

Online Supporting Information B: The accession numbers and sequences of 18 sub-enzyme classes of EC.1(Oxidoreductases), 8 sub-enzyme classes of EC.2(Transferases), 5 sub-enzyme classes of EC.3(Hydrolases), 6 sub-enzyme classes of EC.4(Lyases), 6 sub-enzyme classes of EC.5(Isomerases), 6 sub-enzyme classes of EC.6(Ligases).

EC.1: The accession numbers and sequences of the 1,820 Oxidoreductases sub-enzyme family classified into 18 sub enzyme families: (EC.1.1) Acting on the CH-OH group of donors; (EC.1.2) Acting on the aldehyde or oxo group of donors; (EC.1.3) Acting on the CH-CH group of donors; (EC.1.4) Acting on the CH-NH(2) group of donors; (EC.1.5) Acting on the CH-NH group of donors; (EC.1.6) Acting on NADH or NADPH; (EC.1.7) Acting on other nitrogenous compounds as donors; (EC.1.8) Acting on a sulfur group of donors; (EC.1.9) Acting on a heme group of donors; (EC.1.10) Acting on diphenols and related substances as donors; (EC.1.11) Acting on a peroxide as acceptor (peroxidases); (EC.1.12) Acting on hydrogen as donor; (EC.1.13) Acting on single donors with incorporation of molecular oxygen; (EC.1.14) Acting on paired donors, with incorporation or reduction of molecular oxygen; (EC.1.15) Acting on superoxide as acceptor; (EC.1.16) Oxidizing metal ions; (EC.1.17) Acting on CH or CH(2) groups; (EC.1.18) Acting on iron-sulfur proteins as donors. No pair of proteins in the above 18 sub enzyme families has more than 40% sequence identity. See the text of the paper for further explanation.

(EC.1.1) 449 “Acting on the CH-OH group of donors” proteins

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>O58417

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>P10539

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(EC.1.3) 149 “Acting on the CH-CH group of donors” proteins

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(EC.1.4) 72 “Acting on the CH-NH(2) group of donors ” proteins

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>P41755

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>P19801

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>Q48303

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(EC.1.6) 207 “Acting on NADH or NADPH” proteins

>P15959

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>P50939

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>P08739

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(EC.1.7) 35 “Acting on other nitrogenous compounds as donors” proteins

>Q01537

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>P16163

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>P16164

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>Q05531

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>P45017

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>P32799

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(EC.1.10) 42 “Acting on diphenols and related substances as donors” proteins

>P17489

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>P48502

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>P43311

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>O81755

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(EC.1.13) 47 “Acting on single donors with incorporation of molecular oxygen” proteins

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>Q5U3F8

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>P11122

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>Q53034

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>P29114

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>O42764

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>P0A3V2

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>P93836

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>P17554

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>O16025

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>O33948

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>Q01284

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>Q739J0

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>P42106

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>O15229

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>P19732

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>P19733

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>P19734

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>P79760

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>P05182

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>Q55NZ6

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>O05820

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>P17549

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(EC.1.16) 20 “Oxidizing metal ions” proteins

>P52093

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>Q12333

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(EC.1.17) 67 “Acting on CH or CH(2) groups” proteins

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>Q8EWR9

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>084834

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>O32143

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>O32144

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>P03190

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>Q8KFN9

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(EC.1.18) 26 “Acting on iron-sulfur proteins as donors” proteins

>Q07933

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>Q57118

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 VLQUESTLRRTAEYIEKDWKQETVLLETPLGVKGTDALTAVAKVAEVEVPAELTAERGRL
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 ASKFGE EATVWPEKDLWHLRSLSVFTEPVDLLIGSMQLKYISREANVPLVRVGFPIFDRHH
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>P13452

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>P15333

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>P15334

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 DIADSANALATLALAR YEGATTGEYLEKTFAVPN SLVNTPYGIKNTDDMLRKIAEITGKE
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>P21394

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ISKQKRLAHDIVEMEVVPDKQIAFYPGQYADVECAECASAVRSYSFSAPPQPDGSLSFHVR
 LPVGGVFSGLFGGDRGTATLTLRAPYGQFGLHESNATMVCAGGTGLAPIKCVLQSMTQ
 AQRERDVLLFFGARQQRDLYCLDEIEALQLDWGGRFELIPVLSEESSTSSWKGKRGMVTE
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>P17052

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 SGVCYLRSMEDAKNLRRKLVESAVVVLGGGVIGLEVASAAGVGLGKRVTVIEATPRVMAR
 VVTPAANLVRARLEAEGIEFKLNAKLTSIKGRNGHVEQCVLESSEEIQADLIVVGIGAI
 PELEATEAALEVSNGVVVDDQMCTSRTSIYAIGDCAMARNPFWGTMRLETIHNAVTHA
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>P65528

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 APARRVGRPRIRLAYRLTPRRVLGQRRAGGVQFSVTGTDRLRQLDAGLVLTSIGYRGKPI
 PDLPFDQAALVPNDGGRVIDPGTGEPVPGAYVAGWIKRGPFIGTNKSCSMQTVQALV
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>O07641

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 IYIVTSGEMMAMYAANNIARGILKYAHTGGVRLGGPLICNSRNVNCEAELIEELARRLGTQ
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 LLVDFGILGGEDEYEKALQADKAATKA

>Q52126

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 SKPFEFSPGQYATLQFSPEHARPYSMAGLPDDQEMEFHIRKVPGRVTEYVFEHVREGTS
 IKLSGPLGTAYLRQKHTGPMLCVGGGTGLAPVLSIVRGALKSGMTNPILLYFGVRSQQDL
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>P82861

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 KDVTVEELQEAYHAVVLSYGAEGNRTMGVPGEDLAGVYSAKDFVGWYNGLPRNRELRPDL
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 ACTIKELREMVNLPGTRPEMEASEFEGIAETLKSLARPRKRLTELLKAAIETPGEEL
 KRKAERAWGFRFLRRPLEVLPSTDILIRAAGIRLAVNRLEGDGEGVRAVATGEVEDVECG
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>Q9Z615

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>O05783

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 GSIAAVDFVGWYNAHPHFEQVSPDLGARAVVIGNGNVALDVARILLTDPVLARTDIAD
 HALESLRPRGIQEVVIVGRRGPLQAAFTTLELRELADLDGVVVIDPAELDGITDEAAA
 VGKVCKQNIKVLRGYADREPRPGHRRMVFRFLTSPIEIKGKRKVERIVLGRNELVSDGSG
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>P42454

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>P21890

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>P19066

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 AAEHETTIVPVRCEGFRGVQSLSGHIIANDAIRDWVFDKADGKTDVEFETGPYDVNVIGD
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 KRVMLYVGGLRPRHVITAYEDLGMQIVGTGYFAHNDDYQRTGHYVKTGTLIYDDATSYE
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>P77650

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>O23877

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>O68943

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>Q03304

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>P00467

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>P20620

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>P07771

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GTLARVENLSDSTITFDIQLDDGQPDIHFLAGQYVNVTLPGTETRSYSFSSQPGNRLTG
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>Q44532

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>P71527

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KVEIAKKM GEEAANKIKIVQINTPSFVEHQFKGYDNAIKAI VDTLAEPKDEENGKLNIIIP
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>P11347

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EC.2: The accession numbers and sequences of the 2,847 Transferases sub-enzyme family classified into 8 sub enzyme families: (EC.2.1) Transferring one-carbon groups; (EC.2.2) Transferring aldehyde or ketone residues; (EC.2.3) Acyltransferases; (EC.2.4) Glycosyltransferases; (EC.2.5) Transferring alkyl or aryl groups, other than methyl groups; (EC.2.6) Transferring nitrogenous groups; (EC.2.7) Transferring phosphorous-containing groups; (EC.2.8) Transferring sulfur-containing groups. No pair of proteins in the above 8 sub enzyme families has more than 40% sequence identity. See the text of the paper for further explanation.

(EC.2.1) 529 “Transferring one-carbon groups” proteins

>P11742

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GYLVQYD

>P45975

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>P96172

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>P37725

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>P29568

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>P25283

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>P43422

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(EC.2.5) 277 “Transferring alkyl or aryl groups, other than methyl groups” proteins

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>P32434

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>P38145

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>P55217

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>Q59293

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>P23970

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(EC.2.6) 114 “Transferring nitrogenous groups” proteins

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>O51043

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>O25242

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>O94235

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>O52788

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>Q25197

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>P08922

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>P21343

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>P05306

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>O32193

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>P13186

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>P28976

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>P42356

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EC.3: The accession numbers and sequences of the 3,279 Hydrolases sub-enzyme family classified into 5 sub enzyme families: (EC.3.1) Acting on ester bonds; (EC.3.2) Glycosylases; (EC.3.4) Acting on peptide bonds (peptide hydrolases); (EC.3.5) Acting on carbon-nitrogen bonds, other than peptide bonds; (EC.3.6) Acting on acid anhydrides. No pair of proteins in the above 5 sub enzyme families has more than 40% sequence identity. See the text of the paper for further explanation.

(EC.3.1) 1228 “Acting on ester bonds” proteins

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>P10486

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>O30298

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>P28351

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>Q99024

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>P19531

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>P06864

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>Q59750

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>P54802

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>O60206

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>P23340

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>O13716

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>Q09850

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>P13723

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>P14529

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>Q6BJW6

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>P22134

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>P20533

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>P03480

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>Q8Y2D7

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>P10773

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>P29760

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>P29761

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>Q02834

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>P16084

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>Q92442

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>P07337

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>P29767

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>O75951

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>P29768

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>P11218

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>P18336

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>P18192

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>P00723

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>P61626

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>P18269

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>P21270

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>P84786

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>P54196

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>Q18439

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>P11826

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>P08246

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>P21529

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>P95963

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>P42663

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>P36178

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>P38422

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>P10619

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>P0A3Z1

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>P30432

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EC.4: The accession numbers and sequences of the 892 Lyases sub-enzyme family classified into 6 sub enzyme families: (EC.4.1) Carbon-carbon lyases; (EC.4.2) Carbon-oxygen lyases ; (EC.4.3) Carbon-nitrogen lyases ; (EC.4.4) Carbon-sulfur lyases ; (EC.4.6) Phosphorus-oxygen lyases ; (EC.4.99) Other lyases. No pair of proteins in the above 6 sub enzyme families has more than 40% sequence identity. See the text of the paper for further explanation.

(EC.4.1) 340 “Carbon-carbon lyases” proteins

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>P18417

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(EC.4.6) 56 “Phosphorus-oxygen lyases” proteins

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EC.5: The accession numbers and sequences of the 639 Isomerases sub-enzyme family classified into 6 sub enzyme families: (EC.5.1) Racemases and epimerases; (EC.5.2) Cis-trans-isomerases; (EC.5.3) Intramolecular oxidoreductases; (EC.5.4) Intramolecular transferases (mutases); (EC.5.5) Intramolecular lyases; (EC.5.99) Other isomerases. No pair of proteins in the above 6 sub enzyme families has more than 40% sequence identity. See the text of the paper for further explanation.

(EC.5.1) 111 “Racemases and epimerases” proteins

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>Q83GU5

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>Q1QZC2

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 >P19080

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>P27603

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>Q6AKP0

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>Q12040

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DRPEHEPVDDGRWADAVDALAGTHDFHNLT TDET GTER TWAIDWTRDGQFLVVQLTAGGF
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>Q8ZX6

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>Q57842

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IAEKEI LNE LF DNA VFINDY PDGNFP AHQPD TLKMECL KDI I RAVKNN CEL GLI FDGDG
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>Q5NII8

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>O34676

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RYPDRVLFLVTNQCSMYCRYCTR RRFSGQIGMGVPKKQLDA AIA YIRETPEI RDCL ISGG
DG L LINDQILEYILKELRSI PHLEVIRIGTRAPVVF PQR ITDHLCEILKKYHPVWLNT HF
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>O28544

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>Q49398

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>P23973

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>Q6MDE3

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>P38604

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(EC.5.5) 11 “Intramolecular lyases” proteins

>P38677

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 PPGIPDRDPETGKGLYRADVCALTFSGKYMFASSRANKFELQGYIAGFKLRCGSIEKQL
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>O31385

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>Q38710

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>P46057

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>O81192

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>O65333

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>O33949

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 AVE TALFDAQAO
 RLGVPLSELFGRIRDSVDVAWTLASGDT
 TRDIDEAERVFEAKRHRVFKLKIGS
 RALADD
 VAHVVAI
 QKALQGRGEVRDV
 NQAWTESEAIWAGKRFADAS
 VALIEQPIA
 ENRAGLKRL
 TDLAQVPI
 MADEALHGP
 AD
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 SARAA
 DVFAVKIA
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 ANVAA
 A
 ALA
 NI
 DLY
 GGT
 MLEGAV
 GTIASA
 QLF
 STF
 GELK
 WGTE
 FGPLL
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 TRKG
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>Q9UVY5

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 LPTT
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 IL
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 CH
 AQ
 EPL
 QILD
 VS
 PDEM
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 IEHG
 VTSL
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 STA
 AYL

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>P42428

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>Q9M643

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>O97477

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(EC.5.99) 61 “Other isomerases” proteins

>Q6F598

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 NGHEIKATPDHGLVMREGKLGWVSAKNVREGDYVAFAYNTGHRGRDEYTLKLMIKLGI
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EC.6: The accession numbers and sequences of the 965 Ligases sub-enzyme family classified into 6 sub enzyme families: (EC.6.1) Forming carbon-oxygen bonds; (EC.6.2) Forming carbon-sulfur bonds; (EC.6.3) Forming carbon-nitrogen bonds; (EC.6.4) Forming carbon-carbon bonds; (EC.6.5) Forming phosphoric ester bonds; (EC.6.6) Forming nitrogen-metal bonds. No pair of proteins in the above 6 sub enzyme families has more than 40% sequence identity. See the text of the paper for further explanation.

(EC.6.1) 496 “Forming carbon-oxygen bonds” proteins

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>P41252

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>P78600

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>Q7N8D4

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>Q83G28

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>O51532

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>P39641

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>O29008

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>P32528

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>P49914

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>Q9X844

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>Q06734

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>P43386

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>Q970U8

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>P46363

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>O05272

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>P75216

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>P26046

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 KYVS
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(EC.6.4) 13 “Forming carbon-carbon bonds” proteins

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>O93918

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>O27179

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>P43778

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>P63405

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>P31562

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>Q19842

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>Q06101

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>P18823

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>P57172

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>P97386

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>P28719

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>P44121

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>O25336

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>Q2Y720

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>O74833

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>P00969

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>P32277

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>P43075

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SVS

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>P29929

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