | cytoplasm. |
| Q52LR7 | EPC2_HUMAN | nucleus (by similarity). | nucleus. |
| Q52M93 | Z585B_HUMAN | nucleus (potential). | nucleus. |
| Q52WX2 | SBK1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q53FA7 | QORX_HUMAN | | cytoplasm. |
| Q53FD0 | CN140_HUMAN | | nucleus. |
| Q53FE4 | CD017_HUMAN | | nucleus. |

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| Q53FT3 | CK073_HUMAN | cytoplasm (by similarity). | secreted protein. |
| Q53FV1 | ORML2_HUMAN | membrane; multi-pass membrane protein (potential). endoplasmic reticulum. | endoplasmic reticulum. |
| Q53G59 | KLH12_HUMAN | | cytoplasm. |
| Q53GD3 | CTL4_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q53GI3 | ZN394_HUMAN | nucleus (potential). | nucleus. |
| Q53GT1 | KLH22_HUMAN | | cytoplasm. |
| Q53H80 | CF166_HUMAN | | nucleus. |
| Q53HC5 | KLH26_HUMAN | | cytoplasm. |
| Q53RE8 | ANR39_HUMAN | | cytoplasm. |
| Q53S33 | BOLA3_HUMAN | | nucleus. |
| Q53T03 | RBP22_HUMAN | | cytoplasm. |
| Q53TS8 | AL2SA_HUMAN | | nucleus. |
| Q562E7 | WDR81_HUMAN | | membrane. |
| Q562F6 | SGOL2_HUMAN | nucleus (by similarity). localizes to the centromere (by similarity). | nucleus. |
| Q567U6 | CCD93_HUMAN | | cytoskeleton. |
| Q569K4 | ZN533_HUMAN | nucleus (potential). | nucleus. |
| Q56NI9 | ESCO2_HUMAN | nucleus (probable). | nucleus. |
| Q56UN5 | YSK4_HUMAN | | cytoplasm. |
| Q580R0 | CB027_HUMAN | | cytoplasm. |
| Q587I9 | SFT2C_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. Golgi. |
| Q58EX2 | SDK2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q58EX7 | PKHG4_HUMAN | | cytoplasm. |
| Q58F21 | BRDT_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q58HT5 | AWAT1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q5BJE1 | CR034_HUMAN | | cytoskeleton. |
| Q5BKY1 | LRC10_HUMAN | nucleus (by similarity). | centriole. |
| Q5BKZ1 | ZN326_HUMAN | nucleus; nucleoplasm; nuclear matrix (by similarity). | cytoplasm. nucleus. |
| Q5BN46 | CI116_HUMAN | | membrane. |
| Q5BVD1 | TTMP_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (potential). | endoplasmic reticulum. |
| Q5EBL2 | ZN628_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q5EBL8 | PDZ11_HUMAN | isoform 2: secreted protein (potential). | cytoplasm. |
| Q5EBM4 | ZN542_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q5FYB0 | ARSJ_HUMAN | secreted protein (potential). | endoplasmic reticulum. secreted protein. lysosome. |
| Q5FYB1 | ARSI_HUMAN | secreted protein (potential). | endoplasmic reticulum. secreted protein. lysosome. |
| Q5GAN6 | RNS10_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q5GH70 | XKR9_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5GH72 | XKR7_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5GH73 | XKR6_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5GH76 | XKR4_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |

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| Q5GH77 | XKR3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5H8C1 | FREM1_HUMAN | secreted protein; extracellular space; extracellular matrix; basement membrane (by similarity). localizes at the basement membrane zone of embryonic epidermis and hair follicles (by similarity). | membrane. |
| Q5H9L2 | TCAL5_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q5H9R4 | ARMX4_HUMAN | | endosome; membrane. |
| Q5H9R7 | SAPS3_HUMAN | | cytoplasm. |
| Q5H9T9 | CN155_HUMAN | | cytoplasm. |
| Q5HYA8 | MKS3_HUMAN | cell membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5HYJ3 | FA76B_HUMAN | | nucleus. |
| Q5HYK3 | COQ5_HUMAN | mitochondrion (by similarity). | cytoplasm. mitochondrion. |
| Q5HYR2 | DMRTC_HUMAN | | cytoplasm. |
| Q5I7T1 | AG10B_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q5J5C9 | DB121_HUMAN | secreted protein (potential). | secreted protein. |
| Q5JPB2 | CT174_HUMAN | | nucleus. |
| Q5JPI3 | CC038_HUMAN | | cytoplasm. |
| Q5JRC9 | FA47A_HUMAN | | nucleus. |
| Q5JRK9 | GGEE3_HUMAN | | cytoplasm. |
| Q5JSJ4 | DX26B_HUMAN | | cytoplasm. |
| Q5JSP0 | FGD3_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q5JT25 | RAB41_HUMAN | | Golgi. |
| Q5JT82 | KLF17_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q5JTD0 | TJAP1_HUMAN | Golgi apparatus (by similarity). cell membrane; cell-cell junction; tight junction (by similarity). recruited to tight junctions (tj) during late stages of maturation of the tj complexes. excluded from adherens junctions and desmosomes (by similarity). | cytoplasm. |
| Q5JTV8 | TOIP1_HUMAN | nucleus; nuclear inner membrane; single-pass membrane protein (by similarity). | endosome. |
| Q5JTZ5 | CI152_HUMAN | | cytoplasm. |
| Q5JU69 | TOR2A_HUMAN | secreted protein (probable). | cytoplasm. |
| Q5JU85 | IQEC2_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q5JUK3 | KCNT1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q5JUK9 | GGED1_HUMAN | | cytoplasm. |
| Q5JUW0 | ZN673_HUMAN | | nucleus. |
| Q5JVF3 | PCID2_HUMAN | | nucleus. |
| Q5JVG2 | ZN484_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q5JVG8 | ZN506_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q5JVL4 | EFHC1_HUMAN | | cytoplasm. |
| Q5JW98 | CF078_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5JWF8 | CT134_HUMAN | secreted protein (potential). | cytoplasm. |
| Q5JX69 | CT107_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q5JZY3 | EPHAA_HUMAN | isoform 1, isoform 3: cell membrane; single- pass type i membrane protein (probable). isoform 2: secreted protein | membrane. |
| Q5K4L6 | S27A3_HUMAN | mitochondrion; mitochondrial membrane; multi-pass membrane protein (by | endoplasmic reticulum. mitochondrion. |

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| Q5K651 | SAMD9_HUMAN | | cytoplasm. |
| Q5MAI5 | CDKL4_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q5MCW4 | ZN569_HUMAN | nucleus (by similarity). | nucleus. |
| Q5MJ68 | SPDYC_HUMAN | nucleus (by similarity). | nucleus. |
| Q5MNV8 | FBX47_HUMAN | | cytoplasm. |
| Q5MNZ6 | WIPI3_HUMAN | | nucleus. |
| Q5QGT7 | RTP2_HUMAN | cell membrane; single-pass type iii membrane protein (by similarity). effective cell surface expression depends upon interaction with olfactory receptors (by similarity). | endosome; membrane. |
| Q5QJ74 | TBCEL_HUMAN | | cytoplasm. |
| Q5RI15 | FA36A_HUMAN | | membrane. |
| Q5SGD2 | PP2CL_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q5SQ80 | A20A4_HUMAN | | nucleus. |
| Q5SQH8 | CF136_HUMAN | | membrane. |
| Q5SQQ9 | VAX1_HUMAN | nucleus (by similarity). | nucleus. |
| Q5SQS8 | CJ120_HUMAN | | nucleus. |
| Q5SR56 | HIAL1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q5SR76 | CJ093_HUMAN | | cytoplasm. |
| Q5SRN2 | CF010_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5SSQ6 | G7D_HUMAN | | secreted protein. |
| Q5SVZ6 | ZMYM1_HUMAN | | nucleus. |
| Q5SWW7 | CJ055_HUMAN | | nucleus. |
| Q5SXM1 | ZN678_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q5SXM2 | SNPC4_HUMAN | nucleus (potential). | nucleus. |
| Q5SY85 | CI122_HUMAN | | secreted protein. |
| Q5SZD1 | CF141_HUMAN | | nucleus. |
| Q5SZK8 | FREM2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q5T089 | MORN1_HUMAN | | secreted protein. |
| Q5T1B1 | CJ091_HUMAN | | nucleus. |
| Q5T1J6 | CT091_HUMAN | | nucleus. |
| Q5T200 | ZC313_HUMAN | | cytoplasm. nucleus. |
| Q5T280 | CI114_HUMAN | | nucleus. |
| Q5T2D3 | OTUD3_HUMAN | | cytoplasm. |
| Q5T2R2 | DPS1_HUMAN | | cytoplasm. |
| Q5T2S8 | ARMC4_HUMAN | | cytoplasm. |
| Q5T2W1 | PDZD1_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). associated with peripheral membranes. localizes to the apical compartment of proximal tubular cells and to sinusoidal liver membranes (by | cytoplasm. |
| Q5T447 | HECD3_HUMAN | | cytoplasm. |
| Q5T4F4 | ZFY27_HUMAN | membrane; multi-pass membrane protein (potential). punctate vesicles. | cytoplasm. |
| Q5T4F7 | SFRP5_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q5T4H9 | CJ114_HUMAN | | nucleus. |
| Q5T4W7 | ARTN_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q5T5A8 | LCE3C_HUMAN | | cytoplasm. |
| Q5T5B0 | LCE3E_HUMAN | | cytoplasm. |

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| Q5T5C0 | STXB5_HUMAN | cytoplasm (by similarity). cell membrane; peripheral membrane protein (by similarity). cytoplasmic vesicle; cytoplasmic vesicle membrane; peripheral membrane protein (by similarity). cytoplasmic, and associated with vesicular membranes and the plasma membrane. synapse (by similarity). synaptic vesicle (by similarity). detected at synapses and nucleus (probable). | Golgi. |
| Q5T5D7 | ZN684_HUMAN | | cytoplasm. nucleus. |
| Q5T5N4 | CF118_HUMAN | | nucleus. |
| Q5T6M2 | CF122_HUMAN | | nucleus. |
| Q5T6V5 | CI064_HUMAN | | cytoplasm. |
| Q5T6X5 | GPC6A_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q5T742 | CJ025_HUMAN | | Golgi. |
| Q5T751 | LCE1C_HUMAN | | cytoplasm. |
| Q5T752 | LCE1D_HUMAN | | cytoplasm. |
| Q5T753 | LCE1E_HUMAN | | cytoplasm. |
| Q5T754 | LCE1F_HUMAN | | cytoplasm. |
| Q5T7N3 | ANR38_HUMAN | | cytoplasm. |
| Q5T7P2 | LCE1A_HUMAN | | cytoplasm. |
| Q5T7P3 | LCE1B_HUMAN | | cytoplasm. |
| Q5T7P8 | SYT6_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q5T7W7 | CI097_HUMAN | | cytoplasm. |
| Q5T8I3 | F102B_HUMAN | | cytoplasm. |
| Q5T8R8 | CI066_HUMAN | | nucleus. |
| Q5T9A4 | ATD3B_HUMAN | | cytoplasm. nucleus. |
| Q5T9G4 | CF081_HUMAN | | cytoplasm. |
| Q5TA76 | LCE3A_HUMAN | | cytoplasm. |
| Q5TA77 | LCE3B_HUMAN | | cytoplasm. |
| Q5TA78 | LCE4A_HUMAN | | cytoplasm. |
| Q5TA79 | LCE2A_HUMAN | | cytoplasm. |
| Q5TA81 | LCE2C_HUMAN | | cytoplasm. |
| Q5TA82 | LCE2D_HUMAN | | cytoplasm. |
| Q5TAP6 | UT14C_HUMAN | nucleus; nucleolus (by similarity). | nucleus. |
| Q5TAX3 | ZCH11_HUMAN | | cytoplasm. nucleus. |
| Q5TBB1 | RNH2B_HUMAN | nucleus (probable). | centriole. cytoplasm. |
| Q5TBE3 | CI153_HUMAN | | cytoplasm. |
| Q5TC79 | ZBT37_HUMAN | nucleus (probable). | nucleus. |
| Q5TC82 | RC3H1_HUMAN | cytoplasm (by similarity). localizes to cytosolic rna granules, which are sites for regulating mrna translation and decay, suggesting that it may bind rna (by similarity). | nucleus. |
| Q5TCM9 | LCE5A_HUMAN | | cytoplasm. |
| Q5TCX8 | M3KL4_HUMAN | | cytoplasm. |
| Q5TCY1 | TTBK1_HUMAN | | cytoplasm. |
| Q5TDP6 | GLUD1_HUMAN | | cytoplasm. |
| Q5TEA3 | CT194_HUMAN | | membrane. |
| Q5TEJ8 | ICB1_HUMAN | | cytoplasm. |
| Q5TFE4 | NT5D1_HUMAN | | cytoplasm. |
| Q5TGI0 | CF168_HUMAN | | cytoplasm. |
| Q5TYW2 | A20A1_HUMAN | | cytoplasm; membrane. |

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| Q5TYW8 | PRAM3_HUMAN | | cytoplasm. |
| Q5TYX0 | PRAM5_HUMAN | | cytoplasm. |
| Q5TZF3 | ANR45_HUMAN | | nucleus. |
| Q5U5R9 | HECD2_HUMAN | | cytoplasm. |
| Q5UE93 | PI3R6_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q5VSJ3 | FA23B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5VSL9 | FA40A_HUMAN | | cytoplasm. |
| Q5VT25 | MRCKA_HUMAN | cytoplasm (by similarity). displays a dispersed punctate distribution and concentrates along the cell periphery, especially at the leading edge and cell-cell junction. this concentration is ph-domain dependent (by similarity). | cytoplasm. |
| Q5VT52 | K0460_HUMAN | | nucleus. |
| Q5VTR2 | BRE1A_HUMAN | nucleus (probable). | nucleus. |
| Q5VTY9 | HHAT_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q5VU65 | P210L_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q5VUG0 | SMBT2_HUMAN | | centriole. cytoplasm. |
| Q5VUM1 | CF057_HUMAN | secreted protein (potential). | nucleus. |
| Q5VUR7 | A20A3_HUMAN | | nucleus. |
| Q5VV41 | ARHGG_HUMAN | | cytoplasm. |
| Q5VV43 | K0319_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q5VV52 | ZN691_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q5VVB8 | CF191_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5VVJ2 | MYSM1_HUMAN | nucleus (probable). | nucleus. |
| Q5VVX9 | UBE2U_HUMAN | | cytoplasm. |
| Q5VWM4 | PRAM8_HUMAN | | cytoplasm. |
| Q5VWM5 | PRAM9_HUMAN | | cytoplasm. |
| Q5VWP3 | CF142_HUMAN | | nucleus. |
| Q5VWZ2 | LYPL1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q5VXH4 | PRAM6_HUMAN | | cytoplasm. |
| Q5VXH5 | PRAM7_HUMAN | | cytoplasm. |
| Q5VXT5 | SYPL2_HUMAN | membrane; multi-pass membrane protein (by similarity). triad junction, the junctional complex between the transverse tubule and the sarcoplasmic reticulum (by similarity). | membrane. |
| Q5VXU3 | CHIC1_HUMAN | cell membrane (by similarity). cytoplasmic vesicle (by similarity). also present at a Golgi-like vesicular compartment and at scattered vesicles (by similarity). | cytoplasm. Golgi. |
| Q5VXU9 | CI084_HUMAN | | nucleus. |
| Q5VY09 | IER5_HUMAN | | membrane. |
| Q5VYS8 | ZCHC6_HUMAN | | cytoplasm. nucleus. |
| Q5VYY1 | ANR22_HUMAN | | cytoplasm. |
| Q5VZ03 | CI121_HUMAN | | mitochondrion. |
| Q5VZ18 | SHE_HUMAN | | cytoplasm. |
| Q5VZ89 | DEN4C_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5VZB9 | DMRTA_HUMAN | nucleus (potential). | nucleus. |
| Q5VZQ5 | CJ122_HUMAN | | nucleus. |

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| Q5VZR4 | HIAL2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q5W0B1 | CM007_HUMAN | | cytoplasm. |
| Q5W0B7 | FA23A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q5W0N0 | CI057_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q5W0U4 | BSPRY_HUMAN | cytoplasm. membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q5W0Z9 | ZDH20_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q5W111 | CLLD6_HUMAN | | membrane. |
| Q5XG87 | POLS_HUMAN | nucleus (probable). | nucleus. |
| Q5XG99 | LYSM4_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q5XKK7 | CO017_HUMAN | | cytoplasm. |
| Q5XKL5 | BTBD8_HUMAN | | cytoplasm. |
| Q5XKR9 | F104B_HUMAN | | cytoplasm. |
| Q5XUX0 | FBX31_HUMAN | | cytoplasm. |
| Q5XUX1 | FBXW9_HUMAN | | cytoplasm. |
| Q5YKI7 | GGNB1_HUMAN | cytoplasm. membrane; peripheral membrane protein. Golgi apparatus (by similarity). | cytoplasm. cytoskeleton; membrane. |
| Q5ZPR3 | CD276_HUMAN | membrane; single-pass type i membrane protein (probable). | endosome; membrane. |
| Q63HK3 | ZN694_HUMAN | nucleus (probable). | nucleus. |
| Q63HK5 | TSH3_HUMAN | nucleus (probable). | nucleus. |
| Q63HM1 | AFMID_HUMAN | cytoplasm; cytosol (by similarity). predominantly cytosolic. nucleus. some fraction is nuclear (by similarity). | centriole. cytoplasm. |
| Q63HQ0 | CD016_HUMAN | | Golgi. |
| Q63ZE4 | OAT5_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q63ZY3 | ANR25_HUMAN | | cytoplasm. |
| Q66LE6 | 2ABD_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q676U5 | A16L1_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). localized to preautophagosomal structure (pas) where it is involved in the membrane targeting of atg5 (by similarity). | endosome. |
| Q68CL5 | TPGS2_HUMAN | nucleus (potential). | centriole. cytoplasm. nucleus. |
| Q68D51 | DEN2C_HUMAN | | nucleus. |
| Q68D86 | C102B_HUMAN | | centriole. |
| Q68DC2 | ANKS6_HUMAN | | cytoplasm. |
| Q68DD2 | PA24F_HUMAN | cytoplasm; cytosol. translocates to lysosomal membranes in a calcium-dependent fashion (by similarity). | secreted protein. |
| Q68DN6 | RBP26_HUMAN | | cytoplasm. |
| Q68DU8 | KCD16_HUMAN | | nucleus; membrane. |
| Q69384 | ENK2_HUMAN | | endosome; membrane. |
| Q693B1 | KCD11_HUMAN | | cytoplasm. nucleus; membrane. |
| Q695T7 | S6A19_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q69YI7 | CI090_HUMAN | | nucleus. |
| Q69YQ0 | CYTSA_HUMAN | | cytoplasm. |
| Q6AHZ1 | ZN518_HUMAN | nucleus (probable). | cytoplasm. nucleus. |

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| Q6AI12 | ANR40_HUMAN | | cytoplasm. |
| Q6AI39 | K0240_HUMAN | | nucleus. |
| Q6AZW8 | ZN660_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q6B0I6 | JHD3D_HUMAN | nucleus (by similarity). | nucleus. |
| Q6B9Z1 | IGFL4_HUMAN | secreted protein (potential). | secreted protein. |
| Q6BDS2 | URFB1_HUMAN | | cytoplasm. |
| Q6BEB4 | SP5_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q6DHV5 | CJ130_HUMAN | | membrane. |
| Q6DKI1 | RL7L_HUMAN | | mitochondrion. |
| Q6DKK2 | TTC19_HUMAN | | cytoplasm. |
| Q6EEV6 | SUMO4_HUMAN | | nucleus. |
| Q6EIG7 | CLC6A_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q6EMB2 | TTLL5_HUMAN | | cytoplasm. mitochondrion. |
| Q6EMK4 | VASN_HUMAN | membrane; single-pass type i membrane protein (probable). secreted protein. | membrane. |
| Q6F181 | CPIN1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q6FIF0 | ZFAN6_HUMAN | | nucleus. |
| Q6GMV2 | SMYD5_HUMAN | | nucleus. |
| Q6GYQ0 | GRIPE_HUMAN | cytoplasm (by similarity). nucleus (by similarity). translocated to the nucleus, when associated with tcf3/e12 (by similarity). | centriole. cytoplasm. |
| Q6H8Q1 | ABLM2_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q6HA08 | ASTL_HUMAN | | membrane. |
| Q6IA17 | SIGIR_HUMAN | membrane; single-pass type iii membrane protein (potential). | membrane. |
| Q6IA69 | NADE1_HUMAN | | cytoplasm. |
| Q6IA86 | STTP1_HUMAN | | nucleus. |
| Q6IFS5 | HSN2_HUMAN | secreted protein (potential). | cytoplasm. secreted protein. |
| Q6IMI4 | ST6B1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q6IMI6 | ST1C3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q6IPX3 | TCAL6_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q6IQ26 | RA6I1_HUMAN | | cytoplasm. |
| Q6IQ55 | TTBK2_HUMAN | | cytoplasm. |
| Q6IS24 | GLTL3_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q6IV72 | ZN425_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q6J4K2 | NCKX6_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q6J9G0 | STYK1_HUMAN | cytoplasm (by similarity). | membrane. |
| Q6JEL2 | KLH10_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q6JVE5 | LCN12_HUMAN | secreted protein (potential). | cytoplasm. |
| Q6JVE6 | LCN10_HUMAN | secreted protein (potential). | cytoplasm. |
| Q6JVE9 | LCN8_HUMAN | secreted protein (by similarity). | cytoplasm. |
| Q6KC79 | NIPBL_HUMAN | nucleus (by similarity). | nucleus. |
| Q6KF10 | GDF6_HUMAN | secreted protein (probable). | secreted protein. |
| Q6L8G4 | KR511_HUMAN | | cytoplasm. |
| Q6L8G5 | KR510_HUMAN | | cytoplasm. |
| Q6L8G8 | KRA57_HUMAN | | cytoplasm. |
| Q6L8G9 | KRA56_HUMAN | | cytoplasm. |
| Q6L8H1 | KRA54_HUMAN | | cytoplasm. |
| Q6L8H2 | KRA53_HUMAN | | cytoplasm. |
| Q6L8H4 | KRA51_HUMAN | | cytoplasm. |
| Q6N043 | SUHW4_HUMAN | nucleus (potential). | cytoplasm. nucleus. |

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|--------|-------------|--|-----------------------------------|
| Q6N069 | NARGL_HUMAN | | cytoplasm. |
| Q6NS38 | ALKB2_HUMAN | | nucleus. |
| Q6NSJ2 | PHLB3_HUMAN | | centriole. |
| Q6NSJ5 | LRC8E_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q6NSW5 | FA45B_HUMAN | | cytoplasm. |
| Q6NSX1 | CCD70_HUMAN | secreted protein (potential). | cytoplasm; membrane. |
| Q6NT76 | HMBX1_HUMAN | nucleus (potential). | nucleus. |
| Q6NUJ5 | PWWP2_HUMAN | | nucleus. |
| Q6NUK4 | REEP3_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q6NUM9 | RETST_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; peripheral membrane protein (by similarity). | endoplasmic reticulum. microsome. |
| Q6NUQ1 | RINT1_HUMAN | cytoplasm (probable). endoplasmic reticulum. the snare complex localizes to the endoplasmic reticulum. | endoplasmic reticulum. |
| Q6NUS6 | TECT3_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q6NVV3 | NPAL1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q6NW29 | RWDD4_HUMAN | | nucleus. |
| Q6NW34 | CC017_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q6NW40 | RGMB_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | secreted protein. |
| Q6NX49 | ZN544_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q6NXE6 | ARMC6_HUMAN | | cytoplasm. |
| Q6NXP0 | CC025_HUMAN | | cytoplasm. |
| Q6NY19 | ANR47_HUMAN | | cytoplasm. |
| Q6NYC8 | K1949_HUMAN | | cytoplasm. |
| Q6NZY4 | ZCHC8_HUMAN | | cytoplasm. nucleus. |
| Q6NZY7 | BORG3_HUMAN | intracytoplasmic membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q6P050 | FXL22_HUMAN | | cytoplasm. |
| Q6P0N0 | CV106_HUMAN | nucleus (probable). | nucleus. |
| Q6P158 | DHX57_HUMAN | | nucleus. |
| Q6P1A2 | MBOA5_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q6P1K2 | PMF1_HUMAN | nucleus. associated with the kinetochore (probable). | nucleus. |
| Q6P1L6 | ZN343_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q6P1M0 | S27A4_HUMAN | membrane; multi-pass membrane protein (probable). | endoplasmic reticulum. |
| Q6P1M9 | ARMX5_HUMAN | | cytoplasm. |
| Q6P1R4 | DUS1L_HUMAN | | mitochondrion. |
| Q6P2D0 | ZFP1_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q6P2H3 | CCD21_HUMAN | | cytoplasm. |
| Q6P2M8 | KCC1B_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q6P2Q9 | PRP8_HUMAN | nucleus; nucleoplasm; nuclear speckle (by similarity). | nucleus. |
| Q6P3S6 | FBX42_HUMAN | | cytoplasm. |
| Q6P3V2 | Z585A_HUMAN | nucleus (potential). | nucleus. |
| Q6P3W2 | ZCSL3_HUMAN | | cytoplasm. nucleus. |
| Q6P444 | FA54A_HUMAN | | nucleus. |

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| Q6P499 | NPAL3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q6P4A7 | SFXN4_HUMAN | mitochondrion; mitochondrial membrane; multi-pass membrane protein (by | mitochondrion; membrane. |
| Q6P4R8 | NFRKB_HUMAN | nucleus (probable). | nucleus. |
| Q6P587 | FAHD1_HUMAN | | membrane. |
| Q6P597 | KLC3_HUMAN | elongating spermatid tail midpiece, localized in outer dense fibers (odfs) and associates with mitochondria (by | cytoplasm. |
| Q6P5R6 | RL22L_HUMAN | | mitochondrion. |
| Q6P656 | CO026_HUMAN | | cytoplasm. |
| Q6P6B7 | ANR16_HUMAN | | cytoplasm. |
| Q6P6C2 | ALKB5_HUMAN | | endosome; membrane. |
| Q6P9A2 | GLTL4_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q6P9A3 | ZN549_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q6P9F0 | CCD62_HUMAN | | nucleus. |
| Q6P9F5 | TRI40_HUMAN | | cytoplasm. |
| Q6P9F7 | LRC8B_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q6P9G9 | ZN449_HUMAN | nucleus (potential). | nucleus. |
| Q6PCB7 | S27A1_HUMAN | cell membrane; peripheral membrane protein. intracytoplasmic membrane; peripheral membrane protein. plasma membrane and intracellular membranes, at least in adipocytes. cytoplasm (by similarity). predominantly cytoplasmic in myocytes (by similarity). | endoplasmic reticulum. |
| Q6PCB8 | EMB_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| Q6PCE3 | PGM2L_HUMAN | | cytoplasm. |
| Q6PCT2 | FXL19_HUMAN | | cytoplasm. |
| Q6PD62 | CTR9_HUMAN | nucleus. found in speckles (by similarity). | centriole. nucleus. |
| Q6PD74 | P34_HUMAN | | cytoplasm. |
| Q6PEW1 | ZCH12_HUMAN | | cytoplasm. nucleus. |
| Q6PEX3 | KR261_HUMAN | | cytoplasm. |
| Q6PF04 | ZN613_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q6PF18 | MORN3_HUMAN | | nucleus. |
| Q6PH85 | DCNL2_HUMAN | | cytoplasm. |
| Q6PHR2 | ULK3_HUMAN | | cytoplasm. |
| Q6PI25 | CNIH2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q6PI47 | KCD18_HUMAN | | nucleus; membrane. |
| Q6PI98 | CR037_HUMAN | | nucleus. |
| Q6PIJ6 | FBX38_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | centriole. cytoplasm. nucleus. |
| Q6PIL6 | KCIP4_HUMAN | cell membrane; peripheral membrane protein (by similarity). | membrane. |
| Q6PJ61 | FBX46_HUMAN | | cytoplasm. |
| Q6PJ69 | TRI65_HUMAN | | cytoplasm. |
| Q6PJE2 | POZP3_HUMAN | | centriole. cytoplasm. |
| Q6PJG9 | LRFN4_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q6PKG0 | LARP1_HUMAN | | cytoplasm. |
| Q6PKX4 | DOK6_HUMAN | | centriole. secreted protein. |

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| Q6PL18 | ATAD2_HUMAN | | cytoplasm. nucleus. |
| Q6PL24 | TMED8_HUMAN | | Golgi. |
| Q6PP77 | XKR2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q6PUV4 | CPLX2_HUMAN | cytoplasm; cytosol. enriched at synaptic-releasing sites in mature neurons (by similarity). | cytoplasm. |
| Q6Q6R5 | CRIP3_HUMAN | cytoplasm (by similarity). | cytoplasm. nucleus. |
| Q6QHF9 | PAOX_HUMAN | peroxisome (by similarity). cytoplasm (by similarity). | peroxisome. |
| Q6QHK4 | FIGLA_HUMAN | nucleus (probable). | nucleus. |
| Q6QNY0 | BL1S3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q6R2W3 | ZN452_HUMAN | | nucleus. |
| Q6S545 | POT22_HUMAN | | secreted protein. |
| Q6S5H4 | POT15_HUMAN | | nucleus. |
| Q6S5H5 | POT14_HUMAN | | cytoplasm. |
| Q6S8J7 | POTE8_HUMAN | | nucleus. |
| Q6SA08 | TSSK4_HUMAN | | cytoplasm. |
| Q6SZW1 | SARM1_HUMAN | cytoplasm (by similarity). | membrane. |
| Q6T311 | ARL9_HUMAN | | cytoplasm. |
| Q6T4R5 | NHS_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q6TCH4 | PAQR6_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q6TCH7 | PAQR3_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q6TDP4 | KLH17_HUMAN | postsynaptic density (by similarity). | cytoplasm. |
| Q6TFL3 | CI093_HUMAN | | cytoskeleton. |
| Q6TGC4 | PADI6_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q6U841 | S4A10_HUMAN | cell membrane; multi-pass membrane protein. localizes to the basolateral membrane (by similarity). | membrane. |
| Q6U949 | IG2AS_HUMAN | | nucleus. |
| Q6UVK1 | CSPG4_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). localized at the apical plasma membrane it relocalizes to the lamellipodia of astrocytoma upon phosphorylation by prkca. localizes to the retraction fibers. localizes to the plasma membrane of oligodendrocytes (by similarity). | membrane. |
| Q6UW01 | CBLN3_HUMAN | secreted protein (potential). | membrane. |
| Q6UW10 | SFTPG_HUMAN | secreted protein (probable). | membrane. |
| Q6UW32 | IGFL1_HUMAN | secreted protein (probable). | secreted protein. |
| Q6UW56 | APR3_HUMAN | | membrane. |
| Q6UW60 | PCSK4_HUMAN | membrane; single-pass membrane protein (potential). | secreted protein; membrane. |
| Q6UW63 | KDEL1_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| Q6UW78 | U655_HUMAN | secreted protein (probable). | secreted protein. |
| Q6UW88 | EPGN_HUMAN | membrane; single-pass type i membrane protein (probable). | endosome; membrane. |
| Q6UWE3 | U3045_HUMAN | secreted protein (probable). | membrane. |
| Q6UWF7 | FA55D_HUMAN | secreted protein (potential). | membrane. |
| Q6UWH6 | TX261_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |

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| Q6UWI2 | PARM1_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q6UWI4 | TMM46_HUMAN | membrane; single-pass type i membrane protein (probable). | endosome; membrane. |
| Q6UWJ8 | C16L2_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q6UWL6 | KIRR2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q6UWM7 | LCTL_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q6UWN0 | LYPD4_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | secreted protein. |
| Q6UWN5 | LYPD5_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | secreted protein. |
| Q6UWN8 | ISK6_HUMAN | secreted protein (potential). | secreted protein. |
| Q6UWP2 | DHR11_HUMAN | secreted protein (potential). | cytoplasm. mitochondrion. |
| Q6UWQ5 | LYZL1_HUMAN | secreted protein (probable). | lysosome. |
| Q6UWQ7 | IGFL2_HUMAN | secreted protein (potential). | secreted protein. |
| Q6UWR7 | ENPP6_HUMAN | cell membrane; single-pass type i membrane protein (probable). a minor secreted form also exists. | endosome. |
| Q6UWU4 | CF089_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q6UWV6 | ENPP7_HUMAN | membrane; single-pass type i membrane protein (potential). localized at the surface of the microvillar membrane in small intestine enterocytes, and in endosome-like structures situated beneath the microvillar membrane, and in Golgi | endosome. |
| Q6UWY0 | ARSK_HUMAN | secreted protein (potential). | endoplasmic reticulum. secreted protein. lysosome. |
| Q6UWY5 | OLFL1_HUMAN | secreted protein (potential). | cytoplasm. |
| Q6UX02 | AN36A_HUMAN | | cytoplasm. |
| Q6UX39 | U689_HUMAN | secreted protein (probable). | nucleus. |
| Q6UX41 | BTNL8_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| Q6UX68 | XKR5_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q6UX71 | PXDC2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q6UX98 | ZDH24_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q6UXA7 | CF015_HUMAN | secreted protein (potential). | nucleus. |
| Q6UXB0 | CC040_HUMAN | secreted protein (potential). | nucleus. |
| Q6UXB1 | IGFL3_HUMAN | secreted protein (probable). | secreted protein. |
| Q6UXB3 | LYPD2_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | secreted protein. |
| Q6UXB4 | CLC4G_HUMAN | membrane; single-pass type ii membrane protein (potential). | endosome; membrane. |
| Q6UXE8 | BTNL3_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| Q6UXG8 | BTNL9_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| Q6UXI7 | VITRN_HUMAN | secreted protein (potential). | membrane. |

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|--------|-------------|--|------------------------|
| Q6UXK5 | LRRN1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q6UXL0 | I20RB_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q6UXM1 | LRIG3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q6UXN8 | CLC9A_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q6UXR4 | SPA13_HUMAN | secreted protein (potential). | secreted protein. |
| Q6UXV0 | GFRAL_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q6UXV4 | CX033_HUMAN | secreted protein (potential). | nucleus. |
| Q6UXX9 | RSPO2_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q6UXZ0 | TMIGD_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q6UXZ4 | UNC5D_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q6UY01 | LRC31_HUMAN | | cytoplasm. |
| Q6UY09 | CEA20_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q6UY11 | EGFL9_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q6V0L0 | CP26C_HUMAN | membrane; single-pass membrane protein (potential). | endoplasmic reticulum. |
| Q6V9R5 | ZN562_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q6VAB6 | KSR2_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | cytoplasm. |
| Q6VY07 | PACS1_HUMAN | Golgi apparatus; trans-Golgi network (by similarity). localizes in the paranuclear region, probably the tgn (by similarity). | cytoplasm. |
| Q6W0C5 | DPPA3_HUMAN | nucleus (by similarity). cytoplasm (by similarity). | centriole. cytoplasm. |
| Q6WBX8 | RAD9B_HUMAN | | cytoplasm. |
| Q6WN34 | CRDL2_HUMAN | isoform 1, isoform 2: secreted protein (potential). isoform 3, isoform 4, isoform 5: cytoplasm (potential). | secreted protein. |
| Q6X4W1 | NELF_HUMAN | cell membrane; peripheral membrane protein (by similarity). found on the outside of the lhrh cell membrane and axons projecting from the olfactory pit and epithelium (by similarity). | nucleus. |
| Q6X784 | ZBPB2_HUMAN | secreted protein (potential). | secreted protein. |
| Q6X9E4 | FBW12_HUMAN | | cytoplasm. |
| Q6XPR3 | RPTN_HUMAN | | secreted protein. |
| Q6XUX3 | RIPK5_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q6XXX2 | CU024_HUMAN | | nucleus. |
| Q6XYQ8 | SYT10_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q6XZB0 | LIPI_HUMAN | secreted protein (potential). may associate with lipid draft. | secreted protein. |
| Q6XZF7 | DNMBP_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi stack (by similarity). localized to synapses and Golgi stacks (by similarity). | cytoplasm. |

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| Q6Y2X3 | DCJ14_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q6YFQ2 | CX6B2_HUMAN | mitochondrion; mitochondrial intermembrane space (by similarity). | mitochondrion. |
| Q6ZMG9 | LASS6_HUMAN | nucleus; nuclear membrane; multi-pass membrane protein (potential). endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | nucleus. |
| Q6ZMH5 | S39A5_HUMAN | cell membrane; basolateral cell membrane; multi-pass membrane protein | membrane. |
| Q6ZMI3 | GLDN_HUMAN | cell membrane; single-pass type ii membrane protein. localizes to the nodes of ranvier (by similarity). | membrane. |
| Q6ZMM2 | ATL5_HUMAN | secreted protein (potential). | secreted protein. |
| Q6ZMN7 | PZRN4_HUMAN | | cytoplasm. |
| Q6ZMQ8 | AATK_HUMAN | membrane; single-pass type i membrane protein (by similarity). cytoplasm. predominantly perinuclear. | membrane. |
| Q6ZMR3 | LDH6A_HUMAN | | cytoplasm. |
| Q6ZMT4 | K1718_HUMAN | | nucleus. |
| Q6ZMV9 | KIF6_HUMAN | | cytoplasm. nucleus. |
| Q6ZMY9 | ZN517_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q6ZMZ0 | IBRD3_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. |
| Q6ZN44 | UNC5A_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). the interaction with prkcabp regulates its surface expression and leads to its removal from surface of neurons and growth cones (by similarity). | membrane. |
| Q6ZN79 | Z705A_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q6ZNE5 | K0831_HUMAN | | cytoplasm. |
| Q6ZNG0 | ZN620_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q6ZNG1 | ZN600_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q6ZNH5 | ZN497_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q6ZNL6 | FGD5_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q6ZNR0 | TMM91_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q6ZP36 | AL2SF_HUMAN | | nucleus. |
| Q6ZR52 | ZN493_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q6ZR72 | CU125_HUMAN | | nucleus. |
| Q6ZRH7 | CS015_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q6ZRI6 | CO039_HUMAN | | nucleus. |
| Q6ZRZ4 | CI047_HUMAN | | cytoplasm. |
| Q6ZS30 | NBEAL_HUMAN | | nucleus. |
| Q6ZSA7 | LRC55_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q6ZSB9 | ZN509_HUMAN | nucleus (potential). | nucleus. |
| Q6ZSG1 | RN165_HUMAN | | cytoplasm. |
| Q6ZSI9 | CAN12_HUMAN | | cytoplasm. lysosome. |
| Q6ZSS3 | ZN621_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q6ZSZ6 | TSH1_HUMAN | nucleus (probable). | nucleus. |
| Q6ZT52 | FA43B_HUMAN | | nucleus. |

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| Q6ZT98 | TTLL7_HUMAN | | cytoplasm. mitochondrion. |
| Q6ZTN6 | AN13D_HUMAN | | cytoplasm. |
| Q6ZTQ3 | RASF6_HUMAN | | cytoplasm. |
| Q6ZTR5 | CX022_HUMAN | | membrane. |
| Q6ZTW0 | TPGS1_HUMAN | associated with microtubules from neurites, centrosomes, basal bodies and axonemes (by similarity). | cytoskeleton. |
| Q6ZU52 | K0408_HUMAN | | nucleus. |
| Q6ZU67 | CCDC4_HUMAN | | nucleus. |
| Q6ZU80 | CN145_HUMAN | | cytoplasm. |
| Q6ZUB1 | CI079_HUMAN | | endosome; membrane. |
| Q6ZUJ4 | CC062_HUMAN | | cytoplasm. |
| Q6ZUK4 | TMM26_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q6ZUX7 | LHPL2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q6ZV73 | FGD6_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q6ZV89 | SH2D5_HUMAN | | cytoplasm. |
| Q6ZVD8 | PHLPL_HUMAN | cytoplasm (probable). membrane; peripheral membrane protein (by similarity). may be membrane-associated | cytoplasm. |
| Q6ZVE7 | GOT1A_HUMAN | Golgi apparatus; Golgi membrane; multi-pass membrane protein (by similarity). | Golgi. |
| Q6ZVN8 | RGMC_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | secreted protein. |
| Q6ZVX9 | PAQR9_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q6ZVY0 | ENK19_HUMAN | | centriole. |
| Q6ZW76 | ANKS3_HUMAN | | cytoplasm. |
| Q6ZWJ1 | STXB4_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q6ZWL3 | CP4V2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (potential). | endoplasmic reticulum. |
| Q701N2 | KRA55_HUMAN | | cytoplasm. |
| Q701N4 | KRA52_HUMAN | | cytoplasm. |
| Q70CQ1 | UBP49_HUMAN | | nucleus. |
| Q70CQ2 | UBP34_HUMAN | | nucleus. |
| Q70CQ3 | UBP30_HUMAN | | nucleus. |
| Q70CQ4 | UBP31_HUMAN | | nucleus. |
| Q70EK8 | UBP53_HUMAN | | cytoplasm. |
| Q70EK9 | UBP51_HUMAN | | nucleus. |
| Q70EL1 | UBP54_HUMAN | | nucleus. |
| Q70EL3 | UBP50_HUMAN | | nucleus. |
| Q70EL4 | UBP43_HUMAN | | nucleus. |
| Q70IA8 | MOL2C_HUMAN | | cytoplasm. nucleus. |
| Q70JA7 | CHSS3_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q70YC4 | TALAN_HUMAN | | membrane. |
| Q70YC5 | ZN365_HUMAN | | cytoplasm. |
| Q719H9 | KCTD1_HUMAN | | nucleus; membrane. |
| Q71F56 | T240L_HUMAN | nucleus (by similarity). | nucleus. |
| Q71RC2 | LARP4_HUMAN | | nucleus. |
| Q71RG4 | TMUB2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |

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| Q71RH2 | FA57B_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q71RS6 | NCKX5_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q71U36 | TBA3_HUMAN | | centriole. |
| Q71UI9 | H2AV_HUMAN | nucleus (by similarity). | nucleus. |
| Q71UM5 | RS27L_HUMAN | | mitochondrion. |
| Q765I0 | UTS2B_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q76B58 | FAM5C_HUMAN | secreted protein (potential). | cytoplasm. |
| Q76I76 | SSH2_HUMAN | | cytoplasm. |
| Q7KYR7 | BT2A1_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q7KZ85 | SPT6H_HUMAN | nucleus (probable). | nucleus. |
| Q7L099 | RUFY3_HUMAN | | cytoplasm. |
| Q7L0J3 | SV2A_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein. enriched in chromaffin granules, not present in adrenal microsomes. associated with both insulin granules and synaptic-like microvesicles in insulin-secreting cells of the pancreas (by similarity). | membrane. |
| Q7L190 | DPPA4_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q7L1I2 | SV2B_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein. associated with synaptic-like microvesicles but not with insulin-containing vesicles in insulin-secreting cells of the pancreas. localizes to microvesicles in the pinealocytes. localizes to the acrosome in spermatids (by similarity). | membrane. |
| Q7L1S5 | CHST9_HUMAN | isoform 1: Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). isoform 2: secreted protein (probable). | membrane. |
| Q7L1W4 | LRC8D_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q7L273 | KCTD9_HUMAN | | nucleus; membrane. |
| Q7L2E3 | DHX30_HUMAN | | nucleus. |
| Q7L311 | ARMX2_HUMAN | | endosome. |
| Q7L3T8 | SYPM_HUMAN | mitochondrion; mitochondrial matrix (potential). | cytoplasm. |
| Q7L4S7 | ARMX6_HUMAN | | endosome; membrane. |
| Q7L513 | FCRL_HUMAN | secreted protein (probable). cytoplasm. | secreted protein. |
| Q7L591 | DOK3_HUMAN | cytoplasm (by similarity). | centriole. secreted protein. |
| Q7L5D6 | CG020_HUMAN | | cytoplasm. |
| Q7L5N7 | AYTL1_HUMAN | membrane; single-pass type ii membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q7L5Y6 | DET1_HUMAN | nucleus (probable). | centriole. cytoplasm. nucleus. |
| Q7L622 | K1333_HUMAN | | cytoplasm. |
| Q7L8A9 | VASH_HUMAN | secreted protein (probable). | cytoplasm. |
| Q7L8C5 | SYT13_HUMAN | membrane; single-pass membrane protein (by similarity). | endosome; membrane. |
| Q7L8S5 | OTU6A_HUMAN | | nucleus. |
| Q7LBC6 | JHD2B_HUMAN | nucleus (by similarity). | nucleus. |
| Q7LDI9 | GAK2_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |

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| Q7LFX5 | ST4S6_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). a small fraction may also be present at the cell surface, where it acts as a b-cell receptor. | membrane. |
| Q7LGA3 | HS2ST_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q7LGC8 | CHST3_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q7RTP0 | NIPA1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q7RTP6 | MICA3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q7RTS1 | BHLH8_HUMAN | nucleus (probable). | nucleus. |
| Q7RTS3 | PTF1A_HUMAN | nucleus (by similarity). cytoplasm (by similarity). in chronic pancreatitis associated with pancreas cancer preferentially accumulates in the cytoplasm of acinar/ductular complexes. in the cytoplasm loses its ability to form the ptf1 complex (by similarity). | nucleus. |
| Q7RTS9 | DYM_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q7RTT3 | SSX9_HUMAN | | nucleus. |
| Q7RTT4 | SSX8_HUMAN | | nucleus. |
| Q7RTT5 | SSX7_HUMAN | | nucleus. |
| Q7RTT6 | SSX6_HUMAN | | nucleus. |
| Q7RTU3 | OLIG3_HUMAN | nucleus (by similarity). | nucleus. |
| Q7RTU9 | STRC_HUMAN | cell surface (by similarity). associated with the hair bundle (by similarity). | centriole. cytoplasm. secreted protein. membrane. |
| Q7RTV0 | PHF5A_HUMAN | nucleus (by similarity). | nucleus. |
| Q7RTV2 | GSTA5_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q7RTV5 | CI021_HUMAN | | membrane. |
| Q7RTW8 | OTOAN_HUMAN | cell membrane; apical cell membrane; lipid- anchor; gpi-anchor; extracellular side (probable). at the interface between the apical surface of the epithelia and the overlying acellular gel of the tectorial and otoconial membranes (by similarity). | secreted protein. |
| Q7RTY8 | TMPS7_HUMAN | | secreted protein; membrane. |
| Q7Z2H8 | S36A1_HUMAN | cell membrane; multi-pass membrane protein (by similarity). lysosome; lysosomal membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q7Z2K8 | GRIN1_HUMAN | cell membrane; lipid-anchor (by similarity). highly enriched in growth cone (by similarity). | cytoplasm. Golgi. |
| Q7Z2V1 | TNT_HUMAN | | nucleus. |
| Q7Z2W4 | ZCC2_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q7Z2W7 | TRPM8_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q7Z2X7 | GGEE2_HUMAN | | cytoplasm. |
| Q7Z304 | MAMC2_HUMAN | secreted protein (potential). | endoplasmic reticulum. |
| Q7Z333 | SETX_HUMAN | nucleus (probable). | nucleus. |
| Q7Z340 | ZN551_HUMAN | nucleus (probable). | cytoplasm. nucleus. |

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| Q7Z398 | ZN550_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q7Z3B1 | NEGR1_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| Q7Z3B3 | K1267_HUMAN | | cytoplasm. |
| Q7Z3B4 | NUP54_HUMAN | nucleus; nuclear envelope; nuclear pore complex; cytoplasmic side; nucleoplasmic side. biased towards cytoplasmic side. central region of the nuclear pore complex, within the transporter (by similarity). | cytoplasm. nucleus. |
| Q7Z3C6 | ATG9A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. Golgi. |
| Q7Z3D4 | LYSM3_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q7Z3D6 | CN159_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q7Z3E1 | PARPT_HUMAN | | nucleus. |
| Q7Z3E2 | CJ118_HUMAN | | cytoplasm. |
| Q7Z3G6 | PRIC2_HUMAN | nucleus; nuclear membrane (potential). | cytoplasm. nucleus. |
| Q7Z3H0 | ANR33_HUMAN | | cytoplasm. |
| Q7Z3K3 | POGZ_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q7Z3V5 | ZN571_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q7Z3Z2 | CA036_HUMAN | | cytoplasm. |
| Q7Z3Z3 | PIWL3_HUMAN | | secreted protein. |
| Q7Z3Z4 | PIWL4_HUMAN | | secreted protein. |
| Q7Z401 | MYCPP_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q7Z406 | MYH14_HUMAN | | cytoskeleton. |
| Q7Z407 | CSMD3_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q7Z408 | CSMD2_HUMAN | | endosome; membrane. |
| Q7Z419 | IBRD2_HUMAN | | cytoplasm. |
| Q7Z444 | RASE_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (by similarity). | cytoplasm. |
| Q7Z465 | BNIPL_HUMAN | | cytoplasm. |
| Q7Z478 | DHX29_HUMAN | | nucleus. |
| Q7Z494 | NPHP3_HUMAN | | cytoplasm. |
| Q7Z4B0 | CR020_HUMAN | | nucleus. |
| Q7Z4F1 | LRP10_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q7Z4G1 | COMD6_HUMAN | | cytoplasm. |
| Q7Z4G4 | CF075_HUMAN | | nucleus. |
| Q7Z4H7 | FA29A_HUMAN | | nucleus. |
| Q7Z4H8 | KDEL2_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (potential). | endoplasmic reticulum. |
| Q7Z4K8 | TRI46_HUMAN | | cytoplasm. |
| Q7Z4P5 | GDF7_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q7Z4Q2 | HEAT3_HUMAN | | cytoplasm. |
| Q7Z4S6 | KI21A_HUMAN | | cytoplasm. nucleus. |
| Q7Z4T8 | GLTL5_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q7Z4V0 | ZN438_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q7Z4W1 | DCXR_HUMAN | membrane; peripheral membrane protein (by similarity). probably recruited to membranes via an interaction with phosphatidylinositol (by similarity). | endoplasmic reticulum. mitochondrion. |
| Q7Z4W2 | LYZL2_HUMAN | secreted protein (probable). | lysosome. |
| Q7Z4W3 | KR193_HUMAN | | cytoplasm. |

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| Q7Z4Y8 | AT5L2_HUMAN | | mitochondrion. |
| Q7Z553 | MAMC1_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q7Z570 | CB010_HUMAN | | nucleus. |
| Q7Z5J4 | RAI1_HUMAN | cytoplasm. nucleus. in neurons, localized to neurites (by similarity). | nucleus. |
| Q7Z5L0 | VMO1_HUMAN | secreted protein (probable). | membrane. |
| Q7Z5L2 | GIDRP_HUMAN | | nucleus. |
| Q7Z5M8 | CN029_HUMAN | | microsome. |
| Q7Z5N4 | SDK1_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q7Z5P4 | DHB13_HUMAN | secreted protein (potential). | centriole. cytoplasm. |
| Q7Z5R6 | AB1IP_HUMAN | cell membrane; peripheral membrane protein (by similarity). colocalizes with ena/vasp proteins at lamellipodia tips and focal adhesions, and f-actin at the leading edge. at the membrane surface, associates, via the ph domain, preferentially with the inositol phosphates, ptdins(5)p and ptdins(3)p. this binding appears to be necessary for the efficient interaction of the ra domain to ras-gtpases | cytoplasm. |
| Q7Z5U6 | WDR53_HUMAN | | nucleus. |
| Q7Z628 | ARHG8_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q7Z692 | CEA19_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q7Z698 | SPRE2_HUMAN | membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q7Z699 | SPRE1_HUMAN | membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q7Z6A9 | BTLA_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q7Z6B7 | SRGP1_HUMAN | | cytoplasm. |
| Q7Z6J2 | GRASP_HUMAN | cytoplasm; perinuclear region (by similarity). cell membrane; peripheral membrane protein; cytoplasmic side (by similarity). postsynaptic membrane (by similarity). localized at postsynaptic membranes (by similarity). | cytoplasm. |
| Q7Z6J4 | FGD2_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q7Z6J6 | FRMD5_HUMAN | membrane; single-pass membrane protein (potential). | cytoplasm; membrane. |
| Q7Z6J9 | SEN54_HUMAN | nucleus (probable). nucleus; nucleolus (probable). may be transiently localized in the nucleolus (probable). | centriole. |
| Q7Z6K5 | CO038_HUMAN | | mitochondrion. |
| Q7Z6M2 | FBX33_HUMAN | | cytoplasm. |
| Q7Z6W1 | TMCO2_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q7Z6W7 | DNJB7_HUMAN | | cytoplasm. |
| Q7Z6Z7 | HUWE1_HUMAN | nucleus (by similarity). predominantly cytosolic or perinuclear in some colorectal carcinoma cells. | cytoplasm. |
| Q7Z739 | YTHD3_HUMAN | | cytoplasm. |
| Q7Z794 | K2C1B_HUMAN | | cytoplasm. |

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| Q7Z7B0 | FLIP1_HUMAN | | cytoplasm. |
| Q7Z7B1 | PIGW_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q7Z7B7 | DB132_HUMAN | secreted protein (potential). | secreted protein. |
| Q7Z7B8 | DB128_HUMAN | secreted protein (potential). | secreted protein. |
| Q7Z7E8 | UB2Q1_HUMAN | | cytoplasm. |
| Q7Z7G0 | TARSH_HUMAN | | nucleus. |
| Q7Z7G2 | CPLX4_HUMAN | membrane; lipid-anchor. enriched at the synaptic terminal (by similarity). | endosome. |
| Q7Z7G8 | VP13B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q7Z7H5 | TMED4_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). | endoplasmic reticulum. |
| Q7Z7K2 | ZN467_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q7Z7L7 | ZYGBL_HUMAN | | centriole. cytoplasm. |
| Q7Z7L9 | ZSCA2_HUMAN | nucleus (by similarity). | nucleus. |
| Q7Z7M0 | MEGF8_HUMAN | membrane; single-pass type iii membrane protein (potential). | membrane. |
| Q7Z7M9 | GALT5_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q86SE8 | NPM2_HUMAN | nucleus (by similarity). found in the oocyte nucleus before nuclear membrane breakdown, after which it is redistributed to the cytoplasm (by similarity). | centriole. nucleus. |
| Q86SF2 | GALT7_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q86SG2 | ANR23_HUMAN | nucleus. sarcomeric i-band and some intercalated disks (by similarity). | centriole. cytoplasm. |
| Q86SG4 | DPCA2_HUMAN | | cytoplasm. |
| Q86SG5 | SA7L1_HUMAN | cytoplasm (by similarity). | cytoplasm. nucleus. |
| Q86SG6 | NEK8_HUMAN | cytoplasm (by similarity). predominantly cytoplasmic (by similarity). | cytoplasm. |
| Q86SG7 | LYG2_HUMAN | secreted protein (probable). | centriole. |
| Q86SH2 | ZAR1_HUMAN | cytoplasm (by similarity). | secreted protein. |
| Q86SJ2 | AMGO2_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). nucleus (by similarity). associated with nucleus as well as plasma membrane. restricted to somata of cerebellar as well as hippocampal neurons (by similarity). | membrane. |
| Q86SJ6 | DSG4_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q86SQ9 | DHDDS_HUMAN | | cytoplasm. |
| Q86SR1 | GLT10_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q86SS6 | SYT9_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q86SX6 | GLRX5_HUMAN | | cytoplasm. |
| Q86SZ2 | TPC6B_HUMAN | Golgi apparatus; cis-Golgi network (by similarity). | endoplasmic reticulum. |

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| Q86T13 | CLC14_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q86T20 | CF001_HUMAN | | membrane. |
| Q86T29 | ZN605_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q86T65 | DAAM2_HUMAN | | cytoplasm. |
| Q86T82 | UBP37_HUMAN | | nucleus. |
| Q86TA1 | MOL2B_HUMAN | | cytoplasm. nucleus. |
| Q86TC9 | MYPN_HUMAN | cytoplasm. bound to sarcomere both at the z- line periphery and in the central i-band region. nucleus (by similarity). | centriole. cytoplasm. |
| Q86TD4 | SRCA_HUMAN | sarcoplasmic reticulum; sarcoplasmic reticulum lumen (by similarity). associated through calcium with the membrane (by similarity). | cytoplasm. |
| Q86TH1 | ATL2_HUMAN | secreted protein (potential). | membrane. |
| Q86TI0 | TBCD1_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| Q86TL0 | ATG4D_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q86TL2 | TM110_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q86TV6 | TTC7B_HUMAN | | centriole. |
| Q86TX2 | ACOT1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q86TZ1 | TTC6_HUMAN | | cytoplasm. |
| Q86U06 | RBM23_HUMAN | nucleus (potential). | nucleus. |
| Q86U17 | SPA11_HUMAN | secreted protein (potential). | secreted protein. |
| Q86U38 | CN021_HUMAN | | cytoplasm. |
| Q86U42 | PABP2_HUMAN | nucleus (by similarity). cytoplasm (by similarity). shuttles between the nucleus and the cytoplasm but predominantly found in the nucleus. its nuclear import may involve the nucleocytoplasmic transport receptor transportin and a ran-gtp-sensitive import mechanism. is exported to the cytoplasm by a carrier-mediated pathway that is independent of mrna traffic. nucleus; nucleoplasm; nuclear speckle. colocalizes with skip and poly(a) rna in nuclear speckles (by similarity). | nucleus. |
| Q86U70 | LDB1_HUMAN | nucleus (by similarity). | centriole. nucleus. |
| Q86UD3 | MARH3_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. |
| Q86UE3 | ZN546_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q86UE4 | LYRIC_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (by similarity). nucleus; nuclear membrane; single-pass type ii membrane protein (by similarity). cell membrane; cell-cell junction; tight junction; single-pass type ii membrane protein (by similarity). in epithelial cells, recruited to tight junctions (tj) during the maturation of the tj complexes. nucleus; nucleolus (by similarity). a nucleolar staining may be due to nuclear targeting of an isoform lacking the transmembrane domain (by similarity). | endoplasmic reticulum. |
| Q86UE6 | LRTM1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |

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|--------|-------------|--|-----------------------------------|
| Q86UF2 | CTGE6_HUMAN | | cytoplasm. |
| Q86UK0 | ABCAC_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q86UK5 | LBN_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q86UL3 | PLCF_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q86UL8 | MAGI2_HUMAN | synaptosome; peripheral membrane protein (by similarity). membrane-associated in synaptosomes (by similarity). | cytoplasm. |
| Q86UN6 | AKA28_HUMAN | cytoplasm (probable). | nucleus. |
| Q86UP0 | CAD24_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q86UP6 | CUZD1_HUMAN | membrane; single-pass type i membrane protein (potential). associated with zymogen granule membranes (by | membrane. |
| Q86UP9 | LHPL3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q86UR5 | RIMS1_HUMAN | | cytoplasm. |
| Q86UU1 | PHLB1_HUMAN | | cytoplasm. |
| Q86UU5 | GGN_HUMAN | | cytoplasm. cytoskeleton. nucleus. |
| Q86UV6 | TRI74_HUMAN | | cytoplasm. |
| Q86UV7 | TRI73_HUMAN | | cytoplasm. |
| Q86UW6 | N4BP2_HUMAN | cytoplasm (by similarity). | nucleus. |
| Q86UW8 | HPLN4_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q86UX6 | ST32C_HUMAN | | cytoplasm. |
| Q86UX7 | URP2_HUMAN | cytoplasm (probable). membrane-associated in chronic lymphocytic | cytoplasm; membrane. |
| Q86UZ6 | BTBD4_HUMAN | nucleus (potential). | nucleus. |
| Q86V15 | CASZ1_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q86V20 | FA35A_HUMAN | | nucleus. |
| Q86V25 | VASHL_HUMAN | | cytoplasm. |
| Q86V35 | CABP7_HUMAN | | cytoplasm. |
| Q86V48 | LUZP1_HUMAN | nucleus. also detected in soma and dendrites of neurons (by similarity). | centriole. cytoplasm. |
| Q86V71 | ZN429_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q86V85 | GP180_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q86V86 | PIM3_HUMAN | | cytoplasm. |
| Q86V88 | MGDP1_HUMAN | | cytoplasm. |
| Q86V97 | KBTB6_HUMAN | | cytoplasm. |
| Q86VD1 | MORC1_HUMAN | nucleus (by similarity). | nucleus. |
| Q86VF5 | MOGT3_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q86VH4 | LRTM4_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q86VH5 | LRTM3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q86VI3 | IQGA3_HUMAN | | cytoplasm. |
| Q86VK4 | ZN410_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q86VN1 | VPS36_HUMAN | cytoplasm. endosome; late endosome. colocalizes with ubiquitinated proteins on late endosomes. nucleus (probable). | cytoplasm. nucleus. |

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| Q86VP1 | TAXB1_HUMAN | | cytoplasm. |
| Q86VP6 | CAND1_HUMAN | nucleus (by similarity). | nucleus. |
| Q86VQ0 | CF152_HUMAN | | centriole. |
| Q86VU5 | CMTD1_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q86VX2 | COMD7_HUMAN | | cytoplasm. |
| Q86VZ4 | LRP11_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q86VZ6 | JAZF1_HUMAN | nucleus (potential). | nucleus. |
| Q86W24 | NAL14_HUMAN | | cytoplasm. |
| Q86W25 | NAL13_HUMAN | | cytoplasm. |
| Q86W26 | NAL10_HUMAN | | cytoplasm. |
| Q86W34 | AMZ2_HUMAN | | secreted protein. |
| Q86W42 | THOC6_HUMAN | | cytoplasm. |
| Q86W74 | ANR46_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q86W92 | LIPB1_HUMAN | | membrane. |
| Q86WA6 | BPHL_HUMAN | | mitochondrion. |
| Q86WC4 | OSTM1_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| Q86WD7 | SPA9_HUMAN | secreted protein (potential). | secreted protein. |
| Q86WG3 | ATCAY_HUMAN | | cytoplasm. |
| Q86WI0 | LHPL1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q86WK6 | AMGO1_HUMAN | membrane; single-pass type i membrane protein (potential). associated with axons of neuronal cells (by similarity). | membrane. |
| Q86WK7 | AMGO3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q86WR0 | CCD25_HUMAN | | nucleus. |
| Q86WR7 | CJ047_HUMAN | | nucleus. |
| Q86WZ6 | ZN227_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q86X10 | K1219_HUMAN | | cytoplasm. |
| Q86X29 | LSR_HUMAN | cell membrane; single-pass membrane protein (potential). | membrane. |
| Q86X45 | LRRC6_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q86X52 | CHSS1_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q86X53 | ERIC1_HUMAN | | nucleus. |
| Q86X67 | NUD13_HUMAN | | cytoplasm. |
| Q86X76 | NIT1_HUMAN | | secreted protein. |
| Q86X83 | COMD2_HUMAN | | cytoplasm. |
| Q86XF0 | DYRL1_HUMAN | | microsome. |
| Q86XF7 | ZN575_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q86XJ0 | FA26A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q86XJ1 | GA2L3_HUMAN | | nucleus. |
| Q86XK2 | FBX11_HUMAN | | cytoplasm. |
| Q86XK3 | CJ078_HUMAN | | nucleus. |
| Q86XL7 | ZN539_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q86XN8 | RKHD1_HUMAN | | nucleus. |
| Q86XP0 | PA24D_HUMAN | cytoplasm; cytosol. translocates to membrane vesicles in a calcium-dependent fashion. translocates to perinuclear regions upon ionomycin | secreted protein. |

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| Q86XR5 | PRIMA_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| Q86XR8 | CEP57_HUMAN | nucleus (by similarity). cytoplasm. associates with microtubules and the centrosome. | centriole. |
| Q86XS5 | ANGL5_HUMAN | secreted protein (potential). | secreted protein. |
| Q86XS8 | GOLI_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q86XT4 | TRI50_HUMAN | | cytoplasm. |
| Q86XU0 | ZN677_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q86XW9 | TXND6_HUMAN | cytoplasm (by similarity). cytoskeleton (by similarity). associated with microtubules. detected in cilia of lung epithelium, and associated with the spermatid tail and manchette (by similarity). | cytoplasm. |
| Q86XX4 | FRAS1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q86Y07 | VRK2_HUMAN | membrane; single-pass type iv membrane protein (potential). | cytoplasm. |
| Q86Y22 | CONA1_HUMAN | membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. secreted protein. |
| Q86Y29 | BAGE3_HUMAN | | membrane. |
| Q86Y30 | BAGE2_HUMAN | | membrane. |
| Q86Y37 | CJ046_HUMAN | | cytoplasm. |
| Q86Y38 | XYLT1_HUMAN | or: endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (by similarity). or: Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). secreted protein (probable). some fraction is also found in the extracellular space together with chondroitin sulfate proteoglycans, suggesting that it can be secreted. | endoplasmic reticulum. |
| Q86Y39 | NDUAB_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein; matrix side (by similarity). | mitochondrion. |
| Q86Y56 | HEAT2_HUMAN | | cytoplasm. |
| Q86Y78 | LYPD6_HUMAN | secreted protein (probable). | secreted protein. |
| Q86Y79 | PTH_HUMAN | | cytoplasm. |
| Q86Y82 | STX12_HUMAN | membrane; single-pass membrane protein (potential). | Golgi. |
| Q86YC3 | LRC33_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q86YD3 | TMM25_HUMAN | isoform 1, isoform 4: cell membrane; single-pass type i membrane protein (potential). isoform 2, isoform 3: secreted protein (potential). | endosome; membrane. |
| Q86YH2 | SUHW2_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q86YH6 | DLP1_HUMAN | | cytoplasm. |
| Q86YI8 | PHF13_HUMAN | | nucleus. |
| Q86YJ7 | AN13B_HUMAN | | membrane. |

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| Q86YL7 | PDPN_HUMAN | membrane; single-pass type i membrane protein. localized to actin-rich microvilli and plasma membrane projections such as filopodia, lamellipodia and ruffles (by similarity). | membrane. |
| Q86YM7 | HOME1_HUMAN | cytoplasm (by similarity). postsynaptic density of neuronal cells. isoform 1 inhibits surface expression of grm5 causing it to be retained in the endoplasmic reticulum (by similarity). | cytoplasm. |
| Q86YN1 | DOPP1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q86YQ8 | CPNE8_HUMAN | | membrane. |
| Q86YS6 | RAB43_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| Q86YS7 | K0528_HUMAN | | membrane. |
| Q86YT6 | MIB1_HUMAN | cytoplasm. cell membrane (by similarity). localizes to the plasma membrane (by similarity). according to ref.7, it is mitochondrial, however such localization remains unclear. | cytoplasm. |
| Q86YV9 | HPS6_HUMAN | | cytoplasm. |
| Q86YZ3 | HORN_HUMAN | found in keratohyalin granules in the granular cells of the epidermis (by similarity). | cytoplasm. |
| Q86Z02 | HIPK1_HUMAN | nucleus (by similarity). predominantly nuclear. cytoplasm (by similarity). | nucleus. |
| Q86Z14 | KLOTB_HUMAN | membrane; single-pass type iii membrane protein (potential). | cytoplasm. endosome; membrane. |
| Q8IU57 | I28RA_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q8IU85 | KCC1D_HUMAN | cytoplasm (probable). predominantly cytoplasmic (probable). nucleus (probable). also nuclear upon activation | cytoplasm. |
| Q8IU89 | LASS3_HUMAN | nucleus; nuclear membrane; multi-pass membrane protein (potential). | nucleus. |
| Q8IU99 | FA26C_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8IUA0 | WFDC8_HUMAN | secreted protein (potential). | secreted protein. |
| Q8IUB2 | WFDC3_HUMAN | secreted protein (potential). | secreted protein. |
| Q8IUB3 | WF10B_HUMAN | secreted protein (potential). | membrane. |
| Q8IUB5 | WFD13_HUMAN | secreted protein (potential). | secreted protein. |
| Q8IUB9 | KR191_HUMAN | | cytoplasm. |
| Q8IUC0 | KR131_HUMAN | | cytoplasm. |
| Q8IUC1 | KR111_HUMAN | | cytoplasm. |
| Q8IUC2 | KRA81_HUMAN | | cytoplasm. |
| Q8IUC3 | KRA71_HUMAN | | cytoplasm. |
| Q8IUC8 | GLT13_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |

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|--------|-------------|---|---|
| Q8IUD2 | RB6I2_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein (by similarity). in neurons, localized closed to presynaptic membrane. recruited on Golgi membranes by rab6a in a gtp- | cytoplasm. |
| Q8IUE0 | TF2LY_HUMAN | nucleus (by similarity). | nucleus. |
| Q8IUE1 | TF2LX_HUMAN | nucleus (by similarity). | nucleus. |
| Q8IUG1 | KRA13_HUMAN | | cytoplasm. |
| Q8IUH4 | ZDH13_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8IUH5 | ZDH17_HUMAN | Golgi apparatus; Golgi membrane; multi-pass membrane protein (potential). cytoplasmic vesicle; cytoplasmic vesicle membrane; multi-pass membrane protein (potential). | cytoplasm. endoplasmic reticulum; membrane. |
| Q8IUK5 | PXDC1_HUMAN | isoform 1: membrane; single-pass type i membrane protein (potential). isoform 2, isoform 3: secreted protein (potential). isoform 4: cytoplasm (potential). | membrane. |
| Q8IUK8 | CBLN2_HUMAN | secreted protein (potential). | membrane. |
| Q8IUL8 | CILP2_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q8IUN9 | CLC10_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q8IUR0 | TPPC5_HUMAN | Golgi apparatus; cis-Golgi network (by similarity). | endoplasmic reticulum. |
| Q8IUX4 | ABC3F_HUMAN | | cytoplasm. |
| Q8IUZ0 | LRC49_HUMAN | | cytoplasm. |
| Q8IV01 | SYT12_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q8IV03 | CI150_HUMAN | | cytoplasm. |
| Q8IV20 | CM031_HUMAN | | membrane. |
| Q8IV32 | CCD71_HUMAN | | nucleus. |
| Q8IV36 | CQ028_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8IV38 | ANKY2_HUMAN | | cytoplasm. nucleus. |
| Q8IV42 | PSTK_HUMAN | | cytoplasm. |
| Q8IV48 | THEX1_HUMAN | | nucleus. |
| Q8IV50 | LYSM2_HUMAN | | centriole. |
| Q8IV61 | GRP3_HUMAN | | cytoplasm. |
| Q8IV63 | VRK3_HUMAN | | cytoplasm. |
| Q8IV90 | WASF4_HUMAN | | cytoplasm. |
| Q8IVA1 | PCP2_HUMAN | | cytoplasm. |
| Q8IVB4 | SL9A9_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q8IVB5 | LIX1L_HUMAN | | cytoplasm. |
| Q8IVC4 | ZN584_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8IVD9 | NUDC3_HUMAN | | cytoplasm. |
| Q8IVF6 | ANR18_HUMAN | | nucleus. |
| Q8IVH2 | FOXP4_HUMAN | nucleus (potential). | nucleus. |
| Q8IVH4 | MMAA_HUMAN | mitochondrion (probable). | cytoplasm. mitochondrion. |
| Q8IVH8 | M4K3_HUMAN | | cytoplasm. |
| Q8IVL6 | P3H3_HUMAN | endoplasmic reticulum (by similarity). | endoplasmic reticulum. |

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| Q8IVM0 | CCD50_HUMAN | | nucleus. |
| Q8IVM8 | S22A9_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8IVP9 | ZN547_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8IVQ6 | ZDH21_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q8IVT2 | CS021_HUMAN | | cytoplasm. |
| Q8IVT5 | KSR1_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). in unstimulated cells, where the phosphorylated form is bound to a 14-3-3 protein, sequestration in the cytoplasm occurs. following growth factor treatment, the protein is free for membrane translocation, and it moves from the cytoplasm to the cell periphery (by similarity). | cytoplasm. |
| Q8IVU9 | CJ107_HUMAN | | cytoplasm. |
| Q8IVV7 | CQ039_HUMAN | | cytoplasm. |
| Q8IVV8 | CT058_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8IVW1 | ARL14_HUMAN | | Golgi. |
| Q8IVW4 | CDKL3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8IW19 | CB013_HUMAN | | nucleus. |
| Q8IW36 | ZN695_HUMAN | nucleus (potential). | nucleus. |
| Q8IW41 | MAPK5_HUMAN | cytoplasm (probable). nucleus. also observed in the nucleus. | cytoplasm. |
| Q8IW50 | CI025_HUMAN | | nucleus. |
| Q8IW52 | SLIK4_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q8IW75 | SPA12_HUMAN | secreted protein (potential). | secreted protein. |
| Q8IWA4 | MFN1_HUMAN | mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (probable). isoform 2: cytoplasm. | endosome. mitochondrion. |
| Q8IWA5 | CTL2_HUMAN | membrane; multi-pass membrane protein (by similarity). | Golgi. |
| Q8IWA6 | CCD60_HUMAN | | nucleus. |
| Q8IWB6 | TEX14_HUMAN | cytoplasm. detected in the intercellular bridges that connect male germ cell daughter cells after cell division (by similarity). | cytoplasm. |
| Q8IWB9 | TEX2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8IWG1 | WDR63_HUMAN | | nucleus. |
| Q8IWL3 | HSC20_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q8IWL8 | STH_HUMAN | | membrane. |
| Q8IWN6 | CX052_HUMAN | | cytoplasm. |
| Q8IWN7 | RP1L1_HUMAN | | cytoplasm. |
| Q8IWP9 | CC28A_HUMAN | | cytoplasm. |
| Q8IWQ3 | BRSK2_HUMAN | | cytoplasm. |
| Q8IWR0 | Z3H7A_HUMAN | | nucleus. |
| Q8IWR1 | TRI59_HUMAN | membrane; single-pass membrane protein (potential). | cytoplasm. |
| Q8IWT1 | SCN4B_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |

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| Q8IWT6 | LRC8A_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8IWU5 | SULF2_HUMAN | endoplasmic reticulum (by similarity). Golgi apparatus; Golgi stack (by similarity). cell surface (by similarity). also localized on the cell surface (by similarity). | endoplasmic reticulum. lysosome. |
| Q8IWU6 | SULF1_HUMAN | endoplasmic reticulum (by similarity). Golgi apparatus; Golgi stack (by similarity). cell surface (by similarity). also localized on the cell surface (by similarity). | endoplasmic reticulum. lysosome. |
| Q8IWU9 | TPH2_HUMAN | | endoplasmic reticulum. |
| Q8I WV2 | CNTN4_HUMAN | cell membrane; lipid-anchor; gpi-anchor. secreted protein (by similarity). | membrane. |
| Q8I WV8 | UBR2_HUMAN | nucleus (by similarity). | cytoplasm. |
| Q8I WW6 | RHG12_HUMAN | | cytoplasm. |
| Q8I WX7 | UN45B_HUMAN | | cytoplasm. secreted protein. |
| Q8I WY8 | ZN690_HUMAN | nucleus (probable). | nucleus. |
| Q8I WY9 | CDAN1_HUMAN | | cytoplasm. |
| Q8I WZ4 | TRI48_HUMAN | | cytoplasm. |
| Q8I WZ5 | TRI42_HUMAN | | cytoplasm. |
| Q8I WZ6 | BBS7_HUMAN | | centriole. |
| Q8I WZ8 | SF04_HUMAN | nucleus (probable). | nucleus. |
| Q8I X01 | SFR14_HUMAN | nucleus (potential). | nucleus. |
| Q8I X06 | GOR_HUMAN | | nucleus. |
| Q8I X07 | FOG1_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q8I X21 | CJ006_HUMAN | | nucleus. |
| Q8I X29 | FBX16_HUMAN | | cytoplasm. |
| Q8I X90 | CM003_HUMAN | | nucleus. |
| Q8I X94 | CTGE4_HUMAN | | cytoplasm. |
| Q8I X95 | CTGE3_HUMAN | | cytoplasm. |
| Q8I XF9 | AQP12_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8I XH7 | NELFD_HUMAN | nucleus (probable). | nucleus. |
| Q8I XH8 | CAD26_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q8I XI1 | MIRO2_HUMAN | mitochondrion; mitochondrial outer membrane; single-pass type iv membrane protein (probable). | cytoplasm. endosome. mitochondrion. |
| Q8I XI2 | MIRO1_HUMAN | mitochondrion; mitochondrial outer membrane; single-pass type iv membrane protein (probable). | cytoplasm. endosome. mitochondrion. |
| Q8I XJ9 | ASXL1_HUMAN | nucleus (probable). | nucleus. |
| Q8I XK0 | PHC2_HUMAN | nucleus (probable). | nucleus. |
| Q8I XK2 | GLT12_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q8I XL6 | FA20C_HUMAN | secreted protein (potential). | cytoplasm. |
| Q8I XQ3 | CI040_HUMAN | | nucleus. |
| Q8I XQ5 | KLHL7_HUMAN | | cytoplasm. |
| Q8I XS0 | CF146_HUMAN | | nucleus. |
| Q8I XT2 | DMRTD_HUMAN | nucleus (potential). | nucleus. |
| Q8I XV7 | KLD8B_HUMAN | | membrane. |
| Q8I XX5 | CA037_HUMAN | | endosome; membrane. |
| Q8I XZ2 | ZC3H3_HUMAN | | cytoplasm. nucleus. |
| Q8I XZ3 | SP8_HUMAN | nucleus (potential). | cytoplasm. nucleus. |

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| Q8IY26 | PPAC2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. |
| Q8IY31 | IFT20_HUMAN | Golgi apparatus; cis-Golgi network. centrosome. present at the centrosomes during the cell cycle and associated with the proximal portion of the mother centriole and the lateral aspect of the daughter centriole. associated with basal body at the base of primary cilia (by cytoplasm (by similarity)). | cytoskeleton. |
| Q8IY33 | MILK2_HUMAN | | cytoplasm. nucleus. |
| Q8IY37 | DHX37_HUMAN | | nucleus. |
| Q8IY47 | KBTB2_HUMAN | | cytoplasm. |
| Q8IY49 | PAQRA_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8IY63 | AMOL1_HUMAN | | membrane. |
| Q8IY67 | RAVR1_HUMAN | nucleus (by similarity). cytoplasm (by similarity). nuclear, in perinucleolar structures. shuttles between nucleus and cytoplasm. cytoplasm, at focal contacts and cell-cell contacts. associated with myotubes during muscle differentiation (by | nucleus. |
| Q8IY84 | NIM1_HUMAN | | cytoplasm. |
| Q8IY92 | BTBDC_HUMAN | | cytoplasm. |
| Q8IYB1 | CC059_HUMAN | | cytoplasm. |
| Q8IYB4 | PEX5R_HUMAN | cytoplasm. some fraction is membrane associated via its interaction with rab8b (by similarity). | cytoplasm. |
| Q8IYB5 | SMAP1_HUMAN | cell membrane; peripheral membrane protein; cytoplasmic side (by similarity). | cytoplasm. |
| Q8IYB9 | ZN595_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8IYD2 | KLD8A_HUMAN | | mitochondrion. |
| Q8IYD9 | CR054_HUMAN | secreted protein (potential). | nucleus. |
| Q8IYE1 | CCD13_HUMAN | | cytoplasm. |
| Q8IYF1 | ELOA2_HUMAN | nucleus (probable). | nucleus. |
| Q8IYG6 | LRC56_HUMAN | | cytoplasm. |
| Q8IYI6 | EXOC8_HUMAN | cytoplasm. redistributes to growing neurites and growth cones during cell differentiation (by similarity). binds lipids with phosphatidylinositol-3,4,5-trisphosphate groups. | cytoplasm. |
| Q8IYI8 | ZN440_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8IYJ2 | CJ067_HUMAN | | nucleus. |
| Q8IYK8 | REM2_HUMAN | cell membrane (by similarity). | cytoplasm. |
| Q8IYL2 | U383_HUMAN | | cytoplasm. |
| Q8IYL9 | PSYR_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q8IYM9 | TRI22_HUMAN | | nucleus. |
| Q8IYN0 | ZN100_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8IYN2 | TCAL8_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8IYN6 | F100B_HUMAN | | nucleus. |
| Q8IYP9 | ZDH23_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q8IYQ7 | THNSL_HUMAN | | mitochondrion. |
| Q8IYR0 | CF165_HUMAN | | cytoplasm. |
| Q8IYR2 | SMYD4_HUMAN | | nucleus. |
| Q8IYS8 | FA44C_HUMAN | | cytoplasm. |

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| Q8IYT8 | ULK2_HUMAN | | cytoplasm. |
| Q8IYV9 | IZUM1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8IYW2 | CJ092_HUMAN | | cytoplasm. |
| Q8IYW5 | RN168_HUMAN | | cytoplasm. |
| Q8IYX0 | ZN679_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8IYX1 | TBC21_HUMAN | | cytoplasm. |
| Q8IYX4 | DND1_HUMAN | | nucleus. |
| Q8IYX7 | CI138_HUMAN | | nucleus. |
| Q8IYX8 | CF182_HUMAN | | centriole. cytoplasm. |
| Q8IZ02 | LRC34_HUMAN | | cytoplasm. |
| Q8IZ07 | AN13A_HUMAN | | cytoplasm. |
| Q8IZ20 | ZN683_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8IZ26 | ZNF34_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8IZ40 | RCOR2_HUMAN | nucleus (by similarity). | nucleus. |
| Q8IZ52 | CHSS2_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q8IZ81 | ELMD2_HUMAN | | cytoplasm. |
| Q8IZC7 | ZN101_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8IZD4 | DCP1B_HUMAN | cytoplasm. in discrete foci (by similarity). | cytoplasm. |
| Q8IZD9 | DOCK3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8IZF2 | GP116_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8IZJ1 | UNC5B_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q8IZJ4 | RGDSR_HUMAN | cytoplasmic vesicle (by similarity). | centriole. cytoplasm. |
| Q8IZK6 | MCLN2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8IZN3 | ZDH14_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q8IZP0 | ABI1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). localized to protruding lamellipodia and filopodia tips. also localized to neuronal growth cones and synaptosomes (by similarity). | cytoplasm. nucleus. |
| Q8IZP1 | TBCD3_HUMAN | | cytoplasm. |
| Q8IZP2 | F10A4_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8IZP6 | R113B_HUMAN | | cytoplasm. nucleus. |
| Q8IZP7 | H6ST3_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q8IZQ5 | SELH_HUMAN | | cytoplasm. |
| Q8IZQ8 | MYCD_HUMAN | nucleus (by similarity). | nucleus. |
| Q8IZT6 | ASPM_HUMAN | cytoplasm (by similarity). nucleus (by similarity). the nuclear-cytoplasmic distribution could be regulated by the availability of calmodulin (by similarity). | nucleus. |
| Q8IZT8 | OST5_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q8IZU2 | WDR17_HUMAN | | cytoplasm. |
| Q8IZU3 | SYCP3_HUMAN | nucleus (by similarity). in tripartite segments of synaptonemal complexes, irrespective of whether these are synapsed or unsynapsed (by similarity). | cytoplasm. nucleus. |
| Q8IZU9 | KIRR3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |

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| Q8IZX4 | TAF1L_HUMAN | nucleus (by similarity). | nucleus. |
| Q8J025 | APCD1_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q8N0U8 | VKORL_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N0V1 | CU121_HUMAN | | nucleus. |
| Q8N0V3 | RBFAL_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q8N0V4 | LGI2_HUMAN | secreted protein (potential). | cytoplasm. |
| Q8N0W3 | FUK_HUMAN | | cytoplasm. |
| Q8N0W4 | NLGNX_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8N0X2 | SPG16_HUMAN | cytoplasm (by similarity). flagellum (by similarity). detected on the sperm flagellum. detected in the central apparatus of the axoneme. colocalizes with spag6 on microtubules (by similarity). | centriole. |
| Q8N0X7 | SPG20_HUMAN | | cytoplasm. |
| Q8N0Y2 | ZN444_HUMAN | nucleus (potential). | nucleus. |
| Q8N0Y7 | PGAM4_HUMAN | | cytoplasm. lysosome. |
| Q8N0Z2 | ABRA_HUMAN | localized to the i-band of the sarcomere and to a lesser extent to the sarcomeric structure between z-lines (by similarity). | cytoplasm. |
| Q8N0Z6 | TTC5_HUMAN | | cytoplasm. |
| Q8N0Z8 | PUSL1_HUMAN | | nucleus. |
| Q8N103 | TAGAP_HUMAN | | cytoplasm. |
| Q8N111 | CEND_HUMAN | membrane; single-pass type iv membrane protein (potential). | endosome; membrane. |
| Q8N112 | CC045_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q8N118 | CP4X1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; peripheral membrane protein (by similarity). | endoplasmic reticulum. |
| Q8N119 | MMP21_HUMAN | secreted protein (potential). | secreted protein. |
| Q8N122 | RPTOR_HUMAN | | cytoplasm. |
| Q8N126 | IGS4B_HUMAN | cell membrane; cell-cell junction; single-pass type i membrane protein (by similarity). localized at the cell-cell attached sites of the plasma membrane | membrane. |
| Q8N128 | CN024_HUMAN | | centriole. |
| Q8N131 | PORIM_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8N135 | LGI4_HUMAN | secreted protein (potential). | cytoplasm. |
| Q8N136 | WDR69_HUMAN | | nucleus. |
| Q8N138 | ORML3_HUMAN | membrane; multi-pass membrane protein (potential). endoplasmic reticulum. | endoplasmic reticulum. |
| Q8N141 | ZN545_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N142 | PURA1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8N143 | BCL6B_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q8N145 | LGI3_HUMAN | secreted protein (potential). | cytoplasm. |
| Q8N157 | AHI1_HUMAN | | cytoplasm. |
| Q8N158 | GPC2_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | endoplasmic reticulum. |
| Q8N163 | K1967_HUMAN | | cytoplasm. nucleus. |
| Q8N184 | ZN567_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N187 | AL2S8_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q8N1D0 | BWR1B_HUMAN | | cytoplasm. |

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| Q8N1D5 | CA158_HUMAN | | nucleus. |
| Q8N1E2 | LYG1_HUMAN | secreted protein (probable). | centriole. |
| Q8N1E6 | FXL14_HUMAN | | cytoplasm. |
| Q8N1G0 | ZN687_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N1G4 | LRC47_HUMAN | | cytoplasm. |
| Q8N1H7 | S6OS1_HUMAN | | cytoplasm. |
| Q8N1N0 | CLC4F_HUMAN | membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q8N1N2 | CR026_HUMAN | | endosome; membrane. |
| Q8N1Q1 | CAH13_HUMAN | | cytoplasm. |
| Q8N1W2 | ZN710_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N201 | INT1_HUMAN | nucleus; nuclear membrane; single-pass membrane protein (potential). | endosome. |
| Q8N283 | ANR35_HUMAN | | centriole. |
| Q8N290 | CC053_HUMAN | | nucleus. |
| Q8N2C7 | CB021_HUMAN | | cytoplasm. |
| Q8N2C9 | CU128_HUMAN | | membrane. |
| Q8N2G4 | LYPD1_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | secreted protein. |
| Q8N2G8 | LGP1_HUMAN | endoplasmic reticulum. nucleus; nuclear envelope (by similarity). | centriole. cytoplasm. |
| Q8N2H3 | CJ033_HUMAN | | mitochondrion. |
| Q8N2H9 | PELI3_HUMAN | | cytoplasm. |
| Q8N2I2 | ZN619_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N2K0 | ABD12_HUMAN | | endoplasmic reticulum. |
| Q8N2K1 | UB2J2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type iv membrane protein (by similarity). | endoplasmic reticulum. |
| Q8N2M4 | TM86A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N2M8 | SFR16_HUMAN | nucleus (by similarity). | nucleus. |
| Q8N2N9 | AN36B_HUMAN | | cytoplasm. |
| Q8N2Q7 | NLGN1_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). enriched in synaptic plasma membranes and clustered in synaptic clefts and postsynaptic densities (by similarity). | membrane. |
| Q8N2R0 | OSR2_HUMAN | nucleus (potential). | nucleus. |
| Q8N2R8 | FA43A_HUMAN | | cytoplasm. |
| Q8N2W9 | PIAS4_HUMAN | nucleus; nucleoplasm; nuclear speckle (by similarity). colocalizes with promyelocytic leukemia nuclear bodies (pml nbs) (by similarity). | nucleus. |
| Q8N2Y8 | RUSC2_HUMAN | | nucleus. |
| Q8N2Z3 | NUD9P_HUMAN | | nucleus. |
| Q8N300 | CCD23_HUMAN | | nucleus. |
| Q8N326 | CJ111_HUMAN | | endosome; membrane. |
| Q8N336 | ELMD1_HUMAN | | cytoplasm. |
| Q8N339 | MT1M_HUMAN | | cytoplasm. nucleus. |
| Q8N350 | DOS_HUMAN | | endosome; membrane. |
| Q8N357 | CB018_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N386 | LRC25_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |

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| Q8N387 | MUC15_HUMAN | isoform 1: cell membrane; single-pass type i membrane protein (potential). isoform 2: secreted protein (potential). | endosome; membrane. |
| Q8N392 | RHG18_HUMAN | | cytoplasm. |
| Q8N3C7 | RSNL2_HUMAN | | nucleus. |
| Q8N3F8 | MILK1_HUMAN | cytoplasm (by similarity). | cytoplasm. nucleus. |
| Q8N3I7 | BBS5_HUMAN | localizes to basal bodies (by similarity). | centriole. |
| Q8N3L3 | TXLNB_HUMAN | | cytoplasm. |
| Q8N3R3 | CC023_HUMAN | | mitochondrion. |
| Q8N3T1 | GLTL2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q8N3V7 | SYNPO_HUMAN | cytoplasm (by similarity). cell membrane; cell-cell junction; tight junction (by similarity). localized at the tight junction of cells. cytoplasm; perikarya. in brain, localized to the postsynaptic densities and in the perikarya. associated with dendritic spines of a subset of synapses (by similarity). | cytoplasm. |
| Q8N3Y1 | FBXW8_HUMAN | | cytoplasm. |
| Q8N3Y3 | LARG2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | Golgi. |
| Q8N3Z6 | ZCHC7_HUMAN | | cytoplasm. nucleus. |
| Q8N428 | GLTL1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q8N436 | CPXM2_HUMAN | secreted protein (probable). | secreted protein. |
| Q8N448 | LNK2_HUMAN | | cytoplasm. |
| Q8N456 | LRC18_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8N461 | FXL16_HUMAN | | cytoplasm. |
| Q8N465 | D2HDH_HUMAN | mitochondrion (probable). | mitochondrion. |
| Q8N475 | FSTL5_HUMAN | secreted protein (probable). | cytoplasm. |
| Q8N485 | LIX1_HUMAN | | cytoplasm. |
| Q8N4A0 | GALT4_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | Golgi. |
| Q8N4B4 | FBX39_HUMAN | | cytoplasm. |
| Q8N4C8 | MINK1_HUMAN | | cytoplasm. |
| Q8N4E4 | PDCL2_HUMAN | | cytoplasm. |
| Q8N4F0 | BPIL1_HUMAN | secreted protein (by similarity). | cytoplasm. |
| Q8N4F7 | RN175_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. |
| Q8N4H0 | CI068_HUMAN | | nucleus. |
| Q8N4H5 | CI105_HUMAN | | endoplasmic reticulum. |
| Q8N4J0 | CI041_HUMAN | | cytoplasm. |
| Q8N4L8 | CCD24_HUMAN | | cytoplasm. |
| Q8N4M1 | CTL3_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q8N4Q0 | ZADH2_HUMAN | | cytoplasm. |
| Q8N4S0 | CCD82_HUMAN | | nucleus. |
| Q8N4S7 | PAQR4_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8N4T0 | CBPA6_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q8N4X5 | K1914_HUMAN | | nucleus. |

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| Q8N511 | CQ032_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N531 | FBXL6_HUMAN | | cytoplasm. |
| Q8N554 | ZF276_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N567 | ZCHC9_HUMAN | | cytoplasm. nucleus. |
| Q8N568 | DCAK2_HUMAN | | cytoplasm. |
| Q8N587 | ZN561_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N5A5 | ZGPAT_HUMAN | | nucleus. |
| Q8N5B7 | LASS5_HUMAN | nucleus; nuclear membrane; multi-pass membrane protein (potential). endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | nucleus. |
| Q8N5C1 | CF188_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N5C8 | TAB3_HUMAN | | cytoplasm. |
| Q8N5D0 | WDTC1_HUMAN | | cytoplasm. |
| Q8N5D6 | GBGT1_HUMAN | membrane; single-pass type ii membrane protein (potential). | Golgi. |
| Q8N5G2 | MACOI_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N5H7 | SH2D3_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). associated with the membrane when egf-stimulated. found at ruffling membranes (by similarity). | cytoplasm. |
| Q8N5I2 | ARRD1_HUMAN | | nucleus. |
| Q8N5I3 | KCNRG_HUMAN | | nucleus. |
| Q8N5I4 | DHRXS_HUMAN | | cytoplasm. mitochondrion. |
| Q8N5J4 | SPIC_HUMAN | nucleus (probable). | nucleus. |
| Q8N5M1 | ATPF2_HUMAN | mitochondrion (probable). | mitochondrion. |
| Q8N5N4 | CC022_HUMAN | | nucleus. |
| Q8N5P1 | ZC3H8_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q8N5S9 | KKCC1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q8N5T2 | TBC19_HUMAN | | cytoplasm. |
| Q8N5U0 | CK042_HUMAN | | nucleus. |
| Q8N5V2 | NGEF_HUMAN | cytoplasm. associated with membranes. localizes to axonal growth cones (by similarity). | cytoplasm. |
| Q8N5W8 | FA24B_HUMAN | | membrane. |
| Q8N5Y2 | MS3L1_HUMAN | nucleus (probable). | nucleus. |
| Q8N5Z0 | AADAT_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q8N5Z5 | KCD17_HUMAN | | nucleus. |
| Q8N653 | LZTR1_HUMAN | | cytoplasm. |
| Q8N655 | CJ012_HUMAN | | nucleus. |
| Q8N661 | TM86B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N668 | COMD1_HUMAN | | cytoplasm. |
| Q8N680 | ZBTB2_HUMAN | nucleus (potential). | nucleus. |
| Q8N684 | CPSF7_HUMAN | nucleus (probable). | nucleus. |
| Q8N687 | DB125_HUMAN | secreted protein (potential). | secreted protein. |
| Q8N688 | DB123_HUMAN | secreted protein (potential). | secreted protein. |
| Q8N689 | DB120_HUMAN | secreted protein (potential). | secreted protein. |
| Q8N690 | DB119_HUMAN | secreted protein (potential). | secreted protein. |
| Q8N693 | ESX1L_HUMAN | nucleus (potential). | nucleus. |

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| Q8N6C7 | PGSF1_HUMAN | | nucleus. |
| Q8N6C8 | LIRA3_HUMAN | secreted protein (probable). | membrane. |
| Q8N6D5 | ANR29_HUMAN | | cytoplasm. |
| Q8N6F8 | WBS27_HUMAN | | cytoplasm. |
| Q8N6G5 | CGAT2_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q8N6I4 | CN109_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N6L7 | CI071_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N6M0 | OTU6B_HUMAN | | cytoskeleton. |
| Q8N6M3 | CT142_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N6M5 | ALLC_HUMAN | | cytoplasm. |
| Q8N6M6 | AMPO_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q8N6N7 | ACBD7_HUMAN | | centriole. |
| Q8N6R0 | K0859_HUMAN | | cytoplasm. |
| Q8N6S4 | AN13C_HUMAN | | endoplasmic reticulum. |
| Q8N6T3 | ARFG1_HUMAN | cytoplasm (by similarity). Golgi apparatus (by similarity). associates with the Golgi complex (by similarity). | cytoplasm. |
| Q8N6T7 | SIRT6_HUMAN | nucleus (by similarity). predominantly nuclear. | nucleus. |
| Q8N6V4 | CJ053_HUMAN | | cytoplasm. |
| Q8N6V9 | TEX9_HUMAN | | nucleus. |
| Q8N6Y1 | PCD20_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q8N6Y2 | LRC17_HUMAN | | cytoplasm. |
| Q8N715 | CA065_HUMAN | | centriole. |
| Q8N720 | ZN655_HUMAN | nucleus (probable). | nucleus. |
| Q8N726 | CD2A2_HUMAN | nucleus; nucleolus (by similarity). | nucleus. |
| Q8N752 | KC1AL_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8N755 | PQLC3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N766 | K0090_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q8N782 | ZN525_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N787 | CR030_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N7A1 | KLDC1_HUMAN | | secreted protein. |
| Q8N7C0 | LRC52_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q8N7C4 | CF128_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N7E2 | ZN645_HUMAN | | cytoplasm. nucleus. |
| Q8N7G0 | SPRM1_HUMAN | nucleus (by similarity). | nucleus. |
| Q8N7K0 | ZN433_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N7M2 | ZN283_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N7P3 | CLD22_HUMAN | cell membrane; cell-cell junction; tight junction; multi-pass membrane protein (by similarity). | membrane. |
| Q8N7Q2 | CJ031_HUMAN | secreted protein (potential). | membrane. |
| Q8N7Q3 | ZN676_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N7S2 | DNJ5G_HUMAN | membrane; lipid-anchor (by similarity). | endoplasmic reticulum. |
| Q8N7W2 | CJ030_HUMAN | | nucleus. |

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| Q8N7X0 | CF103_HUMAN | | cytoplasm. |
| Q8N7X4 | MAGB6_HUMAN | | centriole. cytoplasm. |
| Q8N7Y1 | PRR10_HUMAN | | nucleus. |
| Q8N806 | CN130_HUMAN | | cytoplasm. |
| Q8N813 | CC056_HUMAN | | nucleus. |
| Q8N823 | ZN611_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N859 | ZN713_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N865 | CG031_HUMAN | | cytoplasm. |
| Q8N878 | FRMD1_HUMAN | | cytoplasm. |
| Q8N883 | ZN614_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N884 | CF150_HUMAN | | nucleus. |
| Q8N895 | ZN366_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N8A2 | ANR44_HUMAN | | cytoplasm. |
| Q8N8A4 | GAH6_HUMAN | cell membrane (potential). | membrane. |
| Q8N8A6 | DDX51_HUMAN | nucleus; nucleolus (by similarity). | nucleus. |
| Q8N8D1 | PDCD7_HUMAN | nucleus (potential). | nucleus. |
| Q8N8E2 | ZN513_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N8E3 | CCD46_HUMAN | | centriole. |
| Q8N8G2 | VGLL2_HUMAN | nucleus (by similarity). | nucleus. |
| Q8N8J6 | ZN615_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N8K9 | K1958_HUMAN | | nucleus. |
| Q8N8L2 | ZN491_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N8L6 | ARL10_HUMAN | | cytoplasm. |
| Q8N8N0 | RN152_HUMAN | | cytoplasm. |
| Q8N8N7 | ZADH1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8N8Q1 | C56D1_HUMAN | membrane; multi-pass membrane protein (probable). | mitochondrion. |
| Q8N8Q8 | OXAL2_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (potential). | endosome. |
| Q8N8Q9 | NIPA2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8N8R3 | MCATL_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| Q8N8S7 | ENAH_HUMAN | cytoplasm. targeted to the leading edge of lamellipodia and filopodia by mrl family members. colocalizes at filopodial tips with a number of other proteins including vinculin and zyxlin. colocalizes with n-wasp at the leading edge colocalizes with gphn and pfn at synapses (by similarity). | cytoplasm. |
| Q8N8U2 | CDYL2_HUMAN | nucleus (by similarity). | nucleus. |
| Q8N8U3 | ZCHC5_HUMAN | | cytoplasm. nucleus. |
| Q8N8V4 | ANS4B_HUMAN | | nucleus. |
| Q8N8W7 | CU123_HUMAN | | nucleus. |
| Q8N8Z3 | CJ108_HUMAN | | nucleus. |
| Q8N8Z6 | DCBD1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8N8Z8 | ZN441_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N957 | ANKF1_HUMAN | | cytoplasm. |
| Q8N961 | ABTB2_HUMAN | | cytoplasm. nucleus. |
| Q8N966 | ZDH22_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q8N972 | ZN709_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8N988 | ZN557_HUMAN | nucleus (potential). | cytoplasm. nucleus. |

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| Q8N9B4 | ANR42_HUMAN | | cytoplasm. |
| Q8N9F8 | ZN454_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N9I0 | SYT2_HUMAN | synaptic vesicles and chromaffin granules (by similarity). | membrane. |
| Q8N9I9 | DTX3_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8N9K5 | ZN565_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8N9L1 | ZIC4_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q8N9L9 | ACOT4_HUMAN | peroxisome (potential). | peroxisome. |
| Q8N9N7 | LRC57_HUMAN | | cytoplasm. |
| Q8N9R8 | CI126_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q8N9R9 | CN065_HUMAN | | nucleus. |
| Q8N9T2 | CX042_HUMAN | | cytoplasm. |
| Q8N9U0 | TAC2N_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| Q8N9W8 | CN054_HUMAN | | cytoplasm. |
| Q8N9Y4 | CN152_HUMAN | | nucleus. |
| Q8N9Z0 | ZN610_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8NA23 | WDR31_HUMAN | | membrane. |
| Q8NA70 | FA47B_HUMAN | | nucleus. |
| Q8NA75 | WD21C_HUMAN | | membrane. |
| Q8NA92 | THAP8_HUMAN | | cytoplasm. nucleus. |
| Q8NAB2 | KBTB3_HUMAN | | cytoplasm. |
| Q8NAF0 | ZN759_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8NAG6 | ANR41_HUMAN | | membrane. |
| Q8NAJ2 | CI106_HUMAN | | nucleus. |
| Q8NAM6 | ZSCA4_HUMAN | nucleus (potential). | nucleus. |
| Q8NAP3 | ZBT38_HUMAN | nucleus (by similarity). | nucleus. |
| Q8NAP8 | ZBTB8_HUMAN | nucleus (potential). | nucleus. |
| Q8NAT1 | AGO61_HUMAN | | cytoplasm. |
| Q8NAT2 | TDRD5_HUMAN | | cytoplasm. |
| Q8NB12 | SMYD1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. nucleus. |
| Q8NB14 | UBP38_HUMAN | | nucleus. |
| Q8NB15 | ZN511_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8NB16 | MLKL_HUMAN | | cytoplasm. |
| Q8NB25 | CF060_HUMAN | | centriole. |
| Q8NB42 | ZN527_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8NB46 | ANR52_HUMAN | | cytoplasm. |
| Q8NB50 | ZFP62_HUMAN | nucleus (potential). | nucleus. |
| Q8NB54 | CR023_HUMAN | | secreted protein. |
| Q8NB59 | SYT14_HUMAN | membrane; single-pass type iii membrane protein (potential). | endosome; membrane. |
| Q8NB78 | AOF1_HUMAN | | cytoplasm. |
| Q8NBB4 | ZSCA1_HUMAN | nucleus (potential). | nucleus. |
| Q8NBE8 | KLH23_HUMAN | | cytoplasm. |
| Q8NBF1 | GLIS1_HUMAN | nucleus (by similarity). | nucleus. |
| Q8NBF6 | K0241_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q8NBI6 | CC021_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q8NBJ9 | SIDT2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8NBN7 | RDH13_HUMAN | | cytoplasm. mitochondrion. |
| Q8NBP0 | TTC13_HUMAN | | cytoplasm. |
| Q8NBQ5 | DHRS8_HUMAN | secreted protein (potential). | endoplasmic reticulum. |

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| Q8NBQ7 | AQP11_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8NBS9 | TXND5_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| Q8NBT0 | WD51A_HUMAN | | nucleus. |
| Q8NBU5 | ATAD1_HUMAN | | cytoplasm. nucleus. |
| Q8NBV4 | PPAC3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. |
| Q8NBV8 | SYT8_HUMAN | cell membrane; single-pass type iii membrane protein (by similarity). acrosome (by similarity). | endosome; membrane. |
| Q8NBX0 | SCPDH_HUMAN | | mitochondrion. |
| Q8NBZ0 | CCD95_HUMAN | | nucleus. |
| Q8NBZ7 | UXS1_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (by similarity). | endosome; membrane. |
| Q8NC01 | CLC1A_HUMAN | membrane; single-pass type ii membrane protein (probable). | endosome; membrane. |
| Q8NC06 | ACBD4_HUMAN | | centriole. |
| Q8NC24 | CE016_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q8NC44 | CB017_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8NC54 | KCT2_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q8NC60 | CD014_HUMAN | | cytoplasm. |
| Q8NC67 | NETO2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8NC74 | CT151_HUMAN | | nucleus. |
| Q8NC96 | NECP1_HUMAN | cytoplasmic vesicle; clathrin-coated vesicle; clathrin-coated vesicle membrane (by similarity). cell membrane (by similarity). colocalizes with ap-2 at the plasma membrane (by similarity). | endosome. |
| Q8NCA5 | FA98A_HUMAN | | nucleus. |
| Q8NCB2 | CAMKV_HUMAN | membrane; peripheral membrane protein. predominantly observed in association with the plasma membrane of soma and in neurites, both axons and dendrites. may be associated with vesicular structures (by similarity). | cytoplasm. |
| Q8NCC3 | LYPA3_HUMAN | lysosome (by similarity). secreted protein. | mitochondrion. |
| Q8NCH0 | CHSTE_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q8NCJ5 | SPRY3_HUMAN | | nucleus. |
| Q8NCK3 | ZN485_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8NCL4 | GALT6_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q8NCN2 | ZBT34_HUMAN | nucleus (probable). | nucleus. |
| Q8NCN4 | RN169_HUMAN | | cytoplasm. |
| Q8NCQ2 | CS034_HUMAN | | nucleus. |
| Q8NCQ5 | FBX15_HUMAN | | cytoplasm. |

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| Q8NCR0 | B3GL2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | Golgi. |
| Q8NCS7 | CTL5_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q8NCT1 | ARRD4_HUMAN | | membrane. |
| Q8NCU1 | CN048_HUMAN | | cytoplasm. |
| Q8NCW0 | KREM2_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q8NCW6 | GLT11_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q8ND30 | LIPB2_HUMAN | | cytoplasm. |
| Q8ND56 | FA61A_HUMAN | | cytoplasm. |
| Q8ND61 | CC020_HUMAN | membrane; single-pass membrane protein (potential). | endosome. |
| Q8ND76 | CFP1_HUMAN | | centriole. cytoplasm. |
| Q8ND82 | SUHW3_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8ND99 | CS016_HUMAN | | cytoplasm. |
| Q8NDB6 | TMM29_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q8NDC0 | CN032_HUMAN | | nucleus. |
| Q8NDF8 | PAPD5_HUMAN | nucleus (probable). | nucleus. |
| Q8NDH3 | PEPL1_HUMAN | | cytoplasm. |
| Q8NDH6 | AL2SE_HUMAN | | cytoplasm. |
| Q8NDN9 | RCBT1_HUMAN | nucleus (potential). | nucleus. |
| Q8NDP4 | ZN439_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8NDQ6 | ZN540_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8NDT2 | RB15B_HUMAN | nucleus (probable). | nucleus. |
| Q8NDV1 | SIA7C_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |
| Q8NDV3 | SM1L2_HUMAN | nucleus. associates with chromatin. in prophase i stage of meiosis, localizes along the ae of synaptonemal complexes. in late-pachytene-diplotene, the bulk of protein dissociates from the chromosome arms probably because of phosphorylation by plk, except at centromeres, where cohesin complexes remain. remains chromatin associated at the centromeres up to metaphase ii. at anaphase ii, dissociates from centromeres, allowing chromosomes segregation (by similarity). | cytoplasm. nucleus. |
| Q8NDW4 | ZN248_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8NDY6 | BHLH4_HUMAN | nucleus (probable). | nucleus. |
| Q8NDZ0 | CX020_HUMAN | | nucleus. |
| Q8NDZ4 | CC058_HUMAN | | mitochondrion. |
| Q8NE28 | CI096_HUMAN | | cytoplasm. |
| Q8NE31 | FA13C_HUMAN | | nucleus. |
| Q8NE62 | CHDH_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q8NE63 | HIPK4_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8NE71 | ABCF1_HUMAN | | membrane. |
| Q8NE79 | POPD1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8NEA4 | FBX36_HUMAN | | cytoplasm. |

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| Q8NEA5 | CS018_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q8NEA6 | GLIS3_HUMAN | nucleus (by similarity). | nucleus. |
| Q8NEA9 | GMCLL_HUMAN | nucleus; nucleoplasm; nuclear matrix (by similarity). | nucleus. |
| Q8NEB7 | ACRBP_HUMAN | secreted protein (by similarity). acrosome (by similarity). colocalizes with proacrosin in the acrosome of sperm (by similarity). | centriole. |
| Q8NEB9 | PK3C3_HUMAN | | cytoplasm. mitochondrion. nucleus. |
| Q8NEE6 | FXL13_HUMAN | | cytoplasm. |
| Q8NEE8 | TTC16_HUMAN | | cytoplasm. |
| Q8NEG5 | ZSWM2_HUMAN | | cytoplasm. |
| Q8NEJ9 | CN120_HUMAN | | nucleus. |
| Q8NEK5 | ZN548_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8NEL9 | DDHD1_HUMAN | | secreted protein. |
| Q8NEM1 | ZN680_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8NEM2 | SHCBP_HUMAN | | nucleus. |
| Q8NEM7 | FA48A_HUMAN | | cytoplasm. |
| Q8NEN9 | PDZD8_HUMAN | | cytoplasm. |
| Q8NEP3 | LRC50_HUMAN | | cytoplasm. |
| Q8NEP9 | ZN555_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8NEQ6 | CA064_HUMAN | | nucleus. |
| Q8NER1 | TRPV1_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q8NES3 | LFNG_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q8NES8 | DB124_HUMAN | secreted protein (potential). | secreted protein. |
| Q8NET5 | NFAM1_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). partially recruited to lipid rafts upon bcr stimulation (by similarity). | membrane. |
| Q8NET6 | CHSTD_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q8NEV4 | MYO3A_HUMAN | | cytoplasm. |
| Q8NEV8 | SLC2B_HUMAN | | cytoplasm. |
| Q8NEW7 | TMIE_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q8NEX5 | WFDC9_HUMAN | secreted protein (potential). | membrane. |
| Q8NEX6 | WFD11_HUMAN | secreted protein (potential). | secreted protein. |
| Q8NEY3 | SPAT4_HUMAN | | cytoplasm. secreted protein. |
| Q8NEZ3 | WDR19_HUMAN | | cytoplasm. |
| Q8NEZ4 | MLL3_HUMAN | nucleus (probable). | nucleus. |
| Q8NEZ5 | FBX22_HUMAN | | cytoplasm. |
| Q8NF37 | PCAT1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein. Golgi apparatus; Golgi apparatus membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. |
| Q8NF50 | DOCK8_HUMAN | | cytoplasm. |

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| Q8NF91 | SYNE1_HUMAN | nucleus; nuclear membrane; single-pass type iv membrane protein (potential). the largest part of the protein is cytoplasmic, while its c-terminal part is associated with the nuclear envelope, most probably the outer nuclear membrane. in skeletal and smooth muscles, a significant amount is found in the sarcomeres. | membrane. |
| Q8NFA0 | UBP32_HUMAN | | nucleus. |
| Q8NFB2 | FA11A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8NFC6 | FA44A_HUMAN | | nucleus. |
| Q8NFD2 | ANKK1_HUMAN | | cytoplasm. |
| Q8NFG4 | FLCN_HUMAN | | cytoplasm. |
| Q8NFI4 | F10A5_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8NFJ9 | BBS1_HUMAN | | centriole. |
| Q8NFP4 | MDGA1_HUMAN | secreted protein (potential). | cytoplasm. Golgi. |
| Q8NFP7 | NUD10_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8NFP9 | NBEA_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q8NFK5 | BPIL3_HUMAN | secreted protein (by similarity). | cytoplasm. |
| Q8NFK6 | BPIL2_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q8NFR3 | ADMP_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q8NFU9 | CU081_HUMAN | | nucleus. |
| Q8NFW9 | MYRIP_HUMAN | cytoplasm (by similarity). in pre- and post-synaptic areas in photoreceptor cells and in the basal microvilli of retinal pigment epithelium cells. associated with melanosomes. colocalizes with actin filaments (by similarity). | cytoplasm. |
| Q8NFZ0 | FBX18_HUMAN | nucleus (probable). | nucleus. |
| Q8NFZ3 | NLGN1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8NFZ4 | NLGN2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8NG11 | TSN14_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| Q8NG27 | PJA1_HUMAN | | cytoplasm. |
| Q8NG68 | TTL_HUMAN | | cytoplasm. |
| Q8NHG7 | SVIP_HUMAN | endoplasmic reticulum; smooth endoplasmic reticulum; smooth endoplasmic reticulum membrane; peripheral membrane protein (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein (by similarity). cell membrane; peripheral | endoplasmic reticulum. |
| Q8NHH1 | TTL11_HUMAN | | cytoplasm. |
| Q8NHM5 | JHD1B_HUMAN | nucleus (probable). | nucleus. |
| Q8NHP6 | MSPD2_HUMAN | | endosome; membrane. |
| Q8NHQ8 | RASF8_HUMAN | | cytoplasm. |
| Q8NHR9 | PROF4_HUMAN | | cytoplasm. |
| Q8NHS0 | DNJB8_HUMAN | | cytoplasm. |
| Q8NHU2 | CT026_HUMAN | | cytoplasm. |
| Q8NHU6 | TDRD7_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8NHV1 | GIMA7_HUMAN | | cytoplasm. |

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| Q8NHV4 | NEDD1_HUMAN | | cytoplasm. |
| Q8NHW6 | OTOSP_HUMAN | secreted protein (probable). | membrane. |
| Q8NHX4 | SPTA3_HUMAN | | cytoplasm. secreted protein. |
| Q8NHY0 | B4GN2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | Golgi. |
| Q8NHY3 | GA2L2_HUMAN | | cytoplasm. |
| Q8NHY5 | HUS1B_HUMAN | | mitochondrion. |
| Q8NHY6 | ZFP28_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8NHZ7 | MB3L2_HUMAN | | nucleus. |
| Q8NI08 | NCOA7_HUMAN | nucleus (probable). | nucleus. |
| Q8NI27 | THOC2_HUMAN | nucleus (probable). | nucleus. |
| Q8NI28 | CG013_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q8NI29 | FBX27_HUMAN | | cytoplasm. |
| Q8NI35 | INADL_HUMAN | localized in the paranodal region of myelinating schwann cells (by similarity). membrane-associated. localizes to tight junctions in epithelial cells. also found at the apical plasma membrane. | cytoplasm. |
| Q8NI36 | WDR36_HUMAN | | nucleus. |
| Q8NI77 | KI18A_HUMAN | | cytoplasm. nucleus. |
| Q8NI99 | ANGL6_HUMAN | secreted protein (probable). | secreted protein. |
| Q8TA86 | RP9_HUMAN | nucleus (by similarity). | cytoplasm. |
| Q8TA94 | ZN563_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8TAA1 | RNS11_HUMAN | secreted protein (potential). | secreted protein. |
| Q8TAA3 | PSA7L_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q8TAA5 | GRPE2_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | mitochondrion. |
| Q8TAA9 | VANG1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8TAB3 | PCD19_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q8TAB7 | CCD26_HUMAN | | nucleus. |
| Q8TAC2 | JOS2_HUMAN | | cytoplasm. |
| Q8TAC9 | SCAM5_HUMAN | cell membrane; multi-pass membrane protein (by similarity). synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein (by similarity). highly enriched in synaptic vesicles (by similarity). | endosome. Golgi. |
| Q8TAD1 | SPNXE_HUMAN | cytoplasm (by similarity). nucleus (by similarity). associated with nuclear craters (by similarity). | centriole. cytoplasm. |
| Q8TAE6 | PP14C_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). | cytoplasm. Golgi. |
| Q8TAF7 | ZN461_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8TAG5 | IGX1_HUMAN | secreted protein (potential). | nucleus. |
| Q8TAG9 | EXOC6_HUMAN | | cytoplasm. |
| Q8TAK6 | OLIG1_HUMAN | nucleus (by similarity). | nucleus. |
| Q8TAL5 | CI043_HUMAN | | nucleus. |
| Q8TAM1 | BBS10_HUMAN | | cytoplasm. |
| Q8TAP4 | LMO3_HUMAN | | nucleus. |
| Q8TAQ5 | ZN420_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8TAS1 | UHMK1_HUMAN | nucleus. mostly nuclear (by similarity). | cytoplasm. nucleus. |
| Q8TAT2 | FGFP3_HUMAN | secreted protein (probable). | secreted protein. |

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| Q8TAT6 | NPL4_HUMAN | cytoplasm (by similarity). endoplasmic reticulum (by similarity). nucleus (by similarity). associated with the endoplasmic reticulum and nuclear (by similarity). | cytoplasm. |
| Q8TAU0 | NKX23_HUMAN | nucleus (by similarity). | nucleus. |
| Q8TAU3 | ZN417_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8TAV0 | FA76A_HUMAN | | nucleus. |
| Q8TAV3 | CP2W1_HUMAN | membrane; peripheral membrane protein (potential). | endoplasmic reticulum. |
| Q8TAV4 | STML3_HUMAN | cell membrane; single-pass type iii membrane protein (by similarity). detected in lipid rafts (by similarity). | endosome; membrane. |
| Q8TAW3 | ZN671_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8TAX0 | OSR1_HUMAN | nucleus (potential). | nucleus. |
| Q8TB03 | CX038_HUMAN | | cytoplasm. |
| Q8TB05 | F100A_HUMAN | | nucleus. |
| Q8TB24 | RIN3_HUMAN | | cytoplasm. |
| Q8TB36 | GDAP1_HUMAN | cytoplasm (by similarity). | nucleus. |
| Q8TB37 | NUBPL_HUMAN | | cytoplasm. |
| Q8TB40 | ABHD4_HUMAN | | cytoplasm. |
| Q8TB52 | FBX30_HUMAN | | cytoplasm. |
| Q8TB69 | ZN519_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8TB72 | PUM2_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8TB96 | TIP_HUMAN | secreted protein (potential). membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q8TBB0 | THAP6_HUMAN | | cytoplasm. nucleus. |
| Q8TBB1 | LNK1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8TBB5 | KLDC4_HUMAN | | nucleus. |
| Q8TBC4 | UBA3_HUMAN | | cytoplasm. nucleus. |
| Q8TBC5 | ZN447_HUMAN | nucleus (potential). | nucleus. |
| Q8TBE1 | CNIH3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8TBE7 | TMM22_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. |
| Q8TBE9 | NANP_HUMAN | | lysosome. |
| Q8TBF5 | PIGX_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). | endoplasmic reticulum. |
| Q8TBG9 | SYNPR_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q8TBH0 | ARRD2_HUMAN | | membrane. |
| Q8TBK6 | ZCH10_HUMAN | | cytoplasm. nucleus. |
| Q8TBM7 | CJ057_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8TBP0 | TBC16_HUMAN | | cytoplasm. |
| Q8TBR5 | CS023_HUMAN | | nucleus. |
| Q8TBZ3 | WDR20_HUMAN | | membrane. |
| Q8TBZ5 | ZN502_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8TBZ8 | ZN564_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8TC07 | TBC15_HUMAN | | cytoplasm. |
| Q8TC21 | ZN596_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8TC27 | ADA32_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8TC36 | SPA4L_HUMAN | | cytoplasm. secreted protein. |

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| Q8TC41 | IBRD1_HUMAN | | cytoplasm. nucleus. |
| Q8TC44 | WD51B_HUMAN | | cytoplasm. |
| Q8TC59 | PIWL2_HUMAN | cytoplasm (by similarity). | cytoplasm. cytoskeleton. |
| Q8TC84 | FANK1_HUMAN | | cytoplasm. |
| Q8TCA0 | LRC20_HUMAN | | cytoplasm. |
| Q8TCB0 | IFI44_HUMAN | cytoplasm (potential). | secreted protein. |
| Q8TCB7 | METL6_HUMAN | | cytoplasm. |
| Q8TCD6 | PHOP2_HUMAN | | cytoplasm. |
| Q8TCE6 | FA45A_HUMAN | | cytoplasm. |
| Q8TCF1 | ZFAN1_HUMAN | | cytoplasm. nucleus. |
| Q8TCG5 | CPT1C_HUMAN | mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| Q8TCH9 | YV04_HUMAN | | membrane. |
| Q8TCJ0 | FBX25_HUMAN | | cytoplasm. nucleus. |
| Q8TCN5 | ZN507_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8TCT1 | PHOP1_HUMAN | | cytoplasm. |
| Q8TCT6 | PSL4_HUMAN | membrane; multi-pass membrane protein (probable). | endosome. |
| Q8TCT7 | PSL1_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| Q8TCT8 | PSL2_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| Q8TCT9 | HM13_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q8TCU5 | NMD3A_HUMAN | cell membrane; multi-pass membrane protein (by similarity). enriched in post-synaptic plasma membrane and post-synaptic densities. requires the presence of grin1 to be targeted at the plasma membrane (by similarity). | membrane. |
| Q8TCU8 | CH015_HUMAN | | nucleus. |
| Q8TCU9 | CH016_HUMAN | | cytoplasm. |
| Q8TCV5 | WFDC5_HUMAN | secreted protein (potential). | secreted protein. |
| Q8TCX5 | RHPN1_HUMAN | | cytoplasm. |
| Q8TCY5 | MRAP_HUMAN | cytoplasm; perinuclear region. concentrated at the perinuclear membrane region. cytoplasm (by similarity). upon insulin stimulation, it is redistributed into spotty structures throughout the cytoplasm (by similarity). cell membrane. endoplasmic reticulum. localizes both to plasma membrane and endoplasmic | endoplasmic reticulum. |
| Q8TCZ7 | CU074_HUMAN | | cytoplasm. |
| Q8TCZ9 | PKHD1_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q8TD06 | AGR3_HUMAN | secreted protein (probable). | cytoplasm. |
| Q8TD08 | MK15_HUMAN | | cytoplasm. |
| Q8TD10 | MIPO1_HUMAN | | centriole. |
| Q8TD16 | BICD2_HUMAN | Golgi apparatus. in interphase cells mainly localizes to the Golgi complex and colocalizes with dynactin at microtubule plus ends (by similarity). cytoskeleton. its localization is dependent on microtubule morphology. | cytoplasm. |

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|--------|-------------|---|--------------------------|
| Q8TD17 | ZN398_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8TD22 | SFXN5_HUMAN | mitochondrion; mitochondrial membrane; multi-pass membrane protein (by | mitochondrion; membrane. |
| Q8TD26 | CHD6_HUMAN | nucleus (probable). | nucleus. |
| Q8TD30 | ALAT2_HUMAN | | cytoplasm. |
| Q8TD33 | SG1C1_HUMAN | secreted protein (by similarity). | cytoplasm. |
| Q8TD47 | RS4Y2_HUMAN | | cytoplasm. |
| Q8TD88 | CU109_HUMAN | | secreted protein. |
| Q8TD90 | MAGE2_HUMAN | | nucleus. |
| Q8TD91 | MAGC3_HUMAN | | cytoplasm. |
| Q8TD94 | KLF14_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q8TDB6 | DTX3L_HUMAN | | cytoplasm. |
| Q8TDC0 | MYOZ3_HUMAN | localized at the z-line of skeletal muscle (by similarity). | cytoplasm. |
| Q8TDD2 | SP7_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q8TDD5 | MCLN3_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8TDE3 | RNAS8_HUMAN | secreted protein (probable). | secreted protein. |
| Q8TDF5 | NETO1_HUMAN | isoform 2, isoform 3: cell membrane; single-pass type i membrane protein (potential). isoform 1: secreted protein | membrane. |
| Q8TDH9 | MUTED_HUMAN | | cytoplasm. |
| Q8TDI0 | CHD5_HUMAN | nucleus (by similarity). | nucleus. |
| Q8TDI7 | TMC2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8TDI8 | TMC1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8TDL5 | LPLC1_HUMAN | secreted protein (by similarity). | cytoplasm. |
| Q8TDM0 | BCAS4_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q8TDM3 | CB012_HUMAN | | nucleus. |
| Q8TDN4 | CABL1_HUMAN | nucleus (by similarity). cytoplasm (by similarity). located in the cell body and proximal region of the developing axonal shaft of immature neurons. located in axonal growth cone, but not in the distal part of the axon shaft or in dendritic growth cone of mature neurons (by similarity). | cytoplasm. |
| Q8TDN7 | ASAH3_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q8TDN8 | COAS3_HUMAN | | nucleus. |
| Q8TDP1 | RNH2C_HUMAN | nucleus (probable). | centriole. cytoplasm. |
| Q8TDQ0 | TIMD3_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q8TDQ1 | CLM1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8TDR4 | TCP1L_HUMAN | | nucleus. |
| Q8TDW0 | LRC8C_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8TDW5 | SYTL5_HUMAN | membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q8TDX5 | ACMSD_HUMAN | | cytoplasm. |
| Q8TDX6 | CGAT1_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q8TDX7 | NEK7_HUMAN | cytoplasm (by similarity). | cytoplasm. |

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|--------|-------------|---|-----------------------------------|
| Q8TDX9 | PK1L1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8TDY3 | ARPM2_HUMAN | | cytoplasm. |
| Q8TE12 | LMX1A_HUMAN | nucleus (by similarity). | nucleus. |
| Q8TE49 | OTU7A_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. nucleus. |
| Q8TE56 | ATS17_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q8TE57 | ATS16_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q8TE58 | ATS15_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q8TE59 | ATS19_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q8TE60 | ATS18_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q8TE69 | CX04A_HUMAN | | mitochondrion. |
| Q8TE73 | DYH5_HUMAN | | cytoplasm. |
| Q8TE76 | MORC4_HUMAN | | cytoplasm. nucleus. |
| Q8TE77 | SSH3_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q8TE82 | S3TC1_HUMAN | | cytoplasm. |
| Q8TEA8 | DTD1_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q8TEB1 | WDR23_HUMAN | | nucleus. |
| Q8TEH3 | DEN1A_HUMAN | | nucleus. |
| Q8TEK3 | DOT1L_HUMAN | nucleus (probable). | nucleus. |
| Q8TEL6 | TP4AP_HUMAN | | cytoplasm. endoplasmic reticulum. |
| Q8TEQ8 | PIGO_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q8TET4 | GANC_HUMAN | | lysosome. |
| Q8TEW6 | DOK4_HUMAN | | centriole. secreted protein. |
| Q8TEX9 | IPO4_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q8TEY7 | UBP33_HUMAN | | cytoplasm. nucleus. |
| Q8TF05 | PP4R1_HUMAN | | cytoplasm. secreted protein. |
| Q8TF08 | CX7B2_HUMAN | | mitochondrion. |
| Q8TF17 | S3TC2_HUMAN | | cytoplasm. |
| Q8TF32 | ZN431_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8TF39 | ZN483_HUMAN | nucleus (probable). | nucleus. |
| Q8TF42 | STS1_HUMAN | cytoplasm (by similarity). nucleus (potential). | centriole. cytoplasm. |
| Q8TF45 | ZN418_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8TF47 | ZFP90_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q8TF61 | FBX41_HUMAN | | cytoplasm. nucleus. |
| Q8TF64 | GIPC3_HUMAN | | cytoplasm. |
| Q8TF65 | GIPC2_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8TF66 | LRC15_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8TF68 | ZN384_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q8WTP9 | GAGD4_HUMAN | | nucleus. |
| Q8WTR2 | DUS19_HUMAN | | cytoplasm. |
| Q8WTR7 | ZN473_HUMAN | nucleus (probable). | cytoplasm. nucleus. |

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|--------|-------------|---|---|
| Q8WTS1 | ABHD5_HUMAN | cytoplasm (by similarity). colocalized with plin and adrp on the surface of lipid droplets. the localization is dependent upon the metabolic status of the adipocytes and the activity of pka (by | cytoplasm. |
| Q8WTS6 | SETD7_HUMAN | nucleus (probable). | nucleus. |
| Q8WTT0 | CLC4C_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q8WTV1 | THAP3_HUMAN | | cytoplasm. nucleus. |
| Q8WTW4 | TUSC4_HUMAN | | cytoplasm. |
| Q8WTX9 | ZDHC1_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. endoplasmic reticulum. nucleus. |
| Q8WU08 | ST32A_HUMAN | | cytoplasm. |
| Q8WU43 | CB015_HUMAN | | cytoplasm. |
| Q8WU49 | CG033_HUMAN | | membrane. |
| Q8WU66 | TSEAR_HUMAN | secreted protein (probable). | secreted protein. |
| Q8WU76 | SCFD2_HUMAN | | cytoplasm. |
| Q8WU79 | SMP1L_HUMAN | cytoplasm. detected in multiple foci throughout the cytoplasm and in juxtannuclear structures (by similarity). | cytoplasm. |
| Q8WUA2 | PPIL4_HUMAN | nucleus (probable). | nucleus. |
| Q8WUA7 | TB22A_HUMAN | | cytoplasm. |
| Q8WUA8 | TSK_HUMAN | secreted protein (potential). | cytoplasm. |
| Q8WUB8 | PHF10_HUMAN | nucleus (probable). | nucleus. |
| Q8WUD1 | RAB2B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | endoplasmic reticulum. Golgi. |
| Q8WUD4 | CCD12_HUMAN | | nucleus. |
| Q8WUE5 | CX048_HUMAN | | cytoplasm. |
| Q8WUG5 | BOCT_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8WUH1 | CHUR_HUMAN | | nucleus. |
| Q8WUJ0 | STYX_HUMAN | | cytoplasm. |
| Q8WUJ3 | K1199_HUMAN | | centriole. cytoplasm. secreted protein. membrane. |
| Q8WUK0 | PTPM1_HUMAN | mitochondrion; mitochondrial inner membrane (by similarity). associated with the inner membrane (by similarity). | cytoplasm. |
| Q8WUQ7 | CS029_HUMAN | | cytoplasm. |
| Q8WUR7 | U235_HUMAN | | nucleus. |
| Q8WUT4 | CT075_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q8WUU4 | ZN342_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8WUX9 | CHMP7_HUMAN | | cytoplasm. |
| Q8WUY3 | CI065_HUMAN | | nucleus. |
| Q8WUZ0 | BCL7C_HUMAN | | cytoplasm. |
| Q8WV07 | ORAV1_HUMAN | | cytoplasm. |
| Q8WV14 | ZN688_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8WV16 | WD21A_HUMAN | | cytoplasm. |
| Q8WV19 | SFT2A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. Golgi. |
| Q8WV35 | LRC29_HUMAN | | cytoplasm. |
| Q8WV44 | TRI41_HUMAN | | cytoplasm. |
| Q8WV74 | NUDT8_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| Q8WV99 | ZFN2B_HUMAN | | cytoplasm. nucleus. |
| Q8WVD5 | RN141_HUMAN | | cytoplasm. |

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|--------|-------------|---|-----------------------------------|
| Q8WVE6 | TM171_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8WVF2 | CJ049_HUMAN | | cytoplasm. |
| Q8WVF5 | KCTD4_HUMAN | | nucleus; membrane. |
| Q8WVH0 | CPLX3_HUMAN | membrane; lipid-anchor. enriched at the synaptic terminal (by similarity). | endosome. |
| Q8WVJ2 | NUDC2_HUMAN | | cytoplasm. |
| Q8WVK2 | SNUT3_HUMAN | nucleus (potential). | nucleus. |
| Q8WVL7 | ANR49_HUMAN | | cytoplasm. |
| Q8WVM8 | SCFD1_HUMAN | cytoplasm (by similarity). endoplasmic reticulum; endoplasmic reticulum membrane; peripheral membrane protein (by similarity). Golgi apparatus; Golgi stack; Golgi stack membrane; peripheral membrane protein (by similarity). | cytoplasm. endoplasmic reticulum. |
| Q8WVN6 | SCTM1_HUMAN | cell membrane; single-pass type i membrane protein (probable). secreted protein. also found as secreted. | membrane. |
| Q8WVP7 | LMBR1_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q8WVS4 | WDR60_HUMAN | | nucleus. |
| Q8WVT3 | TTC15_HUMAN | | cytoplasm. |
| Q8WVV5 | BT2A2_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q8WVV9 | HNRL1_HUMAN | | nucleus. |
| Q8WVZ1 | ZDH19_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q8WVZ7 | RN133_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q8WVZ9 | KBTB7_HUMAN | | cytoplasm. |
| Q8WW01 | SEN15_HUMAN | nucleus (probable). nucleus; nucleolus (probable). may be transiently localized in the nucleolus (probable). | cytoplasm. nucleus. |
| Q8WW14 | CJ082_HUMAN | | nucleus. |
| Q8WW22 | DNJA4_HUMAN | membrane; lipid-anchor (potential). | cytoplasm. |
| Q8WW27 | ABEC4_HUMAN | | cytoplasm. |
| Q8WW33 | F112B_HUMAN | | centriole. |
| Q8WW36 | ZCH13_HUMAN | | cytoplasm. nucleus. |
| Q8WW38 | FOG2_HUMAN | nucleus (by similarity). | nucleus. |
| Q8WW43 | APH1B_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q8WW59 | SPRY4_HUMAN | | nucleus. |
| Q8WW62 | TMED6_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). | endoplasmic reticulum. |
| Q8WWB3 | DYDC1_HUMAN | | cytoplasm. |
| Q8WWG1 | NRG4_HUMAN | pro-neuregulin-4, membrane-bound isoform: cell membrane; single-pass type i membrane protein (by similarity). neuregulin-4: secreted protein (by similarity). exists also as a proteolytically released soluble growth factor form. the membrane-bound form does not seem to be active (by similarity). | endosome; membrane. |
| Q8WWH4 | ASZ1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8WWI1 | LMO7_HUMAN | | cytoplasm. |

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| Q8WWI5 | CTL1_HUMAN | membrane; multi-pass membrane protein (by similarity). | Golgi. |
| Q8WWM1 | GAGD5_HUMAN | | nucleus. |
| Q8WWM9 | CYGB_HUMAN | cytoplasm (by similarity). | endoplasmic reticulum. |
| Q8WWN8 | CEND3_HUMAN | cytoplasm (by similarity). cell membrane; peripheral membrane protein (by similarity). cytoplasmic, and associated with f-actin-rich membrane ruffles and lamellipodia (by similarity). | cytoplasm. |
| Q8WWP5 | CH006_HUMAN | | membrane. |
| Q8WWP8 | CH005_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8WWR9 | CH022_HUMAN | | cytoplasm. |
| Q8WWU7 | ITLN2_HUMAN | secreted protein (potential). | secreted protein. |
| Q8WWX0 | ASB5_HUMAN | | cytoplasm. |
| Q8WWY7 | WFD12_HUMAN | secreted protein (potential). | secreted protein. |
| Q8WWZ3 | EDAD_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8WX94 | NALP7_HUMAN | | cytoplasm. |
| Q8WXA3 | RUFY2_HUMAN | nucleus (by similarity). | nucleus. |
| Q8WXB4 | ZN606_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q8WXD9 | CSK11_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8WXE0 | CSK12_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8WXF0 | SRR35_HUMAN | nucleus (probable). | nucleus. |
| Q8WXF7 | SPG3A_HUMAN | | cytoplasm. |
| Q8WXG8 | S100Z_HUMAN | | cytoplasm. |
| Q8WXH0 | SYNE2_HUMAN | nucleus; nuclear membrane; single-pass type iv membrane protein (potential). the largest part of the protein is cytoplasmic, while its c-terminal part is associated with the nuclear envelope, most probably the outer nuclear membrane. remains associated with the nuclear envelope during its breakdown in mitotic cells. | membrane. |
| Q8WXH2 | JPH3_HUMAN | cell membrane; peripheral membrane protein (by similarity). endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type iv membrane protein (by similarity). localized predominantly on the plasma membrane. the transmembrane domain is anchored in endoplasmic reticulum membrane, while the n- terminal part associates with the plasma membrane (by similarity). | endoplasmic reticulum. |
| Q8WXH4 | ASB11_HUMAN | | cytoplasm. |
| Q8WXH5 | SOCS4_HUMAN | | cytoplasm. |
| Q8WXH6 | RB40A_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| Q8WXI3 | ASB10_HUMAN | | cytoplasm. |
| Q8WXI8 | CLC4D_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q8WXJ9 | ASB17_HUMAN | | cytoplasm. |
| Q8WXK1 | ASB15_HUMAN | | cytoplasm. |
| Q8WXK2 | ASB14_HUMAN | | cytoplasm. |
| Q8WXK3 | ASB13_HUMAN | | cytoplasm. |
| Q8WXK4 | ASB12_HUMAN | | cytoplasm. |
| Q8WXQ8 | CBPA5_HUMAN | secreted protein (by similarity). | secreted protein. |

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| Q8WXR4 | MYO3B_HUMAN | | cytoplasm. |
| Q8WXS3 | BAALC_HUMAN | cytoplasm. membrane. postsynaptic lipid rafts (by similarity). | cytoplasm. Golgi. |
| Q8WXS4 | CCGL_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q8WXS5 | CCG8_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q8WXS8 | ATS14_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q8WXT5 | FX4L4_HUMAN | nucleus (by similarity). | nucleus. |
| Q8WXW3 | PIBF1_HUMAN | | centriole. |
| Q8WXX5 | DNJC9_HUMAN | | cytoplasm. |
| Q8WXX7 | AUTS2_HUMAN | | centriole. cytoplasm. |
| Q8WY22 | BRI3B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q8WY36 | BBX_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q8WY64 | MYLIP_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8WY91 | THAP4_HUMAN | | cytoplasm. nucleus. |
| Q8WYA0 | IFT81_HUMAN | | cytoplasm. cytoskeleton. |
| Q8WYB5 | MYST4_HUMAN | nucleus (probable). | nucleus. |
| Q8WYH8 | ING5_HUMAN | | nucleus. |
| Q8WYJ6 | SEPT1_HUMAN | | cytoplasm. |
| Q8WYK0 | ACO12_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8WYN0 | ATG4A_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8WYN3 | TAIP2_HUMAN | nucleus (potential). | centriole. nucleus. |
| Q8WYP3 | RIN2_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q8WYP5 | AHTF1_HUMAN | nucleus; nuclear matrix. cytoplasm (by similarity). | nucleus. |
| Q8WYQ4 | CV015_HUMAN | | cytoplasm. |
| Q8WYQ5 | DGCR8_HUMAN | | cytoplasm. |
| Q8WYQ9 | ZCH14_HUMAN | | cytoplasm. nucleus. |
| Q8WYR1 | PI3R5_HUMAN | nucleus (by similarity). cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). | centriole. |
| Q8WYR4 | TSGA2_HUMAN | cytoplasm (by similarity). cytoplasmic in late spermatocytes, secondary spermatocytes and round spermatids. gathered around metaphase chromosomes during meiotic divisions (by similarity). | nucleus. |
| Q8WZ42 | TITIN_HUMAN | cytoplasm (probable). nucleus. | cytoplasm. nucleus. |
| Q8WZ55 | BSND_HUMAN | cell membrane; multi-pass membrane protein (by similarity). cytoplasm (by similarity). a significant amount also observed intracellularly. staining in membranes of the renal tubule and of potassium-secreting epithelia of the inner ear is basolateral (by similarity). | membrane. |
| Q8WZ60 | KLHL6_HUMAN | | cytoplasm. |
| Q8WZ64 | CEND1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q8WZ74 | CTTB2_HUMAN | | nucleus. |
| Q8WZ75 | ROBO4_HUMAN | | membrane. |
| Q8WZ79 | DNS2B_HUMAN | lysosome (probable). | endoplasmic reticulum. nucleus. |
| Q8WZA1 | PMGT1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | Golgi. |

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|--------|-------------|---|---------------------------------------|
| Q8WZA2 | RPGF4_HUMAN | cytoplasm. membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q902F8 | ENK6_HUMAN | | endosome; membrane. |
| Q902F9 | ENK5_HUMAN | | endosome; membrane. |
| Q92185 | SIA8A_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |
| Q92186 | SIA8B_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |
| Q92187 | SIA8D_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |
| Q92478 | CLC2B_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q92481 | AP2B_HUMAN | nucleus (probable). | nucleus. |
| Q92484 | ASM3A_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q92485 | ASM3B_HUMAN | secreted protein (by similarity). | lysosome. |
| Q92499 | DDX1_HUMAN | | cytoplasm. nucleus. |
| Q92502 | STAR8_HUMAN | | cytoplasm. |
| Q92503 | S14L1_HUMAN | | membrane. |
| Q92504 | KE4_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q92506 | DHB8_HUMAN | | cytoplasm. endoplasmic reticulum. |
| Q92508 | FA38A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q92519 | TRIB2_HUMAN | cytoplasm (by similarity). cytoskeleton (by similarity). may associate with the cytoskeleton (by similarity). | cytoplasm. |
| Q92520 | FAM3C_HUMAN | secreted protein (potential). | secreted protein. |
| Q92527 | ANKR7_HUMAN | | centriole. cytoplasm. |
| Q92529 | SHC3_HUMAN | | cytoplasm. |
| Q92530 | PSMF1_HUMAN | | cytoplasm. |
| Q92535 | PIGC_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q92537 | K0247_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q92538 | GBF1_HUMAN | | centriole. cytoplasm. Golgi. nucleus. |
| Q92539 | LPIN2_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q92542 | NICA_HUMAN | membrane; single-pass type i membrane protein (potential). | endoplasmic reticulum. |
| Q92543 | SNX19_HUMAN | | cytoplasm. |
| Q92544 | TM9S4_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q92546 | K0258_HUMAN | | membrane. |

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|--------|-------------|--|------------------------|
| Q92547 | TOPB1_HUMAN | nucleus. detected on unpaired autosomes in meiotic prophase cells. detected on x and y chromosomes during later stages of prophase. colocalizes with atr and h2afx at unsynapsed chromosome cores during prophase (by similarity). has a uniform nuclear distribution during g phase. colocalizes with brca1 at stalled replication forks during s phase. in mitotic cells it colocalizes with brca1 at spindle poles and centrosomes during metaphase and anaphase. detected in discrete foci together with pml and numerous dna repair enzymes after dna damage by alkylating agents, uv or gamma irradiation. | nucleus. |
| Q92561 | PHYIP_HUMAN | | membrane. |
| Q92562 | SAC3_HUMAN | | membrane. |
| Q92563 | TICN2_HUMAN | secreted protein; extracellular space; extracellular matrix (probable). | secreted protein. |
| Q92564 | DCNL4_HUMAN | | centriole. cytoplasm. |
| Q92567 | K0280_HUMAN | | nucleus. |
| Q92569 | P55G_HUMAN | | cytoplasm. |
| Q92570 | NR4A3_HUMAN | nucleus (potential). | nucleus. |
| Q92575 | UBXD2_HUMAN | | cytoplasm. |
| Q92576 | PHF3_HUMAN | | nucleus. |
| Q92599 | SEPT8_HUMAN | | cytoplasm. |
| Q92609 | TBCD5_HUMAN | | cytoplasm. |
| Q92610 | ZN592_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q92611 | EDEM1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (potential). | endoplasmic reticulum. |
| Q92613 | PHF16_HUMAN | | nucleus. |
| Q92614 | MY18A_HUMAN | isoform 1: endoplasmic reticulum-Golgi intermediate compartment (by similarity). isoform 2: cytoplasm (by similarity). isoform 2, which lacks the pdz domain, is diffusely localized in the cytoplasm (by | cytoskeleton. |
| Q92616 | GCN1L_HUMAN | | cytoplasm. |
| Q92618 | ZN516_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q92620 | PRP16_HUMAN | nucleus (by similarity). | nucleus. |
| Q92621 | NU205_HUMAN | nucleus; nuclear envelope; nuclear pore complex (potential). | nucleus. |
| Q92622 | K0226_HUMAN | | cytoplasm. |
| Q92636 | FAN_HUMAN | | cytoplasm. |
| Q92667 | AKAP1_HUMAN | mitochondrion; mitochondrial outer membrane (potential). | endosome. |
| Q92673 | SORL_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q92685 | ALG3_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). | endoplasmic reticulum. |
| Q92686 | NEUG_HUMAN | | secreted protein. |
| Q92688 | AN32B_HUMAN | nucleus. accumulates in the nuclei at the s phase (by similarity). | centriole. |
| Q92696 | PGTA_HUMAN | | cytoplasm. nucleus. |

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|--------|-------------|--|----------------------------------|
| Q92730 | RND1_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| Q92733 | PRCC_HUMAN | | centriole. cytoplasm. |
| Q92734 | TFG_HUMAN | | centriole. cytoplasm. |
| Q92736 | RYR2_HUMAN | membrane; multi-pass membrane protein (probable). | endoplasmic reticulum; membrane. |
| Q92737 | RR22_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| Q92738 | US6NL_HUMAN | | cytoplasm. |
| Q92747 | ARC1A_HUMAN | | cytoplasm. |
| Q92753 | RORB_HUMAN | nucleus (probable). | nucleus. |
| Q92754 | AP2C_HUMAN | nucleus (probable). | nucleus. |
| Q92764 | K1H5_HUMAN | | cytoplasm. |
| Q92765 | SFRP3_HUMAN | secreted protein (probable). | secreted protein. |
| Q92782 | DPF1_HUMAN | cytoplasm (potential). nucleus (potential). | nucleus. |
| Q92783 | STAM1_HUMAN | cytoplasm (probable). | cytoplasm. Golgi. |
| Q92784 | DPF3_HUMAN | cytoplasm (potential). nucleus (potential). | nucleus. |
| Q92786 | PROX1_HUMAN | nucleus (probable). | nucleus. |
| Q92796 | DLG3_HUMAN | | cytoplasm. |
| Q92800 | EZH1_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q92804 | RBP56_HUMAN | nucleus (potential). | nucleus. |
| Q92813 | IOD2_HUMAN | | endosome; membrane. |
| Q92819 | HAS2_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q92823 | NRCAM_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q92824 | PCSK5_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q92828 | COR2A_HUMAN | | cytoplasm. |
| Q92831 | PCAF_HUMAN | nucleus (by similarity). | nucleus. |
| Q92832 | NELL1_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q92833 | JARD2_HUMAN | nucleus (by similarity). | nucleus. |
| Q92834 | RPGR_HUMAN | Golgi apparatus (by similarity). | centriole. |
| Q92839 | HAS1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q92845 | KIFA3_HUMAN | | cytoplasm. |
| Q92851 | CASPA_HUMAN | | cytoplasm. |
| Q92858 | ATOH1_HUMAN | nucleus (probable). | nucleus. |
| Q92870 | APBB2_HUMAN | | cytoplasm. |
| Q92874 | DNLS2_HUMAN | secreted protein (potential). | nucleus. |
| Q92882 | OSTF1_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q92886 | NGN1_HUMAN | nucleus (probable). | nucleus. |
| Q92889 | ERCC4_HUMAN | nucleus (probable). | nucleus. |
| Q92890 | UFD1_HUMAN | | nucleus. |
| Q92901 | RL3L_HUMAN | | cytoplasm. |
| Q92902 | HPS1_HUMAN | | membrane. |
| Q92903 | CDS1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). cytoplasmic aspect of the endoplasmic | endoplasmic reticulum. |
| Q92904 | DAZL_HUMAN | cytoplasm (by similarity). predominantly cytoplasmic. nucleus (by similarity). nuclear in spermatogonia until near the end of the meiotic prophase and cytoplasmic localization from then onward | cytoplasm. |
| Q92913 | FGF13_HUMAN | | secreted protein. |
| Q92914 | FGF11_HUMAN | | secreted protein. |

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|--------|-------------|--|----------------------------------|
| Q92915 | FGF14_HUMAN | nucleus (probable). | secreted protein. |
| Q92917 | GPKOW_HUMAN | | centriole. cytoplasm. |
| Q92918 | M4K1_HUMAN | | cytoplasm. |
| Q92930 | RAB8B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q92932 | PTPR2_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q92935 | EXTL1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. |
| Q92945 | FUBP2_HUMAN | nucleus (by similarity). cytoplasm (by similarity). a small proportion is also found in the cytoplasm of neuronal cell bodies and dendrites (by similarity). | nucleus. |
| Q92956 | TNR14_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q92988 | DLX4_HUMAN | nucleus (potential). | nucleus. |
| Q92990 | GLMN_HUMAN | | cytoplasm. |
| Q92994 | TF3B_HUMAN | nucleus (by similarity). | nucleus. |
| Q92995 | UBP13_HUMAN | | nucleus. |
| Q92997 | DVL3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q93008 | USP9X_HUMAN | | nucleus. |
| Q93034 | CUL5_HUMAN | | nucleus. |
| Q93045 | STMN2_HUMAN | cytoplasm (by similarity). cytoplasm; perinuclear region (by similarity). associated with punctate structures in the perinuclear cytoplasm, axons, and growth cones of developing neurons. scg10 exists in both soluble and membrane- bound forms (by similarity). | cytoplasm. |
| Q93062 | RBPMS_HUMAN | | nucleus. |
| Q93073 | K0256_HUMAN | | nucleus. |
| Q93075 | TATD2_HUMAN | nucleus (potential). | nucleus. |
| Q93083 | MT1L_HUMAN | | cytoplasm. |
| Q93099 | HGD_HUMAN | | endoplasmic reticulum. |
| Q93100 | KPBB_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. mitochondrion. |
| Q969E4 | TCAL3_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q969F0 | FATE1_HUMAN | | endoplasmic reticulum. |
| Q969F1 | CF051_HUMAN | | nucleus. |
| Q969F9 | HPS3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q969G2 | LHX4_HUMAN | nucleus (potential). | nucleus. |
| Q969G6 | RIFK_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q969H0 | FBXW7_HUMAN | nucleus (by similarity). | nucleus. |
| Q969H4 | CNKR1_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). | cytoplasm; membrane. |
| Q969H8 | CS010_HUMAN | secreted protein (by similarity). | cytoplasm. |
| Q969H9 | DIRC1_HUMAN | | nucleus. |
| Q969J2 | ZN307_HUMAN | nucleus (probable). | nucleus. |
| Q969K7 | TMM54_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q969M1 | TM40L_HUMAN | mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| Q969P0 | IGSF8_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |

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|--------|-------------|---|------------------------|
| Q969P5 | FBX32_HUMAN | | cytoplasm. |
| Q969Q0 | RL36L_HUMAN | cytoplasm (probable). | mitochondrion. |
| Q969Q1 | TRI63_HUMAN | cytoplasm (by similarity). nucleus (by similarity). localizes to the m- and z-lines in skeletal muscle. colocalizes with tnai3 in myocytes (by similarity). | cytoplasm. |
| Q969Q4 | ARL11_HUMAN | | Golgi. |
| Q969Q5 | RAB24_HUMAN | endoplasmic reticulum-Golgi intermediate compartment (by similarity). endosome; late endosome (by similarity). | Golgi. |
| Q969R5 | LMBL2_HUMAN | nucleus (probable). | nucleus. |
| Q969S9 | EFG2_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| Q969T3 | SNX21_HUMAN | | cytoplasm. |
| Q969T9 | WBP2_HUMAN | | cytoplasm. |
| Q969U6 | FBXW5_HUMAN | | cytoplasm. |
| Q969V4 | TEKT1_HUMAN | | centriole. |
| Q969V6 | MKL1_HUMAN | nucleus (by similarity). | nucleus. |
| Q969W0 | CN147_HUMAN | | endosome; membrane. |
| Q969W1 | ZDH16_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q969W3 | F104A_HUMAN | | nucleus. |
| Q969W8 | ZN566_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q969W9 | TMEPA_HUMAN | cell membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q969X1 | TMBI1_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q969Z4 | TR19L_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q96A00 | PP14A_HUMAN | cytoplasm (by similarity). | cytoplasm. nucleus. |
| Q96A11 | G3ST3_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q96A23 | CPNE4_HUMAN | | cytoplasm. |
| Q96A25 | T106A_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q96A33 | CCD47_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q96A37 | RN166_HUMAN | | cytoplasm. |
| Q96A44 | SSB4_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q96A47 | ISL2_HUMAN | nucleus (by similarity). | nucleus. |
| Q96A49 | SYAP1_HUMAN | | nucleus. |
| Q96A57 | CT030_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96A61 | TRI52_HUMAN | | cytoplasm. |
| Q96A65 | EXOC4_HUMAN | | cytoplasm. |
| Q96A70 | ADC_HUMAN | | cytoplasm. |
| Q96A72 | MGN2_HUMAN | nucleus (by similarity). | nucleus. |
| Q96AA8 | K0555_HUMAN | | cytoskeleton. |
| Q96AB6 | NTAN1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96AC6 | KIFC2_HUMAN | | cytoplasm. nucleus. |
| Q96AE4 | FUBP1_HUMAN | nucleus (probable). | nucleus. |
| Q96AE7 | TTC17_HUMAN | | cytoplasm. |
| Q96AG4 | LRC59_HUMAN | microsome; microsomal membrane; single-pass type ii membrane protein (by similarity). | membrane. |

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|--------|-------------|--|------------------------|
| Q96AJ9 | VTI1A_HUMAN | | cytoplasm. Golgi. |
| Q96AK3 | ABC3D_HUMAN | | cytoplasm. |
| Q96AP0 | ACD_HUMAN | nucleus (potential). | nucleus. |
| Q96AP4 | CF113_HUMAN | | nucleus. |
| Q96AP7 | ESAM_HUMAN | cell membrane; cell-cell junction; adherens junction; single-pass type i membrane protein (by similarity). cell membrane; cell-cell junction; tight junction; single-pass type i membrane protein (by similarity). | membrane. |
| Q96AT1 | K1143_HUMAN | | nucleus. |
| Q96AT9 | RPE_HUMAN | | cytoplasm. |
| Q96AX1 | VP33A_HUMAN | endosome; late endosome; late endosomal membrane; peripheral membrane protein (by similarity). lysosome; lysosomal membrane; peripheral membrane protein (by similarity). cytoplasmic, peripheral membrane protein associated with late endosomes/lysosomes (by similarity). | endosome. |
| Q96AX2 | RAB37_HUMAN | secretory granules (by similarity). | Golgi. |
| Q96AY3 | FKB10_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| Q96AY4 | TTC28_HUMAN | | cytoplasm. |
| Q96B01 | R51A1_HUMAN | nucleus (by similarity). colocalizes with rad51 to multiple nuclear foci (by | nucleus. |
| Q96B02 | UBE2W_HUMAN | | cytoplasm. |
| Q96B23 | CR025_HUMAN | | nucleus. |
| Q96B26 | EXOS8_HUMAN | cytoplasm (by similarity). nucleus; nucleolus (by similarity). | nucleus. |
| Q96B33 | CLD23_HUMAN | cell membrane; cell-cell junction; tight junction; multi-pass membrane protein (by similarity). | membrane. |
| Q96B45 | CJ032_HUMAN | | cytoplasm. |
| Q96B67 | ARRD3_HUMAN | | membrane. |
| Q96B86 | RGMA_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | secreted protein. |
| Q96BD5 | PF21A_HUMAN | nucleus (by similarity). | nucleus. |
| Q96BD6 | SSB1_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q96BH1 | RNF25_HUMAN | | cytoplasm. |
| Q96BJ8 | ELMO3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96BM0 | FM14B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96BM1 | ANKR9_HUMAN | | secreted protein. |
| Q96BP3 | PPWD1_HUMAN | | cytoplasm. |
| Q96BQ1 | FAM3D_HUMAN | secreted protein (potential). | secreted protein. |
| Q96BQ3 | TRI43_HUMAN | | cytoplasm. |
| Q96BR1 | SGK3_HUMAN | cytoplasmic vesicle (by similarity). endosome; early endosome (by similarity). localized in vesicle- like structures and in the early endosome (by similarity). | cytoplasm. |
| Q96BR6 | ZN669_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96BS2 | TESC_HUMAN | | cytoplasm. |
| Q96BW5 | PTER_HUMAN | | cytoplasm. |
| Q96BW9 | MMP37_HUMAN | mitochondrion (by similarity). | cytoplasm. |
| Q96BX8 | MOL2A_HUMAN | | cytoplasm. |
| Q96BY2 | MOAP1_HUMAN | | cytoplasm. |
| Q96BY6 | DOC10_HUMAN | | cytoplasm. |

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|--------|-------------|--|------------------------|
| Q96BY7 | CN103_HUMAN | | membrane. |
| Q96BY9 | TMM66_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96BZ9 | TBC20_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96C00 | ZBTB9_HUMAN | nucleus (potential). | nucleus. |
| Q96C10 | LGP2_HUMAN | cytoplasm (by similarity). | nucleus. |
| Q96C19 | EFHD2_HUMAN | | cytoplasm. |
| Q96C23 | GALM_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q96C24 | SYTL4_HUMAN | membrane; peripheral membrane protein (by similarity). detected close to the plasma membrane and on secretory granules. in pancreas, interacts with insulin-containing vesicles (by similarity). | cytoplasm. |
| Q96C28 | ZN707_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96C36 | P5CR2_HUMAN | | mitochondrion. |
| Q96C55 | ZN524_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96C90 | PP14B_HUMAN | cytoplasm (probable). | cytoplasm. nucleus. |
| Q96CB8 | PHF22_HUMAN | | nucleus. |
| Q96CD0 | FBXL8_HUMAN | | cytoplasm. |
| Q96CD2 | COAC_HUMAN | | cytoplasm. |
| Q96CE8 | T4S18_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96CG8 | CTHR1_HUMAN | secreted protein; extracellular space (by similarity). | secreted protein. |
| Q96CN5 | LRC45_HUMAN | | cytoplasm. |
| Q96CN9 | GCC1_HUMAN | cytoplasm. Golgi apparatus; Golgi membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q96CS4 | ZN689_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96CS7 | PKHB2_HUMAN | membrane; peripheral membrane protein (potential). | cytoplasm. |
| Q96CT2 | KBTB9_HUMAN | | cytoplasm. |
| Q96CW9 | NTNG2_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| Q96CX2 | KCD12_HUMAN | | nucleus; membrane. |
| Q96CX3 | ZN501_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96D05 | CJ035_HUMAN | | endosome; membrane. |
| Q96D09 | GASP2_HUMAN | | cytoplasm. |
| Q96D15 | RCN3_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (potential). | cytoplasm. |
| Q96D21 | RHES_HUMAN | membrane; lipid-anchor (potential). | cytoplasm. |
| Q96D42 | TIMD1_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q96D59 | RN183_HUMAN | membrane; single-pass membrane protein (potential). | cytoplasm. |
| Q96D71 | REPS1_HUMAN | | cytoplasm. |
| Q96DA0 | U773_HUMAN | secreted protein (potential). | secreted protein. |
| Q96DA2 | RB39B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| Q96DA6 | TIM14_HUMAN | mitochondrion; mitochondrial inner membrane; single-pass membrane protein (probable). | endoplasmic reticulum. |
| Q96DB5 | FA82B_HUMAN | | cytoplasm. |
| Q96DB9 | FXYD5_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |

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|--------|-------------|--|---------------------------|
| Q96DC9 | OTUB2_HUMAN | | cytoplasm. |
| Q96DD0 | LRC39_HUMAN | | cytoplasm. |
| Q96DE0 | NUD16_HUMAN | | cytoplasm. |
| Q96DE5 | CJ104_HUMAN | | cytoplasm. |
| Q96DE9 | CX04B_HUMAN | | mitochondrion. |
| Q96DF8 | DGC14_HUMAN | nucleus (by similarity). | nucleus. |
| Q96DH6 | MSI2H_HUMAN | cytoplasm. associated with polysomes (by similarity). | nucleus. |
| Q96DM3 | MIC1_HUMAN | | cytoplasm. |
| Q96DN5 | WDR67_HUMAN | | nucleus. |
| Q96DN6 | MBD6_HUMAN | | nucleus. |
| Q96DP5 | FMT_HUMAN | mitochondrion (by similarity). | cytoplasm. mitochondrion. |
| Q96DR4 | STAR4_HUMAN | | secreted protein. |
| Q96DR5 | SPLC2_HUMAN | secreted protein (by similarity). | cytoplasm. |
| Q96DR8 | MUCL_HUMAN | secreted protein (probable). membrane (probable). | nucleus. |
| Q96DT5 | DYH11_HUMAN | | cytoplasm. |
| Q96DT6 | ATG4C_HUMAN | cytoplasm (probable). | cytoplasm. lysosome. |
| Q96DT7 | ZBT10_HUMAN | nucleus (potential). | nucleus. |
| Q96DU3 | SLAF6_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q96DU9 | PABP5_HUMAN | cytoplasm (by similarity). | nucleus. |
| Q96DX5 | ASB9_HUMAN | | cytoplasm. |
| Q96DX7 | TRI44_HUMAN | | cytoplasm. |
| Q96DX8 | RTP4_HUMAN | membrane; single-pass type iii membrane protein (potential). | endosome; membrane. |
| Q96DZ1 | XTP3B_HUMAN | secreted protein (potential). | membrane. |
| Q96DZ7 | T4S19_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96E09 | F122A_HUMAN | | nucleus. |
| Q96E11 | RRFM_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| Q96E17 | RAB3C_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q96E40 | CI009_HUMAN | | cytoplasm. |
| Q96E66 | LRC51_HUMAN | | cytoplasm. |
| Q96EC8 | YIPF6_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q96EF6 | FBX17_HUMAN | | cytoplasm. |
| Q96EH3 | CG030_HUMAN | | cytoplasm. |
| Q96EH5 | RL39L_HUMAN | | cytoplasm. mitochondrion. |
| Q96EI5 | TCAL4_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96EK2 | PF21B_HUMAN | | nucleus. |
| Q96EK5 | K1279_HUMAN | | cytoplasm. |
| Q96EK6 | GNA1_HUMAN | Golgi apparatus; Golgi membrane; peripheral membrane protein. endosome; endosomal membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q96EL1 | CC054_HUMAN | | cytoplasm. |
| Q96EN8 | MOCOS_HUMAN | | cytoplasm. |
| Q96EP0 | RNF31_HUMAN | | cytoplasm. |

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|--------|-------------|---|--------------------------|
| Q96EP5 | DAZP1_HUMAN | cytoplasm (by similarity). predominantly cytoplasmic. nucleus (by similarity). nuclear at some stages of spermatozoides development. in midpachytene spermatocytes, it is localized in both the cytoplasm and the nuclei and is clearly excluded from the sex vesicles. in round spermatids, it localizes mainly in the nuclei, whereas in elongated spermatids, it localizes to the cytoplasm (by similarity). | cytoplasm. nucleus. |
| Q96EQ0 | SGTB_HUMAN | | cytoplasm. |
| Q96EQ8 | RN125_HUMAN | | cytoplasm. nucleus. |
| Q96ET8 | F18B2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96EV8 | DTBP1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96EX3 | WDR34_HUMAN | | membrane. |
| Q96EY8 | MMAB_HUMAN | mitochondrion (probable). | mitochondrion. |
| Q96EZ4 | MYEOV_HUMAN | | membrane. |
| Q96F15 | GIMA5_HUMAN | mitochondrion; mitochondrial outer membrane; single-pass type iv membrane protein (by similarity). | endosome. |
| Q96F24 | NRBF2_HUMAN | nucleus (by similarity). cytoplasm (by similarity). | cytoplasm. nucleus. |
| Q96F46 | I17RA_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q96F85 | CB032_HUMAN | | cytoplasm. |
| Q96FA3 | PELI1_HUMAN | | secreted protein. |
| Q96FA7 | CG029_HUMAN | | cytoplasm. |
| Q96FJ0 | STALP_HUMAN | | secreted protein. |
| Q96FJ2 | DYL2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96FL9 | GLT14_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q96FN4 | CPNE2_HUMAN | | membrane. |
| Q96FN5 | KIF12_HUMAN | | cytoplasm. nucleus. |
| Q96FQ6 | S10AG_HUMAN | | cytoplasm. |
| Q96FQ7 | CR018_HUMAN | | cytoplasm. |
| Q96FT7 | ACCN4_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q96FV0 | LRC46_HUMAN | | cytoplasm. |
| Q96FV3 | TSN17_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96FW1 | OTUB1_HUMAN | | cytoplasm. |
| Q96FX7 | TRM61_HUMAN | nucleus (by similarity). | nucleus. |
| Q96FX8 | PERP_HUMAN | cell membrane; cell-cell junction; desmosome; multi-pass membrane protein (by similarity). associated with desmosomes (by similarity). | membrane. |
| Q96FZ2 | DC12_HUMAN | | cytoplasm. |
| Q96G03 | PGM2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96G04 | FA86A_HUMAN | | cytoplasm. |
| Q96G23 | LASS2_HUMAN | nucleus; nuclear membrane; multi-pass membrane protein (potential). endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | nucleus. |

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| Q96G27 | WBP1_HUMAN | | cytoplasm. |
| Q96G30 | CF117_HUMAN | | membrane. |
| Q96G42 | KLD7B_HUMAN | | membrane. |
| Q96G46 | DUS3L_HUMAN | | cytoplasm. |
| Q96G75 | U2508_HUMAN | | cytoplasm. |
| Q96GD0 | PLPP_HUMAN | | membrane. |
| Q96GD3 | SCMH1_HUMAN | nucleus (probable). | nucleus. |
| Q96GE4 | CCD45_HUMAN | | cytoplasm. |
| Q96GE9 | CI123_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96GF1 | RN185_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. |
| Q96GG9 | DCNL1_HUMAN | | nucleus. |
| Q96GM5 | SMRD1_HUMAN | nucleus (probable). | nucleus. |
| Q96GP6 | SREC2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q96GQ7 | DDX27_HUMAN | nucleus (potential). | nucleus. |
| Q96GR4 | ZDH12_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q96GT9 | GAGD3_HUMAN | | nucleus. |
| Q96GU1 | GGEE1_HUMAN | | nucleus. |
| Q96GW9 | SYMM_HUMAN | mitochondrion; mitochondrial matrix (probable). | cytoplasm. |
| Q96GX1 | TECT2_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q96GX5 | MASTL_HUMAN | | cytoplasm. |
| Q96GY3 | ANTOL_HUMAN | | cytoplasm. |
| Q96H15 | TIMD4_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q96H20 | SNF8_HUMAN | cytoplasm (probable). nucleus (probable). | cytoplasm. nucleus. |
| Q96H40 | ZN486_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96H96 | COQ2_HUMAN | mitochondrion; mitochondrial membrane; multi-pass membrane protein (probable). | mitochondrion; membrane. |
| Q96HC4 | PDLI5_HUMAN | | cytoplasm. |
| Q96HD1 | CREL1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96HD9 | ACY3_HUMAN | | lysosome. |
| Q96HE9 | PRR11_HUMAN | | nucleus. |
| Q96HF1 | SFRP2_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q96HH9 | GRAM3_HUMAN | | nucleus. |
| Q96HI0 | SEN5_HUMAN | | lysosome. |
| Q96HJ5 | MS4A3_HUMAN | intracytoplasmic membrane; perinuclear region; multi-pass membrane protein (by similarity). located in the perinuclear area. | membrane. |
| Q96HN2 | SAHH3_HUMAN | | cytoplasm. |
| Q96HP0 | DOCK6_HUMAN | | cytoplasm. |
| Q96HQ0 | ZN419_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96HR3 | TR25_HUMAN | nucleus (potential). | nucleus. |
| Q96HR9 | REEP6_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q96HU8 | DIRA2_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| Q96HW9 | CU124_HUMAN | | membrane. |
| Q96HY6 | CT116_HUMAN | secreted protein (potential). | nucleus. |
| Q96HZ4 | HES6_HUMAN | nucleus (probable). | nucleus. |
| Q96HZ7 | CU119_HUMAN | | secreted protein. |

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|--------|-------------|--|--------------------------------|
| Q96I23 | PREY_HUMAN | mitochondrion (potential). | cytoplasm. |
| Q96I24 | FUBP3_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96I27 | ZN625_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96I34 | PP16A_HUMAN | cell membrane; lipid-anchor (by similarity). | cytoplasm. |
| Q96I51 | WBS16_HUMAN | | cytoplasm. |
| Q96I76 | GPTC3_HUMAN | | centriole. cytoplasm. nucleus. |
| Q96I82 | KAZD1_HUMAN | secreted protein (probable). | secreted protein. |
| Q96ID5 | IGS21_HUMAN | secreted protein (potential). | membrane. |
| Q96IG2 | FXL20_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96IK0 | TM101_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. Golgi. |
| Q96IK1 | FA44B_HUMAN | | cytoplasm. |
| Q96IK5 | GMCL1_HUMAN | nucleus; nucleoplasm; nuclear matrix (by similarity). | nucleus. |
| Q96IL0 | CN153_HUMAN | | nucleus. |
| Q96IM9 | DYDC2_HUMAN | | nucleus. |
| Q96IN3 | CI070_HUMAN | | nucleus. |
| Q96IQ7 | VSIG2_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q96IQ9 | ZN414_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96IT1 | ZN496_HUMAN | nucleus (probable). | nucleus. |
| Q96IT6 | CN128_HUMAN | | mitochondrion. |
| Q96IU2 | ZBED3_HUMAN | | nucleus. |
| Q96IV6 | CE004_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96IW2 | SHD_HUMAN | | cytoplasm. |
| Q96IX5 | USMG5_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q96IZ0 | PAWR_HUMAN | cytoplasm. nucleus. mainly cytoplasmic in absence of apoptosis signal and in normal cells. nuclear in most cancer cell lines. nuclear entry seems to be essential but not sufficient for apoptosis (by similarity). nuclear localization includes nucleoplasm and pml nuclear bodies. | nucleus. |
| Q96IZ2 | CF105_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96IZ6 | METL2_HUMAN | | cytoplasm. |
| Q96IZ7 | RSRC1_HUMAN | | nucleus. |
| Q96J01 | THOC3_HUMAN | nucleus (probable). | nucleus. |
| Q96J02 | ITCH_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q96J84 | KIRR1_HUMAN | membrane; single-pass type i membrane protein (potential). predominantly located at podocyte slit diaphragm. | endosome; membrane. |
| Q96J86 | CYYR1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q96J92 | WNK4_HUMAN | cell membrane; cell-cell junction; tight junction (by similarity). present exclusively in intercellular junctions in the distal convoluted tubule and in both the cytoplasm and intercellular junctions in the cortical collecting duct. wnk4 is part of the tight junction complex (by similarity). | cytoplasm. |
| Q96JB5 | CK5P3_HUMAN | | secreted protein. |

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|--------|-------------|--|---|
| Q96JB6 | LOXL4_HUMAN | secreted protein; extracellular space (potential). | cytoplasm. secreted protein. mitochondrion. |
| Q96JC4 | ZN479_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96JD4 | CF114_HUMAN | | membrane. |
| Q96JG8 | MAGD4_HUMAN | | cytoplasm. |
| Q96JH7 | VCIP1_HUMAN | endoplasmic reticulum (by similarity). Golgi apparatus; Golgi stack (by similarity). associated with Golgi stacks and endoplasmic reticulum (by similarity). | cytoplasm. |
| Q96JH8 | K1849_HUMAN | | cytoplasm. |
| Q96JJ6 | JPH4_HUMAN | membrane; single-pass type iv membrane protein (by similarity). | endosome; membrane. |
| Q96JK2 | WDR22_HUMAN | | nucleus. |
| Q96JL9 | ZN333_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96JM2 | ZN462_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96JM3 | K1802_HUMAN | | nucleus. |
| Q96JM7 | LMBL3_HUMAN | nucleus (potential). | nucleus. |
| Q96JQ0 | PCD16_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q96JQ2 | CLMN_HUMAN | membrane; single-pass type iv membrane protein (potential). | endosome; membrane. |
| Q96JT2 | S45A3_HUMAN | membrane; multi-pass membrane protein (potential). | Golgi. |
| Q96JY0 | MAEL_HUMAN | nucleus (potential). | nucleus. |
| Q96K17 | BT3L4_HUMAN | | cytoplasm. |
| Q96K21 | ZFY19_HUMAN | | cytoplasm. |
| Q96K31 | CH076_HUMAN | | cytoplasm. |
| Q96K37 | SL35E_HUMAN | membrane; multi-pass membrane protein (potential). | Golgi. |
| Q96K62 | ZN499_HUMAN | nucleus (potential). | nucleus. |
| Q96K75 | ZN514_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96K76 | UBP47_HUMAN | | nucleus. |
| Q96KB5 | TOPK_HUMAN | | cytoplasm. |
| Q96KC2 | ARL5B_HUMAN | | Golgi. |
| Q96KC8 | DNJC1_HUMAN | nucleus; nuclear membrane; multi-pass membrane protein. microsome; microsomal membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q96KE9 | BTBD6_HUMAN | | cytoplasm. |
| Q96KF7 | CF162_HUMAN | | endosome; membrane. |
| Q96KH6 | CR012_HUMAN | | membrane. |
| Q96KM6 | K1196_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96KN1 | FA84B_HUMAN | | cytoplasm. |
| Q96KN4 | FA84A_HUMAN | | membrane. |
| Q96KN7 | RPGR1_HUMAN | situated between the axonemal microtubules and the plasma membrane | centriole. |
| Q96KN8 | HRSL5_HUMAN | | nucleus. |
| Q96KP1 | EXOC2_HUMAN | | cytoplasm. |
| Q96KP4 | CNDP2_HUMAN | | secreted protein. |
| Q96KP6 | TNIP3_HUMAN | | centriole. |
| Q96KR6 | CT108_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96KS9 | CH013_HUMAN | | cytoplasm. |
| Q96KT0 | CH012_HUMAN | | nucleus. |
| Q96KT6 | CH014_HUMAN | | nucleus. |

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| Q96KV6 | BT2A3_HUMAN | membrane; single-pass type i membrane protein (by similarity). | endosome; membrane. |
| Q96KX0 | LYZL4_HUMAN | secreted protein (probable). | secreted protein. |
| Q96KX2 | CAZA3_HUMAN | | cytoplasm. |
| Q96L11 | CG034_HUMAN | | membrane. |
| Q96L12 | CALR3_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| Q96L15 | NAR5_HUMAN | secreted protein (potential). | endoplasmic reticulum. |
| Q96L21 | RL10L_HUMAN | | cytoplasm. |
| Q96L34 | MARK4_HUMAN | | cytoplasm. |
| Q96L46 | CPNS2_HUMAN | cytoplasm (by similarity). cell membrane (by similarity). translocates to the plasma membrane upon calcium binding (by similarity). | cytoplasm. |
| Q96L50 | LLR1_HUMAN | | cytoplasm. |
| Q96L62 | ENH4_HUMAN | | endosome; membrane. |
| Q96L91 | EP400_HUMAN | | nucleus. |
| Q96L92 | SNX27_HUMAN | | cytoplasm. |
| Q96L93 | SNX23_HUMAN | | cytoplasm. nucleus. |
| Q96L94 | SNX22_HUMAN | | cytoplasm. |
| Q96LA5 | FCRL2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q96LB4 | VATG3_HUMAN | | mitochondrion. |
| Q96LB8 | PGRP4_HUMAN | membrane; peripheral membrane protein (potential). | cytoplasm. |
| Q96LB9 | PGRP3_HUMAN | membrane; peripheral membrane protein (potential). | cytoplasm. |
| Q96LC9 | BMF_HUMAN | | cytoplasm. |
| Q96LD1 | SGCZ_HUMAN | sarcolemma; sarcolemmal membrane; single- pass type ii membrane protein (by similarity). | endosome; membrane. |
| Q96LD8 | SENP8_HUMAN | | lysosome. |
| Q96LI6 | HSFY1_HUMAN | nucleus (potential). | nucleus. |
| Q96LJ7 | DHRS1_HUMAN | | endoplasmic reticulum. |
| Q96LJ8 | UBXD3_HUMAN | | nucleus. |
| Q96LL9 | WBS18_HUMAN | | cytoplasm. |
| Q96LM9 | CT173_HUMAN | | membrane. |
| Q96LQ0 | CN050_HUMAN | | nucleus. |
| Q96LR5 | UB2E2_HUMAN | | cytoplasm. |
| Q96LR9 | APLD1_HUMAN | cell membrane; multi-pass membrane protein (probable). | membrane. |
| Q96LT4 | SAMD8_HUMAN | membrane; multi-pass membrane protein (potential). | Golgi. |
| Q96LT7 | CI072_HUMAN | | membrane. |
| Q96LW9 | ZN323_HUMAN | nucleus (probable). | nucleus. |
| Q96LX8 | ZN597_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96LZ2 | MAGBA_HUMAN | | cytoplasm. |
| Q96LZ3 | CANB2_HUMAN | | cytoplasm. |
| Q96M20 | CT152_HUMAN | | cytoplasm. |
| Q96M32 | KAD7_HUMAN | | cytoplasm. |
| Q96M42 | CU129_HUMAN | | membrane. |
| Q96M53 | CJ027_HUMAN | | nucleus. |
| Q96M60 | CO033_HUMAN | | cytoplasm. |
| Q96M61 | MAGBI_HUMAN | | cytoplasm. |
| Q96M83 | CCDC7_HUMAN | | nucleus. |
| Q96M91 | CCD11_HUMAN | | centriole. cytoplasm. |

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|--------|-------------|--|----------------------------------|
| Q96M94 | KHL15_HUMAN | | cytoplasm. |
| Q96M95 | CCD42_HUMAN | | cytoskeleton. |
| Q96M96 | FGD4_HUMAN | cytoplasm (by similarity). concentrated in filopodia and poorly detected at lamellipodia. binds along the sides of actin fibers (by similarity). | cytoplasm. |
| Q96M98 | PACRG_HUMAN | | cytoplasm. |
| Q96MC5 | CP045_HUMAN | | nucleus. |
| Q96MC6 | HIAT1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q96MC9 | CA147_HUMAN | | secreted protein. |
| Q96MD7 | CI085_HUMAN | | nucleus. |
| Q96ME1 | FXL18_HUMAN | | cytoplasm. |
| Q96ME7 | ZN512_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96MF2 | STAC3_HUMAN | | cytoplasm. |
| Q96MF6 | CQ10A_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q96MG7 | MAGG1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | centriole. |
| Q96MG8 | PCMD1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96MI6 | PP2CE_HUMAN | nucleus (by similarity). | cytoplasm. |
| Q96MK2 | CT175_HUMAN | | cytoplasm. |
| Q96MK3 | FA20A_HUMAN | secreted protein (potential). | cytoplasm. |
| Q96MM1 | CN081_HUMAN | | mitochondrion. |
| Q96MM6 | HS12B_HUMAN | | cytoplasm. |
| Q96MM7 | H6ST2_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q96MN2 | NALP4_HUMAN | | cytoplasm. |
| Q96MN9 | ZN488_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q96MP5 | ZSWM3_HUMAN | | cytoplasm. nucleus. |
| Q96MR6 | WDR65_HUMAN | membrane; single-pass type iv membrane protein (potential). | endosome; membrane. |
| Q96MR9 | ZN560_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96MS0 | ROBO3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q96MT4 | CF195_HUMAN | | nucleus. |
| Q96MU7 | YTDC1_HUMAN | nucleus (by similarity). localizes to a novel subnuclear structure, the yt bodies (by similarity). | nucleus. |
| Q96MU8 | KREM1_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q96MV8 | ZDH15_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q96MW1 | CCD43_HUMAN | | nucleus. |
| Q96MX3 | ZN553_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96MY1 | CT112_HUMAN | | nucleus. |
| Q96MY7 | CN044_HUMAN | | cytoplasm. |
| Q96MZ0 | GD1L1_HUMAN | | cytoplasm. |
| Q96N03 | CT102_HUMAN | | cytoplasm. |
| Q96N11 | CG026_HUMAN | | nucleus. |
| Q96N20 | ZN75A_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96N35 | TMM83_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q96N46 | TTC14_HUMAN | | cytoplasm. |
| Q96N58 | ZN578_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96N67 | DOCK7_HUMAN | | cytoplasm. |
| Q96N68 | CR015_HUMAN | | endosome; membrane. |

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| Q96N76 | HUTU_HUMAN | | cytoplasm. |
| Q96N87 | S6A18_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q96NA8 | TSNA1_HUMAN | | endosome; membrane. |
| Q96NB2 | SFXN2_HUMAN | mitochondrion; mitochondrial membrane; multi-pass membrane protein (by | mitochondrion; membrane. |
| Q96NB3 | CCD16_HUMAN | nucleus (by similarity). excluded from nucleolus (by similarity). | nucleus. |
| Q96NC0 | ZMAT2_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96ND8 | ZN583_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96NE9 | FRMD6_HUMAN | | cytoplasm. |
| Q96NG5 | ZN558_HUMAN | nucleus (potential). | nucleus. |
| Q96NG8 | ZN582_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96NH3 | CF170_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96NI6 | LRFN5_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q96NI8 | ZN570_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96NJ3 | ZN285_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96NJ5 | BKLH5_HUMAN | | cytoplasm. |
| Q96NJ6 | ZFP3_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96NK8 | NDF6_HUMAN | nucleus (probable). | nucleus. |
| Q96NL3 | ZN599_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96NM4 | GCX1_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q96NR8 | RDH12_HUMAN | | centriole. |
| Q96NS5 | ASB16_HUMAN | | cytoplasm. |
| Q96NT3 | CV013_HUMAN | | cytoplasm. |
| Q96NX5 | KCC1G_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein (by similarity). cell membrane; peripheral membrane protein (by | cytoplasm. |
| Q96NX9 | DACH2_HUMAN | nucleus (probable). | nucleus. |
| Q96NY7 | CLIC6_HUMAN | cytoplasm (by similarity). predominantly cytoplasmic. cell membrane (by similarity). upon chloride ion efflux from the cell, it is translocated to the plasma membrane (by similarity). | membrane. |
| Q96NZ1 | FOXN4_HUMAN | nucleus (potential). | nucleus. |
| Q96P09 | BIRC8_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q96P15 | SPB11_HUMAN | cytoplasm (by similarity). | secreted protein. |
| Q96P26 | 5NT1B_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96P50 | CENB5_HUMAN | | cytoplasm. |
| Q96P53 | WDFY2_HUMAN | | cytoplasm. nucleus. |
| Q96P63 | SPB12_HUMAN | cytoplasm (by similarity). | secreted protein. |
| Q96P70 | IPO9_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q96PB7 | NOE3_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q96PB8 | LRC3B_HUMAN | | cytoplasm. |
| Q96PC5 | MIA2_HUMAN | secreted protein (potential). | nucleus. |
| Q96PD6 | MOGT1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q96PE3 | INP4A_HUMAN | | cytoplasm. |
| Q96PE6 | ZIM3_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96PE7 | MCEE_HUMAN | mitochondrion (probable). | mitochondrion. |
| Q96PF1 | TGM7_HUMAN | | cytoplasm. |

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| Q96PF2 | TSSK2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96PG2 | M4A10_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q96PH1 | NOX5_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q96PI4 | GAK7_HUMAN | | cytoplasm. Golgi. |
| Q96PK2 | MACF4_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96PK6 | RBM14_HUMAN | nucleus (probable). | nucleus. |
| Q96PL2 | TECTB_HUMAN | cell membrane; lipid-anchor; gpi-anchor; extracellular side (probable). found in the non-collagenous matrix of the tectorial membrane (by similarity). | secreted protein. |
| Q96PM5 | ZN363_HUMAN | | cytoplasm. |
| Q96PM8 | CU118_HUMAN | | nucleus. |
| Q96PN8 | TSSK3_HUMAN | | cytoplasm. |
| Q96PP8 | GBP5_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| Q96PP9 | GBP4_HUMAN | | cytoplasm. |
| Q96PQ6 | ZN317_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96PQ7 | KLHL5_HUMAN | | cytoplasm. |
| Q96PS1 | CC024_HUMAN | | cytoplasm. |
| Q96PU8 | QKI_HUMAN | nucleus (by similarity). cytoplasm (by similarity). | cytoplasm. nucleus. |
| Q96PV0 | SYGP1_HUMAN | | cytoplasm. |
| Q96PX8 | SLIK1_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q96PY6 | NEK1_HUMAN | nucleus (probable). | nucleus. |
| Q96PZ0 | PUS7_HUMAN | | nucleus. |
| Q96PZ7 | CSMD1_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q96Q07 | BTBD9_HUMAN | | cytoplasm. |
| Q96Q27 | ASB2_HUMAN | | cytoplasm. |
| Q96Q35 | AL2SB_HUMAN | | cytoplasm. secreted protein. |
| Q96Q40 | AL2S7_HUMAN | | cytoplasm. |
| Q96Q42 | ALS2_HUMAN | | cytoplasm. |
| Q96Q45 | AL2S4_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96Q77 | KIP3_HUMAN | | cytoplasm. |
| Q96Q80 | DERL3_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q96Q83 | ALKB3_HUMAN | | nucleus. |
| Q96QA5 | GSDM1_HUMAN | | cytoplasm. |
| Q96QB1 | RHG07_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q96QE4 | LR37B_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q96QG7 | MTMR9_HUMAN | | cytoplasm. |
| Q96QI5 | OST6_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q96QS1 | TSN32_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96QS3 | ARX_HUMAN | nucleus (by similarity). | nucleus. |
| Q96QS6 | KPSH2_HUMAN | | cytoplasm. |
| Q96QT4 | TRPM7_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |

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| Q96QU1 | PCD15_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q96QV1 | HHIP_HUMAN | cell membrane; peripheral membrane protein (by similarity). secreted protein (by similarity). the last 22 c-terminal amino acids may participate in cell membrane attachment. isoform 2: cytoplasm (probable). | cytoplasm. |
| Q96QZ0 | PANX3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96R05 | RET7_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96RD6 | PANX2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96RD7 | PANX1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96RD9 | FCRL5_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q96RF0 | SNX18_HUMAN | | cytoplasm. |
| Q96RI1 | NR1H4_HUMAN | nucleus (probable). | nucleus. |
| Q96RJ3 | TR13C_HUMAN | membrane; single-pass type iii membrane protein (probable). | membrane. |
| Q96RK0 | CIC_HUMAN | | nucleus. |
| Q96RK4 | BBS4_HUMAN | | centriole. cytoplasm. |
| Q96RL7 | VP13A_HUMAN | | centriole. cytoplasm. |
| Q96RM1 | SPR2F_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96RP7 | G3ST4_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (probable). | Golgi. |
| Q96RQ9 | OXLA_HUMAN | isoform 1, isoform 2: lysosome (by similarity). | mitochondrion. |
| Q96RR4 | KKCC2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q96RT6 | CTGE2_HUMAN | | endosome. |
| Q96RU2 | UBP28_HUMAN | | nucleus. |
| Q96RU7 | TRIB3_HUMAN | | nucleus. |
| Q96RU8 | TRIB1_HUMAN | | cytoplasm. |
| Q96RV3 | PCX1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96RW7 | HMCN1_HUMAN | | secreted protein. |
| Q96RY7 | IF140_HUMAN | | cytoplasm. |
| Q96S15 | WDR24_HUMAN | | membrane. |
| Q96S21 | RB40C_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| Q96S42 | NODAL_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q96S55 | WRIP1_HUMAN | nucleus (by similarity). colocalizes with wrn in granular structures in the nucleus (by similarity). | nucleus. |
| Q96S65 | AXUD1_HUMAN | nucleus (probable). | nucleus. |
| Q96S82 | UBL7_HUMAN | | cytoplasm. |
| Q96S86 | HPLN3_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q96S90 | LYSM1_HUMAN | | centriole. |
| Q96S96 | PEBPL_HUMAN | secreted protein (potential). | membrane. |
| Q96S97 | MYADM_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| Q96SA4 | SERC2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |

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|--------|-------------|---|------------------------------|
| Q96SB3 | NEB2_HUMAN | cytoplasm (by similarity). nucleus (by similarity). enriched at synapse and cadherin-based cell-cell adhesion sites. in neurons, both cytosolic and membrane-associated, and highly enriched in the post-synaptic density apposed to excitatory synapses. colocalizes with ppp1r2 at actin-rich adherens junctions in epithelial cells and in dendritic spines (by similarity). | cytoplasm. nucleus. |
| Q96SE7 | ZN347_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96SF7 | TBX15_HUMAN | nucleus (potential). | nucleus. |
| Q96SI1 | KCD15_HUMAN | | nucleus; membrane. |
| Q96SJ8 | TSN18_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q96SK3 | ZN607_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q96SL4 | GPX7_HUMAN | secreted protein (potential). | cytoplasm. |
| Q96SL8 | FIZ1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. nucleus. |
| Q96SM3 | CPXM1_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q96SN7 | ORAI2_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q96SQ5 | ZN587_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96SR6 | ZN382_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q96ST2 | IWS1_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| Q96ST3 | SIN3A_HUMAN | nucleus (by similarity). | nucleus. |
| Q96SU4 | OSBL9_HUMAN | | secreted protein. |
| Q96SW2 | CRBN_HUMAN | | centriole. |
| Q96SZ4 | ZN206_HUMAN | nucleus (potential). | nucleus. |
| Q96SZ5 | CJ022_HUMAN | | nucleus. |
| Q96SZ6 | CK5P1_HUMAN | | cytoplasm. |
| Q96T21 | SEBP2_HUMAN | nucleus (potential). | nucleus. |
| Q96T25 | ZIC5_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q96T49 | PP16B_HUMAN | cell membrane; lipid-anchor (potential). | membrane. |
| Q96T66 | NMNA3_HUMAN | nucleus (probable). | nucleus. |
| Q96T68 | SETB2_HUMAN | nucleus (potential). | nucleus. |
| Q96T75 | DSCR8_HUMAN | | centriole. cytoplasm. |
| Q96TA1 | NIBL_HUMAN | | nucleus. |
| Q99062 | CSF3R_HUMAN | cell membrane; single-pass type i membrane protein. isoform gcsfr2: secreted protein (probable). isoform gcsfr2, which lacks the transmembrane domain, may represent a soluble form of the | membrane. |
| Q99259 | DCE1_HUMAN | | cytoplasm. |
| Q99418 | CYH2_HUMAN | | cytoplasm. |
| Q99426 | TBCB_HUMAN | | cytoplasm. |
| Q99435 | NELL2_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q99439 | CNN2_HUMAN | | cytoplasm. |
| Q99440 | CD006_HUMAN | | cytoplasm. secreted protein. |
| Q99442 | SEC62_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q99445 | GML_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | secreted protein; membrane. |
| Q99447 | PCY2_HUMAN | | cytoplasm. |
| Q99453 | PHX2B_HUMAN | nucleus (by similarity). | nucleus. |
| Q99456 | K1C12_HUMAN | | cytoplasm. |

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|--------|-------------|---|------------------------------------|
| Q99457 | NP1L3_HUMAN | nucleus (by similarity). | centriole. |
| Q99460 | PSD1_HUMAN | | centriole. cytoplasm. endosome. |
| Q99470 | SDF2_HUMAN | secreted protein (probable). | secreted protein; membrane. |
| Q99471 | PFD5_HUMAN | | nucleus. |
| Q99496 | RING2_HUMAN | nucleus. enriched on inactive x chromosome (xi) in female trophoblast stem (ts) cells as well as differentiating embryonic stem (es) cells. the enrichment on xi is transient during ts and es cell differentiation. the association with xi is mitotically stable in non-differentiated ts | nucleus. |
| Q99501 | GA2L1_HUMAN | | cytoplasm. |
| Q99502 | EYA1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| Q99504 | EYA3_HUMAN | cytoplasm (by similarity). nucleus (by similarity). with decreasing efficiency, translocated to the nucleus by six2 and six5, and six4, respectively (by similarity). | nucleus. |
| Q99526 | FOXE2_HUMAN | nucleus (potential). | nucleus. |
| Q99536 | VAT1_HUMAN | | cytoplasm; membrane. |
| Q99538 | LGMN_HUMAN | lysosome (by similarity). | lysosome. |
| Q99542 | MMP19_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q99543 | ZRF1_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q99550 | MPP9_HUMAN | Golgi apparatus; Golgi membrane; peripheral membrane protein (probable). | cytoplasm. |
| Q99570 | PI3R4_HUMAN | | cytoplasm. |
| Q99576 | T22D3_HUMAN | | nucleus. |
| Q99584 | S10AD_HUMAN | | cytoplasm. |
| Q99593 | TBX5_HUMAN | nucleus (potential). | nucleus. |
| Q99611 | SPS2_HUMAN | | cytoplasm. |
| Q99613 | IF38_HUMAN | | cytoplasm. |
| Q99614 | TTC1_HUMAN | | cytoplasm. |
| Q99615 | DNJC7_HUMAN | | cytoplasm. |
| Q99619 | SPSB2_HUMAN | cytoplasm (probable). | centriole. cytoplasm. |
| Q99622 | C10_HUMAN | | mitochondrion. |
| Q99623 | PHB2_HUMAN | mitochondrion; mitochondrial inner membrane (by similarity). cytoplasm (by similarity). nucleus (by similarity). also cytoplasmic and nuclear (by similarity). | nucleus. |
| Q99624 | S38A3_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q99633 | PRP18_HUMAN | nucleus; nucleoplasm; nuclear speckle (by similarity). colocalizes with spliceosomal snrnps (by similarity). | nucleus. |
| Q99645 | PGLB_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q99653 | CHP1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q99661 | KIF2C_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| Q99666 | RGPD8_HUMAN | | cytoplasm. |
| Q99674 | CGRE1_HUMAN | | cytoplasm. |
| Q99675 | CGRF1_HUMAN | | cytoplasm. |
| Q99676 | ZN184_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q99683 | M3K5_HUMAN | | cytoplasm. |

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|--------|-------------|--|------------------------|
| Q99685 | MGLL_HUMAN | | cytoplasm. |
| Q99687 | MEIS3_HUMAN | nucleus (probable). | nucleus. |
| Q99689 | FEZ1_HUMAN | cytoplasm (by similarity). cell membrane (by similarity). translocated from the plasma membrane to the cytoplasm by activation of the pkc zeta (by similarity). | cytoplasm. |
| Q99698 | LYST_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q99700 | ATX2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q99704 | DOK1_HUMAN | | cytoplasm. |
| Q99714 | HCD2_HUMAN | | mitochondrion. |
| Q99715 | COCA1_HUMAN | | secreted protein. |
| Q99719 | SEPT5_HUMAN | | cytoplasm. |
| Q99726 | ZNT3_HUMAN | synaptic vesicle; synaptic vesicle membrane; multi-pass membrane protein (probable). | membrane. |
| Q99735 | MGST2_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q99742 | NPAS1_HUMAN | nucleus (potential). | nucleus. |
| Q99743 | NPAS2_HUMAN | nucleus (potential). | nucleus. |
| Q99747 | SNAG_HUMAN | membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q99750 | MDFI_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q99755 | PI51A_HUMAN | cell membrane. intracytoplasmic membrane. associated with the plasma membrane and with internal membranes. Golgi apparatus; Golgi stack (by similarity). associated with Golgi stacks (by similarity). detected on rac1-induced plasma membrane ruffles, and on membrane ruffles induced by platelet- derived growth | membrane. |
| Q99758 | ABCA3_HUMAN | | membrane. |
| Q99759 | M3K3_HUMAN | | cytoplasm. |
| Q99765 | YV02_HUMAN | | centriole. cytoplasm. |
| Q99766 | ATP5S_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| Q99767 | APBA2_HUMAN | | cytoplasm. |
| Q99784 | NOE1_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (potential). | membrane. |
| Q99798 | ACON_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| Q99805 | TM9S2_HUMAN | endosome; endosomal membrane; multi-pass membrane protein (probable). | membrane. |
| Q99807 | COQ7_HUMAN | | centriole. |
| Q99811 | PRRX2_HUMAN | nucleus (by similarity). | nucleus. |
| Q99814 | EPAS1_HUMAN | | nucleus. |
| Q99829 | CPNE1_HUMAN | | cytoplasm. |
| Q99832 | TCPH_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q99836 | MYD88_HUMAN | cytoplasm (probable). | membrane. |
| Q99853 | FOXB1_HUMAN | nucleus (by similarity). | nucleus. |
| Q99856 | ARI3A_HUMAN | nucleus (by similarity). | nucleus. |
| Q99865 | SPIN3_HUMAN | | cytoplasm. |
| Q99867 | TBB4Q_HUMAN | | centriole. |
| Q99871 | S1769_HUMAN | | cytoplasm. |
| Q99895 | CLCR_HUMAN | | secreted protein. |
| Q99909 | SSX3_HUMAN | | nucleus. |
| Q99933 | BAG1_HUMAN | | cytoplasm. |

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|--------|-------------|---|------------------------------|
| Q99942 | RNF5_HUMAN | membrane; multi-pass membrane protein (potential). predominantly located in the plasma membrane, with some localization occurring within cytoplasmic organelles. | cytoplasm. |
| Q99943 | PLCA_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q99944 | EGFL8_HUMAN | secreted protein (potential). | membrane. |
| Q99946 | PRRT1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q99952 | PTN18_HUMAN | nucleus (by similarity). cytoplasm (by similarity). | cytoplasm. |
| Q99954 | SMR3A_HUMAN | secreted protein (potential). | secreted protein. |
| Q99958 | FOXC2_HUMAN | nucleus (probable). | nucleus. |
| Q99961 | SH3G1_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). associated with internal membranes. concentrated in presynaptic nerve terminals in neurons (by similarity). | cytoplasm. |
| Q99963 | SH3G3_HUMAN | | cytoplasm. |
| Q99969 | TIG2_HUMAN | secreted protein (potential). | cytoplasm. secreted protein. |
| Q99973 | TEP1_HUMAN | | nucleus. |
| Q99983 | OMD_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q99988 | GDF15_HUMAN | secreted protein (probable). | secreted protein. |
| Q99999 | G3ST1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9BPU6 | DPYL5_HUMAN | cytoplasm (potential). | cytoplasm. secreted protein. |
| Q9BPW4 | APOL4_HUMAN | | secreted protein. |
| Q9BPW8 | NIPS1_HUMAN | | mitochondrion. |
| Q9BPX1 | DHR10_HUMAN | | cytoplasm. mitochondrion. |
| Q9BPY8 | HOP_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BPZ2 | SPIN2_HUMAN | | cytoplasm. |
| Q9BPZ7 | SIN1_HUMAN | | cytoplasm. |
| Q9BQ13 | KCD14_HUMAN | | nucleus; membrane. |
| Q9BQ24 | ZFY21_HUMAN | | cytoplasm. nucleus. |
| Q9BQ48 | RM34_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q9BQ50 | TREX2_HUMAN | nucleus (probable). | nucleus. |
| Q9BQ52 | RNZ2_HUMAN | nucleus (probable). | nucleus. |
| Q9BQ66 | KR412_HUMAN | | cytoplasm. |
| Q9BQ69 | LRP16_HUMAN | | nucleus. |
| Q9BQ75 | CC026_HUMAN | | nucleus. |
| Q9BQ87 | TBL1Y_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9BQ89 | CT055_HUMAN | | nucleus. |
| Q9BQ90 | KLDC3_HUMAN | cytoplasm. also found in meiotic chromatin (by similarity). | cytoplasm. |
| Q9BQA1 | MEP50_HUMAN | | cytoplasm. |
| Q9BQA6 | TTY8_HUMAN | | nucleus. |
| Q9BQB4 | SOST_HUMAN | secreted protein (potential). | cytoplasm. secreted protein. |
| Q9BQD1 | MCHL2_HUMAN | | centriole. cytoplasm. |
| Q9BQE3 | TBA6_HUMAN | | centriole. |
| Q9BQE5 | APOL2_HUMAN | cytoplasm (probable). | secreted protein. |
| Q9BQE9 | BCL7B_HUMAN | | nucleus. |
| Q9BQF6 | SENP7_HUMAN | | nucleus. |

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|--------|-------------|--|--------------------------|
| Q9BQG0 | MBB1A_HUMAN | cytoplasm. nucleus. nucleus; nucleolus. shuttles between the nucleus and cytoplasm. nuclear import may be mediated by kpna2, while export appears to depend partially on xpo1/crm1 (by similarity). predominantly nucleolar. | cytoplasm. nucleus. |
| Q9BQG1 | SYT3_HUMAN | synaptic vesicle; synaptic vesicle membrane; single-pass membrane protein (by similarity). | membrane. |
| Q9BQI0 | IBA2_HUMAN | | cytoplasm. |
| Q9BQI3 | E2AK1_HUMAN | | cytoplasm. |
| Q9BQI4 | CCDC3_HUMAN | | endoplasmic reticulum. |
| Q9BQI5 | SGIP1_HUMAN | | cytoplasm. |
| Q9BQI6 | ANR32_HUMAN | | cytoplasm. |
| Q9BQJ4 | TMM47_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9BQK8 | LPIN3_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q9BQM9 | CT144_HUMAN | | nucleus. |
| Q9BQN1 | CT128_HUMAN | | membrane. |
| Q9BQN2 | CT127_HUMAN | | cytoplasm. nucleus. |
| Q9BQP7 | CT072_HUMAN | | cytoplasm. |
| Q9BQP9 | SPLC3_HUMAN | | cytoplasm. |
| Q9BQQ7 | RTP3_HUMAN | membrane; single-pass type iii membrane protein (potential). | endosome; membrane. |
| Q9BQS2 | SYT15_HUMAN | isoform 1, isoform 2, isoform 4: cell membrane; single-pass type iii membrane protein (by similarity). | endosome; membrane. |
| Q9BQS6 | HSPB9_HUMAN | | cytoplasm. |
| Q9BQS7 | HEPH_HUMAN | membrane; single-pass type i membrane protein (potential). | mitochondrion; membrane. |
| Q9BQS8 | FYCO1_HUMAN | | cytoplasm. nucleus. |
| Q9BQT9 | CSTN3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9BQW3 | COE4_HUMAN | nucleus (potential). | nucleus. |
| Q9BQY4 | PEPP2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BQY6 | WFDC6_HUMAN | secreted protein (potential). | secreted protein. |
| Q9BQY9 | DBND2_HUMAN | | cytoplasm. |
| Q9BR01 | ST4A1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9BR09 | NEUL2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9BR10 | CT165_HUMAN | | endosome; membrane. |
| Q9BR11 | ZSWM1_HUMAN | | cytoplasm. nucleus. |

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|--------|-------------|--|------------------------|
| Q9BR39 | JPH2_HUMAN | cell membrane; peripheral membrane protein (by similarity). endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type iv membrane protein (by similarity). sarcoplasmic reticulum; sarcoplasmic reticulum membrane; single- pass type iv membrane protein (by similarity). localized predominantly on the plasma membrane. the transmembrane domain is anchored in endoplasmic/sarcoplasmic reticulum membrane, while the n-terminal part associates with the plasma membrane. in heart cells, it predominantly associates along z lines within myocytes. in skeletal muscle, it is specifically localized at the junction of a and i bands (by similarity). | endoplasmic reticulum. |
| Q9BR46 | CT078_HUMAN | | nucleus. |
| Q9BR61 | ACBD6_HUMAN | | centriole. |
| Q9BR84 | ZN559_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9BRB3 | PIGQ_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9BRG2 | SH23A_HUMAN | | cytoplasm. |
| Q9BRJ7 | SDOS_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9BRK0 | REEP2_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q9BRK4 | LZTS2_HUMAN | | cytoplasm. |
| Q9BRK5 | CAB45_HUMAN | Golgi apparatus; Golgi lumen (by | cytoplasm. |
| Q9BRP0 | OVOL2_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9BRP4 | WDR71_HUMAN | | nucleus. |
| Q9BRQ0 | PYGO2_HUMAN | nucleus (probable). | nucleus. |
| Q9BRQ5 | ORAI3_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q9BRQ6 | CHCH6_HUMAN | | cytoplasm. |
| Q9BRR0 | ZN306_HUMAN | nucleus (probable). | nucleus. |
| Q9BRR3 | CI125_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9BRR6 | ADPGK_HUMAN | | cytoplasm. |
| Q9BRR9 | RHG09_HUMAN | | cytoplasm. |
| Q9BRS2 | RIOK1_HUMAN | | cytoplasm. |
| Q9BRT2 | CF125_HUMAN | | nucleus. |
| Q9BRT8 | CBWD1_HUMAN | | cytoplasm. |
| Q9BRU2 | TCAL7_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9BRX2 | PELO_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q9BRX8 | CJ058_HUMAN | secreted protein (potential). | mitochondrion. |
| Q9BRX9 | MORG1_HUMAN | cytoplasm (by similarity). predominantly cytoplasmic (by similarity). nucleus (by similarity). partially nuclear (by similarity). | centriole. cytoplasm. |
| Q9BRZ2 | TRI56_HUMAN | | cytoplasm. |
| Q9BS34 | ZN670_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9BS40 | LXN_HUMAN | cytoplasm (probable). | secreted protein. |
| Q9BS86 | ZBP1_HUMAN | secreted protein (potential). | secreted protein. |
| Q9BS92 | NPS3B_HUMAN | | cytoplasm. |
| Q9BSD7 | U334_HUMAN | | cytoplasm. |
| Q9BSG5 | RTBDN_HUMAN | secreted protein (potential). | secreted protein. |
| Q9BSH4 | CCD44_HUMAN | | cytoplasm. |

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| Q9BSJ8 | FA62A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9BSK1 | ZN577_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9BSQ5 | CCM2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9BSR8 | YIPF4_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9BST9 | RTKN_HUMAN | | cytoplasm. |
| Q9BSU1 | CP070_HUMAN | | cytoplasm. |
| Q9BSY4 | CHCH5_HUMAN | | cytoplasm. |
| Q9BSY9 | CA121_HUMAN | | cytoplasm. |
| Q9BT22 | ALG1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (probable). | endoplasmic reticulum. |
| Q9BT30 | ALKB7_HUMAN | | membrane. |
| Q9BT49 | THAP7_HUMAN | | cytoplasm. nucleus. |
| Q9BT56 | CL039_HUMAN | secreted protein (potential). | secreted protein. |
| Q9BT67 | NFIP1_HUMAN | Golgi apparatus; Golgi membrane; multi-pass membrane protein (by similarity). | endosome. Golgi. |
| Q9BT76 | UPK3B_HUMAN | membrane; single-pass type i membrane protein (potential). heterodimer formation with uroplakin ib is a prerequisite to exit out of the endoplasmic reticulum (er). | endoplasmic reticulum. |
| Q9BT81 | SOX7_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9BTA0 | CA090_HUMAN | | cytoplasm. |
| Q9BTC0 | DIDO1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). translocates to the nucleus after pro-apoptotic stimuli (by similarity). | nucleus. |
| Q9BTD5 | YU01_HUMAN | | Golgi. |
| Q9BTE1 | DCTN5_HUMAN | | cytoplasm. |
| Q9BTE3 | CJ119_HUMAN | | cytoplasm. |
| Q9BTF0 | THUM2_HUMAN | | cytoplasm. nucleus. |
| Q9BTJ7 | CU097_HUMAN | | cytoplasm. |
| Q9BTK6 | CP053_HUMAN | | cytoplasm. |
| Q9BTL3 | F103A_HUMAN | | nucleus. |
| Q9BTL4 | IER2_HUMAN | | nucleus. |
| Q9BTM9 | CI074_HUMAN | | cytoplasm. |
| Q9BTN0 | LRFN3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9BTP6 | ZBED2_HUMAN | | nucleus. |
| Q9BTT0 | AN32E_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | centriole. |
| Q9BTV6 | WDR85_HUMAN | | cytoplasm. |
| Q9BTV7 | CABL2_HUMAN | | cytoplasm. nucleus. |
| Q9BTW9 | TBCD_HUMAN | | cytoplasm. |
| Q9BTX7 | CT121_HUMAN | | cytoplasm. |
| Q9BTY2 | FUCO2_HUMAN | secreted protein (probable). | lysosome. |
| Q9BU19 | ZN692_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9BU40 | CRDL1_HUMAN | secreted protein (potential). | membrane. |
| Q9BU76 | MMTA2_HUMAN | | nucleus. |
| Q9BU79 | CG023_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9BU89 | DOHH_HUMAN | | endoplasmic reticulum. |
| Q9BUD6 | SPON2_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9BUF5 | TBB6_HUMAN | | centriole. |
| Q9BUG6 | ZSCA5_HUMAN | nucleus (potential). | nucleus. |

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|--------|-------------|---|------------------------|
| Q9BUH8 | BEGIN_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q9BUI4 | RPC62_HUMAN | nucleus (potential). | nucleus. |
| Q9BUK0 | CHCH7_HUMAN | | nucleus. |
| Q9BUL8 | PDC10_HUMAN | | cytoplasm. |
| Q9BUN5 | CC28B_HUMAN | | cytoplasm. |
| Q9BUP0 | EFHD1_HUMAN | | cytoplasm. |
| Q9BUR4 | WDR79_HUMAN | | membrane. |
| Q9BUV8 | CT024_HUMAN | | centriole. cytoplasm. |
| Q9BUW7 | CI016_HUMAN | | centriole. cytoplasm. |
| Q9BUX1 | CHAC1_HUMAN | | cytoplasm. |
| Q9BUY5 | ZN426_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9BV10 | ALG12_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). | endoplasmic reticulum. |
| Q9BV38 | WDR18_HUMAN | | cytoplasm. |
| Q9BV40 | VAMP8_HUMAN | membrane; single-pass type iv membrane protein (probable). | endosome; membrane. |
| Q9BV68 | RN126_HUMAN | | cytoplasm. |
| Q9BV86 | CI032_HUMAN | | cytoplasm. |
| Q9BV87 | CB024_HUMAN | | endosome; membrane. |
| Q9BV94 | EDEM2_HUMAN | secreted protein (potential). | cytoplasm. |
| Q9BV99 | LRC61_HUMAN | | cytoplasm. |
| Q9BVC6 | TM109_HUMAN | nucleus; nuclear outer membrane; multi-pass membrane protein (by similarity). endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). sarcoplasmic reticulum; sarcoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q9BVG3 | TRI62_HUMAN | | cytoplasm. |
| Q9BVG8 | KIFC3_HUMAN | | cytoplasm. nucleus. |
| Q9BVH7 | SIA7E_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. |
| Q9BVI0 | PHF20_HUMAN | nucleus (probable). | nucleus. |
| Q9BVK2 | ALG8_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9BVK6 | TMED9_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). | endoplasmic reticulum. |
| Q9BVL2 | NUPL1_HUMAN | nucleus; nuclear envelope; nuclear pore complex; cytoplasmic side; nucleoplasmic side. biased towards cytoplasmic side. central region of the nuclear pore complex, within the transporter (by similarity). | centriole. nucleus. |
| Q9BVL4 | SELO_HUMAN | | cytoplasm. |
| Q9BVP2 | GNL3_HUMAN | nucleus (by similarity). nucleus; nucleolus (by similarity). shuttles between the nucleus and nucleolus (by similarity). | nucleus. |
| Q9BVS4 | RIOK2_HUMAN | | cytoplasm. |
| Q9BVT8 | TMUB1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9BVV2 | CT195_HUMAN | | cytoplasm. |

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|--------|-------------|--|---------------------------|
| Q9BVV6 | K0586_HUMAN | | nucleus. |
| Q9BVV7 | TI21L_HUMAN | mitochondrion; mitochondrial membrane; single-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9BVV8 | CS024_HUMAN | | nucleus. |
| Q9BW19 | KIFC1_HUMAN | | cytoplasm. nucleus. |
| Q9BW30 | CG38_HUMAN | | cytoplasm. |
| Q9BW60 | ELOV1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9BW62 | KATL1_HUMAN | | cytoplasm. |
| Q9BW71 | HIRP3_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q9BW72 | HIG2A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9BW83 | RAYL_HUMAN | | cytoplasm. |
| Q9BW85 | CCD94_HUMAN | | nucleus. |
| Q9BWG4 | SSBP4_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BWJ5 | SF3B5_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BWL3 | CA043_HUMAN | | endosome; membrane. |
| Q9BWM5 | ZN416_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9BWQ6 | YIPF2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9BWQ8 | FAIM2_HUMAN | cell membrane; multi-pass membrane protein (by similarity). postsynaptic membrane (by similarity). | membrane. |
| Q9BWT3 | PAPOG_HUMAN | nucleus (probable). | nucleus. |
| Q9BWT7 | CAR10_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9BWU1 | CD2L6_HUMAN | | cytoplasm. |
| Q9BWW7 | TTLL2_HUMAN | | cytoplasm. mitochondrion. |
| Q9BWW4 | SSBP3_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BWW8 | APOL6_HUMAN | cytoplasm (probable). | secreted protein. |
| Q9BWW9 | APOL5_HUMAN | cytoplasm (probable). | secreted protein. |
| Q9BWX5 | GATA5_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BX10 | GTPB2_HUMAN | | cytoplasm. |
| Q9BX26 | SYCP2_HUMAN | nucleus (by similarity). in axial/lateral elements of the tripartite segments of synaptonemal complexes (by similarity). | nucleus. |
| Q9BX40 | FA61B_HUMAN | | cytoplasm. |
| Q9BX51 | GGTL6_HUMAN | | centriole. |
| Q9BX59 | TPSNR_HUMAN | cell membrane; single-pass type i membrane protein (probable). endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (probable). mainly found in endoplasmic reticulum but a minority is found on the cell surface. | endoplasmic reticulum. |
| Q9BX67 | JAM3_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9BX68 | HINT2_HUMAN | | mitochondrion. |
| Q9BX69 | CARD6_HUMAN | | cytoplasm. |
| Q9BX82 | ZN471_HUMAN | nucleus (potential). | cytoplasm. nucleus. |

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|---------|-------------|---|------------------------------|
| Q9BX97 | PLVAP_HUMAN | cell membrane; caveola; single-pass type ii membrane protein (by similarity). membrane-associated protein of caveolae. found in fenestral and stomatal diaphragms in fenestrated endothelia and transendothelial channels. cytoplasm; perinuclear region (by similarity). also colocalized with cav1 in perinuclear region (by similarity). | endosome; membrane. |
| Q9BXA6 | TSSK6_HUMAN | | cytoplasm. |
| Q9BXA7 | TSSK1_HUMAN | | cytoplasm. |
| Q9BXA9 | SALL3_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9BXB4 | OSB11_HUMAN | | secreted protein. |
| Q9BXB5 | OSB10_HUMAN | | secreted protein. |
| Q9BXC9 | BBS2_HUMAN | | centriole. |
| Q9BXF3 | CECR2_HUMAN | | cytoplasm. |
| Q9BXF9 | TEKT3_HUMAN | | centriole. |
| Q9BXH5 | TTY6_HUMAN | | membrane. |
| Q9BXI9 | C1QT6_HUMAN | secreted protein (probable). | secreted protein. |
| Q9BXJ0 | C1QT5_HUMAN | secreted protein (probable). | secreted protein. |
| Q9BXJ1 | C1QT1_HUMAN | secreted protein (potential). | secreted protein. |
| Q9BXJ2 | C1QT7_HUMAN | secreted protein (potential). | secreted protein. |
| Q9BXJ3 | C1QT4_HUMAN | secreted protein (potential). | membrane. |
| Q9BXJ4 | C1QT3_HUMAN | secreted protein (potential). | secreted protein. |
| Q9BXJ5 | C1QT2_HUMAN | secreted protein (potential). | secreted protein. |
| Q9BXJ7 | AMNLS_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9B XK1 | KLF16_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9B XK5 | B2L13_HUMAN | isoform 2: mitochondrion. isoform 2, isoform 3: nucleus (probable). | mitochondrion. |
| Q9B XL8 | CDCA4_HUMAN | | cytoplasm. |
| Q9B XM0 | PRAX_HUMAN | nucleus (by similarity). isoform 1: cell membrane (by similarity). isoform 1 is associated with plasma membrane during myelination. isoform 2: cytoplasm (by similarity). | cytoplasm. nucleus. |
| Q9B XM9 | FSD1L_HUMAN | | centriole. cytoplasm. |
| Q9B XN6 | SPNXD_HUMAN | cytoplasm (by similarity). nucleus (by similarity). associated with nuclear craters (by similarity). | centriole. cytoplasm. |
| Q9B XP5 | ARS2_HUMAN | | cytoplasm. secreted protein. |
| Q9B XP8 | PAP2_HUMAN | secreted protein (by similarity). | membrane. |
| Q9B XQ6 | CECR6_HUMAN | | membrane. |
| Q9B XR0 | TGT_HUMAN | | cytoplasm. nucleus. |
| Q9B XR3 | POK2_HUMAN | | nucleus. |
| Q9B XR5 | TLR10_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9B XS1 | IDI2_HUMAN | peroxisome (by similarity). | cytoplasm. peroxisome. |
| Q9B XS4 | TMM59_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9B XT2 | CCG6_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9B XT4 | TDRD1_HUMAN | | cytoplasm. |
| Q9B XT5 | TEX15_HUMAN | | nucleus. |
| Q9B XT6 | M10L1_HUMAN | | nucleus. |
| Q9B XT8 | RNF17_HUMAN | | cytoplasm. |

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|--------|-------------|---|---|
| Q9BXU0 | TEX12_HUMAN | | nucleus. |
| Q9BXU1 | STK31_HUMAN | | cytoplasm. |
| Q9BXU2 | TX13B_HUMAN | | cytoplasm. |
| Q9BXU3 | TX13A_HUMAN | | cytoplasm. |
| Q9BXU7 | UBP26_HUMAN | | nucleus. |
| Q9BXU8 | FHL17_HUMAN | | secreted protein. |
| Q9BXU9 | CALN1_HUMAN | | cytoplasm. |
| Q9BXV9 | CN142_HUMAN | | cytoplasm. |
| Q9BWX6 | OSBL1_HUMAN | | cytoplasm. secreted protein. |
| Q9BWX7 | CECR5_HUMAN | | lysosome. |
| Q9BXX3 | AN30A_HUMAN | | nucleus. |
| Q9BXY4 | RSPO3_HUMAN | secreted protein (by similarity). | mitochondrion. |
| Q9BXY5 | CAYP2_HUMAN | | cytoplasm. |
| Q9BXY8 | BEX2_HUMAN | cytoplasm (by similarity). nucleus. | centriole. cytoplasm. |
| Q9BY10 | TSCOT_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | Golgi. |
| Q9BY11 | PACN1_HUMAN | cytoplasm (by similarity). colocalizes with dynamin i at vesicular structures in the cell body and neurites (by similarity). | cytoplasm. |
| Q9BY12 | ZN291_HUMAN | | nucleus. |
| Q9BY14 | TX101_HUMAN | cell membrane; lipid-anchor; gpi-anchor. associated with lipid rafts (by similarity). | secreted protein. |
| Q9BY42 | CT043_HUMAN | | nucleus. |
| Q9BY50 | SPC21_HUMAN | microsome; microsomal membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. secreted protein. microsome. membrane. |
| Q9BY60 | GBRL3_HUMAN | | cytoplasm. |
| Q9BY66 | JAD1D_HUMAN | | nucleus. |
| Q9BY71 | LRRC3_HUMAN | | cytoplasm. |
| Q9BY77 | PDIP3_HUMAN | nucleus (probable). | nucleus. |
| Q9BY78 | RNF26_HUMAN | | cytoplasm. |
| Q9BY79 | MFRP_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9BYB4 | GNB1L_HUMAN | | cytoplasm. |
| Q9BYC5 | FUT8_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (by similarity). membrane-bound form in trans cisternae of Golgi (by | Golgi. |
| Q9BYD9 | ARPM1_HUMAN | | cytoplasm. |
| Q9BYE2 | TMPSD_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9BYE3 | LCE3D_HUMAN | | cytoplasm. |
| Q9BYE4 | SPR2G_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9BYG4 | PAR6G_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9BYG5 | PAR6B_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9BYG7 | MSTRO_HUMAN | nucleus; nucleolus (by similarity). | centriole. nucleus. |
| Q9BYH1 | SEZ6L_HUMAN | membrane; single-pass type i membrane protein (probable). | endosome; membrane. |
| Q9BYJ1 | LOXE3_HUMAN | | cytoplasm. |
| Q9BYJ4 | TRI34_HUMAN | | cytoplasm. |
| Q9BYJ9 | YTHD1_HUMAN | | nucleus. |
| Q9BYL1 | SAM10_HUMAN | | nucleus. |
| Q9BYM8 | UB713_HUMAN | | cytoplasm. |
| Q9BYN7 | ZN341_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9BYP7 | WNK3_HUMAN | | cytoplasm. |
| Q9BYP8 | KR171_HUMAN | | cytoplasm. |

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|--------|-------------|---|-------------------------------------|
| Q9BYP9 | KRA99_HUMAN | | cytoplasm. |
| Q9BYQ0 | KRA98_HUMAN | | cytoplasm. |
| Q9BYQ2 | KRA94_HUMAN | | cytoplasm. |
| Q9BYQ3 | KRA93_HUMAN | | cytoplasm. |
| Q9BYQ4 | KRA92_HUMAN | | cytoplasm. |
| Q9BYQ6 | KR414_HUMAN | | cytoplasm. |
| Q9BYQ7 | KR410_HUMAN | | cytoplasm. |
| Q9BYR0 | KRA47_HUMAN | | cytoplasm. |
| Q9BYR2 | KRA45_HUMAN | | cytoplasm. |
| Q9BYR3 | KRA44_HUMAN | | cytoplasm. |
| Q9BYR5 | KRA42_HUMAN | | cytoplasm. |
| Q9BYR6 | KRA33_HUMAN | | cytoplasm. |
| Q9BYR7 | KRA32_HUMAN | | cytoplasm. |
| Q9BYR8 | KRA31_HUMAN | | cytoplasm. |
| Q9BYR9 | KRA24_HUMAN | | cytoplasm. |
| Q9BYS1 | KRA15_HUMAN | | centriole. cytoplasm. |
| Q9BYS8 | LRRC2_HUMAN | | cytoplasm. |
| Q9BYT1 | CT059_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9BYT3 | STK33_HUMAN | | cytoplasm. |
| Q9BYT8 | NEUL_HUMAN | mitochondrion; mitochondrial intermembrane space (by similarity). cytoplasm (by similarity). | cytoplasm. mitochondrion; membrane. |
| Q9BYT9 | TM16C_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| Q9BYU7 | KR413_HUMAN | | cytoplasm. |
| Q9BYV1 | AGT2_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q9BYV2 | TRI54_HUMAN | cytoplasm (by similarity). associates with microtubules. localizes to the z-lines in skeletal muscles (by similarity). | cytoplasm. |
| Q9BYV6 | TRI55_HUMAN | cytoplasm (by similarity). nucleus (by similarity). nuclear under atrophic conditions and upon mechanical signals. localizes to the sarcomeric m-band in cardiomyocytes. colocalizes in part with microtubules (by similarity). | cytoplasm. |
| Q9BYV7 | BCDO2_HUMAN | | cytoplasm. endoplasmic reticulum. |
| Q9BYV9 | BACH2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BYX2 | TBCD2_HUMAN | | cytoplasm. |
| Q9BYX4 | IFIH1_HUMAN | cytoplasm. nucleus (potential). may be found in the nucleus, during apoptosis. | nucleus. |
| Q9BYZ2 | LDH6B_HUMAN | | cytoplasm. |
| Q9BYZ6 | RHBT2_HUMAN | | cytoplasm. |
| Q9BZ01 | FRGL_HUMAN | | nucleus. |
| Q9BZ19 | CT086_HUMAN | | cytoplasm. |
| Q9BZ29 | DOCK9_HUMAN | intracytoplasmic membrane (probable). associated with membranes (probable). | cytoplasm. |
| Q9BZ71 | PITM3_HUMAN | intracytoplasmic membrane; peripheral membrane protein (probable). | cytoplasm. |
| Q9BZ72 | PITM2_HUMAN | intracytoplasmic membrane; peripheral membrane protein (probable). | Golgi; membrane. |
| Q9BZ76 | CNTP3_HUMAN | isoform 1: cell membrane; single-pass type i membrane protein (potential). isoform 2: secreted protein (probable). isoform 2 seems to be secreted. | membrane. |

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| Q9BZ97 | TTY13_HUMAN | | endosome; membrane. |
| Q9BZ98 | TTY12_HUMAN | | Golgi. |
| Q9BZ99 | TTY11_HUMAN | | cytoplasm. |
| Q9BZA0 | TTY10_HUMAN | | secreted protein. |
| Q9BZA1 | TTY9A_HUMAN | | membrane. |
| Q9BZA5 | CY15A_HUMAN | | cytoplasm. |
| Q9BZA7 | PC11X_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9BZA8 | PC11Y_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q9BZC7 | ABCA2_HUMAN | membrane; multi-pass membrane protein (potential). | mitochondrion; membrane. |
| Q9BZE2 | PUS3_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q9BZE3 | BARH1_HUMAN | nucleus (probable). | nucleus. |
| Q9BZE7 | EVG1_HUMAN | | centriole. cytoplasm. |
| Q9BZE9 | ASPC1_HUMAN | cytoplasm. intracellular membrane; peripheral membrane protein (by | secreted protein. |
| Q9BZF1 | OSBL8_HUMAN | | secreted protein. |
| Q9BZF2 | OSBL7_HUMAN | | cytoplasm. secreted protein. |
| Q9BZF3 | OSBL6_HUMAN | | secreted protein. |
| Q9BZG1 | RAB34_HUMAN | cytoplasm (by similarity). Golgi apparatus (by similarity). associates with the Golgi complex (by similarity). | Golgi. |
| Q9BZG9 | LYNX1_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | secreted protein. |
| Q9BZH6 | BRWD2_HUMAN | | endosome; membrane. |
| Q9BZI1 | IRX2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BZJ3 | TRYD_HUMAN | released from the secretory granules upon mast cell activation (by similarity). | secreted protein. |
| Q9BZK7 | TBL1R_HUMAN | nucleus (probable). | nucleus. |
| Q9BZL3 | NID67_HUMAN | | endosome; membrane. |
| Q9BZL6 | KPCD2_HUMAN | | cytoplasm. |
| Q9BZM3 | GSH2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BZP6 | CHIA_HUMAN | secreted protein (probable). isoform 2, isoform 3: cytoplasm. | secreted protein. lysosome. |
| Q9BZQ6 | EDEM3_HUMAN | endoplasmic reticulum (potential). | endoplasmic reticulum. |
| Q9BZQ8 | NIBA_HUMAN | | centriole. cytoplasm. |
| Q9BZR9 | TRIM8_HUMAN | | cytoplasm. |
| Q9BZS1 | FOXP3_HUMAN | nucleus (potential). | nucleus. |
| Q9BZV1 | UBXD1_HUMAN | | centriole. cytoplasm. |
| Q9BZV2 | S19A3_HUMAN | membrane; multi-pass membrane protein (potential). | Golgi. |
| Q9BZV7 | PO5FM_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BZW0 | PO5FN_HUMAN | nucleus (by similarity). | nucleus. |
| Q9BZW5 | TM6S1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9BZW8 | CD244_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9BZX2 | UCK2_HUMAN | | cytoplasm. |
| Q9BZY9 | TRI31_HUMAN | | cytoplasm. |
| Q9C005 | DPY30_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| Q9C009 | FOXQ1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9C010 | IPKB_HUMAN | | cytoplasm. |
| Q9C019 | TRI15_HUMAN | | cytoplasm. |
| Q9C026 | TRIM9_HUMAN | | cytoplasm. |
| Q9C029 | TRIM7_HUMAN | | cytoplasm. |

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|--------|-------------|---|----------------------------------|
| Q9C030 | TRIM6_HUMAN | | cytoplasm. |
| Q9C037 | TRIM4_HUMAN | | cytoplasm. |
| Q9C040 | TRIM2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9C075 | K1C23_HUMAN | | cytoplasm. |
| Q9C0A0 | CNTP4_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9C0A1 | ZFH2_HUMAN | nucleus (probable). | nucleus. |
| Q9C0B0 | ZC3H5_HUMAN | | cytoplasm. nucleus. |
| Q9C0B5 | ZDHC5_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q9C0B6 | FAM5B_HUMAN | secreted protein (potential). | cytoplasm. |
| Q9C0C4 | SEM4C_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9C0D0 | PHAR1_HUMAN | cytoplasm (by similarity). enriched at synapses (by similarity). | cytoplasm. |
| Q9C0D9 | SELI_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9C0E4 | GRIP2_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | cytoplasm. |
| Q9C0E8 | LNP_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9C0F3 | ZN436_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9C0H5 | K1688_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9C0H6 | KLHL4_HUMAN | | cytoplasm. |
| Q9C0I9 | LRC27_HUMAN | | cytoplasm. |
| Q9C0J9 | BHLH3_HUMAN | nucleus (by similarity). | nucleus. |
| Q9C0K0 | BC11B_HUMAN | nucleus (potential). | nucleus. |
| Q9C0K7 | AL2S2_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q9GZL8 | BPEC1_HUMAN | | centriole. cytoplasm. |
| Q9GZM8 | NDEL1_HUMAN | localizes to the cell body of the motor neurons and colocalizes with assembled neurofilaments within axonal processes. localizes to the microtubules of the manchette in elongated spermatids (by similarity). cytoplasm. localizes to the interphase centrosome and the mitotic spindle. | cytoplasm. nucleus. |
| Q9GZN1 | ARP6_HUMAN | | cytoplasm. |
| Q9GZN8 | CT027_HUMAN | | cytoplasm. |
| Q9GZP1 | CT098_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9GZQ3 | COMD5_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| Q9GZR3 | CFC1_HUMAN | | cytoplasm. |
| Q9GZR5 | ELOV4_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9GZR7 | DDX24_HUMAN | | nucleus. |
| Q9GZS0 | DNAI2_HUMAN | | cytoplasm. |
| Q9GZS3 | WDR61_HUMAN | | nucleus. |
| Q9GZS9 | CHST5_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane (probable). Golgi membrane, early secretory pathway. | Golgi. |
| Q9GZT4 | SRR_HUMAN | | cytoplasm. |
| Q9GZT9 | EGLN1_HUMAN | | endoplasmic reticulum. |
| Q9GZU0 | CF062_HUMAN | | nucleus. |
| Q9GZU2 | PEG3_HUMAN | nucleus. cytoplasm (by similarity). | nucleus. |

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|--------|-------------|--|------------------------------|
| Q9GZU5 | NYX_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | centriole. secreted protein. |
| Q9GZU7 | CTDS1_HUMAN | nucleus (potential). | nucleus. |
| Q9GZU8 | NIP30_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q9GZV1 | ANKR2_HUMAN | | centriole. cytoplasm. |
| Q9GZV4 | IF5A2_HUMAN | | cytoplasm. |
| Q9GZV7 | HPLN2_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9GZV8 | PRD14_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9GZW5 | SCND2_HUMAN | nucleus (potential). | nucleus. |
| Q9GZX3 | CHST6_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | Golgi. |
| Q9GZX7 | AICDA_HUMAN | | cytoplasm. |
| Q9GZZ6 | ACH10_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9GZZ9 | UE1D1_HUMAN | | cytoplasm. |
| Q9H000 | MKRN2_HUMAN | | cytoplasm. |
| Q9H063 | MAF1_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9H069 | LRC48_HUMAN | | cytoplasm. |
| Q9H074 | PAIP1_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9H078 | CLPB_HUMAN | | cytoplasm. nucleus. |
| Q9H079 | CO029_HUMAN | | cytoplasm. |
| Q9H082 | RB33B_HUMAN | Golgi apparatus; Golgi membrane; lipid-anchor (by similarity). | Golgi. |
| Q9H091 | ZMY15_HUMAN | | cytoplasm. nucleus. |
| Q9H093 | NUAK2_HUMAN | | cytoplasm. |
| Q9H098 | F107B_HUMAN | | centriole. |
| Q9H0A8 | COMD4_HUMAN | | cytoplasm. |
| Q9H0A9 | CU056_HUMAN | | cytoplasm. |
| Q9H0B3 | K1683_HUMAN | | mitochondrion. |
| Q9H0B6 | KLC2_HUMAN | | cytoplasm. |
| Q9H0B8 | CRLD2_HUMAN | secreted protein (potential). | secreted protein. |
| Q9H0C1 | ZMY12_HUMAN | | cytoplasm. |
| Q9H0C6 | CB031_HUMAN | | membrane. |
| Q9H0D2 | ZN541_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9H0E2 | TOLIP_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9H0E7 | UBP44_HUMAN | | nucleus. |
| Q9H0F5 | RNF38_HUMAN | | cytoplasm. |
| Q9H0F7 | ARL6_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9H0G5 | CCD55_HUMAN | | nucleus. |
| Q9H0H0 | INT2_HUMAN | nucleus; nuclear membrane; single-pass membrane protein (potential). | nucleus. |
| Q9H0H9 | CB014_HUMAN | | membrane. |
| Q9H0J9 | PRP12_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9H0K1 | SN1L2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9H0L4 | CSTFT_HUMAN | nucleus (by similarity). | nucleus. |
| Q9H0M0 | WWP1_HUMAN | | cytoplasm. |
| Q9H0M4 | ZCPW1_HUMAN | | cytoplasm. nucleus. |
| Q9H0M5 | ZN700_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9H0N0 | RAB6C_HUMAN | | Golgi. |
| Q9H0N5 | PHS2_HUMAN | | cytoplasm. |
| Q9H0P0 | 5NT3_HUMAN | cytoplasm (potential). isoform 2: endoplasmic reticulum. | cytoplasm. |
| Q9H0P7 | CF059_HUMAN | | membrane. |
| Q9H0Q0 | FA49A_HUMAN | | cytoplasm. |

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|--------|-------------|---|---|
| Q9H0Q3 | FXVD6_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9H0R1 | CN108_HUMAN | | cytoplasm. |
| Q9H0R3 | CA160_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9H0R8 | GBRL1_HUMAN | | cytoplasm. |
| Q9H0T7 | RAB17_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| Q9H0U3 | IAG2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). | endoplasmic reticulum. |
| Q9H0U4 | RAB1B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q9H0W5 | CCDC8_HUMAN | | cytoplasm; membrane. |
| Q9H0W7 | THAP2_HUMAN | | nucleus. |
| Q9H0X4 | ITFG3_HUMAN | membrane; single-pass type ii membrane protein (potential). | endosome; membrane. |
| Q9H0X9 | OSBL5_HUMAN | | secreted protein. |
| Q9H0Y0 | ATG10_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9H0Z9 | RNPC1_HUMAN | | nucleus. |
| Q9H106 | SIRPD_HUMAN | | nucleus. |
| Q9H114 | CST1L_HUMAN | secreted protein (potential). | secreted protein. |
| Q9H115 | SNAB_HUMAN | membrane; peripheral membrane protein (by similarity). | endoplasmic reticulum. |
| Q9H116 | ZN336_HUMAN | nucleus (probable). | nucleus. |
| Q9H156 | SLIK2_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q9H158 | PCDC1_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9H161 | ALX4_HUMAN | nucleus (by similarity). | nucleus. |
| Q9H169 | STMN4_HUMAN | | cytoplasm. |
| Q9H171 | ZBP1_HUMAN | | cytoplasm. |
| Q9H172 | ABCG4_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9H175 | TAI12_HUMAN | nucleus (potential). | centriole. nucleus. |
| Q9H190 | SDCB2_HUMAN | | cytoplasm. |
| Q9H1A4 | ANC1_HUMAN | | cytoplasm. |
| Q9H1B5 | XYLT2_HUMAN | or: endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (by similarity). or: Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. |
| Q9H1B7 | CN004_HUMAN | nucleus (potential). | nucleus. |
| Q9H1C4 | UN93B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9H1D0 | TRPV6_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9H1D9 | RPO3F_HUMAN | nucleus (potential). | nucleus. |
| Q9H1E5 | TXD13_HUMAN | membrane; single-pass type i membrane protein (potential). | cytoplasm. endoplasmic reticulum. Golgi. mitochondrion. |
| Q9H1F0 | WF10A_HUMAN | secreted protein (potential). | secreted protein. |
| Q9H1H1 | F112A_HUMAN | | nucleus. |
| Q9H1H9 | KI13A_HUMAN | | cytoplasm. nucleus. |
| Q9H1I8 | ASCC2_HUMAN | | cytoplasm. |

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| Q9H1K4 | GHC2_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| Q9H1K6 | MESD1_HUMAN | | membrane. |
| Q9H1L0 | CT166_HUMAN | | secreted protein. |
| Q9H1M3 | DB129_HUMAN | secreted protein (potential). | secreted protein. |
| Q9H1M4 | DB127_HUMAN | secreted protein (potential). | secreted protein. |
| Q9H1N7 | S35B3_HUMAN | membrane; multi-pass membrane protein (potential). | Golgi. |
| Q9H1P3 | OSBL2_HUMAN | | secreted protein. |
| Q9H1P6 | CT085_HUMAN | | cytoplasm. |
| Q9H1Q7 | F113A_HUMAN | | cytoplasm. |
| Q9H1R2 | DUS15_HUMAN | cytoplasm (probable). isoform 3: cell membrane; lipid-anchor; cytoplasmic side. | cytoplasm. |
| Q9H1U4 | MEGF9_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9H1U9 | MCAR1_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (potential). | mitochondrion. |
| Q9H1X1 | CF206_HUMAN | | cytoplasm. |
| Q9H1Z9 | TSN10_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9H201 | EPN3_HUMAN | cytoplasm. cytoplasm; perinuclear region. cytoplasmic vesicle; clathrin-coated vesicle. concentrated in the perinuclear region and associated with clathrin-coated vesicles close to the cell periphery. nucleus (probable). may shuttle to the | cytoplasm. |
| Q9H213 | MAGH1_HUMAN | | cytoplasm. |
| Q9H221 | ABCG8_HUMAN | membrane; multi-pass membrane protein (probable). | Golgi; membrane. |
| Q9H222 | ABCG5_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9H223 | EHD4_HUMAN | | nucleus. |
| Q9H237 | PORCN_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9H246 | CA021_HUMAN | | nucleus. |
| Q9H251 | CAD23_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9H254 | SPTN4_HUMAN | | cytoplasm. |
| Q9H295 | TM7S4_HUMAN | cell membrane; multi-pass membrane protein (potential). localized to the cell surface. | endosome; membrane. |
| Q9H2A3 | NGN2_HUMAN | nucleus (potential). | nucleus. |
| Q9H2A7 | SCYBG_HUMAN | cell membrane; single-pass type i membrane protein (potential). secreted protein. also exists as a soluble form. | membrane. |
| Q9H2A9 | CHST8_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9H2B4 | S26A1_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9H2C0 | GAN_HUMAN | | cytoplasm. |
| Q9H2C1 | LHX5_HUMAN | nucleus (probable). | nucleus. |

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| Q9H2C2 | ARV1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9H2G2 | SLK_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9H2H8 | PPIL3_HUMAN | | cytoplasm. |
| Q9H2I8 | CJ011_HUMAN | | cytoplasm. |
| Q9H2J1 | CI037_HUMAN | | nucleus. |
| Q9H2L4 | TMM60_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9H2L5 | RASF4_HUMAN | | cytoplasm. |
| Q9H2N8 | CI045_HUMAN | | membrane. |
| Q9H2P0 | ADNP_HUMAN | nucleus (potential). | nucleus. |
| Q9H2P9 | DPH5_HUMAN | | cytoplasm. |
| Q9H2R5 | KLK15_HUMAN | secreted protein (probable). | secreted protein. |
| Q9H2S5 | RNF39_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9H2S6 | TNMD_HUMAN | membrane; single-pass type ii membrane protein (potential). | endosome; membrane. |
| Q9H2T7 | RBP17_HUMAN | cytoplasm (by similarity). nucleus (by similarity). nucleus; nuclear envelope; nuclear pore complex (by similarity). | cytoplasm. |
| Q9H2U2 | IPYR2_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q9H2U6 | CO005_HUMAN | | nucleus. |
| Q9H2X0 | CHRD_HUMAN | secreted protein (by similarity). | cytoplasm. |
| Q9H2X3 | CLC4M_HUMAN | isoform 1, isoform 2, isoform 3: cell membrane; single-pass type ii membrane protein (potential). isoform 5, isoform 6, isoform 7, isoform 10: secreted protein (potential). | membrane. |
| Q9H2X8 | FM14A_HUMAN | | endosome; membrane. |
| Q9H2Y7 | ZF106_HUMAN | | nucleus. |
| Q9H2Z4 | NKX24_HUMAN | nucleus (by similarity). | nucleus. |
| Q9H320 | VCX1_HUMAN | | cytoplasm. nucleus. |
| Q9H321 | VCXC_HUMAN | | cytoplasm. nucleus. |
| Q9H322 | VCX2_HUMAN | | centriole. cytoplasm. |
| Q9H329 | E41LB_HUMAN | | cytoplasm. |
| Q9H330 | CI005_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9H334 | FOXP1_HUMAN | nucleus (probable). | nucleus. |
| Q9H336 | CRLD1_HUMAN | secreted protein (potential). | secreted protein. |
| Q9H347 | UBQL3_HUMAN | | cytoplasm. |
| Q9H361 | PABP3_HUMAN | cytoplasm (by similarity). | cytoplasm. nucleus. |
| Q9H367 | CN057_HUMAN | | cytoplasm. |
| Q9H386 | CU116_HUMAN | | endoplasmic reticulum. |
| Q9H3C7 | ZN403_HUMAN | cytoplasmic vesicle (by similarity). associated with vesicular structures (by similarity). | cytoplasm. cytoskeleton. |
| Q9H3E2 | SNX25_HUMAN | | cytoplasm. |
| Q9H3F6 | KCD10_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9H3G5 | CPVL_HUMAN | | secreted protein. |
| Q9H3H1 | MOD5_HUMAN | isoform 1, isoform 4: mitochondrion (potential). isoform 2, isoform 3, isoform 5: cytoplasm (potential). | mitochondrion. |
| Q9H3H3 | CK068_HUMAN | | mitochondrion. |
| Q9H3H9 | TCAL2_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9H3K2 | GHITM_HUMAN | membrane; multi-pass membrane protein (potential). | mitochondrion; membrane. |
| Q9H3K6 | BOLA2_HUMAN | | nucleus. |

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| Q9H3L0 | CB025_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q9H3N1 | TXND1_HUMAN | membrane; single-pass type i membrane protein (potential). predominantly found in the endoplasmic reticulum. | endoplasmic reticulum. |
| Q9H3P7 | GCP60_HUMAN | Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). mitochondrion (by similarity). also mitochondrial (via its interaction with pbr) (by similarity). | mitochondrion. |
| Q9H3Q3 | G3ST2_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9H3R0 | JHD3C_HUMAN | nucleus (by similarity). | nucleus. |
| Q9H3R1 | NDST4_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endosome. |
| Q9H3R2 | MUC13_HUMAN | cell membrane; single-pass type i membrane protein (potential). secreted protein. also exists as a soluble form. | endosome; membrane. |
| Q9H3S1 | SEM4A_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9H3S3 | TMPS5_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9H3S4 | TPK1_HUMAN | | cytoplasm. |
| Q9H3U1 | UN45A_HUMAN | | cytoplasm. secreted protein. |
| Q9H3U7 | SMOC2_HUMAN | secreted protein (probable). | secreted protein. |
| Q9H3W5 | LRRN3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9H3Y6 | SRMS_HUMAN | | cytoplasm. |
| Q9H3Y8 | CT149_HUMAN | | nucleus. |
| Q9H3Z4 | DNJC5_HUMAN | membrane; lipid-anchor (by similarity). | endoplasmic reticulum. |
| Q9H3Z7 | CT135_HUMAN | | nucleus. |
| Q9H426 | RIMS4_HUMAN | | cytoplasm. |
| Q9H446 | RWDD1_HUMAN | | cytoplasm. |
| Q9H469 | FBX37_HUMAN | | cytoplasm. |
| Q9H477 | RBSK_HUMAN | | centriole. cytoplasm. |
| Q9H478 | KCQ1D_HUMAN | | centriole. cytoplasm. |
| Q9H479 | FN3K_HUMAN | | cytoplasm. |
| Q9H488 | OFUT1_HUMAN | endoplasmic reticulum (by similarity). | endoplasmic reticulum. Golgi. |
| Q9H489 | TSYL3_HUMAN | | centriole. cytoplasm. |
| Q9H490 | PIGU_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9H493 | CT051_HUMAN | | cytoplasm. |
| Q9H4A4 | AMPB_HUMAN | secreted protein (by similarity). | cytoplasm; membrane. |
| Q9H4A9 | DPEP2_HUMAN | membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| Q9H4B6 | SAV1_HUMAN | | cytoplasm. |
| Q9H4B7 | TBB1_HUMAN | | centriole. |
| Q9H4B8 | DPEP3_HUMAN | membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| Q9H4D0 | CSTN2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9H4E5 | RHOJ_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |

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| Q9H4F1 | SIA7D_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. |
| Q9H4G0 | E41L1_HUMAN | | cytoplasm. |
| Q9H4G1 | CST9_HUMAN | secreted protein (potential). | secreted protein. |
| Q9H4G8 | ZCSL1_HUMAN | | cytoplasm. nucleus. |
| Q9H4H8 | CT129_HUMAN | | cytoplasm. |
| Q9H4I3 | TRABD_HUMAN | | cytoplasm. |
| Q9H4I8 | SERHL_HUMAN | cytoplasm; perinuclear region (by similarity). concentrated in perinuclear vesicles. peroxisome (by similarity). may be located in peroxisomes (by similarity). | peroxisome. |
| Q9H4K7 | GTPB5_HUMAN | | cytoplasm. |
| Q9H4L4 | SENP3_HUMAN | nucleus; nucleolus (by similarity). | nucleus. |
| Q9H4L5 | OSBL3_HUMAN | | cytoplasm. secreted protein. |
| Q9H4L7 | SMRCD_HUMAN | nucleus (by similarity). | nucleus. |
| Q9H4M3 | FBX44_HUMAN | | cytoplasm. |
| Q9H4M7 | PKHA4_HUMAN | cytoplasm (probable). membrane; peripheral membrane protein (probable). | cytoplasm. |
| Q9H4P4 | RNF41_HUMAN | | cytoplasm. |
| Q9H4Q3 | PRD13_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9H4Q4 | PRD12_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9H4S2 | GSHI_HUMAN | nucleus (by similarity). | nucleus. |
| Q9H4T2 | ZN435_HUMAN | nucleus (probable). | nucleus. |
| Q9H4W6 | COE3_HUMAN | nucleus (potential). | nucleus. |
| Q9H4Y5 | GSTO2_HUMAN | | cytoplasm. |
| Q9H4Z2 | ZN335_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9H501 | ABTAP_HUMAN | nucleus; nucleolus (by similarity). nucleus; nucleoplasm (by similarity). | cytoplasm. nucleus. |
| Q9H553 | ALG2_HUMAN | | endoplasmic reticulum. |
| Q9H560 | ANR19_HUMAN | | cytoplasm. |
| Q9H579 | CT132_HUMAN | | nucleus. |
| Q9H598 | VIAAT_HUMAN | cytoplasmic vesicle; cytoplasmic vesicle membrane; multi-pass membrane protein (by similarity). | Golgi. |
| Q9H5I1 | SUV92_HUMAN | nucleus (by similarity). associates with constitutive heterochromatin (by similarity). | nucleus. |
| Q9H5I5 | FA38B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9H5J0 | ZBTB3_HUMAN | nucleus (potential). | nucleus. |
| Q9H5P4 | PDZD7_HUMAN | | cytoplasm. |
| Q9H5U6 | ZCHC4_HUMAN | | cytoplasm. |
| Q9H5U8 | CX045_HUMAN | | nucleus. |
| Q9H5V8 | CDCP1_HUMAN | cell membrane; single-pass membrane protein (potential). its shedding may lead to a soluble peptide. | endosome; membrane. |
| Q9H5X1 | FA96A_HUMAN | | nucleus. |
| Q9H5Y7 | SLIK6_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q9H5Z1 | DHX35_HUMAN | | nucleus. |
| Q9H609 | ZN576_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9H611 | CO020_HUMAN | | cytoplasm. |
| Q9H668 | OBFC1_HUMAN | | centriole. cytoplasm. |
| Q9H672 | ASB7_HUMAN | | cytoplasm. |
| Q9H6A0 | DEN2D_HUMAN | | cytoplasm. |
| Q9H6B1 | ZN659_HUMAN | nucleus (potential). | nucleus. |

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| Q9H6D3 | XKR8_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9H6D5 | WBS23_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q9H6D7 | CN094_HUMAN | | cytoplasm. |
| Q9H6H4 | REEP4_HUMAN | membrane; multi-pass membrane protein (by similarity). | endosome; membrane. |
| Q9H6I2 | SOX17_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9H6K1 | CF106_HUMAN | | membrane. |
| Q9H6K4 | OPA3_HUMAN | | centriole. |
| Q9H6L4 | ARMC7_HUMAN | | cytoplasm. |
| Q9H6P5 | TASP1_HUMAN | | cytoplasm. |
| Q9H6R4 | NOL6_HUMAN | nucleus; nucleolus. localizes to condensed chromosomes in mitosis (by similarity). | nucleus. |
| Q9H6R6 | ZDHC6_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q9H6S0 | YTDC2_HUMAN | | nucleus. |
| Q9H6U6 | BCAS3_HUMAN | nucleus (potential). | centriole. cytoplasm. |
| Q9H6U8 | ALG9_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). | endoplasmic reticulum. |
| Q9H6X2 | ANTR1_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q9H6Y2 | WDR55_HUMAN | | nucleus. |
| Q9H6Y7 | RN167_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q9H714 | CM018_HUMAN | | cytoplasm. |
| Q9H756 | LRC19_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9H761 | CN139_HUMAN | | cytoplasm. |
| Q9H772 | GREM2_HUMAN | secreted protein (probable). | secreted protein. |
| Q9H777 | RNZ1_HUMAN | nucleus (probable). | nucleus. |
| Q9H788 | SH24A_HUMAN | | cytoplasm. |
| Q9H7C9 | CK067_HUMAN | | cytoplasm. |
| Q9H7D7 | WDR26_HUMAN | | cytoplasm. |
| Q9H7E2 | TDRD3_HUMAN | | nucleus. |
| Q9H7F0 | AT133_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9H7F4 | FA11B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9H7H1 | CU096_HUMAN | | membrane. |
| Q9H7L9 | SDS3_HUMAN | | nucleus. |
| Q9H7M6 | ZSWM4_HUMAN | | cytoplasm. nucleus. |
| Q9H7M9 | GI24_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9H7P6 | F125B_HUMAN | | membrane. |
| Q9H7R0 | ZN442_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9H7S9 | ZN703_HUMAN | | nucleus. |
| Q9H7T0 | CN161_HUMAN | | endosome; membrane. |
| Q9H7T3 | CJ095_HUMAN | | nucleus. |
| Q9H7V2 | CT039_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9H7X3 | ZN696_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9H7Y0 | CX036_HUMAN | | nucleus. |
| Q9H7Z3 | CN102_HUMAN | | cytoplasm. |
| Q9H808 | TLE6_HUMAN | nucleus (by similarity). | nucleus. |

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| Q9H819 | DJC18_HUMAN | membrane; single-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9H841 | NPAL2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9H845 | ACAD9_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q9H857 | NT5D2_HUMAN | | Golgi. |
| Q9H867 | CN138_HUMAN | | mitochondrion. |
| Q9H892 | TTC12_HUMAN | | cytoplasm. |
| Q9H8G1 | ZN430_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9H8G2 | CI082_HUMAN | | nucleus. |
| Q9H8J5 | MANS1_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9H8M1 | CQ10B_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q9H8M2 | BRD9_HUMAN | | cytoplasm. |
| Q9H8S9 | MOL1B_HUMAN | | cytoplasm. nucleus. |
| Q9H8U3 | ZFAN3_HUMAN | | nucleus. |
| Q9H8V3 | ECT2_HUMAN | | cytoplasm. |
| Q9H8W2 | CF155_HUMAN | | nucleus. |
| Q9H8W3 | CJ084_HUMAN | | nucleus. |
| Q9H8X2 | IPPK_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. nucleus. |
| Q9H8X3 | CF208_HUMAN | | cytoplasm. |
| Q9H8X9 | ZDH11_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q9H8Y1 | CN115_HUMAN | | nucleus. |
| Q9H8Y5 | ANKZ1_HUMAN | | nucleus. |
| Q9H936 | GHC1_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (potential). | mitochondrion. |
| Q9H939 | PIIP2_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | cytoplasm. |
| Q9H943 | CJ068_HUMAN | | cytoplasm. |
| Q9H944 | TRFP_HUMAN | nucleus (probable). | nucleus. |
| Q9H963 | ZN702_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9H967 | WDR76_HUMAN | | membrane. |
| Q9H972 | CN093_HUMAN | | nucleus. |
| Q9H977 | WDR54_HUMAN | | cytoplasm. |
| Q9H981 | ARP8_HUMAN | | cytoplasm. |
| Q9H987 | SYP2L_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9H993 | CF211_HUMAN | | cytoplasm. |
| Q9H999 | PANK3_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9H9A6 | LRC40_HUMAN | | cytoplasm. |
| Q9H9A7 | RMI1_HUMAN | | cytoplasm. |
| Q9H9B4 | SFXN1_HUMAN | mitochondrion; mitochondrial membrane; multi-pass membrane protein (by | mitochondrion; membrane. |
| Q9H9C1 | CN133_HUMAN | | nucleus. |
| Q9H9D4 | ZN408_HUMAN | nucleus (potential). | nucleus. |
| Q9H9E1 | ANRA2_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by | cytoplasm. |
| Q9H9F9 | ARP5_HUMAN | | nucleus. |
| Q9H9G7 | I2C3_HUMAN | | cytoplasm. |
| Q9H9J4 | UBP42_HUMAN | | nucleus. |
| Q9H9P2 | CHODL_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |

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| Q9H9S3 | S61A2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9H9S4 | CB39L_HUMAN | | cytoplasm. |
| Q9H9S5 | FKRP_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9H9V4 | RN122_HUMAN | Golgi apparatus. endoplasmic reticulum. membrane; single-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9H9Y3 | T142B_HUMAN | | secreted protein. |
| Q9H9Y4 | ATPB_B_HUMAN | | cytoplasm. |
| Q9HA47 | UCK1_HUMAN | | cytoplasm. |
| Q9HA64 | KT3K_HUMAN | | cytoplasm. |
| Q9HA65 | TBC17_HUMAN | | cytoplasm. |
| Q9HA72 | FA26B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9HA82 | LASS4_HUMAN | nucleus; nuclear membrane; multi-pass membrane protein (potential). endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | nucleus. |
| Q9HA90 | CCD48_HUMAN | | cytoplasm. |
| Q9HAB8 | PPCS_HUMAN | | cytoplasm. mitochondrion. |
| Q9HAC7 | CG010_HUMAN | | cytoplasm. |
| Q9HAC8 | UBTD1_HUMAN | | cytoplasm. |
| Q9HAD4 | WDR41_HUMAN | | cytoplasm. |
| Q9HAF5 | CO028_HUMAN | | secreted protein. |
| Q9HAH7 | FBSH_HUMAN | | nucleus. |
| Q9HAP6 | LIN7B_HUMAN | cell membrane; peripheral membrane protein. cell membrane; cell-cell junction; peripheral membrane protein (by similarity). localized to postsynaptic densities of neurons and epithelial cell-cell junctions (by similarity). mainly basolateral in renal epithelial cells. | Golgi. |
| Q9HAQ2 | KIF9_HUMAN | | cytoplasm. nucleus. |
| Q9HAS0 | NJMU_HUMAN | | cytoplasm. secreted protein. |
| Q9HAT1 | LMA1L_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9HAT2 | SIAE_HUMAN | lysosome (probable). | lysosome. |
| Q9HAT8 | PELI2_HUMAN | | cytoplasm. |
| Q9HAU0 | PKHA5_HUMAN | | cytoplasm. |
| Q9HAU6 | FKSG2_HUMAN | | cytoplasm. |
| Q9HAU8 | RNPL1_HUMAN | | cytoplasm. |
| Q9HAV0 | GBB4_HUMAN | | nucleus. |
| Q9HAY2 | MAGF1_HUMAN | | cytoplasm. |
| Q9HAY6 | BCDO1_HUMAN | | cytoplasm. endoplasmic reticulum. |
| Q9HAZ1 | CLK4_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9HAZ2 | PRD16_HUMAN | nucleus (potential). | nucleus. |
| Q9HB03 | ELOV3_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9HB07 | MYG1_HUMAN | | cytoplasm. |
| Q9HB09 | B2L12_HUMAN | | cytoplasm. |

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| Q9HB14 | KCNKD_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9HB15 | KCNKC_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9HB19 | PKHA2_HUMAN | cytoplasm (by similarity). cell membrane; peripheral membrane protein (by similarity). nucleus (by similarity). locates to the plasma membrane after treatments that stimulate the production of ptdins3,4p2 (by similarity). | nucleus. |
| Q9HB20 | PKHA3_HUMAN | cytoplasm (probable). membrane; peripheral membrane protein (probable). | cytoplasm. |
| Q9HB40 | RISC_HUMAN | secreted protein (potential). | secreted protein. |
| Q9HBA0 | TRPV4_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9HBB8 | MUCDL_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9HBH0 | RHOF_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| Q9HBH1 | DEFM_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q9HBH5 | RDH14_HUMAN | | endoplasmic reticulum. |
| Q9HBH7 | BEX1_HUMAN | nucleus. cytoplasm. shuttles between the cytoplasm and the nucleus (by similarity). | nucleus. |
| Q9HBH9 | MKNK2_HUMAN | | cytoplasm. |
| Q9HBI0 | PARVG_HUMAN | cell membrane; cell-matrix junction; focal adhesion; cytoplasmic side (by similarity). constituent of focal adhesions (by | cytoplasm. |
| Q9HBI5 | CC014_HUMAN | | cytoplasm. |
| Q9HBI6 | CP4FB_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9HBJ7 | UBP29_HUMAN | | nucleus. |
| Q9HBJ8 | TMM27_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9HBK9 | AS3MT_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9HBL7 | CI046_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9HBM0 | VEZA_HUMAN | membrane; single-pass type iii membrane protein (probable). | endosome; membrane. |
| Q9HBM6 | TAF9B_HUMAN | nucleus (by similarity). | nucleus. |
| Q9HBT7 | ZN287_HUMAN | nucleus (probable). | nucleus. |
| Q9HBT8 | ZN286_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9HBU1 | BARX1_HUMAN | nucleus (probable). | nucleus. |
| Q9HBU9 | POPD2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9HBV1 | POPD3_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9HBV2 | SACA1_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9HBW1 | LRRC4_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9HBY0 | NOX3_HUMAN | membrane; multi-pass membrane protein (potential). | mitochondrion. |
| Q9HBY8 | SGK2_HUMAN | | cytoplasm. |
| Q9HBZ2 | ARNT2_HUMAN | nucleus (potential). | nucleus. |

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| Q9HC07 | TM165_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9HC10 | OTOF_HUMAN | membrane; single-pass type ii membrane protein (by similarity). | endosome; membrane. |
| Q9HC24 | TMBI4_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9HC47 | CTGE1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9HC56 | PCDH9_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9HC57 | WFDC1_HUMAN | secreted protein (potential). | secreted protein. |
| Q9HC73 | CRLF2_HUMAN | isoform 1: cell membrane; single-pass type i membrane protein (probable). isoform 2, isoform 3: secreted protein (probable). | membrane. |
| Q9HC78 | ZBT20_HUMAN | nucleus (probable). | nucleus. |
| Q9HC96 | CAN10_HUMAN | | cytoplasm. |
| Q9HC97 | GPR35_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9HCB6 | SPON1_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9HCC6 | HES4_HUMAN | | nucleus. |
| Q9HCC9 | ZFY28_HUMAN | | cytoplasm. nucleus. |
| Q9HCE1 | MOV10_HUMAN | | cytoplasm. nucleus. |
| Q9HCE7 | SMUF1_HUMAN | | cytoplasm. |
| Q9HCE9 | TM16H_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| Q9HCG1 | KR18_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9HCH3 | CPNE5_HUMAN | | membrane. |
| Q9HCH5 | SYTL2_HUMAN | intracytoplasmic membrane; peripheral membrane protein (by similarity). bound to melanosomes (by similarity). | cytoplasm. |
| Q9HCI5 | MAGE1_HUMAN | | cytoplasm. nucleus. |
| Q9HCJ1 | ANKH_HUMAN | membrane; multi-pass membrane protein (probable). | Golgi. |
| Q9HCJ3 | RAVR2_HUMAN | nucleus (by similarity). cytoplasm (by similarity). may shuttle between the nucleus and the cytoplasm (by similarity). | nucleus. |
| Q9HCJ6 | K1576_HUMAN | | mitochondrion. |
| Q9HCK0 | ZBT26_HUMAN | nucleus (probable). | nucleus. |
| Q9HCK4 | ROBO2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9HCK5 | I2C4_HUMAN | | cytoplasm. |
| Q9HCK8 | CHD8_HUMAN | nucleus (probable). | nucleus. |
| Q9HCL0 | PCD18_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9HCL2 | PLSB_HUMAN | mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9HCL3 | ZFP14_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9HCM2 | PLXA4_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9HCM4 | E41L5_HUMAN | | cytoplasm. |
| Q9HCM9 | TRI39_HUMAN | | cytoplasm. |
| Q9HCN3 | TMEM8_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |

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| Q9HCN8 | SDF2L_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (potential). | endoplasmic reticulum. |
| Q9HCP6 | GUP1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9HCQ5 | GALT9_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | secreted protein. Golgi. |
| Q9HCS2 | CP4FC_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9HCS5 | E41LA_HUMAN | | cytoplasm. |
| Q9HCS7 | XAB2_HUMAN | nucleus (by similarity). detected in the splicing complex carrying pre-mrna (by similarity). | nucleus. |
| Q9HCT0 | FGF22_HUMAN | secreted protein (potential). | secreted protein. |
| Q9HCU0 | CD248_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q9HCU5 | PREB_HUMAN | nucleus (by similarity). endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9HCU9 | BRMS1_HUMAN | | cytoplasm. |
| Q9HCX3 | ZN304_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9HCX4 | TRPC7_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9HCZ1 | ZN334_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9HD20 | AT131_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9HD23 | MRS2L_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9HD26 | GOPC_HUMAN | cytoplasm. Golgi apparatus; Golgi membrane; peripheral membrane protein. Golgi apparatus; trans-Golgi network; trans-Golgi network membrane; peripheral membrane protein. enriched in synaptosomal and postsynaptic densities (psd) fractions (by similarity). expressed in cell bodies and dendrites of purkinje cells (by similarity). localized at the trans-Golgi network (tgn) of spermatids and the medulla of round spermatides (by | Golgi. |
| Q9HD34 | CF149_HUMAN | | mitochondrion. |
| Q9HD45 | TM9S3_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9HD64 | GAGD2_HUMAN | | secreted protein. |
| Q9HD67 | MYO10_HUMAN | | cytoplasm. |
| Q9HD90 | NDF4_HUMAN | nucleus (potential). | nucleus. |
| Q9HDB5 | NRX3B_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9HDB8 | ENK8_HUMAN | | centriole. |
| Q9HDB9 | GAK8_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |
| Q9HDC0 | ENK15_HUMAN | | centriole. |

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|--------|-------------|--|---|
| Q9HDC5 | JPH1_HUMAN | cell membrane; peripheral membrane protein (by similarity). endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type iv membrane protein (by similarity). sarcoplasmic reticulum; sarcoplasmic reticulum membrane; single- pass type iv membrane protein (by similarity). localized predominantly on the plasma membrane. the transmembrane domain is anchored in endoplasmic/sarcoplasmic reticulum membrane, while the n-terminal part associates with the plasma membrane. in skeletal muscle cells, it is predominantly localized at the junction of the a and i bands (by similarity). | endoplasmic reticulum. |
| Q9HDC9 | APMAP_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9HDD0 | HRSL1_HUMAN | | membrane. |
| Q9N2J8 | ENH3_HUMAN | | endosome; membrane. |
| Q9N2J9 | ENH2_HUMAN | | endosome; membrane. |
| Q9N2K0 | ENH1_HUMAN | | membrane. |
| Q9NNW5 | WDR6_HUMAN | | centriole. cytoplasm. |
| Q9NNX1 | TUFT1_HUMAN | secreted protein (by similarity). secreted at a very early stage of enamel formation, concentrated at the dentin-enamel junction and tightly bound to the surface of the growing crystallites (by similarity). | secreted protein. |
| Q9NNX6 | CD209_HUMAN | isoform 1, isoform 2, isoform 3, isoform 4, isoform 5: cell membrane; single-pass type ii membrane protein (probable). isoform 6, isoform 7, isoform 8, isoform 9, isoform 10, isoform 11, isoform 12, isoform 13: secreted protein (probable). | membrane. |
| Q9NNX9 | VCX3_HUMAN | | cytoplasm. |
| Q9NNZ3 | DNJC4_HUMAN | membrane; single-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9NP50 | FA60A_HUMAN | | nucleus. |
| Q9NP55 | PLUNC_HUMAN | secreted protein (by similarity). found in the nasal mucus (by similarity). apical side of airway epithelial cells. detected in nasal mucus (by similarity). | secreted protein. |
| Q9NP56 | PDE7B_HUMAN | | cytoplasm. |
| Q9NP58 | ABCB6_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (potential). | mitochondrion; membrane. |
| Q9NP62 | GCM1_HUMAN | nucleus (potential). | nucleus. |
| Q9NP66 | HM20A_HUMAN | nucleus (by similarity). | nucleus. |
| Q9NP72 | RAB18_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. endosome. Golgi. |
| Q9NP78 | ABCB9_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (probable). lysosome; lysosomal membrane; multi-pass | endoplasmic reticulum. mitochondrion; membrane. |
| Q9NP79 | CF055_HUMAN | | cytoplasm. |
| Q9NP81 | SYSM_HUMAN | mitochondrion (by similarity). | cytoplasm. |

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| Q9NP86 | CABP5_HUMAN | | cytoplasm. |
| Q9NP87 | DPOLM_HUMAN | | nucleus. |
| Q9NP90 | RAB9B_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q9NP91 | S6A20_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9NP99 | TREM1_HUMAN | isoform 1: cell membrane; single-pass type i membrane protein (potential). isoform 2: secreted protein (potential). | membrane. |
| Q9NPA0 | CO024_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome. |
| Q9NPA3 | M1IP1_HUMAN | nucleus (by similarity). cytoplasm (by similarity). associated with microtubules (by similarity). | cytoplasm. nucleus. |
| Q9NPA5 | ZF64A_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9NPB0 | CF064_HUMAN | | endosome; membrane. |
| Q9NPB3 | CABP2_HUMAN | | cytoplasm. |
| Q9NPB8 | K1434_HUMAN | | lysosome. |
| Q9NPC2 | KCNK9_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9NPC4 | A4GAT_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | Golgi. |
| Q9NPC6 | MYOZ2_HUMAN | colocalizes with actn1 and ppp3ca at the z-line of heart and skeletal muscle (by similarity). | cytoplasm. |
| Q9NPD0 | PART1_HUMAN | | centriole. cytoplasm. |
| Q9NPD8 | UBE2T_HUMAN | | cytoplasm. |
| Q9NPE6 | SPAG4_HUMAN | membrane; multi-pass membrane protein (potential). axoneme (by similarity). in spermatids, it is localized in the transient manchette and in the axoneme of elongating spermatids and epididymal sperm (by similarity). | membrane. |
| Q9NPF0 | CD320_HUMAN | membrane; single-pass type i membrane protein (potential). | endoplasmic reticulum. |
| Q9NPF2 | CHSTB_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9NPF4 | GCP_HUMAN | | secreted protein. |
| Q9NPG2 | NGB_HUMAN | | cytoplasm. mitochondrion. |
| Q9NPG8 | ZDHC4_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. mitochondrion; membrane. |
| Q9NPH0 | PPA6_HUMAN | secreted protein (probable). | cytoplasm. |
| Q9NPH5 | NOX4_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein. cell membrane; multi-pass membrane protein (probable). cell membrane; cell-matrix junction; focal adhesion (probable). may localize to plasma membrane and focal adhesions. nucleus (probable). may also localize to the nucleus (ref.14). | endoplasmic reticulum. |
| Q9NPH6 | OBP2B_HUMAN | secreted protein (probable). | secreted protein. |
| Q9NPI5 | NRK2_HUMAN | | cytoplasm. |
| Q9NPJ1 | MKKS_HUMAN | | cytoplasm. |
| Q9NPJ3 | THEM2_HUMAN | | cytoplasm. |

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| Q9NPJ4 | PNRC2_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9NPQ8 | RIC8A_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NPU4 | CN132_HUMAN | | endosome; membrane. |
| Q9NQ03 | SCRT2_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9NQ11 | AT132_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9NQ29 | LUC7L_HUMAN | | cytoplasm. nucleus. |
| Q9NQ32 | CK016_HUMAN | | nucleus. |
| Q9NQ34 | TMM9B_HUMAN | membrane; single-pass membrane protein (potential). | endosome. Golgi. |
| Q9NQ35 | NRIP3_HUMAN | | cytoplasm. |
| Q9NQ40 | CT054_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NQ50 | RM40_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| Q9NQ66 | PLCB1_HUMAN | | cytoplasm. nucleus. |
| Q9NQ69 | LHX9_HUMAN | nucleus (probable). | nucleus. |
| Q9NQ75 | CT032_HUMAN | | Golgi. |
| Q9NQ79 | CRAC1_HUMAN | secreted protein (probable). | cytoplasm. |
| Q9NQ86 | TRI36_HUMAN | cytoplasm (by similarity). found in the acrosomal region of elongated spermatids and mature sperm (by similarity). | cytoplasm. |
| Q9NQ88 | CL005_HUMAN | | centriole. cytoplasm. mitochondrion. |
| Q9NQ89 | CL004_HUMAN | | nucleus. |
| Q9NQ90 | TM16B_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| Q9NQ94 | ACF_HUMAN | nucleus. predominantly nuclear where it localizes to heterochromatin. endoplasmic reticulum (by similarity). also cytoplasmic where it is found at the outer surface of the endoplasmic reticulum (by similarity). cytoplasm. shuttles between the nucleus and cytoplasm. may be transported into the nucleus by the nuclear import protein tnp2/trn2 or by apobec1. | nucleus. |
| Q9NQA5 | TRPV5_HUMAN | cell membrane; apical cell membrane; multi-pass membrane protein. colocalized with s100a10 and anax2 along the apical domain of kidney distal tubular cells (by | membrane. |
| Q9NQC1 | PHF15_HUMAN | | nucleus. |
| Q9NQC8 | CK060_HUMAN | | cytoplasm. nucleus. |
| Q9NQF3 | SEHL2_HUMAN | | cytoplasm. |
| Q9NQG1 | MANBL_HUMAN | | endosome; membrane. |
| Q9NQG5 | CT077_HUMAN | | nucleus. |
| Q9NQG7 | HPS4_HUMAN | | cytoplasm. |
| Q9NQM4 | CX041_HUMAN | | nucleus. |
| Q9NQP4 | PFD4_HUMAN | | cytoplasm. |
| Q9NQQ0 | GUSL1_HUMAN | | lysosome. |
| Q9NQQ7 | S35C2_HUMAN | membrane; multi-pass membrane protein (potential). | Golgi. |
| Q9NQR1 | SETD8_HUMAN | nucleus (by similarity). associates with silent chromatin on euchromatic arms. no association with constitutive heterochromatin (by similarity). | nucleus. |
| Q9NQR7 | CN162_HUMAN | | membrane. |

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| Q9NQS3 | PVRL3_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q9NQT6 | FSCN3_HUMAN | | cytoplasm. |
| Q9NQT8 | KI13B_HUMAN | | cytoplasm. nucleus. |
| Q9NQU5 | PAK6_HUMAN | | cytoplasm. |
| Q9NQV5 | PRD11_HUMAN | | centriole. cytoplasm. |
| Q9NQV6 | PRD10_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9NQV7 | PRDM9_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9NQV8 | PRDM8_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9NQW5 | PRDM7_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9NQW7 | XPP1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NQX1 | PRDM5_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9NQX3 | GEPH_HUMAN | | cytoplasm. |
| Q9NQX4 | MYO5C_HUMAN | | cytoskeleton. |
| Q9NQX5 | NPDC1_HUMAN | | endosome; membrane. |
| Q9NQX6 | ZN331_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9NQX7 | ITM2C_HUMAN | membrane; single-pass type ii membrane protein (potential). | endosome. |
| Q9NQY0 | BIN3_HUMAN | | centriole. cytoplasm. |
| Q9NQZ2 | SAS10_HUMAN | nucleus (by similarity). | nucleus. |
| Q9NQZ5 | STAR7_HUMAN | | nucleus. |
| Q9NQZ6 | HC127_HUMAN | | cytoplasm. |
| Q9NQZ8 | ZNF71_HUMAN | nucleus (potential). | nucleus. |
| Q9NR00 | CH004_HUMAN | | nucleus. |
| Q9NR09 | BIRC6_HUMAN | | cytoplasm. |
| Q9NR11 | ZN302_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9NR12 | PDLI7_HUMAN | cytoplasm (by similarity). cytoskeleton (by similarity). colocalizes with ret to the cell periphery and in some cytoskeletal components. colocalizes with tpm2 near the z line in muscle. co-localizes with tbx4 and tbx5 to actin filaments (by similarity). | nucleus. |
| Q9NR16 | C163B_HUMAN | isoform 1, isoform 2: cell membrane; single- pass type i membrane protein (probable). isoform 3: secreted protein | endosome; membrane. |
| Q9NR20 | DYRK4_HUMAN | | cytoplasm. |
| Q9NR22 | ANM8_HUMAN | | nucleus. |
| Q9NR23 | GDF3_HUMAN | secreted protein (probable). | secreted protein. |
| Q9NR31 | SAR1A_HUMAN | | Golgi. |
| Q9NR33 | DPOE4_HUMAN | nucleus (potential). | nucleus. |
| Q9NR45 | SIAS_HUMAN | | cytoplasm. nucleus. |
| Q9NR50 | EI2BG_HUMAN | | cytoplasm. |
| Q9NR61 | DLL4_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q9NR63 | CP26B_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; peripheral membrane protein (by similarity). | endoplasmic reticulum. |
| Q9NR80 | ARHG4_HUMAN | isoform 3: cytoplasm. associated with membrane ruffles (probable). | cytoplasm. |
| Q9NR81 | ARHG3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NR96 | TLR9_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9NR97 | TLR8_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9NRA0 | SPHK2_HUMAN | | cytoplasm. |

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| Q9NRA2 | S17A5_HUMAN | lysosome; lysosomal membrane; multi-pass membrane protein (potential). | membrane. |
| Q9NRB3 | CHSTC_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | Golgi. |
| Q9NRC9 | OTOR_HUMAN | secreted protein (potential). | centriole. cytoplasm. secreted protein. membrane. |
| Q9NRD0 | FBX8_HUMAN | | cytoplasm. nucleus. |
| Q9NRD1 | FBX6_HUMAN | | cytoplasm. |
| Q9NRE2 | TSH2_HUMAN | nucleus (probable). | nucleus. |
| Q9NRF8 | PYRG2_HUMAN | | cytoplasm. |
| Q9NRF9 | DPOE3_HUMAN | nucleus (potential). | nucleus. |
| Q9NRG0 | CHRC1_HUMAN | nucleus (potential). | nucleus. |
| Q9NRG4 | SMYD2_HUMAN | | nucleus. |
| Q9NRG9 | AAAS_HUMAN | | cytoplasm. |
| Q9NRH1 | CG036_HUMAN | | cytoplasm. |
| Q9NRH2 | SNRK_HUMAN | nucleus (by similarity). | cytoplasm. |
| Q9NRH3 | TBG2_HUMAN | | centriole. |
| Q9NRI5 | DISC1_HUMAN | localizes to neurites (by similarity). cytoplasm. localizes to punctate cytoplasmic foci which overlap in part with mitochondria. also localizes to the centrosome. | cytoplasm. |
| Q9NRJ5 | PAPOB_HUMAN | nucleus (probable). | nucleus. |
| Q9NRJ7 | PCDBG_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9NRL3 | STRN4_HUMAN | cytoplasm (by similarity). membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q9NRM2 | ZN277_HUMAN | nucleus (potential). | nucleus. |
| Q9NRP2 | CP061_HUMAN | | cytoplasm. |
| Q9NRP4 | ACN9_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q9NRQ2 | PLS4_HUMAN | membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9NRQ5 | CK075_HUMAN | | centriole. cytoplasm. |
| Q9NRR2 | TRYG1_HUMAN | membrane; single-pass membrane protein (potential). | secreted protein; membrane. |
| Q9NRR5 | UBQL4_HUMAN | nucleus. cytoplasm. endoplasmic reticulum (probable). may associate with the endoplasmic reticulum. | nucleus. |
| Q9NRR6 | INP5E_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; peripheral membrane protein; cytoplasmic side (by similarity). peripheral membrane protein associated with Golgi stacks (by similarity). | cytoplasm. |
| Q9NRS4 | TMPS4_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9NRS6 | SNX15_HUMAN | | cytoplasm. Golgi. |
| Q9NRT5 | SMG64_HUMAN | | cytoplasm. secreted protein. |
| Q9NRW3 | ABC3C_HUMAN | | cytoplasm. |
| Q9NRW4 | DUS22_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. nucleus. |

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| Q9NRW7 | VPS45_HUMAN | Golgi apparatus; Golgi membrane; peripheral membrane protein (by similarity). endosome; endosomal membrane; peripheral membrane protein (by similarity). associated with Golgi/endosomal vesicles and the trans- | endosome. |
| Q9NRX2 | RM17_HUMAN | mitochondrion (potential). | mitochondrion. |
| Q9NRX3 | NUA4L_HUMAN | | mitochondrion. |
| Q9NRX4 | PHP14_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NRX5 | SERC1_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9NRY4 | GRLF1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| Q9NRY6 | PLS3_HUMAN | membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9NRY7 | PLS2_HUMAN | membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9NRZ4 | GAW2_HUMAN | cell membrane (potential). | secreted protein. |
| Q9NRZ5 | PLCD_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q9NRZ7 | PLCC_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q9NS15 | LTBP3_HUMAN | secreted protein (by similarity). secretion occurs after coexpression with tgfb1 and requires complexing with cys-33 of the tgfb1 propeptide (by similarity). | cytoplasm. |
| Q9NS28 | RGS18_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NS61 | KCIP2_HUMAN | isoform 1, isoform 2, isoform 6: cell membrane; lipid-anchor (by similarity). | membrane. |
| Q9NS62 | THSD1_HUMAN | isoform 1 and isoform 2: membrane; single- pass type i membrane protein (potential). isoform 3: secreted protein. | endosome; membrane. |
| Q9NS66 | GP173_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9NS67 | GPR27_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9NS68 | TNR19_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q9NS69 | TOM22_HUMAN | mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9NS71 | GKN1_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q9NS80 | TRI49_HUMAN | | cytoplasm. |
| Q9NS82 | AAA1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9NS84 | CHST7_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9NS85 | CAH10_HUMAN | | cytoplasm. |
| Q9NS91 | RAD18_HUMAN | nucleus (by similarity). | nucleus. |
| Q9NSA1 | FGF21_HUMAN | secreted protein (potential). | secreted protein. |
| Q9NSA3 | CNBP1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| Q9NSB2 | KRHB4_HUMAN | | cytoplasm. |
| Q9NSB4 | KRHB2_HUMAN | | cytoplasm. |
| Q9NSC2 | SALL1_HUMAN | nucleus (by similarity). | nucleus. |

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| Q9NSC5 | HOME3_HUMAN | cytoplasm (by similarity). postsynaptic density of neuronal cells (by similarity). | cytoplasm. |
| Q9NSC7 | SIA7A_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |
| Q9NSD4 | ZN275_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9NSD9 | SYFB_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NSE2 | CISH_HUMAN | | cytoplasm. |
| Q9NSE4 | SYIM_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | cytoplasm. |
| Q9NSE7 | ABCCD_HUMAN | | Golgi; membrane. |
| Q9NSG0 | RHG08_HUMAN | | cytoplasm. |
| Q9NSI2 | CU070_HUMAN | | nucleus. |
| Q9NSI6 | BRWD1_HUMAN | cytoplasm. nucleus (by similarity). | nucleus. |
| Q9NSI8 | SAMN1_HUMAN | | centriole. |
| Q9NSK0 | KLC4_HUMAN | | cytoplasm. |
| Q9NSP4 | CENPM_HUMAN | nucleus. cytoplasm. nuclear in non-confluent cells and cytoplasmic in confluent or dividing cells (by similarity). localizes in the kinetochore domain of | nucleus. |
| Q9NSU2 | TREX1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9NSV0 | PRR8_HUMAN | | nucleus. |
| Q9NSV4 | DIAP3_HUMAN | | cytoplasm. |
| Q9NSY0 | NRBP2_HUMAN | | cytoplasm. |
| Q9NSY2 | STAR5_HUMAN | | secreted protein. |
| Q9NT22 | EMIL3_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9NT99 | LRC4B_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q9NTG1 | PKDRE_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9NTJ4 | MA2C1_HUMAN | | lysosome. |
| Q9NTK1 | DEPP_HUMAN | | mitochondrion. |
| Q9NTK5 | PTD4_HUMAN | | cytoplasm. |
| Q9NTM9 | CUTC_HUMAN | | secreted protein. |
| Q9NTU7 | CBLN4_HUMAN | secreted protein (potential). | membrane. |
| Q9NTW7 | ZF64B_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9NTX7 | RN146_HUMAN | | cytoplasm. |
| Q9NTX9 | CT177_HUMAN | | nucleus. |
| Q9NU02 | ANKR5_HUMAN | | cytoplasm. |
| Q9NU19 | TB22B_HUMAN | | cytoplasm. |
| Q9NU22 | MDN1_HUMAN | nucleus (by similarity). | cytoplasm. |
| Q9NU39 | FX4L1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9NU53 | CF072_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9NUA8 | ZBT40_HUMAN | nucleus (potential). | nucleus. |
| Q9NUB1 | ACS2L_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | mitochondrion. |
| Q9NUB4 | CT141_HUMAN | | endosome; membrane. |
| Q9NUD5 | ZCHC3_HUMAN | | cytoplasm. nucleus. |
| Q9NUD7 | CT096_HUMAN | | nucleus. |
| Q9NUE0 | ZDH18_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q9NUG4 | CT160_HUMAN | | membrane. |
| Q9NUG6 | PDRG1_HUMAN | | cytoplasm. |
| Q9NUH8 | TM14B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |

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| Q9NUI1 | DECR2_HUMAN | peroxisome (by similarity). | peroxisome. |
| Q9NUL3 | STAU2_HUMAN | cytoplasm. nucleus. nucleus; nucleolus. shuttles between the nucleolus, nucleus and the cytoplasm. nuclear export of isoform 1 is independent of xpo1/crm1 and requires the exportin xpo5. nuclear export of isoform 2 and isoform 3 can occur by both xpo1/crm1-dependent and xpo1/crm1-independent pathways. found in large cytoplasmic ribonucleoprotein (rnp) granules which are present in the actin rich regions of myelinating processes and associated with microtubules, polysomes and the endoplasmic reticulum. also recruited to stress granules (sgs) upon inhibition of translation or oxidative stress. these structures are thought to harbor housekeeping mrnas when translation is aborted (by similarity). | nucleus. |
| Q9NUM4 | T106B_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q9NUP1 | CNO_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NUP7 | CCD76_HUMAN | | cytoplasm. |
| Q9NUP9 | LIN7C_HUMAN | cell membrane; peripheral membrane protein. cell membrane; cell-cell junction; peripheral membrane protein (by similarity). localized to postsynaptic densities of neurons and epithelial cell-cell junctions (by similarity). mainly basolateral in renal epithelial cells. | Golgi. |
| Q9NUQ2 | PLCE_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q9NUQ3 | TXLNG_HUMAN | | cytoskeleton. |
| Q9NUQ8 | ABCF3_HUMAN | | cytoplasm. |
| Q9NUQ9 | FA49B_HUMAN | | endoplasmic reticulum. |
| Q9NUR3 | CT046_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NUS5 | CT029_HUMAN | | nucleus. |
| Q9NUU7 | DD19A_HUMAN | cytoplasm (by similarity). nucleus; nuclear envelope; nuclear pore complex; cytoplasmic side (by similarity). nuclear pore complex cytoplasmic fibrils. | nucleus. |
| Q9NUV7 | CT038_HUMAN | | endosome; membrane. |
| Q9NUV9 | GIMA4_HUMAN | | cytoplasm. |
| Q9NUW8 | TYDP1_HUMAN | nucleus (probable). | nucleus. |
| Q9NUY9 | TDRD4_HUMAN | | cytoplasm. |
| Q9NV23 | SAST_HUMAN | | cytoplasm. |
| Q9NV29 | TM100_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NV35 | NUD15_HUMAN | | cytoplasm. |
| Q9NV44 | CU077_HUMAN | | membrane. |
| Q9NV70 | EXOC1_HUMAN | | cytoplasm. |
| Q9NV72 | ZN701_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9NV79 | PCMD2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NV96 | CC50A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |

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| Q9NVA1 | BZFB_HUMAN | cytoplasmic vesicle (by similarity). cytoplasmic vesicular structures (by similarity). | nucleus. |
| Q9NVA2 | SEP11_HUMAN | | cytoplasm. |
| Q9NVD3 | SETD4_HUMAN | | centriole. cytoplasm. |
| Q9NVE5 | UBP40_HUMAN | | nucleus. |
| Q9NVE7 | PANK4_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9NVF7 | FBX28_HUMAN | | cytoplasm. |
| Q9NVF9 | EKI2_HUMAN | | cytoplasm. |
| Q9NVG8 | TBC13_HUMAN | | cytoplasm. |
| Q9NVH0 | EXDL2_HUMAN | | nucleus. |
| Q9NVH1 | DCJ11_HUMAN | | cytoplasm. |
| Q9NVH6 | TMLH_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | mitochondrion. |
| Q9NVI1 | K1794_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NVI7 | ATD3A_HUMAN | | cytoplasm. nucleus. |
| Q9NVL8 | CN105_HUMAN | | nucleus. |
| Q9NVM4 | ANM7_HUMAN | | nucleus. |
| Q9NVM6 | DJC17_HUMAN | | nucleus. |
| Q9NVM9 | CL011_HUMAN | | cytoplasm. |
| Q9NVP1 | DDX18_HUMAN | | nucleus. |
| Q9NVP4 | CT012_HUMAN | | cytoplasm. |
| Q9NVQ4 | FAIM1_HUMAN | | cytoplasm. |
| Q9NVR0 | KLH11_HUMAN | secreted protein (potential). | cytoplasm. |
| Q9NVR5 | CN104_HUMAN | | cytoplasm. |
| Q9NVS9 | PNPO_HUMAN | | endoplasmic reticulum. peroxisome. |
| Q9NVT9 | ARMC1_HUMAN | | cytoplasm. |
| Q9NVU1 | MC3AP_HUMAN | | nucleus. |
| Q9NVV5 | AIG1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NVW2 | RNF12_HUMAN | | nucleus. |
| Q9NVX7 | KBTB4_HUMAN | | cytoplasm. |
| Q9NVZ3 | NECP2_HUMAN | cytoplasmic vesicle; clathrin-coated vesicle; clathrin-coated vesicle membrane (by similarity). cell membrane (by similarity). colocalizes with ap-2 at the plasma membrane (by similarity). | Golgi. |
| Q9NW07 | ZN358_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9NW75 | GPTC2_HUMAN | | centriole. nucleus. |
| Q9NW97 | TMM51_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NWB1 | A2BP1_HUMAN | | nucleus. |
| Q9NWC5 | TM45A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NWD9 | BEX4_HUMAN | nucleus (by similarity). cytoplasm (by similarity). | centriole. cytoplasm. |
| Q9NWM0 | SMOX_HUMAN | cytoplasm (by similarity). | mitochondrion. |
| Q9NWM3 | CUED1_HUMAN | | nucleus. |
| Q9NWM8 | FKB14_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (potential). | endoplasmic reticulum. |
| Q9NWN3 | FBX34_HUMAN | | cytoplasm. |
| Q9NWQ4 | CN118_HUMAN | | endosome; membrane. |
| Q9NWQ9 | CN119_HUMAN | | nucleus. |
| Q9NWS6 | F118A_HUMAN | | endosome; membrane. |

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| Q9NWS8 | CF096_HUMAN | | nucleus. |
| Q9NWS9 | ZN446_HUMAN | nucleus (potential). | nucleus. |
| Q9NWT6 | HIF1N_HUMAN | nucleus (potential). | nucleus. |
| Q9NWW6 | NRK1_HUMAN | | cytoplasm. |
| Q9NWW9 | HRSL2_HUMAN | | membrane. |
| Q9NWX5 | ASB6_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NWX3 | IRAK4_HUMAN | | cytoplasm. |
| Q9NWX5 | UCKL1_HUMAN | | cytoplasm. |
| Q9NWX8 | F51A1_HUMAN | | nucleus. |
| Q9NX05 | F120C_HUMAN | | cytoplasm. |
| Q9NX08 | COMD8_HUMAN | | mitochondrion. |
| Q9NX14 | NDUBB_HUMAN | mitochondrion; mitochondrial inner membrane (potential). | mitochondrion. |
| Q9NX31 | CT111_HUMAN | | nucleus. |
| Q9NX36 | CU055_HUMAN | | cytoplasm. |
| Q9NX57 | RAB20_HUMAN | highly enriched on apical endocytic structures in polarized epithelial cells of kidney proximal tubules (by similarity). | Golgi. |
| Q9NX63 | CHCH3_HUMAN | | cytoplasm. |
| Q9NX65 | ZN434_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9NX77 | ENK14_HUMAN | | endosome; membrane. |
| Q9NX78 | CN101_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NX94 | CJ026_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q9NXA8 | SIRT5_HUMAN | | nucleus. |
| Q9NXB0 | MKS1_HUMAN | | cytoplasm. |
| Q9NXB9 | ELOV2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9NXF8 | ZDHC7_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. endoplasmic reticulum; membrane. |
| Q9NXG0 | CI039_HUMAN | | Golgi. |
| Q9NXG2 | THUM1_HUMAN | | nucleus. |
| Q9NXH3 | PP14D_HUMAN | cytoplasm (probable). | cytoplasm. nucleus. |
| Q9NXH9 | TRM1_HUMAN | | nucleus. |
| Q9NXJ5 | PGPI_HUMAN | cytoplasm (by similarity). | lysosome. |
| Q9NXK6 | MPRG_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9NXK8 | FXL12_HUMAN | | cytoplasm. |
| Q9NXL6 | SIDT1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NXL9 | CF061_HUMAN | | nucleus. |
| Q9NXR1 | NDE1_HUMAN | cytoplasm. centrosome. kinetochore. spindle. concentrates at the plus ends of microtubules coincident with kinetochores in metaphase and anaphase. also localizes to the cleavage furrow during cytokinesis (by similarity). localizes to the interphase centrosome and to the mitotic spindle. | cytoplasm. |
| Q9NXR5 | ANR10_HUMAN | | cytoplasm. |
| Q9NXS3 | BTBD5_HUMAN | | cytoplasm. |
| Q9NXT0 | ZN586_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9NXV2 | KCTD5_HUMAN | | nucleus; membrane. |
| Q9NXV8 | CN113_HUMAN | | secreted protein. |

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| Q9NXW2 | DNJBC_HUMAN | membrane; single-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9NXW9 | ALKB4_HUMAN | | nucleus. |
| Q9NXX6 | CJ086_HUMAN | | cytoplasm. |
| Q9NXZ2 | DDX43_HUMAN | | nucleus. |
| Q9NY15 | STAB1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9NY25 | CLC5A_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9NY28 | GALT8_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | Golgi. |
| Q9NY30 | BTG4_HUMAN | | nucleus. |
| Q9NY35 | CLDND_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NY43 | BARH2_HUMAN | nucleus (probable). | nucleus. |
| Q9NY56 | OBP2A_HUMAN | secreted protein (probable). | secreted protein. |
| Q9NY57 | ST32B_HUMAN | | cytoplasm. |
| Q9NY64 | GTR8_HUMAN | membrane; multi-pass membrane protein. principally intracellular. may move between intracellular vesicles and the plasma membrane. the dileucine internalization motif is critical for intracellular sequestration (by similarity). | membrane. |
| Q9NY65 | TBA8_HUMAN | | centriole. |
| Q9NY84 | VNN3_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | cytoplasm. |
| Q9NY97 | B3GN1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | Golgi. |
| Q9NYA1 | SPHK1_HUMAN | | cytoplasm. |
| Q9NYA3 | GOGA6_HUMAN | | Golgi. |
| Q9NYB9 | ABI2_HUMAN | cytoplasm (by similarity). isoform 1 but not isoform 3 is localized to protruding lamellipodia and filopodia tips (by | cytoplasm. nucleus. |
| Q9NYC9 | DYH9_HUMAN | | cytoplasm. |
| Q9NYF0 | DACT1_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | centriole. secreted protein. nucleus. |
| Q9NYF3 | FA53C_HUMAN | | nucleus. |
| Q9NYF5 | CE005_HUMAN | | cytoplasm. |
| Q9NYG2 | ZDHC3_HUMAN | Golgi apparatus; Golgi membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum; membrane. |
| Q9NYG8 | KCNK4_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9NYI0 | EFA6R_HUMAN | | cytoplasm. |
| Q9NYJ7 | DLL3_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q9NYK1 | TLR7_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9NYK6 | EURL_HUMAN | | nucleus. |
| Q9NYL2 | MLTK_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NYL4 | FKB11_HUMAN | | cytoplasm. |
| Q9NYL9 | TMOD3_HUMAN | | cytoplasm. |
| Q9NYM9 | BET1L_HUMAN | Golgi apparatus; Golgi membrane; single-pass type iv membrane protein (by similarity). | endosome. Golgi. |
| Q9NYP3 | DONS_HUMAN | nucleus (by similarity). | nucleus. |

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| Q9NYP8 | CU062_HUMAN | | centriole. cytoplasm. |
| Q9NYP9 | CU045_HUMAN | | centriole. cytoplasm. |
| Q9NYQ8 | FAT2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9NYR9 | KBRS2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9NYS7 | WSB2_HUMAN | | cytoplasm. |
| Q9NYT0 | PLEK2_HUMAN | | cytoplasm. |
| Q9NYT6 | ZN226_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9NYU1 | UGGG2_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). endoplasmic reticulum-Golgi intermediate compartment (by similarity). | endoplasmic reticulum. |
| Q9NYV4 | CD2L7_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9NYV6 | RRN3_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9NYX4 | D1IP_HUMAN | membrane; single-pass type iii membrane protein (potential). | membrane. |
| Q9NYY3 | PLK2_HUMAN | | cytoplasm. |
| Q9NYY8 | FAKD2_HUMAN | | cytoplasm. |
| Q9NYZ1 | FA18B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NYZ2 | MFRN1_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| Q9NZ01 | GPSN2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9NZ09 | UBAP1_HUMAN | | cytoplasm. |
| Q9NZ32 | ARP10_HUMAN | | cytoplasm. |
| Q9NZ38 | CJ110_HUMAN | | nucleus. |
| Q9NZ43 | USE1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (by similarity). Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. |
| Q9NZ45 | ZCD1_HUMAN | | mitochondrion. |
| Q9NZ56 | FMN2_HUMAN | | cytoplasm. |
| Q9NZ63 | CI078_HUMAN | | nucleus. |
| Q9NZ71 | RTEL1_HUMAN | nucleus (potential). | nucleus. |
| Q9NZ72 | STMN3_HUMAN | | cytoplasm. |
| Q9NZ81 | PRR13_HUMAN | | nucleus. |
| Q9NZ94 | NLGN3_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9NZB2 | F120A_HUMAN | | cytoplasm. |
| Q9NZB8 | MOS1B_HUMAN | | mitochondrion. |
| Q9NZC2 | TREM2_HUMAN | isoform 1: cell membrane; single-pass type i membrane protein (potential). isoform 2, isoform 3: secreted protein (potential). | membrane. |
| Q9NZC7 | WVOX_HUMAN | cytoplasm. nucleus. mitochondrion. partially localizes to the mitochondria. ref.16: Golgi apparatus. translocates to the nucleus upon genotoxic stress or tnf stimulation (by similarity). isoform 5 and isoform 6 may localize in the nucleus. | cytoplasm. mitochondrion. nucleus. |
| Q9NZC9 | SMAL1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9NZD2 | GLTP_HUMAN | cytoplasm (by similarity). | endosome. |
| Q9NZF1 | PLAC8_HUMAN | | secreted protein. |
| Q9NZH7 | IL1F8_HUMAN | secreted protein (potential). | secreted protein. |

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| Q9NZI2 | KCIP1_HUMAN | cell membrane; peripheral membrane protein (by similarity). | membrane. |
| Q9NZJ4 | SACS_HUMAN | | cytoplasm. |
| Q9NZJ6 | COQ3_HUMAN | mitochondrion; mitochondrial matrix (probable). | cytoplasm. mitochondrion. |
| Q9NZJ7 | MTCH1_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (potential). | mitochondrion. |
| Q9NZK5 | CECR1_HUMAN | | secreted protein. |
| Q9NZL4 | HPBP1_HUMAN | | cytoplasm. |
| Q9NZL6 | RGL1_HUMAN | | cytoplasm. |
| Q9NZM1 | MYOF_HUMAN | cell membrane; single-pass type ii membrane protein (by similarity). nucleus; nuclear membrane; single-pass type ii membrane protein (by similarity). found at nuclear and plasma membranes. | membrane. |
| Q9NZM4 | GSCR1_HUMAN | | nucleus. |
| Q9NZM6 | PK2L2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9NZN3 | EHD3_HUMAN | | nucleus. |
| Q9NZN4 | EHD2_HUMAN | | nucleus. |
| Q9NZN5 | ARHGC_HUMAN | cytoplasm (probable). membrane (probable). translocated to the membrane upon stimulation (probable). | cytoplasm. |
| Q9NZN8 | CNOT2_HUMAN | | nucleus. |
| Q9NZP6 | CO002_HUMAN | | cytoplasm. |
| Q9NZQ9 | TMOD4_HUMAN | | cytoplasm. |
| Q9NZR1 | TMOD2_HUMAN | | cytoplasm. |
| Q9NZR2 | LRP1B_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9NZR4 | VSX1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9NZS2 | KLRF1_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9NZT1 | CALL5_HUMAN | | cytoplasm. |
| Q9NZT2 | OGFR_HUMAN | cytoplasm (by similarity). nucleus (by similarity). the ogf/ogfr complex is probably translocated to the nucleus. | membrane. |
| Q9NZU0 | FLRT3_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q9NZU1 | FLRT1_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q9NZU5 | LMCD1_HUMAN | | cytoplasm. |
| Q9NZU6 | CABP3_HUMAN | | cytoplasm. |
| Q9NZV5 | SEPNI_HUMAN | isoform 2: endoplasmic reticulum; endoplasmic reticulum membrane (probable). probably membrane- | endoplasmic reticulum. secreted protein. |
| Q9NZV6 | MSRX_HUMAN | | cytoplasm. |
| Q9NZV7 | ZIM2_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9NZW5 | MPP6_HUMAN | membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q9NZY2 | FM30A_HUMAN | | membrane. |
| Q9P000 | COMD9_HUMAN | | cytoplasm. |
| Q9P003 | CNIH4_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9P032 | CF066_HUMAN | | nucleus. |

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| Q9P055 | CN100_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9P0G3 | KLK14_HUMAN | secreted protein (probable). | secreted protein. |
| Q9P0I2 | TM111_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9P0J1 | PDP1_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | cytoplasm. |
| Q9P0K7 | RAI14_HUMAN | cytoplasm; cell cortex (by similarity). cytoskeleton (by similarity). associated with the cortical actin cytoskeleton structures in terminal web and cell-cell adhesion sites and stress fibers (by similarity). | cytoplasm. |
| Q9P0K9 | CI004_HUMAN | membrane; single-pass membrane protein (potential). | membrane. |
| Q9P0L1 | ZN167_HUMAN | nucleus (potential). | nucleus. |
| Q9P0L2 | MARK1_HUMAN | cytoplasm (by similarity). appears to localize to an intracellular network (by similarity). | cytoplasm. |
| Q9P0L9 | PK2L1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9P0N8 | MARH2_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. |
| Q9P0N9 | TBCD7_HUMAN | | cytoplasm. |
| Q9P0P8 | CF203_HUMAN | | nucleus. |
| Q9P0R6 | CN129_HUMAN | | nucleus. |
| Q9P0S2 | CN112_HUMAN | | nucleus. |
| Q9P0S3 | ORML1_HUMAN | membrane; multi-pass membrane protein (potential). endoplasmic reticulum. | endoplasmic reticulum. |
| Q9P0S9 | TM14C_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9P0T4 | ZN581_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9P0W8 | SPAT7_HUMAN | | nucleus. |
| Q9P104 | DOK5_HUMAN | | centriole. secreted protein. |
| Q9P107 | GMIP_HUMAN | | cytoplasm. |
| Q9P144 | CY014_HUMAN | | endosome; membrane. |
| Q9P1A2 | PP4RL_HUMAN | | cytoplasm. |
| Q9P1F3 | CF115_HUMAN | | cytoplasm. |
| Q9P1J3 | CN167_HUMAN | | secreted protein. |
| Q9P1N6 | CN091_HUMAN | | cytoplasm. |
| Q9P1Q0 | VPS54_HUMAN | | cytoplasm. |
| Q9P1U0 | RPA12_HUMAN | nucleus (potential). | nucleus. |
| Q9P1W9 | PIM2_HUMAN | | cytoplasm. |
| Q9P1Y5 | K1543_HUMAN | | centriole. cytoplasm. |
| Q9P1Z0 | ZBTB4_HUMAN | nucleus (potential). | nucleus. |
| Q9P202 | WHRN_HUMAN | cytoplasm. detected at the level of stereocilia in inner hair cells and in outer hair cells in the cochlea. co-localizes with the growing ends of actin filaments (by similarity). | cytoplasm. |
| Q9P203 | BTBD7_HUMAN | | cytoplasm. |
| Q9P215 | POGK_HUMAN | nucleus (probable). | nucleus. |
| Q9P217 | ZSWM5_HUMAN | | cytoplasm. nucleus. |
| Q9P218 | COKA1_HUMAN | secreted protein; extracellular space (probable). | secreted protein. |
| Q9P232 | CNTN3_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |

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| Q9P241 | AT10D_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9P243 | ZN406_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9P246 | STIM2_HUMAN | cell membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9P267 | MBD5_HUMAN | nucleus (probable). | nucleus. |
| Q9P286 | PAK7_HUMAN | | cytoplasm. |
| Q9P291 | ARMX1_HUMAN | | endosome. |
| Q9P298 | HIG1B_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9P299 | COPZ2_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). the coatomer is cytoplasmic or polymerized on the cytoplasmic side of the Golgi, as well as on the vesicles/buds originating from it (by similarity). | endosome. |
| Q9P2B2 | FPRP_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). Golgi apparatus; trans-Golgi network; trans-Golgi network membrane; single-pass type i membrane protein (by similarity). | endoplasmic reticulum. |
| Q9P2B4 | CT2NL_HUMAN | | cytoplasm. |
| Q9P2C4 | GP178_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9P2E2 | KIF17_HUMAN | | cytoplasm. nucleus. |
| Q9P2E3 | ZNFX1_HUMAN | | nucleus. |
| Q9P2E5 | CHGUT_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9P2E7 | PCD10_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9P2E8 | MARH4_HUMAN | membrane; multi-pass membrane protein (potential). | cytoplasm. |
| Q9P2E9 | RRBP1_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type iii membrane protein (by similarity). | endoplasmic reticulum. |
| Q9P2F8 | SI1L2_HUMAN | | cytoplasm. |
| Q9P2F9 | ZN319_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9P2G3 | KLH14_HUMAN | | cytoplasm. |
| Q9P2G4 | K1383_HUMAN | | nucleus. |
| Q9P2G9 | KLHL8_HUMAN | | cytoplasm. |
| Q9P2H0 | K1377_HUMAN | | cytoplasm. |
| Q9P2H3 | IFT80_HUMAN | | membrane. |
| Q9P2H5 | UBP35_HUMAN | | nucleus. |
| Q9P2I0 | CPSF2_HUMAN | nucleus (potential). | nucleus. |
| Q9P2J3 | KLHL9_HUMAN | | cytoplasm. |
| Q9P2J5 | SYLC_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9P2J8 | ZN624_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9P2J9 | PDP2_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | cytoplasm. |
| Q9P2K3 | RCOR3_HUMAN | nucleus (by similarity). | nucleus. |
| Q9P2K6 | KLDC5_HUMAN | | cytoplasm. |
| Q9P2K8 | E2AK4_HUMAN | | cytoplasm. |
| Q9P2L0 | WDR35_HUMAN | | cytoplasm. |

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|--------|-------------|--|---------------------------|
| Q9P2M4 | TBC14_HUMAN | | cytoplasm. |
| Q9P2M7 | CING_HUMAN | | cytoplasm. |
| Q9P2N4 | ATS9_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9P2N7 | KLH13_HUMAN | | cytoplasm. |
| Q9P2Q2 | FRM4A_HUMAN | | cytoplasm. |
| Q9P2S2 | NRX2A_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9P2S5 | WDR8_HUMAN | cytoplasm (potential). | membrane. |
| Q9P2S6 | ANKY1_HUMAN | | cytoplasm. nucleus. |
| Q9P2T1 | GMPR2_HUMAN | | cytoplasm. mitochondrion. |
| Q9P2V4 | LRC21_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). | endoplasmic reticulum. |
| Q9P2W3 | GBG13_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| Q9P2W9 | STX18_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type iv membrane protein (probable). | endoplasmic reticulum. |
| Q9P2X7 | DEC1_HUMAN | | cytoplasm. |
| Q9P2X8 | CI027_HUMAN | | centriole. cytoplasm. |
| Q9P2Z0 | THA10_HUMAN | | cytoplasm. nucleus. |
| Q9QC07 | POK7_HUMAN | | nucleus. |
| Q9UBA6 | G8_HUMAN | | cytoplasm. |
| Q9UBB4 | ATX10_HUMAN | | cytoplasm. |
| Q9UBB9 | TFP11_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | nucleus. |
| Q9UBC1 | IKBL_HUMAN | | centriole. |
| Q9UBC2 | EP15R_HUMAN | cell membrane; peripheral membrane protein (by similarity). nucleus (by similarity). localized to plasma membrane coated pits (by similarity). | centriole. cytoplasm. |
| Q9UBC5 | MYO1A_HUMAN | | cytoskeleton. |
| Q9UBD9 | CLCF1_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q9UBE0 | ULE1A_HUMAN | | nucleus. |
| Q9UBE8 | NLK_HUMAN | nucleus (by similarity). predominantly nuclear. cytoplasm (by similarity). a smaller fraction is cytoplasmic (by similarity). | cytoplasm. nucleus. |
| Q9UBF2 | COPG2_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). the coatomer is cytoplasmic or polymerized on the cytoplasmic side of the Golgi, as well as on the vesicles/buds originating from it (by similarity). | endosome. |
| Q9UBG7 | RBPSL_HUMAN | nucleus (probable). | nucleus. |
| Q9UBI1 | COMD3_HUMAN | | membrane. |
| Q9UBI4 | STML1_HUMAN | membrane; single-pass type iii membrane protein (potential). | membrane. |
| Q9UBI6 | GBG12_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| Q9UBI9 | HDC_HUMAN | | cytoplasm. |
| Q9UBK7 | RBL2A_HUMAN | | cytoplasm. |
| Q9UBK8 | MTRR_HUMAN | isoform b, isoform c: cytoplasm. isoform a: mitochondrion (probable). | mitochondrion. |
| Q9UBK9 | UXT_HUMAN | | cytoplasm. |

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| Q9UBL0 | ARP21_HUMAN | cytoplasm (by similarity). | centriole. cytoplasm. |
| Q9UBL3 | ASH2L_HUMAN | nucleus (probable). | nucleus. |
| Q9UBL6 | CPNE7_HUMAN | | secreted protein. |
| Q9UBM1 | PEMT_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). mitochondrion; mitochondrial membrane; multi-pass membrane protein (potential). found in endoplasmic reticulum where most pemt activity is generated and in mitochondria (by similarity). | endoplasmic reticulum. |
| Q9UBM4 | OPT_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9UBN1 | CCG4_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9UBN4 | TRPC4_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9UBP5 | HEY2_HUMAN | nucleus (probable). | nucleus. |
| Q9UBP6 | TRMB_HUMAN | | cytoplasm. |
| Q9UBQ6 | EXTL2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (by similarity). processed form: secreted protein. a soluble form is found in the serum. | endoplasmic reticulum. |
| Q9UBQ7 | GRHPR_HUMAN | | cytoplasm. |
| Q9UBR4 | LHX3_HUMAN | nucleus (probable). | nucleus. |
| Q9UBR5 | CKLF_HUMAN | isoform ck1f1: secreted protein. isoform ck1f2, isoform ck1f4: membrane; multi-pass membrane protein (potential). | membrane. |
| Q9UBS0 | KS6B2_HUMAN | | cytoplasm. |
| Q9UBS3 | DNJB9_HUMAN | cytoplasm (by similarity). nucleus (by similarity). stress induces its translocation to the nucleus (by similarity). | cytoplasm. |
| Q9UBS5 | GABR1_HUMAN | cell membrane; multi-pass membrane protein. moreover coexpression of gaba-b-r1 and gaba-b-r2 appears to be a prerequisite for maturation and transport of gaba-b-r1 to the plasma membrane. isoform 1e: secreted protein (probable). | membrane. |
| Q9UBT2 | ULE1B_HUMAN | | cytoplasm. |
| Q9UBU6 | FA8A1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9UBV2 | SEL1L_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9UBW7 | ZN198_HUMAN | nucleus (potential). | nucleus. |
| Q9UBX0 | HESX1_HUMAN | nucleus (probable). | nucleus. |
| Q9UBY9 | HSPB7_HUMAN | | cytoplasm. |
| Q9UBZ4 | APEX2_HUMAN | nucleus (probable). mitochondrion (probable). co-localized partly with pcna in nuclear foci. | nucleus. |
| Q9UBZ9 | REV1_HUMAN | nucleus (probable). | nucleus. |
| Q9UC06 | ZNF70_HUMAN | nucleus (potential). | nucleus. |
| Q9UDP3 | S11Y_HUMAN | | cytoplasm. |
| Q9UDR5 | AASS_HUMAN | mitochondrion (by similarity). | mitochondrion. |

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| Q9UDT6 | CYLN2_HUMAN | cytoplasm (by similarity). cytoskeleton (by similarity). associated with the cytoskeleton (by similarity). | cytoplasm. |
| Q9UDV6 | ZN212_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9UDV7 | ZN282_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UDW1 | UCR10_HUMAN | mitochondrion; mitochondrial inner membrane (by similarity). | mitochondrion. |
| Q9UDX3 | S14L4_HUMAN | | membrane. |
| Q9UDX4 | S14L3_HUMAN | | membrane. |
| Q9UDY2 | ZO2_HUMAN | cell membrane; cell-cell junction; adherens junction; peripheral membrane protein; cytoplasmic side (by similarity). cell membrane; cell-cell junction; tight junction; peripheral membrane protein; cytoplasmic side (by similarity). also nuclear under environmental stress conditions and in migratory endothelial cells and subconfluent epithelial cell cultures (by | cytoplasm. |
| Q9UDY4 | DNJB4_HUMAN | | endoplasmic reticulum. |
| Q9UEE9 | CFDP1_HUMAN | | secreted protein. |
| Q9UET6 | RRMJ1_HUMAN | | nucleus. |
| Q9UEU5 | GAGE8_HUMAN | | cytoplasm. |
| Q9UEW3 | MARCO_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9UEW8 | STK39_HUMAN | cytoplasm (probable). nucleus (probable). nucleus when caspase-cleaved (probable). | cytoplasm. |
| Q9UEY8 | ADDG_HUMAN | | cytoplasm. |
| Q9UF02 | CCG5_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9UF33 | EPHA6_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UF47 | DNJ5B_HUMAN | membrane; lipid-anchor (by similarity). | endoplasmic reticulum. |
| Q9UFE4 | CCD39_HUMAN | | mitochondrion. |
| Q9UFM2 | CU030_HUMAN | | nucleus. |
| Q9UG22 | GIMA2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. |
| Q9UG56 | PISD_HUMAN | mitochondrion (by similarity). | cytoplasm. |
| Q9UG63 | ABCF2_HUMAN | | mitochondrion; membrane. |
| Q9UGB7 | MIOX_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9UGC6 | RGS17_HUMAN | membrane; lipid-anchor (potential). | Golgi; membrane. |
| Q9UGI8 | TES_HUMAN | | cytoplasm. nucleus. |
| Q9UGI9 | AAKG3_HUMAN | | mitochondrion. |
| Q9UGJ0 | AAKG2_HUMAN | | cytoplasm. secreted protein. |
| Q9UGL9 | NICE1_HUMAN | | nucleus. |
| Q9UGM1 | ACHA9_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9UGM3 | DMBT1_HUMAN | secreted protein (by similarity). some isoforms may be membrane-bound. localized to the luminal aspect of crypt cells in the small intestine. in the colon, seen in the luminal aspect of surface epithelial cells. formed in the ducts of von ebner gland, and released into the fluid bathing the taste buds contained in the taste papillae (by similarity). | cytoplasm. |
| Q9UGM5 | FETUB_HUMAN | secreted protein (potential). | secreted protein. |

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| Q9UGN4 | CM35H_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9UGP4 | LIMD1_HUMAN | | cytoplasm. nucleus. |
| Q9UGQ2 | CI007_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9UGQ3 | GTR6_HUMAN | cell membrane; multi-pass membrane protein (by similarity). the dileucine internalization motif is critical for intracellular sequestration (by similarity). | membrane. |
| Q9UGR2 | Z3H7B_HUMAN | nucleus (probable). | nucleus. |
| Q9UGT4 | SUSD2_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9UGU5 | HM2L1_HUMAN | nucleus (potential). | nucleus. |
| Q9UGV2 | NDRG3_HUMAN | | centriole. cytoplasm. nucleus. |
| Q9UGV6 | HMG1X_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UH03 | SEPT3_HUMAN | | cytoplasm. |
| Q9UH17 | ABC3B_HUMAN | | cytoplasm. |
| Q9UH36 | SRR1L_HUMAN | | centriole. secreted protein. nucleus. |
| Q9UH62 | ARMX3_HUMAN | | endosome. |
| Q9UH64 | NSGX_HUMAN | | centriole. |
| Q9UH73 | COE1_HUMAN | nucleus (potential). | nucleus. |
| Q9UH77 | KLHL3_HUMAN | | cytoplasm. |
| Q9UH90 | FBX40_HUMAN | | cytoplasm. |
| Q9UH99 | UN84B_HUMAN | nucleus; nuclear membrane; single-pass membrane protein. endosome; endosomal membrane; single-pass membrane protein (probable). | nucleus. |
| Q9UHA2 | S18L2_HUMAN | | cytoplasm. |
| Q9UHA4 | MK111_HUMAN | endosome; late endosome; late endosomal membrane; peripheral membrane protein; cytoplasmic side (by similarity). | cytoplasm. Golgi. |
| Q9UHA7 | IL1F6_HUMAN | secreted protein (potential). | secreted protein. |
| Q9UHC1 | MLH3_HUMAN | nucleus (potential). | nucleus. |
| Q9UHC3 | ACCN3_HUMAN | cell membrane; multi-pass membrane protein (by similarity). cell surface expression may be stabilized by interaction with lin7b and cytoplasmic retention by interaction with dlg4. cytoplasm (by similarity). in part cytoplasmic in cochlea cells (by similarity). | membrane. |
| Q9UHC6 | CNTP2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9UHC7 | MKRN1_HUMAN | | cytoplasm. nucleus. |

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| Q9UHC9 | NPCL1_HUMAN | cell membrane; apical cell membrane; multi-pass membrane protein. cell membrane; multi-pass membrane protein (by similarity). subfractionation of brush border membranes from proximal enterocytes suggests considerable association with the apical membrane fraction. exists as a predominantly cell surface membrane expressed protein (by similarity). ref.3: cytoplasmic vesicle; cytoplasmic vesicle membrane; multi-pass membrane protein. according to ref.3 localizes in a subcellular vesicular | endoplasmic reticulum. |
| Q9UHD4 | CIDEB_HUMAN | | cytoplasm. |
| Q9UHD8 | SEPT9_HUMAN | cell membrane (by similarity). | cytoplasm. |
| Q9UHE8 | STEA1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9UHF1 | EGFL7_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q9UHF4 | I20RA_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UHF7 | TRPS1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UHG0 | DCDC2_HUMAN | | cytoplasm. |
| Q9UHI5 | LAT2_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9UHI8 | ATS1_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9UHJ3 | SMBT1_HUMAN | | centriole. cytoplasm. |
| Q9UHJ6 | CARKL_HUMAN | cytoplasm (potential). | centriole. cytoplasm. |
| Q9UHM2 | AD13_HUMAN | | endosome; membrane. |
| Q9UHM6 | OPN4_HUMAN | membrane; multi-pass membrane protein. found in soma, dendrites and proximal part of axons of certain retinal ganglion cells (by similarity). | membrane. |
| Q9UHP3 | UBP25_HUMAN | | nucleus. |
| Q9UHP6 | RTDR1_HUMAN | | centriole. cytoplasm. |
| Q9UHP9 | SMPX_HUMAN | | cytoplasm. cytoskeleton. |
| Q9UHQ4 | BAP29_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9UHR4 | BI2L1_HUMAN | | centriole. cytoplasm. |
| Q9UHR6 | ZNHI2_HUMAN | | cytoplasm. nucleus. |
| Q9UHT4 | P1854_HUMAN | | membrane. |
| Q9UHT9 | P1768_HUMAN | | nucleus. |
| Q9UHV2 | SRTD1_HUMAN | | cytoplasm. nucleus. |
| Q9UHV8 | PP13_HUMAN | | secreted protein. |
| Q9UHV9 | PFD2_HUMAN | | cytoplasm. |
| Q9UHY8 | FEZ2_HUMAN | | cytoplasm. |
| Q9UHZ2 | MALAT_HUMAN | | membrane. |
| Q9UI08 | EVL_HUMAN | cytoplasm. targeted to the leading edge of lamellipodia and the distal tip of stress fibers through interaction with a number of proteins. in activated t-cells, localizes to the f-actin collar and the distal tip of microspikes (by similarity). | cytoplasm. |
| Q9UI10 | EI2BD_HUMAN | | cytoplasm. |
| Q9UI12 | VATH_HUMAN | | mitochondrion. |

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| Q9UI14 | PRAF1_HUMAN | cell membrane; multi-pass membrane protein (by similarity). cytoplasm (by similarity). Golgi apparatus (by similarity). synaptic vesicle (by similarity). according to some authors, it is an integral membrane protein, while others showed that it is cytoplasmic and membrane-associated to Golgi and synaptic vesicles | endosome; membrane. |
| Q9UI15 | TAGL3_HUMAN | | secreted protein. |
| Q9UI25 | P461_HUMAN | | nucleus. |
| Q9UI30 | TR112_HUMAN | | cytoplasm. |
| Q9UI33 | SC11A_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9UI38 | TSP50_HUMAN | | secreted protein. |
| Q9UI42 | CBPA4_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q9UI46 | DNAI1_HUMAN | | cytoplasm. |
| Q9UI54 | P628_HUMAN | | membrane. |
| Q9UI56 | YYY4_HUMAN | | cytoplasm. |
| Q9UI66 | CI038_HUMAN | | nucleus. |
| Q9UI95 | MD2L2_HUMAN | | centriole. cytoplasm. |
| Q9UIA0 | CYH4_HUMAN | | cytoplasm. |
| Q9UIA9 | XPO7_HUMAN | cytoplasm. nucleus. nucleus; nuclear envelope; nuclear pore complex. shuttles between the nucleus and the cytoplasm (probable). | cytoplasm. |
| Q9UIC8 | LCMT1_HUMAN | | cytoplasm. |
| Q9UID3 | CK002_HUMAN | | membrane. |
| Q9UIE0 | ZN230_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UIF8 | BAZ2B_HUMAN | | nucleus. |
| Q9UIG0 | BAZ1B_HUMAN | nucleus (potential). accumulates in pericentromeric heterochromatin during replication. | nucleus. |
| Q9UIG4 | PS1C2_HUMAN | secreted protein (potential). | centriole. cytoplasm. |
| Q9UIG5 | PS1C1_HUMAN | | nucleus. |
| Q9UII5 | ZN588_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9UII6 | DUS13_HUMAN | | cytoplasm. |
| Q9UIJ5 | ZDHC2_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q9UIJ7 | KAD3_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | mitochondrion. |
| Q9UIK5 | TEFF2_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9UIL4 | KIF25_HUMAN | | cytoplasm. nucleus. |
| Q9UIL8 | PHF11_HUMAN | | nucleus. |
| Q9UIR0 | BTNL2_HUMAN | secreted protein; extracellular space (potential). | membrane. |
| Q9UIU6 | SIX4_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UIW0 | VAX2_HUMAN | nucleus (probable). | nucleus. |
| Q9UIW2 | PLXA1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9UIY3 | RWDD2_HUMAN | | cytoplasm. |
| Q9UJ04 | TSYL4_HUMAN | | centriole. cytoplasm. |
| Q9UJ14 | GGT4_HUMAN | membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9UJ37 | SIA7B_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |

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| Q9UJ55 | MAGL2_HUMAN | | centriole. cytoplasm. |
| Q9UJ68 | MSRA_HUMAN | | cytoplasm. |
| Q9UJ70 | NAGK_HUMAN | | cytoplasm. |
| Q9UJ72 | ANX10_HUMAN | | cytoplasm. |
| Q9UJ78 | ZMYM5_HUMAN | nucleus (potential). | nucleus. |
| Q9UJ99 | CAD22_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UJA5 | TRM6_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9UJA9 | ENPP5_HUMAN | secreted protein (potential). | endosome. |
| Q9UJC3 | HOOK1_HUMAN | cytoplasm (by similarity). does not associate with the Golgi complex. during spermiogenesis, it localizes to the manchette in spermatids from steps 8-10. it is also present between the microtubule manchette and the nucleus. during manchette elongation, it is preferentially localized to the nuclear ring of the manchette, whereas the strong localization to the manchette decreases. in more mature spermatids, while the manchette migrates posteriorly, it localizes to punctuates spots. at later stages of spermatid differentiation, the punctuate expression pattern is found at both the attachment site and the proximal end of the elongated manchette. in contrast, it is not present in mature spermatozoa (by | cytoplasm. |
| Q9UJD0 | RIMS3_HUMAN | | cytoplasm. |
| Q9UJF2 | NGAP_HUMAN | | cytoplasm. |
| Q9UJG1 | MSPD1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9UJH8 | METRN_HUMAN | secreted protein (by similarity). | membrane. |
| Q9UJJ9 | GNPTG_HUMAN | secreted protein (by similarity). Golgi apparatus (by similarity). | cytoplasm. |
| Q9UJL9 | ZN643_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9UJP4 | KLH21_HUMAN | | cytoplasm. |
| Q9UJQ1 | CT103_HUMAN | membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9UJQ4 | SALL4_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UJQ7 | CT079_HUMAN | | mitochondrion. peroxisome. |
| Q9UJT2 | TSKS_HUMAN | | cytoplasm. |
| Q9UJT9 | FBXL7_HUMAN | | cytoplasm. |
| Q9UJU2 | LEF1_HUMAN | nucleus (by similarity). found in nuclear bodies upon piag binding (by similarity). | nucleus. |
| Q9UJU3 | ZN228_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UJU5 | FOXD3_HUMAN | nucleus (probable). | nucleus. |
| Q9UJV9 | DDX41_HUMAN | nucleus (potential). | nucleus. |
| Q9UJW3 | DNM3L_HUMAN | nucleus (by similarity). | cytoplasm. |
| Q9UJW8 | ZN180_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UJX0 | OKL38_HUMAN | | secreted protein. |
| Q9UJX2 | CDC23_HUMAN | | nucleus. |
| Q9UJX3 | APC7_HUMAN | | nucleus. |
| Q9UJX4 | ANC5_HUMAN | | nucleus. |
| Q9UJX5 | ANC4_HUMAN | | cytoplasm. nucleus. |
| Q9UJX6 | ANC2_HUMAN | | cytoplasm. |
| Q9UJY1 | HSPB8_HUMAN | | cytoplasm. |

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| Q9UK00 | CC018_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q9UK08 | GBG8_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | membrane. |
| Q9UK10 | ZN225_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UK11 | ZN223_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UK12 | ZN222_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UK13 | ZN221_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UK22 | FBX2_HUMAN | | cytoplasm. |
| Q9UK28 | BSMAP_HUMAN | membrane (potential). could be associated with an intracellular organelle or membrane (potential). | endosome; membrane. |
| Q9UK32 | KS6A6_HUMAN | | cytoplasm. |
| Q9UK33 | ZN580_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UK39 | NOCT_HUMAN | | nucleus. |
| Q9UK41 | VPS28_HUMAN | | cytoplasm. |
| Q9UK45 | LSM7_HUMAN | nucleus (potential). | nucleus. |
| Q9UK53 | ING1_HUMAN | nucleus (probable). | nucleus. |
| Q9UK58 | CCNL1_HUMAN | nucleus; nucleoplasm; nuclear speckle (by similarity). more specifically found in nuclear intrachromatin granules clusters (igc), also called nuclear speckles, which are storage compartments for nuclear proteins involved in mrna processing (by nucleus (probable)). | nucleus. |
| Q9UK59 | DBR1_HUMAN | | nucleus. |
| Q9UK80 | UBP21_HUMAN | | nucleus. |
| Q9UK96 | FBX10_HUMAN | | cytoplasm. nucleus. |
| Q9UK97 | FBX9_HUMAN | | cytoplasm. |
| Q9UK99 | FBX3_HUMAN | | cytoplasm. |
| Q9UKA1 | FBXL5_HUMAN | | cytoplasm. nucleus. |
| Q9UKA8 | CCP3_HUMAN | | nucleus. |
| Q9UKA9 | PTBP2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UKB1 | FBW1B_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q9UKB3 | DJC12_HUMAN | | cytoplasm. |
| Q9UKC9 | FBXL2_HUMAN | | cytoplasm. |
| Q9UKE5 | TNIK_HUMAN | | cytoplasm. |
| Q9UKF6 | CPSF3_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UKG9 | OCTC_HUMAN | peroxisome (potential). | mitochondrion. |
| Q9UKH3 | ENK4_HUMAN | | endosome; membrane. |
| Q9UKH7 | ENK9_HUMAN | | centriole. |
| Q9UKH8 | GAK9_HUMAN | | cytoplasm. Golgi. |
| Q9UKI3 | VPRE3_HUMAN | | endoplasmic reticulum. |
| Q9UKI8 | TLK1_HUMAN | nucleus. isoform 3: cytoplasm (probable). | nucleus. |
| Q9UKJ0 | PILRB_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9UKJ1 | PILRA_HUMAN | isoform 1, isoform 2: cell membrane; single- pass type i membrane protein (potential). isoform 3, isoform 4: secreted protein (potential). | membrane. |
| Q9UKJ3 | GPTC8_HUMAN | | centriole. cytoplasm. |
| Q9UKK9 | NUDT5_HUMAN | | cytoplasm. |
| Q9UKL6 | PPCT_HUMAN | cytoplasm (by similarity). | secreted protein. |
| Q9UKM9 | RALY_HUMAN | nucleus (probable). | nucleus. |
| Q9UKN5 | PRDM4_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9UKP3 | ITBP2_HUMAN | | cytoplasm. |

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|--------|-------------|---|------------------------------------|
| Q9UKP4 | ATS7_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9UKP5 | ATS6_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9UKQ9 | KLK9_HUMAN | secreted protein (probable). | secreted protein. |
| Q9UKR0 | KLK12_HUMAN | secreted protein (probable). | secreted protein. |
| Q9UKR3 | KLK13_HUMAN | secreted protein (probable). | secreted protein. |
| Q9UKR5 | ERG28_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9UKR8 | TSN16_HUMAN | membrane; multi-pass membrane protein (probable). | endosome; membrane. |
| Q9UKS7 | HELI_HUMAN | nucleus (by similarity). | cytoplasm. nucleus. |
| Q9UKT4 | FBX5_HUMAN | | cytoplasm. |
| Q9UKT5 | FBX4_HUMAN | | cytoplasm. |
| Q9UKT6 | FXL21_HUMAN | | cytoplasm. nucleus. |
| Q9UKT8 | FBXW2_HUMAN | | cytoplasm. |
| Q9UKT9 | AIOL_HUMAN | nucleus (probable). | nucleus. |
| Q9UKU0 | ACSL6_HUMAN | mitochondrion; mitochondrial outer membrane; single-pass type iii membrane protein (by similarity). peroxisome; peroxisomal membrane; single-pass type iii membrane protein (by similarity). microsome; microsomal membrane; single-pass type iii membrane protein (by similarity). | endoplasmic reticulum. peroxisome. |
| Q9UKV0 | HDAC9_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UKW4 | VAV3_HUMAN | | cytoplasm. |
| Q9UKW6 | ELF5_HUMAN | | nucleus. |
| Q9UKX7 | NUP50_HUMAN | nucleus; nuclear envelope; nuclear pore complex; nucleoplasmic side (by similarity). localizes to the nucleoplasmic fibrils of the nuclear pore complex (by similarity). | nucleus. |
| Q9UKY0 | PRND_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | secreted protein. |
| Q9UKZ1 | CB029_HUMAN | | cytoplasm. |
| Q9UKZ9 | PCOC2_HUMAN | secreted protein (probable). | secreted protein. |
| Q9UL01 | SART2_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9UL12 | SARDH_HUMAN | mitochondrion; mitochondrial matrix (by similarity). | mitochondrion. |
| Q9UL15 | BAG5_HUMAN | | cytoplasm. |
| Q9UL16 | CCD19_HUMAN | | secreted protein. |
| Q9UL18 | I2C1_HUMAN | | cytoplasm. |
| Q9UL19 | TIG3_HUMAN | | cytoplasm. |
| Q9UL25 | RAB21_HUMAN | Golgi apparatus; Golgi membrane; lipid-anchor (by similarity). | Golgi. |
| Q9UL26 | RB22A_HUMAN | endosome; endosomal membrane; lipid-anchor (by similarity). cell membrane; lipid-anchor (by similarity). | endosome. |
| Q9UL36 | ZN236_HUMAN | nucleus (probable). | nucleus. |
| Q9UL46 | PSME2_HUMAN | | cytoplasm. |
| Q9UL52 | TM11E_HUMAN | cell membrane; single-pass type ii membrane protein (by similarity). processed form: secreted protein (by similarity). activated by cleavage and secreted (by similarity). | secreted protein; membrane. |

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|--------|-------------|---|---------------------------------------|
| Q9UL54 | TAOK2_HUMAN | cytoplasmic vesicle; cytoplasmic vesicle membrane; multi-pass membrane protein (potential). found to be perinuclear and localized to vesicular compartment. | cytoplasm. nucleus. |
| Q9UL58 | ZN215_HUMAN | nucleus (probable). | nucleus. |
| Q9UL59 | ZN214_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UL62 | TRPC5_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9UL63 | MKLN1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9UL68 | MYT1L_HUMAN | nucleus (by similarity). | nucleus. |
| Q9ULB1 | NRX1A_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9ULB4 | CADH9_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9ULC0 | MUCEN_HUMAN | isoform 1: cell membrane; single-pass type i membrane protein. isoform 1: membrane; single-pass type i membrane protein. consistently localized to the plasma membrane and less abundantly to Golgi-like perinuclear stacks. isoform 2: secreted protein (potential). | endosome; membrane. |
| Q9ULC3 | RAB23_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | Golgi. |
| Q9ULC5 | ACSL5_HUMAN | mitochondrion; mitochondrial outer membrane; single-pass type iii membrane protein (by similarity). peroxisome; peroxisomal membrane; single-pass type iii membrane protein (by similarity). microsome; microsomal membrane; single-pass type iii membrane protein (by similarity). | endoplasmic reticulum. peroxisome. |
| Q9ULC8 | ZDHC8_HUMAN | cytoplasmic vesicle; cytoplasmic vesicle membrane; multi-pass membrane protein (potential). | endoplasmic reticulum; membrane. |
| Q9ULD4 | BRPF3_HUMAN | | nucleus. |
| Q9ULD6 | PDZD6_HUMAN | | cytoplasm. |
| Q9ULE0 | WWC3_HUMAN | | cytoplasm. |
| Q9ULE3 | DEN2A_HUMAN | | nucleus. |
| Q9ULH4 | LRFN2_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9ULH7 | MKL2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9ULJ3 | ZN295_HUMAN | nucleus (potential). | nucleus. |
| Q9ULJ7 | ANR50_HUMAN | | cytoplasm. |
| Q9ULJ8 | NEB1_HUMAN | cytoplasm. synaptosome (by similarity). also found in the soluble synaptosomal fractions (by similarity). | cytoplasm. nucleus. |
| Q9ULK0 | GRID1_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9ULK5 | VANG2_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9ULL0 | K1210_HUMAN | | nucleus. |
| Q9ULL1 | PKHG1_HUMAN | | cytoplasm. |
| Q9ULL5 | PRR12_HUMAN | | nucleus. |
| Q9ULM2 | ZN490_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9ULM3 | YETS2_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9ULM6 | CNOT6_HUMAN | cytoplasm (probable). nucleus (probable). | cytoplasm. nucleus. |

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|--------|-------------|---|-----------------------|
| Q9ULP0 | NDRG4_HUMAN | | cytoplasm. nucleus. |
| Q9ULQ0 | FA40B_HUMAN | | nucleus. |
| Q9ULR0 | ISY1_HUMAN | nucleus (potential). | nucleus. |
| Q9ULS5 | TMCC3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9ULS6 | KCNS2_HUMAN | cell membrane; multi-pass membrane protein (by similarity). membrane; multi-pass membrane protein (by similarity). may not reach the plasma membrane but remain in an intracellular compartment in the absence of kcnb1 (by similarity). | membrane. |
| Q9ULT0 | TTC7A_HUMAN | | cytoplasm. |
| Q9ULT8 | HECD1_HUMAN | | cytoplasm. |
| Q9ULU4 | PKCB1_HUMAN | | nucleus. |
| Q9ULU8 | CAPS1_HUMAN | cytoplasmic vesicle (by similarity). membrane-associated to vesicles. strongly enriched in synaptic fractions. preferentially binds to dcvs but not to svcs. binds phosphoinositides, with a strong selectivity for ptdins(4,5)p2 over ptdins(3,4,5)p3. probably localizes to different vesicles compared to cadps2 (by | endosome. |
| Q9ULV0 | MYO5B_HUMAN | | cytoskeleton. |
| Q9ULV4 | COR1C_HUMAN | | cytoplasm. |
| Q9ULV8 | CBLC_HUMAN | nucleus (potential). | cytoplasm. |
| Q9ULW3 | ABT1_HUMAN | nucleus (by similarity). nucleus; nucleolus (by similarity). | nucleus. |
| Q9ULW5 | RAB26_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| Q9ULW6 | NP1L2_HUMAN | nucleus (probable). | centriole. |
| Q9ULX5 | ZN179_HUMAN | | cytoplasm. |
| Q9ULX7 | CAH14_HUMAN | membrane; single-pass type i membrane protein (potential). | cytoplasm. |
| Q9ULY5 | CLC4E_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9ULZ2 | STAP1_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q9UM01 | YLA1_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9UM13 | APC10_HUMAN | | cytoplasm. nucleus. |
| Q9UM19 | HPCL4_HUMAN | | centriole. cytoplasm. |
| Q9UM44 | HHLA2_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UM54 | MYO6_HUMAN | | cytoskeleton. |
| Q9UM63 | PLAL1_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UMN6 | MLL4_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UMQ6 | CAN11_HUMAN | | cytoplasm. lysosome. |
| Q9UMR7 | CLC4A_HUMAN | membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9UMS0 | HIRP5_HUMAN | | cytoplasm. |
| Q9UMS5 | PHTF1_HUMAN | nucleus (potential). | nucleus. |
| Q9UMS6 | SYNP2_HUMAN | nucleus (by similarity). cytoplasm (by similarity). shuttles between the nucleus and the cytoplasm in a differentiation-dependent and stress-induced fashion. localizes to the z-disk in mature striated muscle. the nuclear export is xpo1- | cytoplasm. nucleus. |

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| Q9UMW8 | UBP18_HUMAN | | nucleus. |
| Q9UMX2 | OAZ3_HUMAN | nucleus. cytoplasm (by similarity). | centriole. cytoplasm. |
| Q9UMX3 | BOK_HUMAN | | cytoplasm. |
| Q9UMX5 | NENF_HUMAN | secreted protein; extracellular space (by similarity). | cytoplasm. |
| Q9UMX9 | S45A2_HUMAN | melanosome; melanosomal membrane; multi-pass membrane protein (by | membrane. |
| Q9UMY4 | SNX12_HUMAN | | cytoplasm. |
| Q9UMZ2 | SYNG_HUMAN | cytoplasm (by similarity). Golgi apparatus; trans-Golgi network; trans-Golgi network membrane; peripheral membrane protein (by similarity). associates with membranes via the adapter protein complex ap-1. colocalizes with ap1g1 (by similarity). | endosome. |
| Q9UN11 | CT040_HUMAN | | membrane. |
| Q9UN30 | SCML1_HUMAN | nucleus (probable). | nucleus. |
| Q9UN42 | AT1B4_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9UN66 | PCDB8_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UN67 | PCDBA_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UN70 | PCDGK_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UN71 | PCDGG_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UN72 | PCDA7_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UN73 | PCDA6_HUMAN | isoform 1: cell membrane; single-pass type i membrane protein (by similarity). isoform 2: secreted protein (potential). | membrane. |
| Q9UN74 | PCDA4_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UN75 | PCDAC_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9UN79 | SOX13_HUMAN | nucleus (potential). | nucleus. |
| Q9UNA0 | ATS5_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9UNA1 | RHG26_HUMAN | | cytoplasm. |
| Q9UNA3 | A4GCT_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | Golgi. |
| Q9UNE0 | EDAR_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q9UNF0 | PACN2_HUMAN | cytoplasmic vesicle (by similarity). vesicle-like cytoplasmic distribution (by similarity). | cytoplasm. |
| Q9UNF1 | MAGD2_HUMAN | | nucleus. |
| Q9UNG2 | TNF18_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9UNH6 | SNX7_HUMAN | | cytoplasm. |
| Q9UNH7 | SNX6_HUMAN | | cytoplasm. endosome. |
| Q9UNK0 | STX8_HUMAN | membrane; single-pass type iv membrane protein (by similarity). preferentially associated with the early endosome. to lesser extends, also present in late endosome, the plasma membrane and coated pits (by similarity). | endoplasmic reticulum. |

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|--------|-------------|---|--------------------------|
| Q9UNK4 | PA2GD_HUMAN | secreted protein (potential). | secreted protein. |
| Q9UNK9 | ANGE1_HUMAN | | cytoplasm. |
| Q9UNM6 | PSD13_HUMAN | | cytoplasm. |
| Q9UNN5 | FAF1_HUMAN | nucleus (potential). | cytoplasm. |
| Q9UNP4 | SIAT9_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. Golgi. |
| Q9UNS1 | TIM_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UNT1 | RBL2B_HUMAN | | cytoplasm. |
| Q9UNW1 | MINP1_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| Q9UNW8 | GP132_HUMAN | membrane; multi-pass membrane protein (probable). internalized and accumulated in endosomal compartments. lpc triggers the relocalization from the endosomal compartment to the cell surface (by similarity). | membrane. |
| Q9UNW9 | NOVA2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UNX3 | RL26L_HUMAN | | mitochondrion. |
| Q9UNZ2 | NSF1C_HUMAN | nucleus (by similarity). predominantly nuclear in interphase cells. bound to the axial elements of sex chromosomes in pachytene spermatocytes. Golgi apparatus; Golgi stack (by similarity). a small proportion of the protein is cytoplasmic, associated with Golgi stacks | centriole. cytoplasm. |
| Q9UNZ5 | L10K_HUMAN | | nucleus. |
| Q9UP52 | TFR2_HUMAN | cell membrane; single-pass type ii membrane protein. isoform beta: cytoplasm (probable). the beta isoform lacks the transmembrane domain and is probably intracellular. | membrane. |
| Q9UP79 | ATS8_HUMAN | secreted protein; extracellular space; extracellular matrix (by similarity). | secreted protein. |
| Q9UPA5 | BSN_HUMAN | cytoplasm (by similarity). localized to the active zone of presynaptic density (by similarity). | nucleus. |
| Q9UPC5 | GPR34_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9UPE1 | STK23_HUMAN | | cytoplasm. |
| Q9UPG8 | PLAL2_HUMAN | nucleus (probable). | nucleus. |
| Q9UPM6 | LHX61_HUMAN | nucleus (probable). | nucleus. |
| Q9UPN6 | RBM16_HUMAN | | nucleus. |
| Q9UPN7 | SAPS1_HUMAN | | cytoplasm. |
| Q9UPN9 | TIF1G_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UPP1 | PHF8_HUMAN | | nucleus. |
| Q9UPP2 | IQEC3_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q9UPQ4 | TRI35_HUMAN | cytoplasm (by similarity). nucleus (by similarity). found predominantly in cytoplasm with a granular distribution. found in punctuate nuclear bodies (by | cytoplasm. |
| Q9UPQ7 | PZRN3_HUMAN | | cytoplasm. |
| Q9UPQ8 | TMM15_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9UPQ9 | TNC6B_HUMAN | | cytoplasm. |
| Q9UPS8 | ANR26_HUMAN | | centriole. |

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|--------|-------------|--|------------------------------|
| Q9UPT5 | EXOC7_HUMAN | cytoplasm; cytosol (by similarity). cell membrane; peripheral membrane protein (by similarity). translocates, as a preformed complex with sec6 and sec8, to the plasma membrane in response to insulin through the activation of arhq (by | cytoplasm. |
| Q9UPT6 | JIP3_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9UPT8 | CS007_HUMAN | | cytoplasm. nucleus. |
| Q9UPT9 | UBP22_HUMAN | | nucleus. |
| Q9UPU5 | UBP24_HUMAN | | nucleus. |
| Q9UPU6 | ZN409_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9UPU9 | SAMD4_HUMAN | | centriole. |
| Q9UPV7 | K1045_HUMAN | | cytoplasm. nucleus. |
| Q9UPW0 | FOXJ3_HUMAN | nucleus (potential). | nucleus. |
| Q9UPW6 | SATB2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9UPW8 | UN13A_HUMAN | cell membrane; peripheral membrane protein (by similarity). localized to the active zone of presynaptic density. translocated to the plasma membrane as response to phorbol ester binding (by | cytoplasm. |
| Q9UPX6 | K1024_HUMAN | | endosome; membrane. |
| Q9UPX8 | SHAN2_HUMAN | cytoplasm; postsynaptic density of neuronal cells (by similarity). | cytoplasm. |
| Q9UPY3 | DICER_HUMAN | | nucleus. |
| Q9UPY5 | XCT_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9UPZ3 | HPS5_HUMAN | | cytoplasm. |
| Q9UPZ9 | ICK_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9UQ03 | COR2B_HUMAN | | cytoplasm. |
| Q9UQ07 | MOK_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9UQ13 | SHOC2_HUMAN | | cytoplasm. |
| Q9UQ16 | DYN3_HUMAN | cytoplasm (potential). cytoskeleton (potential). microtubule-associated (potential). | cytoskeleton. mitochondrion. |
| Q9UQ26 | RIMS2_HUMAN | | cytoplasm. |
| Q9UQ52 | CNTN6_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| Q9UQ72 | PSG11_HUMAN | secreted protein (potential). | secreted protein. |
| Q9UQ74 | PSG8_HUMAN | secreted protein (potential). | membrane. |
| Q9UQB3 | CTND2_HUMAN | nucleus (by similarity). cell membrane; cell-cell junction; adherens junction (by similarity). | nucleus. |
| Q9UQC2 | GAB2_HUMAN | | nucleus. |
| Q9UQF0 | ENW1_HUMAN | tm: cell membrane; single-pass type i membrane protein. su: cell membrane; peripheral membrane protein. the surface protein is not anchored to the membrane, but localizes to the extracellular surface through its binding to tm (by similarity). | endosome; membrane. |

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| Q9UQF2 | JIP1_HUMAN | cytoplasm (by similarity). accumulates in cell surface projections. cytoplasm; perinuclear region (by similarity). under certain stress conditions, translocates to the perinuclear region of neurons. nucleus (by similarity). in insulin-secreting cells, detected in both the cytoplasm and nucleus (by similarity). | cytoplasm. |
| Q9UQG0 | POK20_HUMAN | | nucleus. |
| Q9UQM7 | KCC2A_HUMAN | postsynaptic lipid rafts (by similarity). | cytoplasm. |
| Q9UQP3 | TENN_HUMAN | | cytoplasm. secreted protein. |
| Q9UQQ2 | LNK_HUMAN | | cytoplasm. |
| Q9UQR0 | SCML2_HUMAN | nucleus (probable). | nucleus. |
| Q9WJR5 | POK3_HUMAN | | nucleus. |
| Q9Y210 | TRPC6_HUMAN | membrane; multi-pass membrane protein (probable). | membrane. |
| Q9Y215 | COLQ_HUMAN | | secreted protein. |
| Q9Y216 | MTMR7_HUMAN | | cytoplasm. |
| Q9Y217 | MTMR6_HUMAN | | cytoplasm. |
| Q9Y223 | GLCNE_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9Y225 | RNF24_HUMAN | | cytoplasm. |
| Q9Y226 | ORCT3_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9Y227 | ENP4_HUMAN | membrane; multi-pass membrane protein (potential). localizes in the Golgi and autophagic vacuoles/lysosomes. | endosome. secreted protein. |
| Q9Y228 | T3JAM_HUMAN | membrane; single-pass type iv membrane protein (potential). | endosome; membrane. |
| Q9Y231 | FUT9_HUMAN | Golgi apparatus; Golgi stack; Golgi stack membrane; single-pass type ii membrane protein (by similarity). membrane-bound form in trans cisternae of Golgi (by similarity). | Golgi. |
| Q9Y232 | CDYL1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y235 | ABEC2_HUMAN | | nucleus. |
| Q9Y236 | CH001_HUMAN | | cytoplasm. secreted protein. |
| Q9Y237 | PIN4_HUMAN | | cytoplasm. |
| Q9Y241 | HIG1A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9Y242 | TCF19_HUMAN | nucleus (potential). | nucleus. |
| Q9Y247 | FA50B_HUMAN | | centriole. cytoplasm. |
| Q9Y248 | PSF2_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| Q9Y250 | LZTS1_HUMAN | cytoplasm (by similarity). cell membrane (by similarity). associated with the plasma membrane and with microtubules. detected in dendritic spines, especially in the postsynaptic density (by similarity). | nucleus. |
| Q9Y252 | RNF6_HUMAN | | cytoplasm. |
| Q9Y255 | PX19_HUMAN | | secreted protein. |
| Q9Y256 | FACE2_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9Y257 | KCNK6_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9Y259 | CHKB_HUMAN | | cytoplasm. |
| Q9Y262 | IF3I_HUMAN | | cytoplasm. |
| Q9Y263 | PLAP_HUMAN | | cytoplasm. |

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| Q9Y264 | ANGP4_HUMAN | secreted protein (probable). | cytoplasm. |
| Q9Y267 | ORCT4_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9Y272 | RASD1_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. Golgi. |
| Q9Y274 | SIA10_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | secreted protein. |
| Q9Y278 | OST2_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9Y279 | VSIG4_HUMAN | membrane; single-pass type i membrane protein (probable). | membrane. |
| Q9Y281 | COF2_HUMAN | nucleus (by similarity). cytoplasm (by similarity). | cytoplasm. |
| Q9Y283 | INVS_HUMAN | cytoplasm (by similarity). cytoskeleton (by similarity). spindle (by similarity). associates with several components of the cytoskeleton including ciliary, random and polarized microtubules. during mitosis, it is recruited to mitotic spindle. membrane; peripheral membrane protein (by similarity). frequently membrane-associated, membrane localization is dependent upon cell-cell contacts and is redistributed when cell adhesion is disrupted after incubation of the cell monolayer with low- calcium/egta medium. | nucleus. |
| Q9Y284 | CS056_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9Y285 | SYFA_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9Y287 | ITM2B_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9Y296 | TPPC4_HUMAN | Golgi apparatus; cis-Golgi network (by similarity). | endoplasmic reticulum. |
| Q9Y2A4 | ZN443_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9Y2A7 | NCKP1_HUMAN | | membrane. |
| Q9Y2A9 | B3GN3_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | Golgi. |
| Q9Y2B0 | MSAP_HUMAN | | cytoplasm. |
| Q9Y2B1 | TMEM5_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9Y2B2 | PIGL_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9Y2B5 | CP007_HUMAN | | mitochondrion. |
| Q9Y2B9 | IPKG_HUMAN | | cytoplasm. |
| Q9Y2C2 | UST_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (by similarity). | membrane. |
| Q9Y2C3 | B3GT5_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | Golgi. |
| Q9Y2C4 | NUCGL_HUMAN | | secreted protein. mitochondrion. |
| Q9Y2C9 | TLR6_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). phagosome; phagosomal membrane; single- pass type i membrane protein (by | membrane. |

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|--------|-------------|--|---------------------------------|
| Q9Y2D4 | SC15B_HUMAN | | cytoplasm. |
| Q9Y2D5 | AKAP2_HUMAN | | cytoplasm. |
| Q9Y2D8 | ADIP_HUMAN | cell membrane; cell-cell junction; adherens junction (by similarity). not found at cell-matrix ajs (by similarity). nucleus. | cytoplasm. nucleus. |
| Q9Y2E4 | DIP2C_HUMAN | | centriole. |
| Q9Y2E5 | MA2B2_HUMAN | secreted protein (potential). | lysosome. |
| Q9Y2F9 | BTBD3_HUMAN | | cytoplasm. |
| Q9Y2G4 | ANKR6_HUMAN | | cytoplasm. |
| Q9Y2G5 | OFUT2_HUMAN | endoplasmic reticulum (by similarity). | Golgi. |
| Q9Y2G7 | ZFP30_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9Y2G8 | DJC16_HUMAN | membrane; single-pass type iv membrane protein (potential). | endoplasmic reticulum. |
| Q9Y2H0 | DLGP4_HUMAN | membrane; peripheral membrane protein (by similarity). | cytoplasm. |
| Q9Y2H5 | PKHA6_HUMAN | | nucleus. |
| Q9Y2H6 | FNDC3_HUMAN | | membrane. |
| Q9Y2H8 | ZN510_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9Y2H9 | MAST1_HUMAN | cell membrane; peripheral membrane protein (by similarity). colocalizes with syntrophins at the cell membrane (by similarity). | cytoplasm. |
| Q9Y2I2 | NTNG1_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| Q9Y2I8 | WDR37_HUMAN | | cytoplasm. |
| Q9Y2J0 | RP3A_HUMAN | | cytoplasm. |
| Q9Y2J2 | E41L3_HUMAN | | cytoplasm. |
| Q9Y2J4 | AMOL2_HUMAN | | centriole. |
| Q9Y2K1 | ZBTB1_HUMAN | nucleus (potential). | nucleus. |
| Q9Y2K6 | UBP20_HUMAN | | nucleus. |
| Q9Y2K7 | JHD1A_HUMAN | nucleus (probable). | nucleus. |
| Q9Y2K9 | STB5L_HUMAN | cytoplasm (potential). cell membrane; peripheral membrane protein (potential). membrane; peripheral membrane protein (potential). cytoplasmic, and associated with vesicular membranes and the plasma membrane (potential). | Golgi. |
| Q9Y2L5 | TRS85_HUMAN | Golgi apparatus; cis-Golgi network (by similarity). | cytoplasm. |
| Q9Y2L6 | FRM4B_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9Y2L8 | ZFP95_HUMAN | nucleus (potential). | nucleus. |
| Q9Y2L9 | LRCH1_HUMAN | | cytoplasm. |
| Q9Y2M5 | KLH20_HUMAN | | cytoplasm. |
| Q9Y2N3 | PO121_HUMAN | nucleus; nuclear envelope; nuclear pore complex; single-pass type ii membrane protein (by similarity). endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (by similarity). stably associated with the npc throughout interphase and the endoplasmic reticulum during metaphase (by similarity). | endoplasmic reticulum. nucleus. |
| Q9Y2P4 | S27A6_HUMAN | membrane; multi-pass membrane protein (potential). in heart is exclusively located on the sarcolemma in areas juxtaposed with small blood vessels where it colocalizes cd36 (by similarity). | endoplasmic reticulum. |

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|--------|-------------|---|---------------------|
| Q9Y2P7 | ZN256_HUMAN | nucleus (probable). | nucleus. |
| Q9Y2P8 | RCL1_HUMAN | nucleus; nucleolus (by similarity). | nucleus. |
| Q9Y2Q0 | AT8A1_HUMAN | integral membrane protein. chromaffin granules (by similarity). | membrane. |
| Q9Y2Q1 | ZN257_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9Y2Q5 | MAPIP_HUMAN | endosome; late endosome; late endosomal membrane; peripheral membrane protein; cytoplasmic side (by similarity). lysosome; lysosomal membrane; peripheral membrane protein; cytoplasmic side (by | lysosome. |
| Q9Y2R0 | CCD56_HUMAN | | endosome; membrane. |
| Q9Y2R2 | PTN22_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9Y2S2 | CRYL1_HUMAN | | mitochondrion. |
| Q9Y2S6 | CCD72_HUMAN | | nucleus. |
| Q9Y2S7 | PDIP2_HUMAN | nucleus (probable). | centriole. |
| Q9Y2T2 | AP3M1_HUMAN | | Golgi. |
| Q9Y2T3 | GUAD_HUMAN | | cytoplasm. |
| Q9Y2T4 | 2ABG_HUMAN | | cytoplasm. |
| Q9Y2T7 | YBOX2_HUMAN | cytoplasm (by similarity). nucleus (by similarity). | cytoplasm. |
| Q9Y2U2 | KCNK7_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9Y2U9 | KLDC2_HUMAN | | secreted protein. |
| Q9Y2V2 | CHSP1_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9Y2V7 | COG6_HUMAN | Golgi apparatus (by similarity). | endosome. |
| Q9Y2W1 | TR150_HUMAN | nucleus (potential). | nucleus. |
| Q9Y2W3 | S45A1_HUMAN | membrane; multi-pass membrane protein (potential). | Golgi; membrane. |
| Q9Y2W6 | TDRKH_HUMAN | | cytoplasm. |
| Q9Y2W7 | CSEN_HUMAN | cytoplasm. cell membrane; lipid-anchor (by similarity). also membrane-bound, associated with the plasma membrane (by similarity). endoplasmic reticulum. Golgi apparatus. in the presence of psen2 associated with the endoplasmic reticulum and Golgi. nucleus (probable). | membrane. |
| Q9Y2Y1 | RPC11_HUMAN | nucleus (potential). | nucleus. |
| Q9Y2Z0 | SUGT1_HUMAN | | cytoplasm. |
| Q9Y2Z2 | MTO1_HUMAN | mitochondrion (by similarity). | mitochondrion. |
| Q9Y2Z4 | SYYM_HUMAN | mitochondrion (potential). | cytoplasm. |
| Q9Y2Z9 | COQ6_HUMAN | | mitochondrion. |
| Q9Y302 | GPR89_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. Golgi. |
| Q9Y305 | ACOT9_HUMAN | | cytoplasm. |
| Q9Y312 | CT004_HUMAN | | cytoplasm. |
| Q9Y315 | DEOC_HUMAN | | cytoplasm. |
| Q9Y316 | CB004_HUMAN | | cytoplasm. |
| Q9Y324 | CN111_HUMAN | | nucleus. |
| Q9Y328 | NSG2_HUMAN | intracytoplasmic membrane; single-pass type ii membrane protein (by similarity). mainly Golgi stack, but also found in small vacuolar organelles and multivesicular bodies (by similarity). | endosome; membrane. |
| Q9Y330 | ZBT12_HUMAN | nucleus (potential). | nucleus. |
| Q9Y333 | LSM2_HUMAN | nucleus (potential). | nucleus. |
| Q9Y334 | G7C_HUMAN | | membrane. |

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| Q9Y343 | SNX24_HUMAN | | cytoplasm. |
| Q9Y365 | PCTL_HUMAN | | nucleus. |
| Q9Y366 | IFT52_HUMAN | | endoplasmic reticulum. |
| Q9Y375 | CIA30_HUMAN | mitochondrion (probable). | mitochondrion. |
| Q9Y376 | CAB39_HUMAN | cytoplasm (potential). | cytoplasm. |
| Q9Y383 | LC7L2_HUMAN | | cytoplasm. nucleus. |
| Q9Y388 | RBMX2_HUMAN | | nucleus. |
| Q9Y394 | DHRS7_HUMAN | | cytoplasm. mitochondrion. |
| Q9Y3A0 | COQ4_HUMAN | | mitochondrion. |
| Q9Y3A2 | UTP11_HUMAN | nucleus; nucleolus (by similarity). | nucleus. |
| Q9Y3A4 | ZG14_HUMAN | | nucleus. |
| Q9Y3A5 | SBDS_HUMAN | | cytoplasm. |
| Q9Y3A6 | TMED5_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). | endoplasmic reticulum. |
| Q9Y3B1 | CT045_HUMAN | | nucleus. |
| Q9Y3B3 | TMED7_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type i membrane protein (by similarity). | endoplasmic reticulum. |
| Q9Y3B4 | PM14_HUMAN | nucleus (potential). | nucleus. |
| Q9Y3B6 | CN122_HUMAN | | cytoplasm. |
| Q9Y3C0 | CCD53_HUMAN | | centriole. cytoplasm. |
| Q9Y3C5 | RNF11_HUMAN | | cytoplasm. |
| Q9Y3C6 | PPIL1_HUMAN | | cytoplasm. |
| Q9Y3C7 | MED31_HUMAN | nucleus (potential). | nucleus. |
| Q9Y3C8 | UFC1_HUMAN | | cytoplasm. |
| Q9Y3D0 | FA96B_HUMAN | | cytoplasm. |
| Q9Y3D2 | MSRB2_HUMAN | | mitochondrion. |
| Q9Y3D7 | TIM16_HUMAN | mitochondrion; mitochondrial inner membrane; peripheral membrane protein (by similarity). | cytoplasm. endosome. |
| Q9Y3E2 | BOLA1_HUMAN | | nucleus. |
| Q9Y3F1 | TA6P_HUMAN | | membrane. |
| Q9Y3F4 | STRAP_HUMAN | | cytoplasm. |
| Q9Y3I1 | FBX7_HUMAN | | cytoplasm. |
| Q9Y3K8 | GBG5L_HUMAN | | secreted protein. |
| Q9Y3L3 | 3BP1_HUMAN | | cytoplasm. |
| Q9Y3L5 | RAP2C_HUMAN | cell membrane; lipid-anchor; cytoplasmic side (potential). | cytoplasm. |
| Q9Y3M9 | ZN337_HUMAN | nucleus (probable). | cytoplasm. nucleus. |
| Q9Y3P4 | RHBD3_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9Y3Q0 | NALD2_HUMAN | membrane; single-pass type ii membrane protein (potential). | membrane. |
| Q9Y3Q8 | T22D4_HUMAN | nucleus (potential). | nucleus. |
| Q9Y3R0 | GRIP1_HUMAN | cytoplasmic and membrane-associated with vesicles, peri-Golgi complexes and endoplasmic reticulum. enriched in post-synaptic plasma membrane and post-synaptic densities (by similarity). | nucleus. |
| Q9Y3R4 | NEUR2_HUMAN | cytoplasm (by similarity). | lysosome. |
| Q9Y3R5 | CU005_HUMAN | | cytoplasm. |
| Q9Y3S1 | WNK2_HUMAN | | cytoplasm. |
| Q9Y3U8 | RL36_HUMAN | | cytoplasm. mitochondrion. |
| Q9Y3V2 | RWDD3_HUMAN | | cytoplasm. |
| Q9Y3X0 | CCDC9_HUMAN | | cytoplasm. |

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| Q9Y3Y2 | CA077_HUMAN | | cytoplasm. |
| Q9Y3Y4 | PYGO1_HUMAN | nucleus (probable). | nucleus. |
| Q9Y3Z3 | SAMH1_HUMAN | | cytoplasm. nucleus. |
| Q9Y421 | FA32A_HUMAN | | nucleus. |
| Q9Y426 | CU025_HUMAN | | nucleus. |
| Q9Y442 | CV024_HUMAN | | endosome; membrane. |
| Q9Y450 | HBS1L_HUMAN | | cytoplasm. |
| Q9Y458 | TBX22_HUMAN | nucleus (potential). | nucleus. |
| Q9Y462 | ZN711_HUMAN | nucleus (probable). | nucleus. |
| Q9Y466 | NR2E1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y467 | SALL2_HUMAN | nucleus (probable). | nucleus. |
| Q9Y471 | CMAH_HUMAN | | cytoplasm. |
| Q9Y478 | AAKB1_HUMAN | | cytoplasm. |
| Q9Y483 | MTF2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y484 | WIPI4_HUMAN | | nucleus. |
| Q9Y485 | DMXL1_HUMAN | | cytoplasm. |
| Q9Y487 | VPP2_HUMAN | membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9Y490 | TLN1_HUMAN | | cytoplasm. cytoskeleton. |
| Q9Y493 | ZAN_HUMAN | cell membrane; single-pass type i membrane protein. exclusively on the apical region of the sperm head (by | membrane. |
| Q9Y496 | KIF3A_HUMAN | | cytoplasm. nucleus. |
| Q9Y4A0 | JERKL_HUMAN | | centriole. |
| Q9Y4A8 | NF2L3_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y4C0 | NRX3A_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9Y4C1 | JHD2A_HUMAN | nucleus (probable). | nucleus. |
| Q9Y4C5 | CHST2_HUMAN | Golgi apparatus; trans-Golgi network; trans- Golgi network membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9Y4D1 | DAAM1_HUMAN | | cytoplasm. |
| Q9Y4D7 | PLXD1_HUMAN | | membrane. |
| Q9Y4E1 | FA21C_HUMAN | | nucleus. |
| Q9Y4E5 | ZN451_HUMAN | nucleus (potential). | cytoplasm. nucleus. |
| Q9Y4E6 | WDR7_HUMAN | | membrane. |
| Q9Y4E8 | UBP15_HUMAN | | nucleus. |
| Q9Y4F1 | FARP1_HUMAN | | cytoplasm. |
| Q9Y4G8 | RPGF2_HUMAN | cell membrane (by similarity). associated with the synaptic plasma membrane (by similarity). | cytoplasm. |
| Q9Y4H2 | IRS2_HUMAN | | centriole. nucleus. |
| Q9Y4I1 | MYO5A_HUMAN | | cytoskeleton. |
| Q9Y4I5 | MTL5_HUMAN | | cytoplasm. cytoskeleton. |
| Q9Y4K0 | LOXL2_HUMAN | secreted protein; extracellular space (potential). | secreted protein. |
| Q9Y4K1 | AIM1_HUMAN | | centriole. cytoplasm. |
| Q9Y4K3 | TRAF6_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9Y4L5 | ZN364_HUMAN | | cytoplasm. |
| Q9Y4P1 | ATG4B_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9Y4P3 | TBL2_HUMAN | | centriole. cytoplasm. |
| Q9Y4P8 | WIPI2_HUMAN | | cytoplasm. |
| Q9Y4P9 | SPEF1_HUMAN | cytoplasm. present in the tails of developing and epididymal sperm, internal to the fibrous sheath and around the outer dense fibers of the sperm flagellum (by | centriole. cytoplasm. |

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| Q9Y4R7 | TTLL3_HUMAN | | cytoplasm. mitochondrion. |
| Q9Y4U1 | MMAC_HUMAN | | cytoplasm. secreted protein. |
| Q9Y4X0 | AMER1_HUMAN | | centriole. cytoplasm. |
| Q9Y4X1 | UDA1_HUMAN | | microsome. |
| Q9Y4X3 | CCL27_HUMAN | | secreted protein. |
| Q9Y4Y9 | LSM5_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y4Z0 | LSM4_HUMAN | nucleus (potential). | nucleus. |
| Q9Y4Z2 | NGN3_HUMAN | nucleus (potential). | nucleus. |
| Q9Y508 | ZN313_HUMAN | | cytoplasm. |
| Q9Y512 | SAM50_HUMAN | mitochondrion; mitochondrial outer membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| Q9Y519 | CV005_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9Y530 | CF130_HUMAN | | nucleus. |
| Q9Y534 | CSDC2_HUMAN | nucleus (by similarity). cytoplasm (by similarity). pippin-rna complexes are located to the nucleus (by similarity). | cytoplasm. |
| Q9Y543 | HES2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y546 | LRC42_HUMAN | | cytoplasm. |
| Q9Y547 | PP25_HUMAN | | cytoplasm. secreted protein. |
| Q9Y548 | YIPF1_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9Y566 | SHAN1_HUMAN | cytoplasm (by similarity). postsynaptic density of neuronal cells (by similarity). | cytoplasm. |
| Q9Y570 | PPME1_HUMAN | | cytoplasm. |
| Q9Y572 | RIPK3_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9Y573 | IPP_HUMAN | | cytoplasm. |
| Q9Y574 | ASB4_HUMAN | | cytoplasm. |
| Q9Y575 | ASB3_HUMAN | | cytoplasm. |
| Q9Y576 | ASB1_HUMAN | | cytoplasm. |
| Q9Y577 | TRI17_HUMAN | | cytoplasm. |
| Q9Y580 | RBM7_HUMAN | | nucleus. |
| Q9Y581 | INSL6_HUMAN | secreted protein (by similarity). | secreted protein. |
| Q9Y584 | TIM22_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (by similarity). | mitochondrion. |
| Q9Y592 | CCD41_HUMAN | | cytoplasm. |
| Q9Y597 | KCTD3_HUMAN | | nucleus; membrane. |
| Q9Y5A6 | ZFP38_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y5A9 | YTHD2_HUMAN | | membrane. |
| Q9Y5B6 | GCFC_HUMAN | nucleus (potential). | nucleus. |
| Q9Y5B8 | NDK7_HUMAN | | cytoplasm. |
| Q9Y5C1 | ANGL3_HUMAN | secreted protein (by similarity). | cytoplasm. secreted protein. |
| Q9Y5E1 | PCDB9_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5E2 | PCDB7_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5E3 | PCDB6_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5E4 | PCDB5_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5E5 | PCDB4_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5E6 | PCDB3_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |

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| Q9Y5E7 | PCDB2_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5E8 | PCDBF_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5E9 | PCDBE_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5F0 | PCDBD_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5F1 | PCDBC_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5F2 | PCDBB_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5F3 | PCDB1_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5F6 | PCDGM_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5F7 | PCDGL_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5F8 | PCDGJ_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5F9 | PCDGI_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G0 | PCDGH_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G1 | PCDGF_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G2 | PCDGE_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G3 | PCDGD_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G4 | PCDG9_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G5 | PCDG8_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G6 | PCDG7_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G7 | PCDG6_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G8 | PCDG5_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5G9 | PCDG4_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5H0 | PCDG3_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5H1 | PCDG2_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5H2 | PCDGB_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5H3 | PCDGA_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5H4 | PCDG1_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5H5 | PCDA9_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5H6 | PCDA8_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |

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|--------|-------------|---|------------------------------|
| Q9Y5H7 | PCDA5_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5H8 | PCDA3_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5H9 | PCDA2_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5I0 | PCDAD_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5I1 | PCDAB_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5I2 | PCDAA_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). isoform 2: secreted protein (potential). | membrane. |
| Q9Y5I3 | PCDA1_HUMAN | isoform 1: cell membrane; single-pass type i membrane protein (by similarity). isoform 2: secreted protein (potential). | membrane. |
| Q9Y5I4 | PCDC2_HUMAN | membrane; single-pass type i membrane protein (by similarity). | membrane. |
| Q9Y5J3 | HEY1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y5J5 | PHLA3_HUMAN | | secreted protein. |
| Q9Y5J9 | TIM8B_HUMAN | mitochondrion; mitochondrial inner membrane; peripheral membrane protein; intermembrane side (by similarity). | mitochondrion. |
| Q9Y5K1 | SPO11_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y5K5 | UCHL5_HUMAN | | nucleus. |
| Q9Y5K6 | CD2AP_HUMAN | cytoplasm (by similarity). colocalizes with f-actin and bcar1/p130cas in membrane ruffles. located at podocyte slit diaphragm between podocyte foot processes (by similarity). | cytoplasm. |
| Q9Y5K8 | VATD_HUMAN | | mitochondrion. |
| Q9Y5L2 | HIG2_HUMAN | | endosome; membrane. |
| Q9Y5L3 | ENP2_HUMAN | membrane; multi-pass membrane protein (potential). | endosome. |
| Q9Y5L5 | LENEP_HUMAN | | nucleus. |
| Q9Y5M1 | APR2_HUMAN | | centriole. cytoplasm. |
| Q9Y5M8 | SRPRB_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass membrane protein (by similarity). | endoplasmic reticulum. |
| Q9Y5N5 | HEMK2_HUMAN | | cytoplasm. |
| Q9Y5P2 | CSAG2_HUMAN | | cytoplasm. nucleus. |
| Q9Y5P3 | RAI2_HUMAN | | cytoplasm. secreted protein. |
| Q9Y5P4 | C43BP_HUMAN | | cytoplasm. |
| Q9Y5Q3 | MAFB_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y5Q6 | INSL5_HUMAN | secreted protein (probable). | secreted protein. |
| Q9Y5R2 | MMP24_HUMAN | cell membrane; single-pass type i membrane protein (by similarity). processed form: secreted protein (by similarity). also shed from cell surface as soluble proteinase, by a proteolytic | membrane. |
| Q9Y5R4 | HEMK1_HUMAN | | cytoplasm. |
| Q9Y5R5 | DMRT2_HUMAN | nucleus (potential). | nucleus. |
| Q9Y5R6 | DMRT1_HUMAN | nucleus (potential). | nucleus. |
| Q9Y5R8 | TPPC1_HUMAN | Golgi apparatus; cis-Golgi network (by similarity). | endoplasmic reticulum. |

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|--------|-------------|--|-----------------------------|
| Q9Y5S1 | TRPV2_HUMAN | cell membrane; multi-pass membrane protein (by similarity). cytoplasm (by similarity). translocates from the cytoplasm to the plasma membrane upon ligand stimulation (by similarity). | membrane. |
| Q9Y5S2 | MRCKB_HUMAN | cytoplasm (by similarity). displays a dispersed punctate distribution and concentrates along the cell periphery, especially at the leading edge and cell-cell junction. this concentration is ph-domain dependent (by similarity). | cytoplasm. |
| Q9Y5S8 | NOX1_HUMAN | membrane; multi-pass membrane protein (potential). | membrane. |
| Q9Y5T4 | DCJ15_HUMAN | membrane; single-pass membrane protein (potential). | endosome; membrane. |
| Q9Y5T5 | UBP16_HUMAN | | nucleus. |
| Q9Y5U8 | BR44L_HUMAN | | cytoplasm. |
| Q9Y5V0 | ZN706_HUMAN | | nucleus. |
| Q9Y5V3 | MAGD1_HUMAN | cytoplasm. cell membrane; peripheral membrane protein. expression shifts from the cytoplasm to the plasma membrane upon stimulation with ngf (by similarity). | cytoplasm. |
| Q9Y5W3 | KLF2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y5W7 | SNX14_HUMAN | | cytoplasm. |
| Q9Y5W8 | SNX13_HUMAN | | cytoplasm. |
| Q9Y5W9 | SNX11_HUMAN | | cytoplasm. |
| Q9Y5X0 | SNX10_HUMAN | | cytoplasm. |
| Q9Y5X1 | SNX9_HUMAN | | cytoplasm. |
| Q9Y5X2 | SNX8_HUMAN | | cytoplasm. |
| Q9Y5X3 | SNX5_HUMAN | | cytoplasm. |
| Q9Y5Y0 | FLVC1_HUMAN | cell membrane; multi-pass membrane protein (by similarity). | membrane. |
| Q9Y5Y2 | NUBP2_HUMAN | | cytoplasm. |
| Q9Y5Y6 | ST14_HUMAN | membrane; single-pass type ii membrane protein (probable). | secreted protein; membrane. |
| Q9Y5Y9 | SC10A_HUMAN | membrane; multi-pass membrane protein (by similarity). it can be translocated to the extracellular membrane through association with s100a10 (by similarity). | membrane. |
| Q9Y5Z4 | HEBP2_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9Y5Z6 | B3GT1_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | Golgi. |
| Q9Y600 | CSAD_HUMAN | | cytoplasm. |
| Q9Y606 | TRUA_HUMAN | nucleus (by similarity). | centriole. cytoplasm. |
| Q9Y608 | LRRF2_HUMAN | | cytoplasm. |
| Q9Y614 | ACL7B_HUMAN | | cytoplasm. |
| Q9Y615 | ACL7A_HUMAN | | cytoskeleton. |
| Q9Y616 | IRAK3_HUMAN | | cytoplasm. |
| Q9Y617 | SERC_HUMAN | | cytoplasm. |
| Q9Y620 | RA54B_HUMAN | nucleus (probable). | nucleus. |
| Q9Y625 | GPC6_HUMAN | cell membrane; lipid-anchor; gpi-anchor (by similarity). | membrane. |
| Q9Y639 | NPTN_HUMAN | cell membrane; single-pass type i membrane protein (potential). | endosome; membrane. |
| Q9Y644 | RFNG_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (potential). | membrane. |

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| Q9Y651 | SOX21_HUMAN | nucleus (potential). | nucleus. |
| Q9Y661 | OST4_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9Y662 | OST3B_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9Y663 | OST3A_HUMAN | Golgi apparatus; Golgi membrane; single-pass type ii membrane protein (probable). | membrane. |
| Q9Y672 | ALG6_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9Y673 | ALG5_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; single-pass type ii membrane protein (by similarity). | endoplasmic reticulum. |
| Q9Y678 | COPG_HUMAN | cytoplasm (by similarity). Golgi apparatus; Golgi membrane; peripheral membrane protein; cytoplasmic side (by similarity). the coatomer is cytoplasmic or polymerized on the cytoplasmic side of the Golgi, as well as on the vesicles/buds originating from it (by similarity). | endosome. |
| Q9Y679 | AUP1_HUMAN | | cytoplasm. |
| Q9Y680 | FKBP7_HUMAN | endoplasmic reticulum; endoplasmic reticulum lumen (by similarity). | endoplasmic reticulum. |
| Q9Y689 | ARL5A_HUMAN | | Golgi. |
| Q9Y693 | LHFP_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9Y6A5 | TACC3_HUMAN | cytoplasm (by similarity). | nucleus. |
| Q9Y6A9 | SPCS1_HUMAN | membrane; multi-pass membrane protein (potential). | endoplasmic reticulum. |
| Q9Y6B6 | SAR1B_HUMAN | endoplasmic reticulum; endoplasmic reticulum membrane; peripheral membrane protein (by similarity). Golgi apparatus; Golgi stack; Golgi stack membrane; peripheral membrane protein (by similarity). associated with the endoplasmic reticulum and Golgi stacks, in particular in the juxta-nuclear Golgi region | endoplasmic reticulum. Golgi. |
| Q9Y6C7 | NAG7_HUMAN | | membrane. |
| Q9Y6C9 | MTCH2_HUMAN | mitochondrion; mitochondrial inner membrane; multi-pass membrane protein (potential). | mitochondrion. |
| Q9Y6D0 | SELK_HUMAN | | cytoplasm. |
| Q9Y6D5 | BIG2_HUMAN | | cytoplasm. |
| Q9Y6D6 | BIG1_HUMAN | | cytoplasm. |
| Q9Y6E0 | STK24_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9Y6E7 | SIRT4_HUMAN | | nucleus. |
| Q9Y6F7 | CDY2_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y6F8 | CDY1_HUMAN | nucleus (by similarity). | nucleus. |
| Q9Y6G1 | TM14A_HUMAN | membrane; multi-pass membrane protein (potential). | endosome; membrane. |
| Q9Y6G5 | COMDA_HUMAN | | cytoplasm. |
| Q9Y6G9 | DC1L1_HUMAN | | cytoplasm. |
| Q9Y6H1 | CHCH2_HUMAN | | nucleus. |
| Q9Y6H5 | SNCAP_HUMAN | | cytoplasm. |
| Q9Y6I0 | VPK2_HUMAN | | secreted protein. |

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| Q9Y6I3 | EPN1_HUMAN | cytoplasm (by similarity). cell membrane; peripheral membrane protein (by similarity). nucleus (by similarity). cell membrane; coated pit; clathrin-coated pit (by similarity). associated with the cytoplasmic membrane at sites where clathrin-coated pits are forming. colocalizes with clathrin and ap-2 in a punctate pattern on the plasma membrane. detected in presynaptic nerve terminals and in Golgi stacks. may shuttle to the nucleus when associated with | cytoplasm. |
| Q9Y6I4 | UBP3_HUMAN | | nucleus. |
| Q9Y6I7 | WSB1_HUMAN | | nucleus. |
| Q9Y6I8 | PXMP4_HUMAN | peroxisome; peroxisomal membrane; multi-pass membrane protein (by | peroxisome. |
| Q9Y6I9 | TX264_HUMAN | secreted protein (potential). | membrane. |
| Q9Y6J3 | DAMS_HUMAN | | cytoplasm. |
| Q9Y6J8 | STYL1_HUMAN | | cytoplasm. |
| Q9Y6K1 | DNM3A_HUMAN | nucleus (probable). | nucleus. |
| Q9Y6K5 | OAS3_HUMAN | | microsome. |
| Q9Y6L7 | TLL2_HUMAN | secreted protein (probable). | secreted protein. |
| Q9Y6M0 | TEST_HUMAN | cell membrane; lipid-anchor; gpi-anchor (potential). | secreted protein. |
| Q9Y6M1 | IF2B2_HUMAN | cytoplasm (probable). | cytoplasm. |
| Q9Y6M5 | ZNT1_HUMAN | cell membrane; multi-pass membrane protein (probable). | membrane. |
| Q9Y6M7 | S4A7_HUMAN | cell membrane; basolateral cell membrane; multi-pass membrane protein. also described at the apical cell membrane. localizes to the stereocilia of cochlear outer hair cells and to the lateral membrane of cochlear inner hair cells (by | membrane. |
| Q9Y6N5 | SQRD_HUMAN | mitochondrion (probable). | mitochondrion. |
| Q9Y6N7 | ROBO1_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9Y6N8 | CAD10_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9Y6N9 | USH1C_HUMAN | | centriole. |
| Q9Y6Q1 | CAN6_HUMAN | | cytoplasm. |
| Q9Y6Q3 | ZFP37_HUMAN | nucleus (probable). | nucleus. |
| Q9Y6Q6 | TNR11_HUMAN | membrane; single-pass type i membrane protein (potential). | membrane. |
| Q9Y6R4 | M3K4_HUMAN | | cytoplasm. |
| Q9Y6S9 | RPKL1_HUMAN | | cytoplasm. |
| Q9Y6T7 | DGKB_HUMAN | cytoplasm (by similarity). | cytoplasm. |
| Q9Y6U3 | ADSV_HUMAN | | cytoplasm. |
| Q9Y6V0 | PCLO_HUMAN | concentrated at the presynaptic side of synaptic junctions (by similarity). | cytoplasm. synapse. |
| Q9Y6V7 | DDX49_HUMAN | | nucleus. |
| Q9Y6W5 | WASF2_HUMAN | cytoplasm (by similarity). at the interface between the lamellipodial actin meshwork and the membrane (by similarity). | cytoplasm. |
| Q9Y6W8 | ICOS_HUMAN | isoform 1: cell membrane; single-pass type i membrane protein (potential). isoform 2: secreted protein (potential). | membrane. |

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| Q9Y6X2 | PIAS3_HUMAN | nucleus; nucleoplasm; nuclear speckle (by similarity). mostly (by similarity). | nucleus. |
| Q9Y6X9 | MORC2_HUMAN | | cytoplasm. nucleus. |
| Q9Y6Y1 | CMTA1_HUMAN | nucleus (probable). cytoplasm. | cytoplasm. nucleus. |
| Q9Y6Z2 | CF123_HUMAN | | centriole. cytoplasm. |
| Q9Y6Z4 | CF054_HUMAN | | centriole. cytoplasm. |
| Q9YNA8 | GAK3_HUMAN | cell membrane. cytoplasmic membrane (in a transfection system) (by similarity). | cytoplasm. |