

Protein Name

Cluster of Glial fibrillary acidic protein OS=Homo sapiens GN=GFAP PE=1 SV=1 (P14
Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2
Cluster of Isoform 3 of Plectin OS=Homo sapiens GN=PLEC (Q15149-3)
Cluster of Hemoglobin subunit beta OS=Homo sapiens GN=HBB PE=1 SV=2 (P68871)
Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4
Cluster of Tubulin beta-3 chain OS=Homo sapiens GN=TUBB3 PE=1 SV=2 (Q13509)
Cluster of Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1 (P60709)
Cluster of Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1 (P68363)
Cluster of Isoform 2 of Spectrin alpha chain, non-erythrocytic 1 OS=Homo sapiens
Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2
Cluster of Spectrin beta chain, non-erythrocytic 1 OS=Homo sapiens GN=SPTBN1 PE=1
Cluster of Pyruvate kinase isozymes M1/M2 OS=Homo sapiens GN=PKM PE=1 SV=4 (P1461
Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3
Clathrin heavy chain 1 OS=Homo sapiens GN=CLTC PE=1 SV=5
Filamin-A OS=Homo sapiens GN=FLNA PE=1 SV=4
Cytoplasmic dynein 1 heavy chain 1 OS=Homo sapiens GN=DYNC1H1 PE=1 SV=5
Cluster of ATPase, Na⁺/K⁺ transporting, alpha 2 (+) polypeptide OS=Homo sapiens G
Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1 SV=2
Fibrinogen alpha chain OS=Homo sapiens GN=FGA PE=1 SV=2
Dihydropyrimidinase-related protein 2 OS=Homo sapiens GN=DPYSL2 PE=1 SV=1
Cluster of Alpha-actinin-1 OS=Homo sapiens GN=ACTN1 PE=1 SV=2 (P12814)
60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2
ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 PE=1 SV=1
Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1
2',3'-cyclic-nucleotide 3'-phosphodiesterase OS=Homo sapiens GN=CNP PE=1 SV=2
Creatine kinase B-type OS=Homo sapiens GN=CKB PE=1 SV=1
Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2
Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2
Glutamate dehydrogenase 1, mitochondrial OS=Homo sapiens GN=GLUD1 PE=1 SV=2
Cluster of Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1 (P6810
Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3
Cluster of Myelin basic protein OS=Homo sapiens GN=MBP PE=1 SV=3 (P02686)
Neurofilament light polypeptide OS=Homo sapiens GN=NEFL PE=1 SV=3
Isoform V2 of Versican core protein OS=Homo sapiens GN=VCAN
Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5
Annexin OS=Homo sapiens GN=ANXA6 PE=3 SV=3
Cluster of Fibrinogen gamma chain OS=Homo sapiens GN=FGG PE=1 SV=3 (P02679)
Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2
Neuroblast differentiation-associated protein AHNAK OS=Homo sapiens GN=AHNAK PE=1
14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1
Nestin OS=Homo sapiens GN=NES PE=1 SV=2
Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1
Neurofilament medium polypeptide OS=Homo sapiens GN=NEFM PE=3 SV=1
Tenascin OS=Homo sapiens GN=TNC PE=1 SV=3
Microtubule-associated protein 1B OS=Homo sapiens GN=MAP1B PE=1 SV=2
ATP synthase subunit beta OS=Homo sapiens GN=ATP5B PE=3 SV=1

Alpha-2-macroglobulin OS=Homo sapiens GN=A2M PE=1 SV=3
Cluster of Complement component C4B (Childo blood group) OS=Homo sapiens GN=C4B-1
Cluster of Isoform 2 of Gelsolin OS=Homo sapiens GN=GSN (P06396-2)
Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3
Cluster of Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1 (P01857)
Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2
Cluster of Histone H2B type 1-0 OS=Homo sapiens GN=HIST1H2B0 PE=1 SV=3 (P23527)
Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4
Aconitase 2, mitochondrial OS=Homo sapiens GN=ACO2 PE=4 SV=1
Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2
Malate dehydrogenase, mitochondrial OS=Homo sapiens GN=MDH2 PE=1 SV=3
Cluster of Isoform 2 of Dynamin-1 OS=Homo sapiens GN=DNM1 (Q05193-2)
Alpha-crystallin B chain OS=Homo sapiens GN=CRYAB PE=1 SV=2
Ubiquitin carboxyl-terminal hydrolase isozyme L1 OS=Homo sapiens GN=UCHL1 PE=1 SV=1
Cluster of Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=1
Filamin-C OS=Homo sapiens GN=FLNC PE=1 SV=3
Brevican core protein OS=Homo sapiens GN=BCAN PE=1 SV=2
Transketolase OS=Homo sapiens GN=TKT PE=4 SV=1
Protein disulfide isomerase family A, member 3, isoform CRA_b OS=Homo sapiens GN=PDIA3 PE=1 SV=1
Gamma-enolase OS=Homo sapiens GN=ENO2 PE=1 SV=3
Carbonyl reductase [NADPH] 1 OS=Homo sapiens GN=CBR1 PE=1 SV=3
Myelin proteolipid protein OS=Homo sapiens GN=PLP1 PE=1 SV=2
Alpha-internexin OS=Homo sapiens GN=INA PE=1 SV=2
Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2
Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4
Fructose-bisphosphate aldolase C OS=Homo sapiens GN=ALDOC PE=1 SV=2
Heat shock 70 kDa protein 1A/1B OS=Homo sapiens GN=HSPA1A PE=1 SV=5
Hexokinase-1 OS=Homo sapiens GN=HK1 PE=4 SV=1
Endoplasmic reticulum chaperone protein OS=Homo sapiens GN=HSP90B1 PE=1 SV=1
Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2
Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1 SV=3
Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2
Cluster of Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV=1
Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1
Septin-7 OS=Homo sapiens GN=SEPT7 PE=1 SV=2
Amine oxidase [flavin-containing] B OS=Homo sapiens GN=MAOB PE=2 SV=1
Galectin-1 OS=Homo sapiens GN=LGALS1 PE=1 SV=2
78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2
Lactoferrin OS=Homo sapiens GN=LTF PE=4 SV=1
Rab GDP dissociation inhibitor alpha OS=Homo sapiens GN=GDI1 PE=1 SV=2
L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2
Cluster of Ig lambda-2 chain C regions OS=Homo sapiens GN=IGLC2 PE=1 SV=1 (P0CG05)
Tenascin-R OS=Homo sapiens GN=TNR PE=1 SV=3
Cluster of Neural cell adhesion molecule 1 OS=Homo sapiens GN=NCAM1 PE=1 SV=3 (P1)
Heat shock protein beta-1 OS=Homo sapiens GN=HSPB1 PE=1 SV=2
Cluster of Isoform 2 of Ankyrin-2 OS=Homo sapiens GN=ANK2 (Q01484-2)
Talin-1 OS=Homo sapiens GN=TLN1 PE=1 SV=3

Phosphatidylethanolamine-binding protein 1 OS=Homo sapiens GN=PEBP1 PE=1 SV=3
14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1
Phosphoserine aminotransferase OS=Homo sapiens GN=PSAT1 PE=1 SV=2
Cluster of Calnexin OS=Homo sapiens GN=CANX PE=2 SV=1 (B4DGP8)
Cluster of Phosphoglycerate mutase 1 OS=Homo sapiens GN=PGAM1 PE=1 SV=2 (P18669)
Receptor-type tyrosine-protein phosphatase zeta OS=Homo sapiens GN=PTPRZ1 PE=1 SV=1
Cluster of Plasma membrane calcium-transporting ATPase 4 OS=Homo sapiens GN=ATP2B
Triosephosphate isomerase OS=Homo sapiens GN=TPI1 PE=1 SV=3
N(G),N(G)-dimethylarginine dimethylaminohydrolase 1 OS=Homo sapiens GN=DDAH1 PE=1
Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4
Vesicle-fusing ATPase OS=Homo sapiens GN=NSF PE=1 SV=3
Isoform LCRMP-4 of Dihydropyrimidinase-related protein 3 OS=Homo sapiens GN=DPYSL
Microtubule-associated protein OS=Homo sapiens GN=MAP4 PE=4 SV=1
Isoform 2 of Syntaxin-binding protein 1 OS=Homo sapiens GN=STXBP1
Cofilin-1 OS=Homo sapiens GN=CFL1 PE=4 SV=1
Cluster of Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 (P
Isocitrate dehydrogenase [NADP], mitochondrial OS=Homo sapiens GN=IDH2 PE=1 SV=2
Moesin OS=Homo sapiens GN=MSN PE=1 SV=3
Transitional endoplasmic reticulum ATPase OS=Homo sapiens GN=VCP PE=1 SV=4
NAD(P) transhydrogenase, mitochondrial OS=Homo sapiens GN=NNT PE=1 SV=3
Ubiquitin OS=Homo sapiens GN=UBB PE=2 SV=1
Band 4.1-like protein 2 OS=Homo sapiens GN=EPB41L2 PE=4 SV=1
Cluster of Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4 (P15311)
Septin-2 OS=Homo sapiens GN=SEPT2 PE=3 SV=1
14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1 SV=1
Cluster of Mitogen-activated protein kinase 1 OS=Homo sapiens GN=MAPK1 PE=1 SV=3
WD repeat-containing protein 1 OS=Homo sapiens GN=WDR1 PE=1 SV=4
Ig kappa chain C region OS=Homo sapiens GN=IGKC PE=1 SV=1
4-trimethylaminobutyraldehyde dehydrogenase OS=Homo sapiens GN=ALDH9A1 PE=1 SV=3
Synemin OS=Homo sapiens GN=SYNM PE=1 SV=2
Reticulon-4 OS=Homo sapiens GN=RTN4 PE=1 SV=2
Malate dehydrogenase, cytoplasmic OS=Homo sapiens GN=MDH1 PE=3 SV=1
Filamin-B OS=Homo sapiens GN=FLNB PE=1 SV=2
Trifunctional enzyme subunit alpha, mitochondrial OS=Homo sapiens GN=HADHA PE=1 S
Cluster of Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3 (P07237)
Cluster of Histone H2A.Z OS=Homo sapiens GN=H2AFZ PE=1 SV=2 (POCOS5)
Cluster of Glycogen phosphorylase, brain form OS=Homo sapiens GN=PYGB PE=1 SV=5 (Alpha-aminoadipic semialdehyde dehydrogenase OS=Homo sapiens GN=ALDH7A1 PE=1 SV=5
Cytosolic non-specific dipeptidase OS=Homo sapiens GN=CNDP2 PE=1 SV=2
Microtubule-associated protein 2 OS=Homo sapiens GN=MAP2 PE=1 SV=4
Peroxioredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5
Major vault protein OS=Homo sapiens GN=MVP PE=1 SV=4
Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 PE=4 SV=1
Apolipoprotein E OS=Homo sapiens GN=APOE PE=1 SV=1
Voltage-dependent anion-selective channel protein 1 OS=Homo sapiens GN=VDAC1 PE=1
Cluster of AP-2 complex subunit beta OS=Homo sapiens GN=AP2B1 PE=1 SV=1 (P63010)
V-type proton ATPase catalytic subunit A OS=Homo sapiens GN=ATP6V1A PE=2 SV=1

D-3-phosphoglycerate dehydrogenase OS=Homo sapiens GN=PHGDH PE=1 SV=4
Contactin-1 OS=Homo sapiens GN=CNTN1 PE=1 SV=1
Chaperonin containing TCP1, subunit 8 (Theta), isoform CRA_a OS=Homo sapiens GN=C
Methylmalonate-semialdehyde dehydrogenase [acylating], mitochondrial OS=Homo sapi
Haptoglobin OS=Homo sapiens GN=HP PE=1 SV=1
Prelamin-A/C OS=Homo sapiens GN=LMNA PE=1 SV=1
Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2
Guanine nucleotide-binding protein G(o) subunit alpha OS=Homo sapiens GN=GNAO1 PE
Peroxisome oxidoreductin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3
Cluster of 1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase delta-1 OS=H
Laminin subunit beta-2 OS=Homo sapiens GN=LAMB2 PE=1 SV=2
Fibronectin OS=Homo sapiens GN=FN1 PE=4 SV=1
Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4
Ubiquitin-like modifier-activating enzyme 1 OS=Homo sapiens GN=UBA1 PE=1 SV=3
Cluster of Isoform 2 of Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB (Q1
Puromycin-sensitive aminopeptidase OS=Homo sapiens GN=NPEPPS PE=4 SV=1
4-aminobutyrate aminotransferase, mitochondrial OS=Homo sapiens GN=ABAT PE=3 SV=1
Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2
Fatty acid synthase OS=Homo sapiens GN=FASN PE=1 SV=3
Aldehyde dehydrogenase, mitochondrial OS=Homo sapiens GN=ALDH2 PE=3 SV=1
Cluster of Putative adenosylhomocysteinase 2 OS=Homo sapiens GN=AHCYL1 PE=1 SV=2
Cluster of Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Homo sapien
Isoform Cytoplasmic+peroxisomal of Peroxisome oxidoreductin-5, mitochondrial OS=Homo sapiens
Heat shock-related 70 kDa protein 2 OS=Homo sapiens GN=HSPA2 PE=1 SV=1
Alpha-1-antichymotrypsin His-Pro-less OS=Homo sapiens GN=SERPINA3 PE=3 SV=1
Adenylate kinase isoenzyme 1 OS=Homo sapiens GN=AK1 PE=1 SV=3
Neurofilament heavy polypeptide OS=Homo sapiens GN=NEFH PE=1 SV=4
Protein disulfide-isomerase A4 OS=Homo sapiens GN=PDIA4 PE=1 SV=2
V-type proton ATPase subunit B, brain isoform OS=Homo sapiens GN=ATP6V1B2 PE=1 SV
Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1
T-complex protein 1 subunit eta OS=Homo sapiens GN=CCT7 PE=1 SV=2
Dihydropyrimidinase-related protein 5 OS=Homo sapiens GN=DPYSL5 PE=1 SV=1
Cluster of Band 4.1-like protein 3 OS=Homo sapiens GN=EPB41L3 PE=1 SV=2 (Q9Y2J2)
Catenin beta-1 OS=Homo sapiens GN=CTNNB1 PE=2 SV=1
Cluster of Isoform IB of Synapsin-1 OS=Homo sapiens GN=SYN1 (P17600-2)
Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3
Cluster of Synaptosomal-associated protein 25 OS=Homo sapiens GN=SNAP25 PE=1 SV=1
Adenylyl cyclase-associated protein 1 OS=Homo sapiens GN=CAP1 PE=1 SV=5
Fascin OS=Homo sapiens GN=FSCN1 PE=1 SV=3
Trifunctional enzyme subunit beta, mitochondrial OS=Homo sapiens GN=HADHB PE=2 SV
T-complex protein 1 subunit epsilon OS=Homo sapiens GN=CCT5 PE=2 SV=1
T-complex protein 1 subunit beta OS=Homo sapiens GN=CCT2 PE=3 SV=2
Protein S100-A9 OS=Homo sapiens GN=S100A9 PE=1 SV=1
Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1
6-phosphogluconate dehydrogenase, decarboxylating OS=Homo sapiens GN=PGD PE=2 SV=2
Plasminogen OS=Homo sapiens GN=PLG PE=1 SV=2
Keratin, type II cytoskeletal 1 OS=Homo sapiens GN=KRT1 PE=1 SV=6

Cluster of ADP/ATP translocase 1 OS=Homo sapiens GN=SLC25A4 PE=1 SV=4 (P12235)
Isoform 1 of Four and a half LIM domains protein 1 OS=Homo sapiens GN=FHL1
Protein DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2
Tubulin polymerization-promoting protein family member 3 OS=Homo sapiens GN=TPPP3
Myosin-10 OS=Homo sapiens GN=MYH10 PE=4 SV=1
Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1
Cluster of Protein kinase C alpha type OS=Homo sapiens GN=PRKCA PE=1 SV=4 (P17252)
Nicotinamide phosphoribosyltransferase OS=Homo sapiens GN=NAMPT PE=1 SV=1
Prolow-density lipoprotein receptor-related protein 1 OS=Homo sapiens GN=LRP1 PE=1
Cullin-associated NEDD8-dissociated protein 1 OS=Homo sapiens GN=CAND1 PE=1 SV=2
Neuronal cell adhesion molecule OS=Homo sapiens GN=NRCAM PE=4 SV=1
Cluster of Hypoxia up-regulated protein 1 OS=Homo sapiens GN=HYOU1 PE=3 SV=1 (E9P
ATP synthase subunit d, mitochondrial OS=Homo sapiens GN=ATP5H PE=1 SV=3
Superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD1 PE=1 SV=2
T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1
Heat shock 70 kDa protein 4 OS=Homo sapiens GN=HSPA4 PE=1 SV=4
Voltage-dependent anion-selective channel protein 2 OS=Homo sapiens GN=VDAC2 PE=1
Isoform 2 of Mitochondrial inner membrane protein OS=Homo sapiens GN=IMMT
Superoxide dismutase OS=Homo sapiens GN=SOD2 PE=2 SV=1
Calmodulin OS=Homo sapiens GN=CALM2 PE=4 SV=1
Aldehyde dehydrogenase 5 family, member A1 (Succinate-semialdehyde dehydrogenase)
L-lactate dehydrogenase A chain OS=Homo sapiens GN=LDHA PE=1 SV=2
Carbonic anhydrase 2 OS=Homo sapiens GN=CA2 PE=1 SV=2
Vitamin D-binding protein OS=Homo sapiens GN=GC PE=1 SV=1
Alanine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=AARS PE=1 SV=2
Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU PE=1 SV=6
Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2
Stress-induced-phosphoprotein 1 OS=Homo sapiens GN=STIP1 PE=4 SV=1
Aspartate aminotransferase, mitochondrial OS=Homo sapiens GN=GOT2 PE=1 SV=3
Vinculin OS=Homo sapiens GN=VCL PE=1 SV=4
Acetyl-CoA acetyltransferase, mitochondrial OS=Homo sapiens GN=ACAT1 PE=1 SV=1
Gamma-adducin OS=Homo sapiens GN=ADD3 PE=1 SV=1
Cluster of Ras-related protein Rab-5A OS=Homo sapiens GN=RAB5A PE=2 SV=1 (B4DJA5)
Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo
Leukotriene A-4 hydrolase OS=Homo sapiens GN=LTA4H PE=1 SV=2
Inter-alpha-trypsin inhibitor heavy chain H2 OS=Homo sapiens GN=ITIH2 PE=1 SV=2
NADH-ubiquinone oxidoreductase 75 kDa subunit, mitochondrial OS=Homo sapiens GN=N
Cytosol aminopeptidase OS=Homo sapiens GN=LAP3 PE=1 SV=3
Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3
Succinyl-CoA:3-ketoacid coenzyme A transferase 1, mitochondrial OS=Homo sapiens G
Pre-B-cell leukemia transcription factor-interacting protein 1 OS=Homo sapiens GN
Citrate synthase OS=Homo sapiens GN=CS PE=2 SV=1
Dihydropyrimidinase-related protein 1 OS=Homo sapiens GN=CRMP1 PE=4 SV=1
Thioredoxin-dependent peroxide reductase, mitochondrial OS=Homo sapiens GN=PRDX3
Very long-chain-specific acyl-CoA dehydrogenase, mitochondrial OS=Homo sapiens GN
AP-2 complex subunit alpha-1 OS=Homo sapiens GN=AP2A1 PE=1 SV=3
Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1 PE=1 SV=2

Epoxide hydrolase 1 OS=Homo sapiens GN=EPHX1 PE=1 SV=1
Fumarate hydratase, mitochondrial OS=Homo sapiens GN=FH PE=1 SV=3
Laminin subunit gamma-1 OS=Homo sapiens GN=LAMC1 PE=1 SV=3
Electron transfer flavoprotein subunit alpha, mitochondrial OS=Homo sapiens GN=ET
Cytochrome b-c1 complex subunit 2, mitochondrial OS=Homo sapiens GN=UQCRC2 PE=1 S
Cluster of Kinesin-1 heavy chain OS=Homo sapiens GN=KIF5B PE=1 SV=1 (P33176)
T-complex protein 1 subunit gamma OS=Homo sapiens GN=CCT3 PE=1 SV=4
Protein-arginine deiminase type-2 OS=Homo sapiens GN=PADI2 PE=1 SV=2
Cytoplasmic dynein 1 light intermediate chain 2 OS=Homo sapiens GN=DYNC1LI2 PE=2
S-phase kinase-associated protein 1 OS=Homo sapiens GN=SKP1 PE=4 SV=1
Protein-L-isoaspartate O-methyltransferase OS=Homo sapiens GN=PCMT1 PE=3 SV=1
NADH-cytochrome b5 reductase 3 OS=Homo sapiens GN=CYB5R3 PE=1 SV=3
Carbonic anhydrase 1 OS=Homo sapiens GN=CA1 PE=1 SV=2
Pyruvate dehydrogenase E1 component subunit beta, mitochondrial OS=Homo sapiens G
Cluster of ADP-ribosylation factor 4 OS=Homo sapiens GN=ARF4 PE=1 SV=3 (P18085)
Lamin-B1 OS=Homo sapiens GN=LMNB1 PE=1 SV=2
Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPQ PE=1 SV=2
Coronin-1A OS=Homo sapiens GN=CORO1A PE=1 SV=4
Aquaporin-4 OS=Homo sapiens GN=AQP4 PE=1 SV=2
14-3-3 protein gamma OS=Homo sapiens GN=YWHAG PE=1 SV=2
Ras-related protein Rab-11B OS=Homo sapiens GN=RAB11B PE=1 SV=4
Staphylococcal nuclease domain-containing protein 1 OS=Homo sapiens GN=SND1 PE=1
RuvB-like 2 OS=Homo sapiens GN=RUVBL2 PE=1 SV=3
Chloride intracellular channel protein 4 OS=Homo sapiens GN=CLIC4 PE=1 SV=4
Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC PE=2 SV=
MAP1A heavy chain OS=Homo sapiens GN=MAP1A PE=4 SV=1
Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha isoform
Isoform IIB of Myc box-dependent-interacting protein 1 OS=Homo sapiens GN=BIN1
Myeloperoxidase OS=Homo sapiens GN=MPO PE=1 SV=1
Ribonuclease inhibitor OS=Homo sapiens GN=RNH1 PE=1 SV=2
Isoform 3 of Nucleoside diphosphate kinase B OS=Homo sapiens GN=NME2
Cytochrome b-c1 complex subunit 1, mitochondrial OS=Homo sapiens GN=UQCRC1 PE=1 S
Peroxisomal multifunctional enzyme type 2 OS=Homo sapiens GN=HSD17B4 PE=1 SV=3
Cluster of 2-oxoglutarate dehydrogenase, mitochondrial OS=Homo sapiens GN=OGDH PE
Cluster of Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2 (Q15365)
Chondroitin sulfate proteoglycan 4 OS=Homo sapiens GN=CSPG4 PE=1 SV=2
Calcium/calmodulin-dependent protein kinase type II subunit alpha OS=Homo sapiens
Myelin oligodendrocyte glycoprotein OS=Homo sapiens GN=MOG PE=4 SV=1
Erln-2 (Fragment) OS=Homo sapiens GN=ERLIN2 PE=4 SV=1
Rab GDP dissociation inhibitor beta OS=Homo sapiens GN=GDI2 PE=4 SV=1
Neurofascin OS=Homo sapiens GN=NFASC PE=4 SV=1
Cluster of Ferritin OS=Homo sapiens GN=FTH1 PE=3 SV=1 (G3V192)
Prohibitin-2 OS=Homo sapiens GN=PHB2 PE=4 SV=1
Coagulation factor XIII A chain OS=Homo sapiens GN=F13A1 PE=1 SV=4
Band 3 anion transport protein OS=Homo sapiens GN=SLC4A1 PE=1 SV=3
Sodium/potassium-transporting ATPase subunit beta-1 OS=Homo sapiens GN=ATP1B1 PE=
Cluster of Histone H1.4 OS=Homo sapiens GN=HIST1H1E PE=1 SV=2 (P10412)

Stathmin OS=Homo sapiens GN=STMN1 PE=1 SV=3
Glutathione S-transferase Mu 3 OS=Homo sapiens GN=GSTM3 PE=1 SV=3
Peptidyl-prolyl cis-trans isomerase B OS=Homo sapiens GN=PIIB PE=1 SV=2
Adenosylhomocysteinase OS=Homo sapiens GN=AHCY PE=1 SV=4
Delta-1-pyrroline-5-carboxylate dehydrogenase, mitochondrial OS=Homo sapiens GN=A
Cluster of Serine/threonine-protein phosphatase 2A catalytic subunit alpha isoform
Collagen alpha-1(XIV) chain OS=Homo sapiens GN=COL14A1 PE=1 SV=3
Cytoplasmic FMR1-interacting protein 1 OS=Homo sapiens GN=CYFIP1 PE=1 SV=1
Cluster of Endophilin-A1 OS=Homo sapiens GN=SH3GL2 PE=1 SV=1 (Q99962)
Cluster of Histone H3 OS=Homo sapiens GN=H3F3A PE=2 SV=1 (B4DEB1)
Cluster of Septin-11 OS=Homo sapiens GN=SEPT11 PE=3 SV=1 (D6RER5)
Phosphoribosylaminoimidazole-succinocarboxamide synthase (Fragment) OS=Homo sapiens
Phosphoribosylaminoimidazolecarboxamide formyltransferase OS=Homo sapiens GN=ATIC
Dynactin subunit 2 OS=Homo sapiens GN=DCTN2 PE=4 SV=1
Neurocan core protein OS=Homo sapiens GN=NCAN PE=2 SV=3
6-phosphogluconolactonase OS=Homo sapiens GN=PGLS PE=1 SV=2
Creatine kinase U-type, mitochondrial OS=Homo sapiens GN=CKMT1A PE=1 SV=1
Sarcoplasmic/endoplasmic reticulum calcium ATPase 2 OS=Homo sapiens GN=ATP2A2 PE=
Aspartate aminotransferase, cytoplasmic OS=Homo sapiens GN=GOT1 PE=1 SV=3
Inter-alpha-trypsin inhibitor heavy chain H1 OS=Homo sapiens GN=ITIH1 PE=1 SV=3
40S ribosomal protein S3 OS=Homo sapiens GN=RPS3 PE=1 SV=2
Phosphoglucomutase-1 OS=Homo sapiens GN=PGM1 PE=1 SV=3
Ras GTPase-activating-like protein IQGAP1 OS=Homo sapiens GN=IQGAP1 PE=1 SV=1
10 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPE1 PE=1 SV=2
Basement membrane-specific heparan sulfate proteoglycan core protein OS=Homo sapiens
Serine/threonine-protein phosphatase 2B catalytic subunit alpha isoform OS=Homo sapiens
ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4
NAD-dependent protein deacetylase sirtuin-2 OS=Homo sapiens GN=SIRT2 PE=1 SV=2
Adducin 1 (Alpha) OS=Homo sapiens GN=ADD1 PE=4 SV=1
Sorcin OS=Homo sapiens GN=SRI PE=4 SV=2
Cluster of Tropomyosin 1 (Alpha), isoform CRA_m OS=Homo sapiens GN=TPM1 PE=3 SV=1
N(G),N(G)-dimethylarginine dimethylaminohydrolase 2 OS=Homo sapiens GN=DDAH2 PE=1
Hemopexin OS=Homo sapiens GN=HPX PE=1 SV=2
Isoform 2 of HLA class II histocompatibility antigen gamma chain OS=Homo sapiens
Cathepsin D OS=Homo sapiens GN=CTSD PE=1 SV=1
Dihydropteridine reductase OS=Homo sapiens GN=QDPR PE=1 SV=2
X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3
Kinectin OS=Homo sapiens GN=KTN1 PE=1 SV=1
X-ray repair complementing defective repair in Chinese hamster cells 6 (Ku autoan
Neuronal-specific septin-3 OS=Homo sapiens GN=SEPT3 PE=4 SV=1
Cluster of Probable ATP-dependent RNA helicase DDX17 OS=Homo sapiens GN=DDX17 PE=
Prohibitin (Fragment) OS=Homo sapiens GN=PHB PE=4 SV=2
Flotillin-2 OS=Homo sapiens GN=FLOT2 PE=4 SV=1
Alpha-synuclein OS=Homo sapiens GN=SNCA PE=4 SV=1
Synapsin-2 OS=Homo sapiens GN=SYN2 PE=4 SV=1
Phosphate carrier protein, mitochondrial OS=Homo sapiens GN=SLC25A3 PE=3 SV=1
Ceruloplasmin OS=Homo sapiens GN=CP PE=1 SV=1

Glutamine synthetase OS=Homo sapiens GN=GLUL PE=1 SV=4
Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3
14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB PE=1 SV=3
Transaldolase OS=Homo sapiens GN=TALDO1 PE=1 SV=2
T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3
Glycerol-3-phosphate dehydrogenase, mitochondrial OS=Homo sapiens GN=GPD2 PE=1 SV=1
Ubiquitin carboxyl-terminal hydrolase 5 OS=Homo sapiens GN=USP5 PE=1 SV=2
Hsc70-interacting protein OS=Homo sapiens GN=ST13 PE=1 SV=2
T-complex protein 1 subunit delta OS=Homo sapiens GN=CCT4 PE=1 SV=4
Heterogeneous nuclear ribonucleoprotein A3 OS=Homo sapiens GN=HNRNPA3 PE=1 SV=2
Alpha-soluble NSF attachment protein OS=Homo sapiens GN=NAPA PE=1 SV=3
6-phosphofructokinase type C OS=Homo sapiens GN=PFKP PE=1 SV=2
Calcyphosin OS=Homo sapiens GN=CAPS PE=1 SV=1
Synaptic vesicle membrane protein VAT-1 homolog OS=Homo sapiens GN=VAT1 PE=1 SV=2
Adipocyte plasma membrane-associated protein OS=Homo sapiens GN=APMAP PE=1 SV=2
GTP-binding nuclear protein Ran OS=Homo sapiens GN=RAN PE=4 SV=1
Programmed cell death 6-interacting protein OS=Homo sapiens GN=PDCD6IP PE=4 SV=1
Rap1 GTPase-GDP dissociation stimulator 1 OS=Homo sapiens GN=RAP1GDS1 PE=4 SV=1
Neuronal membrane glycoprotein M6-a OS=Homo sapiens GN=GPM6A PE=4 SV=1
Cytosolic acyl coenzyme A thioester hydrolase OS=Homo sapiens GN=ACOT7 PE=1 SV=3
Heat shock 70 kDa protein 12A OS=Homo sapiens GN=HSPA12A PE=1 SV=2
Extracellular matrix protein 2 OS=Homo sapiens GN=ECM2 PE=2 SV=1
Protein S100-B OS=Homo sapiens GN=S100B PE=1 SV=2
Cluster of Isoform Fetal-tau of Microtubule-associated protein tau OS=Homo sapiens
Cluster of cAMP-dependent protein kinase type I-alpha regulatory subunit OS=Homo sapiens
Actin-related protein 3 OS=Homo sapiens GN=ACTR3 PE=1 SV=3
Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK PE=1 SV=1
Tropomyosin alpha-4 chain OS=Homo sapiens GN=TPM4 PE=1 SV=3
Cytoskeleton-associated protein 4 OS=Homo sapiens GN=CKAP4 PE=1 SV=2
CAP-Gly domain-containing linker protein 2 OS=Homo sapiens GN=CLIP2 PE=1 SV=1
Dynactin subunit 1 OS=Homo sapiens GN=DCTN1 PE=4 SV=2
Cluster of Capping protein (Actin filament) muscle Z-line, beta OS=Homo sapiens GN=CLIP2 PE=1 SV=1
Protein disulfide-isomerase A6 OS=Homo sapiens GN=PDIA6 PE=3 SV=1
Succinyl-CoA ligase [ADP-forming] subunit beta, mitochondrial OS=Homo sapiens GN=4F2 cell-surface antigen heavy chain OS=Homo sapiens GN=SLC3A2 PE=4 SV=1
Voltage-dependent anion-selective channel protein 3 OS=Homo sapiens GN=VDAC3 PE=4 SV=1
Methylenetetrahydrofolate dehydrogenase OS=Homo sapiens GN=MTHFD1 PE=3 SV=1
Heterogeneous nuclear ribonucleoprotein Q OS=Homo sapiens GN=SYNCRIP PE=1 SV=2
Perilipin-3 OS=Homo sapiens GN=PLIN3 PE=1 SV=3
Glutathione reductase, mitochondrial OS=Homo sapiens GN=GSR PE=1 SV=2
Ig gamma-2 chain C region OS=Homo sapiens GN=IGHG2 PE=1 SV=2
Ig mu chain C region OS=Homo sapiens GN=IGHM PE=1 SV=3
Collagen alpha-1(VI) chain OS=Homo sapiens GN=COL6A1 PE=1 SV=3
Cluster of Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6 (P1
Cluster of HLA class I histocompatibility antigen, A-11 alpha chain OS=Homo sapiens
ADP-ribosyl cyclase 1 OS=Homo sapiens GN=CD38 PE=1 SV=2
26S protease regulatory subunit 7 OS=Homo sapiens GN=PSMC2 PE=1 SV=3

V-type proton ATPase subunit E 1 OS=Homo sapiens GN=ATP6V1E1 PE=1 SV=1
Glutathione S-transferase omega-1 OS=Homo sapiens GN=GSTO1 PE=1 SV=2
Complement component 1 Q subcomponent-binding protein, mitochondrial OS=Homo sapiens GN=CC1Q1 PE=1 SV=1
Lethal(2) giant larvae protein homolog 1 OS=Homo sapiens GN=LLGL1 PE=1 SV=3
tRNA-splicing ligase RtcB homolog OS=Homo sapiens GN=C22orf28 PE=1 SV=1
Leucine zipper transcription factor-like 1, isoform CRA_b OS=Homo sapiens GN=LZTF1 PE=1 SV=1
CD166 antigen OS=Homo sapiens GN=ALCAM PE=2 SV=1
Acid ceramidase subunit beta OS=Homo sapiens GN=ASAH1 PE=4 SV=1
Calcium/calmodulin-dependent protein kinase type II subunit delta OS=Homo sapiens GN=PKCDELTA PE=1 SV=1
Thrombin light chain OS=Homo sapiens GN=F2 PE=3 SV=1
Tubulin polymerization-promoting protein OS=Homo sapiens GN=TPPP PE=1 SV=1
AP-2 complex subunit alpha-2 OS=Homo sapiens GN=AP2A2 PE=1 SV=2
Spectrin alpha chain, erythrocytic 1 OS=Homo sapiens GN=SPTA1 PE=1 SV=5
Catalase OS=Homo sapiens GN=CAT PE=1 SV=3
Retinol-binding protein 1 OS=Homo sapiens GN=RBP1 PE=1 SV=2
Annexin A4 OS=Homo sapiens GN=ANXA4 PE=1 SV=4
Poly [ADP-ribose] polymerase 1 OS=Homo sapiens GN=PARP1 PE=1 SV=4
Alcohol dehydrogenase class-3 OS=Homo sapiens GN=ADH5 PE=1 SV=4
Glucosidase 2 subunit beta OS=Homo sapiens GN=PRKCSH PE=1 SV=2
6-phosphofructokinase, liver type OS=Homo sapiens GN=PFKL PE=1 SV=6
Myristoylated alanine-rich C-kinase substrate OS=Homo sapiens GN=MARCKS PE=1 SV=4
ES1 protein homolog, mitochondrial OS=Homo sapiens GN=C21orf33 PE=1 SV=3
Electron transfer flavoprotein subunit beta OS=Homo sapiens GN=ETFB PE=1 SV=3
Cluster of Guanine nucleotide-binding protein G(q) subunit alpha OS=Homo sapiens GN=GNAS1 PE=1 SV=1
Ras-related protein Rab-7a OS=Homo sapiens GN=RAB7A PE=1 SV=1
Hepatocyte cell adhesion molecule OS=Homo sapiens GN=HEPACAM PE=1 SV=1
Microtubule-associated protein RP/EB family member 1 OS=Homo sapiens GN=MAPRE1 PE=1 SV=1
Tropomyosin 3 OS=Homo sapiens GN=TPM3 PE=2 SV=1
ATP-dependent RNA helicase DDX1 OS=Homo sapiens GN=DDX1 PE=1 SV=2
Heat shock protein 105 kDa OS=Homo sapiens GN=HSPH1 PE=1 SV=1
Protein kinase C and casein kinase substrate in neurons protein 1 OS=Homo sapiens GN=CKS1B PE=1 SV=1
Ankyrin repeat and FYVE domain-containing protein 1 OS=Homo sapiens GN=ANKFY1 PE=1 SV=1
Drebrin OS=Homo sapiens GN=DBN1 PE=4 SV=1
Proactivator polypeptide OS=Homo sapiens GN=PSAP PE=4 SV=1
Dihydrolipoyl dehydrogenase OS=Homo sapiens GN=DLD PE=2 SV=1
Complement factor B Ba fragment OS=Homo sapiens GN=CFB PE=2 SV=1
Secernin-1 OS=Homo sapiens GN=SCRN1 PE=4 SV=2
CD44 antigen OS=Homo sapiens GN=CD44 PE=4 SV=2
Cluster of Isocitrate dehydrogenase [NAD] subunit alpha, mitochondrial OS=Homo sapiens GN=IDH3A PE=1 SV=1
Microtubule-actin cross-linking factor 1, isoforms 1/2/3/5 OS=Homo sapiens GN=MAC1 PE=1 SV=1
Eukaryotic translation initiation factor 5A-1 (Fragment) OS=Homo sapiens GN=EIF5A PE=1 SV=1
Synaptotagmin I, isoform CRA_b OS=Homo sapiens GN=SYT1 PE=4 SV=1
Lamin B2, isoform CRA_a OS=Homo sapiens GN=LMNB2 PE=4 SV=1
BAG family molecular chaperone regulator 3 OS=Homo sapiens GN=BAG3 PE=1 SV=3
Cystatin-B OS=Homo sapiens GN=CSTB PE=1 SV=2
Aldose reductase OS=Homo sapiens GN=AKR1B1 PE=1 SV=3
ATP synthase subunit b, mitochondrial OS=Homo sapiens GN=ATP5F1 PE=1 SV=2

3-ketoacyl-CoA thiolase, mitochondrial OS=Homo sapiens GN=ACAA2 PE=1 SV=2
Dextrin OS=Homo sapiens GN=DSTN PE=1 SV=3
Sorbitol dehydrogenase OS=Homo sapiens GN=SORD PE=1 SV=4
Cluster of Aldo-keto reductase family 1 member C1 OS=Homo sapiens GN=AKR1C1 PE=1
14-3-3 protein eta OS=Homo sapiens GN=YWHAH PE=1 SV=4
Nuclear mitotic apparatus protein 1 OS=Homo sapiens GN=NUMA1 PE=1 SV=2
Early endosome antigen 1 OS=Homo sapiens GN=EEA1 PE=1 SV=2
Succinyl-CoA ligase [GDP-forming] subunit beta, mitochondrial OS=Homo sapiens GN=
Isoform 5 of Sorbin and SH3 domain-containing protein 1 OS=Homo sapiens GN=SORBS1
Septin-9 OS=Homo sapiens GN=SEPT9 PE=1 SV=2
Quinone oxidoreductase OS=Homo sapiens GN=CRYZ PE=4 SV=1
Protein NDRG2 OS=Homo sapiens GN=NDRG2 PE=2 SV=1
Cluster of N-acetyl-D-glucosamine kinase OS=Homo sapiens GN=NAGK PE=2 SV=1 (B4DLZ
2,4-dienoyl-CoA reductase, mitochondrial OS=Homo sapiens GN=DECR1 PE=2 SV=1
40S ribosomal protein SA (Fragment) OS=Homo sapiens GN=RPSA PE=3 SV=1
26S protease regulatory subunit 6A OS=Homo sapiens GN=PSMC3 PE=3 SV=1
Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 PE=4 SV=1
60S acidic ribosomal protein P0 (Fragment) OS=Homo sapiens GN=RPLP0 PE=3 SV=1
Actin-related protein 2/3 complex subunit 2 OS=Homo sapiens GN=ARPC2 PE=1 SV=1
Protein S100-A8 OS=Homo sapiens GN=S100A8 PE=1 SV=1
Complement factor H OS=Homo sapiens GN=CFH PE=1 SV=4
Heterogeneous nuclear ribonucleoprotein L OS=Homo sapiens GN=HNRNPL PE=1 SV=2
Aspartate--tRNA ligase, cytoplasmic OS=Homo sapiens GN=DARS PE=1 SV=2
Laminin subunit alpha-2 OS=Homo sapiens GN=LAMA2 PE=1 SV=4
Glutathione S-transferase Mu 2 OS=Homo sapiens GN=GSTM2 PE=1 SV=2
Tropomodulin-1 OS=Homo sapiens GN=TMOD1 PE=1 SV=1
Cluster of Interferon-induced guanylate-binding protein 1 OS=Homo sapiens GN=GBP1
Lon protease homolog, mitochondrial OS=Homo sapiens GN=LONP1 PE=1 SV=2
Excitatory amino acid transporter 2 OS=Homo sapiens GN=SLC1A2 PE=1 SV=2
Cluster of F-actin-capping protein subunit alpha-2 OS=Homo sapiens GN=CAPZA2 PE=1
ATP-citrate synthase OS=Homo sapiens GN=ACLY PE=1 SV=3
Tyrosine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=YARS PE=1 SV=4
Cluster of Heterogeneous nuclear ribonucleoprotein H2 OS=Homo sapiens GN=HNRNPH2
Eukaryotic initiation factor 4A-I OS=Homo sapiens GN=EIF4A1 PE=1 SV=1
Cluster of Ras-related C3 botulinum toxin substrate 1 OS=Homo sapiens GN=RAC1 PE=
Tyrosine-protein phosphatase non-receptor type substrate 1 OS=Homo sapiens GN=SIR
Calponin-3 OS=Homo sapiens GN=CNN3 PE=1 SV=1
Cluster of EH domain-containing protein 1 OS=Homo sapiens GN=EHD1 PE=1 SV=2 (Q9H4
Phenylalanine--tRNA ligase beta subunit OS=Homo sapiens GN=FARSB PE=1 SV=3
Guanine deaminase OS=Homo sapiens GN=GDA PE=1 SV=1
Macrophage migration inhibitory factor OS=Homo sapiens GN=MIF PE=1 SV=4
Septin-8 OS=Homo sapiens GN=SEPT8 PE=3 SV=1
Astrocytic phosphoprotein PEA-15 OS=Homo sapiens GN=PEA15 PE=2 SV=1
Thioredoxin OS=Homo sapiens GN=TXN PE=4 SV=1
UTP--glucose-1-phosphate uridylyltransferase OS=Homo sapiens GN=UGP2 PE=4 SV=1
DNA-dependent protein kinase catalytic subunit OS=Homo sapiens GN=PRKDC PE=4 SV=1
Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Homo

Cysteine and glycine-rich protein 1 (Fragment) OS=Homo sapiens GN=CSRP1 PE=4 SV=1
Cluster of V-type proton ATPase subunit d 1 OS=Homo sapiens GN=ATP6V0D1 PE=4 SV=1
Cluster of Vesicle-associated membrane protein 2 OS=Homo sapiens GN=VAMP2 PE=4 SV=1
Interleukin enhancer binding factor 3, 90kDa, isoform CRA_b OS=Homo sapiens GN=IL
Proteasome subunit alpha type (Fragment) OS=Homo sapiens GN=PSMA4 PE=3 SV=1
Exportin-1 OS=Homo sapiens GN=XPO1 PE=1 SV=1
Copine-3 OS=Homo sapiens GN=CPNE3 PE=1 SV=1
Cluster of Paralemmin-1 OS=Homo sapiens GN=PALM PE=1 SV=2 (075781)
Argininosuccinate synthase OS=Homo sapiens GN=ASS1 PE=1 SV=2
Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2
Glutathione peroxidase 1 OS=Homo sapiens GN=GPX1 PE=1 SV=4
Calpain-1 catalytic subunit OS=Homo sapiens GN=CAPN1 PE=1 SV=1
Bifunctional glutamate/proline--tRNA ligase OS=Homo sapiens GN=EPRS PE=1 SV=5
Cytochrome c oxidase subunit 5B, mitochondrial OS=Homo sapiens GN=COX5B PE=1 SV=2
Alcohol dehydrogenase [NADP(+)] OS=Homo sapiens GN=AKR1A1 PE=1 SV=3
40S ribosomal protein S12 OS=Homo sapiens GN=RPS12 PE=1 SV=3
Catenin alpha-2 OS=Homo sapiens GN=CTNNA2 PE=1 SV=5
DnaJ homolog subfamily A member 1 OS=Homo sapiens GN=DNAJA1 PE=1 SV=2
Dihydrolipoyllysine-residue succinyltransferase component of 2-oxoglutarate dehyd
Arginine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=RARS PE=1 SV=2
Cluster of Cell division control protein 42 homolog OS=Homo sapiens GN=CDC42 PE=1
Actin-related protein 2 OS=Homo sapiens GN=ACTR2 PE=1 SV=1
A-kinase anchor protein 12 OS=Homo sapiens GN=AKAP12 PE=1 SV=4
Selenium-binding protein 1 OS=Homo sapiens GN=SELENBP1 PE=1 SV=2
Eukaryotic initiation factor 4A-II OS=Homo sapiens GN=EIF4A2 PE=1 SV=2
Non-POU domain-containing octamer-binding protein OS=Homo sapiens GN=NONO PE=1 SV
Cluster of Reticulon-1 OS=Homo sapiens GN=RTN1 PE=1 SV=1 (Q16799)
Protein NDRG1 OS=Homo sapiens GN=NDRG1 PE=1 SV=1
3-hydroxybutyrate dehydrogenase type 2 OS=Homo sapiens GN=BDH2 PE=1 SV=2
Coronin-2B OS=Homo sapiens GN=CORO2B PE=2 SV=4
HLA class II histocompatibility antigen, DR alpha chain OS=Homo sapiens GN=HLA-DR
Plastin-3 OS=Homo sapiens GN=PLS3 PE=2 SV=1
ATP synthase gamma chain OS=Homo sapiens GN=ATP5C1 PE=2 SV=1
Elongation factor 1-gamma OS=Homo sapiens GN=EEF1G PE=2 SV=1
Cluster of Serine/arginine-rich-splicing factor 3 OS=Homo sapiens GN=SRSF3 PE=2 S
Myosin light polypeptide 6 OS=Homo sapiens GN=MYL6 PE=2 SV=1
Cluster of Ras-related protein Rab-1A OS=Homo sapiens GN=RAB1A PE=2 SV=1 (B7Z8M7)
Apolipoprotein D (Fragment) OS=Homo sapiens GN=APOD PE=4 SV=1
26S proteasome non-ATPase regulatory subunit 2 OS=Homo sapiens GN=PSMD2 PE=4 SV=1
Elongation factor 1-delta (Fragment) OS=Homo sapiens GN=EEF1D PE=3 SV=1
Myosin-14 OS=Homo sapiens GN=MYH14 PE=4 SV=1
Cluster of Spliceosome RNA helicase DDX39B OS=Homo sapiens GN=DDX39B PE=4 SV=1 (F
Adenylate kinase 2, mitochondrial OS=Homo sapiens GN=AK2 PE=3 SV=1
NADH dehydrogenase (Ubiquinone) flavoprotein 1, 51kDa, isoform CRA_c OS=Homo sapi
Paraoxonase 2, isoform CRA_a OS=Homo sapiens GN=PON2 PE=4 SV=1
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 5 (Fragment) OS=Homo s
6-phosphofructokinase OS=Homo sapiens GN=PFKM PE=3 SV=1

Asparagine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=NARS PE=1 SV=1
Gamma-synuclein OS=Homo sapiens GN=SNCG PE=1 SV=2
Glutaminase kidney isoform, mitochondrial OS=Homo sapiens GN=GLS PE=1 SV=1
Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1
Cluster of Ras-related protein Ral-A OS=Homo sapiens GN=RALA PE=1 SV=1 (P11233)
Neuromodulin OS=Homo sapiens GN=GAP43 PE=1 SV=1
Trifunctional purine biosynthetic protein adenosine-3 OS=Homo sapiens GN=GART PE=
Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1 PE=1 SV=1
Catenin alpha-1 OS=Homo sapiens GN=CTNNA1 PE=1 SV=1
Serpins B6 OS=Homo sapiens GN=SERPINB6 PE=1 SV=3
ATP synthase subunit O, mitochondrial OS=Homo sapiens GN=ATP50 PE=1 SV=1
Myotrophin OS=Homo sapiens GN=MTPN PE=1 SV=2
CD81 antigen OS=Homo sapiens GN=CD81 PE=1 SV=1
Nuclear transport factor 2 OS=Homo sapiens GN=NUTF2 PE=1 SV=1
Brain acid soluble protein 1 OS=Homo sapiens GN=BASP1 PE=1 SV=2
116 kDa U5 small nuclear ribonucleoprotein component OS=Homo sapiens GN=EFTUD2 PE
3-hydroxyacyl-CoA dehydrogenase type-2 OS=Homo sapiens GN=HSD17B10 PE=1 SV=3
Endoplasmic reticulum resident protein 44 OS=Homo sapiens GN=ERP44 PE=1 SV=1
Band 4.1-like protein 1 OS=Homo sapiens GN=EPB41L1 PE=1 SV=2
CDGSH iron-sulfur domain-containing protein 1 OS=Homo sapiens GN=CISD1 PE=1 SV=1
Nck-associated protein 1 OS=Homo sapiens GN=NCKAP1 PE=1 SV=1
Talin-2 OS=Homo sapiens GN=TLN2 PE=1 SV=4
Chromobox protein homolog 3 OS=Homo sapiens GN=CBX3 PE=1 SV=4
Cluster of Ras-related protein Rap-1b-like protein OS=Homo sapiens PE=2 SV=1 (A6N
Biglycan OS=Homo sapiens GN=BGN PE=2 SV=1
Valine--tRNA ligase OS=Homo sapiens GN=VAR5 PE=3 SV=1
Calcium-binding mitochondrial carrier protein Aralar1 OS=Homo sapiens GN=SLC25A12
Nucleosome assembly protein 1-like 1 OS=Homo sapiens GN=NAP1L1 PE=2 SV=1
Annexin OS=Homo sapiens GN=ANXA7 PE=2 SV=1
Annexin OS=Homo sapiens GN=ANXA11 PE=2 SV=1
ITIH4 protein OS=Homo sapiens GN=ITIH4 PE=2 SV=1
Profilin-2 OS=Homo sapiens GN=PFN2 PE=4 SV=1
Tripartite motif-containing 2, isoform CRA_a OS=Homo sapiens GN=TRIM2 PE=4 SV=1
NADH dehydrogenase [ubiquinone] flavoprotein 2, mitochondrial OS=Homo sapiens GN=
Nesprin-1 OS=Homo sapiens GN=SYNE1 PE=4 SV=2
Myelin-associated glycoprotein OS=Homo sapiens GN=MAG PE=4 SV=1
Inosine-5'-monophosphate dehydrogenase 2 (Fragment) OS=Homo sapiens GN=IMPDH2 PE=
Hydroxyacyl-coenzyme A dehydrogenase, mitochondrial OS=Homo sapiens GN=HADH PE=4
Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1
Heterogeneous nuclear ribonucleoprotein R OS=Homo sapiens GN=HNRNPR PE=1 SV=1
Antithrombin-III OS=Homo sapiens GN=SERPINC1 PE=1 SV=1
Propionyl-CoA carboxylase alpha chain, mitochondrial OS=Homo sapiens GN=PCCA PE=1
Protein S100-A6 OS=Homo sapiens GN=S100A6 PE=1 SV=1
Spectrin beta chain, erythrocytic OS=Homo sapiens GN=SPTB PE=1 SV=5
Pyruvate carboxylase, mitochondrial OS=Homo sapiens GN=PC PE=1 SV=2
Sodium/potassium-transporting ATPase subunit beta-2 OS=Homo sapiens GN=ATP1B2 PE=
Calpain-2 catalytic subunit OS=Homo sapiens GN=CAPN2 PE=1 SV=6

Cadherin-2 OS=Homo sapiens GN=CDH2 PE=1 SV=4
Tryptophan--tRNA ligase, cytoplasmic OS=Homo sapiens GN=WARS PE=1 SV=2
14-3-3 protein sigma OS=Homo sapiens GN=SFN PE=1 SV=1
Beta-adducin OS=Homo sapiens GN=ADD2 PE=1 SV=3
Platelet-activating factor acetylhydrolase IB subunit alpha OS=Homo sapiens GN=PA
Ribose-phosphate pyrophosphokinase 1 OS=Homo sapiens GN=PRPS1 PE=1 SV=2
Ras-related protein Rab-10 OS=Homo sapiens GN=RAB10 PE=1 SV=1
Syntaxin-1B OS=Homo sapiens GN=STX1B PE=2 SV=1
Peptidyl-prolyl cis-trans isomerase FKBP4 OS=Homo sapiens GN=FKBP4 PE=1 SV=3
Isoform 2 of Src substrate cortactin OS=Homo sapiens GN=CTTN
Immunoglobulin superfamily member 8 OS=Homo sapiens GN=IGSF8 PE=1 SV=1
Cluster of Aminopeptidase B OS=Homo sapiens GN=RNPEP PE=1 SV=2 (Q9H4A4)
Sideroflexin-1 OS=Homo sapiens GN=SFXN1 PE=1 SV=4
3-hydroxyacyl-CoA dehydratase 3 OS=Homo sapiens GN=PTPLAD1 PE=1 SV=2
Glyoxylate reductase/hydroxypyruvate reductase OS=Homo sapiens GN=GRHPR PE=1 SV=1
Nuclear migration protein nudC OS=Homo sapiens GN=NUDC PE=1 SV=1
Cofilin-2 OS=Homo sapiens GN=CFL2 PE=1 SV=1
Lactoylglutathione lyase OS=Homo sapiens GN=GL01 PE=1 SV=4
Acyl-CoA-binding protein OS=Homo sapiens GN=DBI PE=1 SV=2
Reticulon-3 OS=Homo sapiens GN=RTN3 PE=4 SV=1
Coronin OS=Homo sapiens GN=COR01C PE=2 SV=1
Excitatory amino acid transporter 1 OS=Homo sapiens GN=SLC1A3 PE=2 SV=1
3' (2'), 5'-bisphosphate nucleotidase 1 OS=Homo sapiens GN=BPNT1 PE=2 SV=1
Medium-chain-specific acyl-CoA dehydrogenase, mitochondrial OS=Homo sapiens GN=AC
FERM, RhoGEF and pleckstrin domain-containing protein 1 OS=Homo sapiens GN=FARP1
Actin-related protein 2/3 complex subunit 4 OS=Homo sapiens GN=ARPC4 PE=4 SV=1
Catenin delta-1 OS=Homo sapiens GN=CTNND1 PE=4 SV=2
Dystonin OS=Homo sapiens GN=DST PE=4 SV=1
Phosphorylase OS=Homo sapiens GN=PYGL PE=3 SV=1
Ubiquitin-conjugating enzyme E2 N OS=Homo sapiens GN=UBE2N PE=3 SV=1
ATPase, H⁺ transporting, lysosomal 50/57kDa, V1 subunit H, isoform CRA_c OS=Homo
ATP synthase-coupling factor 6, mitochondrial OS=Homo sapiens GN=ATP5J PE=3 SV=1
Fatty aldehyde dehydrogenase OS=Homo sapiens GN=ALDH3A2 PE=3 SV=1
Na(+)/H(+) exchange regulatory cofactor NHE-RF1 OS=Homo sapiens GN=SLC9A3R1 PE=1
Laminin subunit alpha-5 OS=Homo sapiens GN=LAMA5 PE=1 SV=8
Neural cell adhesion molecule 2 OS=Homo sapiens GN=NCAM2 PE=1 SV=2
Fatty acid-binding protein, brain OS=Homo sapiens GN=FABP7 PE=1 SV=3
Isocitrate dehydrogenase [NADP] cytoplasmic OS=Homo sapiens GN=IDH1 PE=1 SV=2
Apoptosis-inducing factor 1, mitochondrial OS=Homo sapiens GN=AIFM1 PE=1 SV=1
Beta-2-glycoprotein 1 OS=Homo sapiens GN=APOH PE=1 SV=3
Alpha-1B-glycoprotein OS=Homo sapiens GN=A1BG PE=1 SV=4
Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2 OS=Homo
Glucose-6-phosphate 1-dehydrogenase OS=Homo sapiens GN=G6PD PE=1 SV=4
Cytochrome c oxidase subunit 4 isoform 1, mitochondrial OS=Homo sapiens GN=COX4I1
Cytochrome c oxidase subunit 5A, mitochondrial OS=Homo sapiens GN=COX5A PE=1 SV=2
Proteasome subunit alpha type-1 OS=Homo sapiens GN=PSMA1 PE=1 SV=1
Flavin reductase (NADPH) OS=Homo sapiens GN=BLVRB PE=1 SV=3

ATP synthase subunit delta, mitochondrial OS=Homo sapiens GN=ATP5D PE=1 SV=2
Cytochrome b-c1 complex subunit Rieske, mitochondrial OS=Homo sapiens GN=UQCRCFS1
Serine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=SARS PE=1 SV=3
Glycine amidinotransferase, mitochondrial OS=Homo sapiens GN=GATM PE=1 SV=1
Cluster of Alpha-actinin OS=Homo sapiens GN=ACTR1A PE=1 SV=1 (P61163)
Interleukin enhancer-binding factor 2 OS=Homo sapiens GN=ILF2 PE=1 SV=2
Delta(3,5)-Delta(2,4)-dienoyl-CoA isomerase, mitochondrial OS=Homo sapiens GN=ECH
DNA damage-binding protein 1 OS=Homo sapiens GN=DDB1 PE=1 SV=1
Cell adhesion molecule 2 OS=Homo sapiens GN=CADM2 PE=2 SV=1
Gamma-soluble NSF attachment protein OS=Homo sapiens GN=NAPG PE=1 SV=1
GTP:AMP phosphotransferase, mitochondrial OS=Homo sapiens GN=AK3 PE=1 SV=4
RuvB-like 1 OS=Homo sapiens GN=RUVBL1 PE=1 SV=1
SAM domain and HD domain-containing protein 1 OS=Homo sapiens GN=SAMHD1 PE=1 SV=2
S-formylglutathione hydrolase OS=Homo sapiens GN=ESD PE=1 SV=2
Alpha-2-HS-glycoprotein chain A OS=Homo sapiens GN=AHSG PE=4 SV=1
Alpha-1-syntrophin OS=Homo sapiens GN=SNTA1 PE=1 SV=1
26S protease regulatory subunit 8 OS=Homo sapiens GN=PSMC5 PE=2 SV=1
HCG2002594, isoform CRA_a OS=Homo sapiens GN=SEPT5 PE=2 SV=1
Heterogeneous nuclear ribonucleoprotein D0 OS=Homo sapiens GN=HNRNPD PE=2 SV=1
Coatomer protein complex, subunit beta 2 (Beta prime), isoform CRA_b OS=Homo sapiens
Cytochrome c (Fragment) OS=Homo sapiens GN=CYCS PE=3 SV=1
Polyadenylate-binding protein 1 OS=Homo sapiens GN=PABPC1 PE=4 SV=1
Xaa-Pro dipeptidase OS=Homo sapiens GN=PEPD PE=3 SV=1
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 8 OS=Homo sapiens GN=N
Methyltransferase-like protein 7A (Fragment) OS=Homo sapiens GN=METTTL7A PE=4 SV=1
Endoplasmic reticulum resident protein 29 OS=Homo sapiens GN=ERP29 PE=4 SV=1
Peripheral plasma membrane protein CASK OS=Homo sapiens GN=CASK PE=4 SV=1
ATP-dependent dihydroxyacetone kinase (Fragment) OS=Homo sapiens GN=DAK PE=4 SV=1
NADH dehydrogenase [ubiquinone] iron-sulfur protein 2, mitochondrial OS=Homo sapiens
Obg-like ATPase 1 OS=Homo sapiens GN=OLA1 PE=4 SV=1
Cluster of Tyrosine-protein kinase Yes OS=Homo sapiens GN=YES1 PE=4 SV=1 (J3QRU1)
Protein NipSnap homolog 2 OS=Homo sapiens GN=GBAS PE=1 SV=1
NADH dehydrogenase [ubiquinone] iron-sulfur protein 6, mitochondrial OS=Homo sapiens
Flotillin-1 OS=Homo sapiens GN=FLOT1 PE=1 SV=3
Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1
C4b-binding protein alpha chain OS=Homo sapiens GN=C4BPA PE=1 SV=2
Cluster of 60S acidic ribosomal protein P2 OS=Homo sapiens GN=RPLP2 PE=1 SV=1 (P0
Apolipoprotein A-IV OS=Homo sapiens GN=APOA4 PE=1 SV=3
Clathrin light chain A OS=Homo sapiens GN=CLTA PE=1 SV=1
Nidogen-1 OS=Homo sapiens GN=NID1 PE=1 SV=3
Proteasome subunit beta type-1 OS=Homo sapiens GN=PSMB1 PE=1 SV=2
Signal transducer and activator of transcription 1-alpha/beta OS=Homo sapiens GN=
Lumican OS=Homo sapiens GN=LUM PE=1 SV=2
Coatomer subunit beta OS=Homo sapiens GN=COPB1 PE=1 SV=3
Exportin-2 OS=Homo sapiens GN=CSE1L PE=1 SV=3
Hemoglobin subunit gamma-1 OS=Homo sapiens GN=HBG1 PE=1 SV=2
Proteasome activator complex subunit 1 OS=Homo sapiens GN=PSME1 PE=1 SV=1

Serine/threonine-protein kinase PAK 2 OS=Homo sapiens GN=PAK2 PE=1 SV=3
Isoform 2B of Cytoplasmic dynein 1 intermediate chain 2 OS=Homo sapiens GN=DYNC11
Coactosin-like protein OS=Homo sapiens GN=COTL1 PE=1 SV=3
Guanine nucleotide-binding protein subunit alpha-13 OS=Homo sapiens GN=GNA13 PE=1
Protein phosphatase 1 regulatory subunit 7 OS=Homo sapiens GN=PPP1R7 PE=1 SV=1
Guanine nucleotide-binding protein G(s) subunit alpha isoforms XLas OS=Homo sapie
Hydroxysteroid dehydrogenase-like protein 2 OS=Homo sapiens GN=HSDL2 PE=1 SV=1
Isoform 3 of Protein AHNAK2 OS=Homo sapiens GN=AHNAK2
Ermin OS=Homo sapiens GN=ERMN PE=2 SV=1
Cluster of Transportin-1 OS=Homo sapiens GN=TNP01 PE=1 SV=2 (Q92973)
Vacuolar protein sorting-associated protein 35 OS=Homo sapiens GN=VPS35 PE=1 SV=2
Beta-soluble NSF attachment protein OS=Homo sapiens GN=NAPB PE=1 SV=2
Isoform 2 of Glyoxalase domain-containing protein 4 OS=Homo sapiens GN=GLOD4
Electrogenic sodium bicarbonate cotransporter 1 OS=Homo sapiens GN=SLC4A4 PE=1 SV
Prostaglandin reductase 1 OS=Homo sapiens GN=PTGR1 PE=1 SV=2
Cluster of 40S ribosomal protein S15a OS=Homo sapiens GN=RPS15A PE=4 SV=1 (I3L3P7
Matrin-3 OS=Homo sapiens GN=MATR3 PE=4 SV=1
Cluster of Fibulin 1 (Fragment) OS=Homo sapiens GN=FBLN1 PE=4 SV=1 (B1AHM9)
NAD(P)H dehydrogenase [quinone] 1 OS=Homo sapiens GN=NQ01 PE=2 SV=1
Dynamin-like 120 kDa protein, mitochondrial OS=Homo sapiens GN=OPA1 PE=3 SV=1
Transcription elongation factor B polypeptide 1 (Fragment) OS=Homo sapiens GN=TCE
Catechol O-methyltransferase (Fragment) OS=Homo sapiens GN=COMT PE=4 SV=1
Dystrophin OS=Homo sapiens GN=DMD PE=4 SV=1
Microtubule-associated protein RP/EB family member 2 OS=Homo sapiens GN=MAPRE2 PE
Copine-6 OS=Homo sapiens GN=CPNE6 PE=4 SV=1
V-type proton ATPase 116 kDa subunit a isoform 1 OS=Homo sapiens GN=ATP6V0A1 PE=4
Heat shock protein 75 kDa, mitochondrial OS=Homo sapiens GN=TRAP1 PE=3 SV=1
Isoleucine--tRNA ligase, mitochondrial OS=Homo sapiens GN=IARS2 PE=3 SV=1
Proteasome subunit alpha type OS=Homo sapiens GN=PSMA6 PE=3 SV=1
Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=4 SV=1
Transforming growth factor-beta-induced protein ig-h3 OS=Homo sapiens GN=TGFBI PE
Inositol monophosphatase 1 (Fragment) OS=Homo sapiens GN=IMPA1 PE=4 SV=1
Isovaleryl Coenzyme A dehydrogenase, isoform CRA_a OS=Homo sapiens GN=IVD PE=3 SV
ATP-dependent RNA helicase DDX3X OS=Homo sapiens GN=DDX3X PE=1 SV=3
Dihydropyrimidinase-related protein 4 OS=Homo sapiens GN=DPYSL4 PE=1 SV=2
Heterogeneous nuclear ribonucleoprotein D-like OS=Homo sapiens GN=HNRPDL PE=1 SV=
Plexin-B1 OS=Homo sapiens GN=PLXNB1 PE=1 SV=3
Aflatoxin B1 aldehyde reductase member 2 OS=Homo sapiens GN=AKR7A2 PE=1 SV=3
Cluster of Core histone macro-H2A.1 OS=Homo sapiens GN=H2AFY PE=1 SV=4 (075367)
Cytosolic 10-formyltetrahydrofolate dehydrogenase OS=Homo sapiens GN=ALDH1L1 PE=1
Apolipoprotein A-II OS=Homo sapiens GN=APOA2 PE=1 SV=1
Integrin beta-1 OS=Homo sapiens GN=ITGB1 PE=1 SV=2
Synaptophysin OS=Homo sapiens GN=SYP PE=1 SV=3
Pyruvate dehydrogenase E1 component subunit alpha, somatic form, mitochondrial OS
Collagen alpha-2(VI) chain OS=Homo sapiens GN=COL6A2 PE=1 SV=4
V-type proton ATPase subunit C 1 OS=Homo sapiens GN=ATP6V1C1 PE=1 SV=4
Cytoplasmic aconitate hydratase OS=Homo sapiens GN=ACO1 PE=1 SV=3

Cluster of NAD-dependent malic enzyme, mitochondrial OS=Homo sapiens GN=ME2 PE=1
Laminin subunit alpha-1 OS=Homo sapiens GN=LAMA1 PE=1 SV=2
Glycogen debranching enzyme OS=Homo sapiens GN=AGL PE=1 SV=3
60S ribosomal protein L4 OS=Homo sapiens GN=RPL4 PE=1 SV=5
Eukaryotic initiation factor 4A-III OS=Homo sapiens GN=EIF4A3 PE=1 SV=4
Leucine-rich PPR motif-containing protein, mitochondrial OS=Homo sapiens GN=LRPPR
60S ribosomal protein L5 OS=Homo sapiens GN=RPL5 PE=1 SV=3
Serpine H1 OS=Homo sapiens GN=SERPINH1 PE=1 SV=2
Isocitrate dehydrogenase [NAD] subunit gamma, mitochondrial OS=Homo sapiens GN=ID
B-cell receptor-associated protein 31 OS=Homo sapiens GN=BCAP31 PE=1 SV=3
Ribonuclease UK114 OS=Homo sapiens GN=HRSP12 PE=1 SV=1
Succinyl-CoA ligase [ADP/GDP-forming] subunit alpha, mitochondrial OS=Homo sapien
Voltage-dependent calcium channel subunit alpha-2/delta-1 OS=Homo sapiens GN=CACN
Ras-related protein Rab-14 OS=Homo sapiens GN=RAB14 PE=1 SV=4
Cluster of Serine/threonine-protein phosphatase PP1-beta catalytic subunit OS=Hom
Single-stranded DNA-binding protein, mitochondrial OS=Homo sapiens GN=SSBP1 PE=1
Cytoskeleton-associated protein 5 OS=Homo sapiens GN=CKAP5 PE=1 SV=3
Inorganic pyrophosphatase OS=Homo sapiens GN=PPA1 PE=1 SV=2
Protein NipSnap homolog 1 OS=Homo sapiens GN=NIPSNAP1 PE=1 SV=1
Hyaluronan and proteoglycan link protein 2 OS=Homo sapiens GN=HAPLN2 PE=1 SV=1
Cluster of Ethylmalonyl-CoA decarboxylase OS=Homo sapiens GN=ECHDC1 PE=1 SV=2 (Q9
Endoplasmic reticulum aminopeptidase 1 OS=Homo sapiens GN=ERAP1 PE=1 SV=3
Cluster of Microtubule-associated protein RP/EB family member 3 OS=Homo sapiens G
Calpain-5 OS=Homo sapiens GN=CAPN5 PE=4 SV=2
Mannose-6-phosphate isomerase OS=Homo sapiens GN=MPI PE=3 SV=1
Cytochrome c oxidase subunit 2 OS=Homo sapiens GN=MT-CO2 PE=1 SV=1
Plastin-2 OS=Homo sapiens GN=LCP1 PE=1 SV=6
Oligodendrocyte-myelin glycoprotein OS=Homo sapiens GN=OMG PE=1 SV=2
Receptor expression-enhancing protein 5 OS=Homo sapiens GN=REEP5 PE=1 SV=3
Lin-7 homolog C (C. elegans), isoform CRA_b OS=Homo sapiens GN=LIN7C PE=4 SV=1
High mobility group protein B1 OS=Homo sapiens GN=HMGB1 PE=1 SV=3
Translationally-controlled tumor protein OS=Homo sapiens GN=TPT1 PE=1 SV=1
Rab3 GTPase-activating protein non-catalytic subunit OS=Homo sapiens GN=RAB3GAP2
40S ribosomal protein S20 OS=Homo sapiens GN=RPS20 PE=1 SV=1
Fumarylacetoacetate hydrolase domain-containing protein 2B OS=Homo sapiens GN=FAH
Lysosome membrane protein 2 OS=Homo sapiens GN=SCARB2 PE=1 SV=2
Monocarboxylate transporter 1 OS=Homo sapiens GN=SLC16A1 PE=1 SV=3
Peptidyl-prolyl cis-trans isomerase-like 3 OS=Homo sapiens GN=PPIL3 PE=3 SV=1
Ubiquitin-like modifier-activating enzyme 6 OS=Homo sapiens GN=UBA6 PE=1 SV=1
Drebrin-like protein OS=Homo sapiens GN=DBNL PE=2 SV=1
Macrophage-capping protein OS=Homo sapiens GN=CAPG PE=2 SV=1
Purine nucleoside phosphorylase A OS=Homo sapiens GN=MTAP PE=2 SV=1
Erythrocyte band 7 integral membrane protein OS=Homo sapiens GN=STOM PE=2 SV=1
Acetyl-CoA acetyltransferase, cytosolic OS=Homo sapiens GN=ACAT2 PE=2 SV=1
Aldehyde dehydrogenase, dimeric NADP-preferring (Fragment) OS=Homo sapiens GN=ALD
Guanine nucleotide-binding protein subunit beta-2-like 1 OS=Homo sapiens GN=GNB2L
Dihydrolipoyllysine-residue acetyltransferase component of pyruvate dehydrogenase

General vesicular transport factor p115 OS=Homo sapiens GN=USO1 PE=4 SV=1
Cluster of Ras-related protein Rab-6A OS=Homo sapiens GN=RAB6A PE=3 SV=1 (F5H668)
Proliferation-associated protein 2G4 OS=Homo sapiens GN=PA2G4 PE=4 SV=1
Unconventional myosin-Va OS=Homo sapiens GN=MYO5A PE=4 SV=1
X-prolyl aminopeptidase (Aminopeptidase P) 1, soluble, isoform CRA_b OS=Homo sapiens GN=AMPEP1 PE=4 SV=1
Cluster of Importin-5 (Fragment) OS=Homo sapiens GN=IPO5 PE=4 SV=1 (HOY8C6)
Fatty acid-binding protein, epidermal OS=Homo sapiens GN=FABP5 PE=4 SV=1
26S proteasome non-ATPase regulatory subunit 3 OS=Homo sapiens GN=PSMD3 PE=1 SV=2
DnaJ homolog subfamily A member 2 OS=Homo sapiens GN=DNAJA2 PE=1 SV=1
NADH dehydrogenase [ubiquinone] iron-sulfur protein 3, mitochondrial OS=Homo sapiens GN=ND3 PE=1 SV=1
Isoform 2 of Nebulette OS=Homo sapiens GN=NEBL
Purine nucleoside phosphorylase OS=Homo sapiens GN=PNP PE=1 SV=2
Cystatin-C OS=Homo sapiens GN=CST3 PE=1 SV=1
Alpha-1-acid glycoprotein 1 OS=Homo sapiens GN=ORM1 PE=1 SV=1
Cytochrome c1, heme protein, mitochondrial OS=Homo sapiens GN=CYC1 PE=1 SV=3
cAMP-dependent protein kinase type II-alpha regulatory subunit OS=Homo sapiens GN=PKRII PE=1 SV=1
Gap junction alpha-1 protein OS=Homo sapiens GN=GJA1 PE=1 SV=2
5'-nucleotidase OS=Homo sapiens GN=NT5E PE=1 SV=1
Enoyl-CoA hydratase, mitochondrial OS=Homo sapiens GN=ECHS1 PE=1 SV=4
Keratin, type I cytoskeletal 9 OS=Homo sapiens GN=KRT9 PE=1 SV=3
Chitinase-3-like protein 1 OS=Homo sapiens GN=CHI3L1 PE=1 SV=2
Acyl-coenzyme A thioesterase 2, mitochondrial OS=Homo sapiens GN=ACOT2 PE=1 SV=6
Heterogeneous nuclear ribonucleoprotein F OS=Homo sapiens GN=HNRNPF PE=1 SV=3
NEDD8-conjugating enzyme Ubc12 OS=Homo sapiens GN=UBE2M PE=1 SV=1
Cluster of Transforming protein RhoA OS=Homo sapiens GN=RHOA PE=1 SV=1 (P61586)
Dual specificity mitogen-activated protein kinase kinase 1 OS=Homo sapiens GN=MAP2K1 PE=1 SV=1
Rho GTPase-activating protein 1 OS=Homo sapiens GN=ARHGAP1 PE=1 SV=1
Leukocyte surface antigen CD47 OS=Homo sapiens GN=CD47 PE=1 SV=1
EGF-containing fibulin-like extracellular matrix protein 1 OS=Homo sapiens GN=EFE1 PE=1 SV=1
Transcription intermediary factor 1-beta OS=Homo sapiens GN=TRIM28 PE=1 SV=5
Calcium/calmodulin-dependent protein kinase type II subunit beta OS=Homo sapiens GN=PKCII PE=1 SV=1
Eukaryotic translation initiation factor 3 subunit A OS=Homo sapiens GN=EIF3A PE=1 SV=1
RAB18, member RAS oncogene family OS=Homo sapiens GN=RAB18 PE=2 SV=1
Isoform 2 of Choline transporter-like protein 2 OS=Homo sapiens GN=SLC44A2 PE=1 SV=1
Far upstream element-binding protein 3 OS=Homo sapiens GN=FUBP3 PE=1 SV=2
Extended synaptotagmin-1 OS=Homo sapiens GN=ESYT1 PE=1 SV=1
Guanine nucleotide-binding protein subunit beta-4 OS=Homo sapiens GN=GNB4 PE=1 SV=1
Heme-binding protein 1 OS=Homo sapiens GN=HEBP1 PE=1 SV=1
Ribosome-binding protein 1 OS=Homo sapiens GN=RRBP1 PE=1 SV=4
Alpha-aminoadipic semialdehyde synthase, mitochondrial OS=Homo sapiens GN=AASS PE=1 SV=1
Cluster of Ubiquilin-2 OS=Homo sapiens GN=UBQLN2 PE=1 SV=2 (Q9UHD9)
Pre-mRNA-processing factor 19 OS=Homo sapiens GN=PRPF19 PE=1 SV=1
IgG Fc-binding protein OS=Homo sapiens GN=FCGBP PE=1 SV=3
Clathrin light chain B OS=Homo sapiens GN=CLTB PE=1 SV=1
Septin 10, isoform CRA_c OS=Homo sapiens GN=SEPT10 PE=3 SV=2
Electron transfer flavoprotein-ubiquinone oxidoreductase, mitochondrial OS=Homo sapiens GN=ETF1B PE=1 SV=1
Shootin-1 OS=Homo sapiens GN=KIAA1598 PE=1 SV=4

Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2
Cathepsin B OS=Homo sapiens GN=CTSB PE=1 SV=3
HCG2044799 OS=Homo sapiens GN=hCG_2044799 PE=4 SV=1
26S proteasome non-ATPase regulatory subunit 11 OS=Homo sapiens GN=PSMD11 PE=1 SV=1
Prostaglandin E synthase 3 OS=Homo sapiens GN=PTGES3 PE=2 SV=1
CD9 antigen OS=Homo sapiens GN=CD9 PE=4 SV=1
Protein S100-A10 OS=Homo sapiens GN=S100A10 PE=1 SV=2
Cysteine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=CARS PE=2 SV=1
Syntaxin 12 (Fragment) OS=Homo sapiens GN=STX12 PE=4 SV=1
Protein kinase, cAMP-dependent, catalytic, beta OS=Homo sapiens GN=PRKACB PE=2 SV=1
Non-specific lipid-transfer protein OS=Homo sapiens GN=SCP2 PE=2 SV=1
ATP-dependent RNA helicase DDX19A OS=Homo sapiens GN=DDX19A PE=2 SV=1
Activator of 90 kDa heat shock protein ATPase homolog 1 OS=Homo sapiens GN=AHSA1
Eukaryotic translation initiation factor 3 subunit C OS=Homo sapiens GN=EIF3CL PE=1
Delta-aminolevulinic acid dehydratase OS=Homo sapiens GN=ALAD PE=2 SV=1
Cysteine-rich protein 2 OS=Homo sapiens GN=CRIP2 PE=2 SV=1
3-ketoacyl-CoA thiolase, peroxisomal OS=Homo sapiens GN=ACAA1 PE=3 SV=1
Coiled-coil-helix-coiled-coil-helix domain-containing protein 3, mitochondrial OS=Homo sapiens GN=EIF4G1 PE=4 SV=1
Eukaryotic translation initiation factor 4 gamma 1 OS=Homo sapiens GN=EIF4G1 PE=4 SV=1
Eukaryotic translation initiation factor 4 gamma 2 (Fragment) OS=Homo sapiens GN=EIF4G2 PE=4 SV=1
Cluster of CLIP-associating protein 2 OS=Homo sapiens GN=CLASP2 PE=4 SV=2 (E3W994)
Cluster of Kinesin light chain 1 OS=Homo sapiens GN=KLC1 PE=4 SV=1 (E7EVH7)
Integrin alpha-V light chain OS=Homo sapiens GN=ITGAV PE=3 SV=1
Protein arginine N-methyltransferase 1 OS=Homo sapiens GN=PRMT1 PE=4 SV=1
TOM1-like protein 2 OS=Homo sapiens GN=TOM1L2 PE=4 SV=2
SPARC (Fragment) OS=Homo sapiens GN=SPARC PE=4 SV=1
Cysteine and glycine-rich protein 2 OS=Homo sapiens GN=CSRP2 PE=4 SV=1
UPF0317 protein C14orf159, mitochondrial OS=Homo sapiens GN=C14orf159 PE=4 SV=1
Sorting nexin-1 OS=Homo sapiens GN=SNX1 PE=4 SV=1
Thiomorpholine-carboxylate dehydrogenase OS=Homo sapiens GN=CRYM PE=4 SV=1
Eukaryotic peptide chain release factor GTP-binding subunit ERF3A OS=Homo sapiens GN=ERF3A PE=4 SV=1
PDZ and LIM domain protein 1 OS=Homo sapiens GN=PDLIM1 PE=1 SV=4
LanC-like protein 1 OS=Homo sapiens GN=LANCL1 PE=1 SV=1
Sorting nexin-3 OS=Homo sapiens GN=SNX3 PE=1 SV=3
Vesicle-trafficking protein SEC22b OS=Homo sapiens GN=SEC22B PE=1 SV=4
Mitochondrial import receptor subunit TOM70 OS=Homo sapiens GN=TOMM70A PE=1 SV=1
LETM1 and EF-hand domain-containing protein 1, mitochondrial OS=Homo sapiens GN=LETM1 PE=1 SV=1
Alpha-2-antiplasmin OS=Homo sapiens GN=SERPINF2 PE=1 SV=3
Hyaluronan and proteoglycan link protein 1 OS=Homo sapiens GN=HAPLN1 PE=2 SV=2
40S ribosomal protein S2 OS=Homo sapiens GN=RPS2 PE=1 SV=2
Histone H1.5 OS=Homo sapiens GN=HIST1H1B PE=1 SV=3
Cluster of Ras-related protein Rab-3A OS=Homo sapiens GN=RAB3A PE=1 SV=1 (P20336)
Succinate dehydrogenase [ubiquinone] iron-sulfur subunit, mitochondrial OS=Homo sapiens GN=SDHA PE=1 SV=1
Proteasome subunit alpha type-3 OS=Homo sapiens GN=PSMA3 PE=1 SV=2
Threonine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=TARS PE=1 SV=3
Proteasome subunit alpha type-5 OS=Homo sapiens GN=PSMA5 PE=1 SV=3
60S ribosomal protein L12 OS=Homo sapiens GN=RPL12 PE=1 SV=1

Leukocyte elastase inhibitor OS=Homo sapiens GN=SERPINB1 PE=1 SV=1
Adenylyl cyclase-associated protein 2 OS=Homo sapiens GN=CAP2 PE=1 SV=1
Carnitine O-palmitoyltransferase 1, liver isoform OS=Homo sapiens GN=CPT1A PE=1 SV=1
Biliverdin reductase A OS=Homo sapiens GN=BLVRA PE=1 SV=2
26S protease regulatory subunit 4 OS=Homo sapiens GN=PSMC1 PE=1 SV=1
Growth factor receptor-bound protein 2 OS=Homo sapiens GN=GRB2 PE=1 SV=1
Contactin-associated protein 1 OS=Homo sapiens GN=CNTNAP1 PE=1 SV=1
Transcriptional activator protein Pur-alpha OS=Homo sapiens GN=PURA PE=1 SV=2
Transgelin OS=Homo sapiens GN=TAGLN PE=1 SV=4
Peptidyl-prolyl cis-trans isomerase D OS=Homo sapiens GN=PPID PE=1 SV=3
Heterogeneous nuclear ribonucleoprotein A0 OS=Homo sapiens GN=HNRNPA0 PE=1 SV=1
Cluster of Rho-associated protein kinase 1 OS=Homo sapiens GN=ROCK1 PE=1 SV=1 (Q1
Calcium/calmodulin-dependent protein kinase type II subunit gamma OS=Homo sapiens
Beta-1-syntrophin OS=Homo sapiens GN=SNTB1 PE=1 SV=3
Platelet-activating factor acetylhydrolase IB subunit gamma OS=Homo sapiens GN=PA
Beta-synuclein OS=Homo sapiens GN=SNCB PE=1 SV=1
Heterochromatin protein 1-binding protein 3 OS=Homo sapiens GN=HP1BP3 PE=1 SV=1
Twinfilin-2 OS=Homo sapiens GN=TWF2 PE=1 SV=2
Cell cycle exit and neuronal differentiation protein 1 OS=Homo sapiens GN=CEND1 P
Saccharopine dehydrogenase-like oxidoreductase OS=Homo sapiens GN=SCCPDH PE=1 SV=
NAD(P)H-hydrate epimerase OS=Homo sapiens GN=APOA1BP PE=1 SV=2
Cell adhesion molecule 4 OS=Homo sapiens GN=CADM4 PE=1 SV=1
BTB/POZ domain-containing protein KCTD12 OS=Homo sapiens GN=KCTD12 PE=1 SV=1
CB1 cannabinoid receptor-interacting protein 1 OS=Homo sapiens GN=CNRI1 PE=1 SV=
Phytanoyl-CoA hydroxylase-interacting protein-like OS=Homo sapiens GN=PHYHIPL PE=
Phosphoglucomutase-2 OS=Homo sapiens GN=PGM2 PE=1 SV=4
Tubulin-folding cofactor B OS=Homo sapiens GN=TBCB PE=1 SV=2
Proteasome subunit beta type-7 OS=Homo sapiens GN=PSMB7 PE=1 SV=1
Redox-regulatory protein FAM213A OS=Homo sapiens GN=FAM213A PE=1 SV=3
Transmembrane protein 43 OS=Homo sapiens GN=TMEM43 PE=1 SV=1
Mitochondrial glutamate carrier 1 OS=Homo sapiens GN=SLC25A22 PE=1 SV=1
Methylcrotonoyl-CoA carboxylase beta chain, mitochondrial OS=Homo sapiens GN=MCCC
Isoform 2 of ATPase family AAA domain-containing protein 3A OS=Homo sapiens GN=AT
LIM and cysteine-rich domains protein 1 OS=Homo sapiens GN=LMCD1 PE=1 SV=1
Vesicle-associated membrane protein-associated protein A OS=Homo sapiens GN=VAPA
NADH-cytochrome b5 reductase 1 OS=Homo sapiens GN=CYB5R1 PE=1 SV=1
Lambda-crystallin homolog OS=Homo sapiens GN=CRYL1 PE=1 SV=3
Cytochrome b-c1 complex subunit 7 OS=Homo sapiens GN=UQCRB PE=2 SV=1
Lupus La protein OS=Homo sapiens GN=SSB PE=4 SV=1
NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial (Fragment) O
Transthyretin OS=Homo sapiens GN=TTR PE=1 SV=1
Cluster of 6-phosphofructo-2-kinase/fructose-2, 6-bisphosphatase 2 transcript var
Thioredoxin domain-containing protein 12 OS=Homo sapiens GN=TXNDC12 PE=1 SV=1
Glycogenin-1 OS=Homo sapiens GN=GYG1 PE=1 SV=4
Histamine N-methyltransferase OS=Homo sapiens GN=HNMT PE=1 SV=1
Fermitin family homolog 2 (Fragment) OS=Homo sapiens GN=FERMT2 PE=4 SV=1
Histidine triad nucleotide-binding protein 1 OS=Homo sapiens GN=HINT1 PE=1 SV=2

Tripeptidyl-peptidase 1 OS=Homo sapiens GN=TPP1 PE=1 SV=2
Ubiquitin-like protein ISG15 OS=Homo sapiens GN=ISG15 PE=1 SV=5
60S ribosomal protein L23a OS=Homo sapiens GN=RPL23A PE=3 SV=1
Heterogeneous nuclear ribonucleoprotein A/B OS=Homo sapiens GN=HNRNPAB PE=4 SV=1
Programmed cell death protein 6 OS=Homo sapiens GN=PDCD6 PE=1 SV=1
26S proteasome non-ATPase regulatory subunit 14 OS=Homo sapiens GN=PSMD14 PE=1 SV=1
Dynein light chain 1, cytoplasmic OS=Homo sapiens GN=DYNLL1 PE=1 SV=1
Calpain small subunit 1 OS=Homo sapiens GN=CAPNS1 PE=1 SV=1
Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-7 OS=Homo sapiens
SH3 domain binding glutamic acid-rich protein like 3 OS=Homo sapiens GN=SH3BGRL3
AP-3 complex subunit delta-1 OS=Homo sapiens GN=AP3D1 PE=4 SV=1
Nucleosome assembly protein 1-like 4 (Fragment) OS=Homo sapiens GN=NAP1L4 PE=3 SV=1
Archain 1, isoform CRA_a OS=Homo sapiens GN=ARCN1 PE=4 SV=1
Cluster of Histidine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=HARS PE=2 SV=1
Sorting nexin-2 OS=Homo sapiens GN=SNX2 PE=2 SV=1
Solute carrier family 25 (Mitochondrial carrier; citrate transporter), member 1,
ELAV-like protein 1 OS=Homo sapiens GN=ELAVL1 PE=2 SV=1
Plasma protease C1 inhibitor OS=Homo sapiens GN=SERPING1 PE=2 SV=1
Stomatin-like protein 2 OS=Homo sapiens GN=STOML2 PE=2 SV=1
Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP7 PE=2 SV=1
Propionyl-CoA carboxylase beta chain, mitochondrial OS=Homo sapiens GN=PCCB PE=4
Catenin delta-2 OS=Homo sapiens GN=CTNND2 PE=4 SV=1
Heat shock 70 kDa protein 4L OS=Homo sapiens GN=HSPA4L PE=3 SV=2
Acylglycerol kinase, mitochondrial OS=Homo sapiens GN=AGK PE=4 SV=1
AP-2 complex subunit mu OS=Homo sapiens GN=AP2M1 PE=4 SV=1
Pyridoxal kinase OS=Homo sapiens GN=PDXK PE=4 SV=1
Doublecortin and CaM kinase-like 2, isoform CRA_c OS=Homo sapiens GN=DCLK2 PE=4 S
Heme oxygenase 2 OS=Homo sapiens GN=HMOX2 PE=4 SV=1
Cytochrome c oxidase subunit 6B1 OS=Homo sapiens GN=COX6B1 PE=4 SV=1
Serine/arginine-rich-splicing factor 1 OS=Homo sapiens GN=SRSF1 PE=4 SV=1
Chloride intracellular channel protein 1 OS=Homo sapiens GN=CLIC1 PE=1 SV=4
Plexin-B2 OS=Homo sapiens GN=PLXNB2 PE=1 SV=3
Syntaxin-7 OS=Homo sapiens GN=STX7 PE=1 SV=4
Actin-related protein 2/3 complex subunit 5 OS=Homo sapiens GN=ARPC5 PE=1 SV=3
Isocitrate dehydrogenase [NAD] subunit beta, mitochondrial OS=Homo sapiens GN=IDH
Periplakin OS=Homo sapiens GN=PPL PE=1 SV=4
Enoyl-CoA delta isomerase 2, mitochondrial OS=Homo sapiens GN=ECI2 PE=1 SV=4
U5 small nuclear ribonucleoprotein 200 kDa helicase OS=Homo sapiens GN=SNRNP200 P
ATP synthase subunit g, mitochondrial OS=Homo sapiens GN=ATP5L PE=1 SV=3
SEC14-like protein 2 OS=Homo sapiens GN=SEC14L2 PE=1 SV=1
Endonuclease domain-containing 1 protein OS=Homo sapiens GN=ENDOD1 PE=1 SV=2
Collagen alpha-1(IV) chain OS=Homo sapiens GN=COL4A1 PE=1 SV=3
60S ribosomal protein L7 OS=Homo sapiens GN=RPL7 PE=1 SV=1
Keratin, type II cytoskeletal 2 epidermal OS=Homo sapiens GN=KRT2 PE=1 SV=2
Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4
Glutamate--cysteine ligase catalytic subunit OS=Homo sapiens GN=GCLC PE=1 SV=2
Activated RNA polymerase II transcriptional coactivator p15 OS=Homo sapiens GN=SU

Epiplakin OS=Homo sapiens GN=EPPK1 PE=1 SV=2
Cluster of Liprin-alpha-1 OS=Homo sapiens GN=PPFIA1 PE=1 SV=1 (Q13136)
Nidogen-2 OS=Homo sapiens GN=NID2 PE=1 SV=3
Dystroglycan OS=Homo sapiens GN=DAG1 PE=1 SV=2
LIM and SH3 domain protein 1 OS=Homo sapiens GN=LASP1 PE=1 SV=2
Lysine--tRNA ligase OS=Homo sapiens GN=KARS PE=1 SV=3
Coiled-coil domain-containing protein 6 OS=Homo sapiens GN=CCDC6 PE=1 SV=2
GRIP1-associated protein 1 OS=Homo sapiens GN=GRIPAP1 PE=1 SV=1
Glucose 1,6-bisphosphate synthase OS=Homo sapiens GN=PGM2L1 PE=1 SV=3
Cluster of DBIRD complex subunit KIAA1967 OS=Homo sapiens GN=KIAA1967 PE=1 SV=2 (Actin-related protein 2/3 complex subunit 1A OS=Homo sapiens GN=ARPC1A PE=1 SV=2
26S proteasome non-ATPase regulatory subunit 1 OS=Homo sapiens GN=PSMD1 PE=1 SV=2
UBX domain-containing protein 6 OS=Homo sapiens GN=UBXN6 PE=1 SV=1
Prenylcysteine oxidase 1 OS=Homo sapiens GN=PCYOX1 PE=1 SV=3
Enolase-phosphatase E1 OS=Homo sapiens GN=ENOPH1 PE=1 SV=1
Transgelin-3 OS=Homo sapiens GN=TAGLN3 PE=1 SV=2
F-box only protein 2 OS=Homo sapiens GN=FBXO2 PE=1 SV=2
Cluster of Probable phospholipid-transporting ATPase IA OS=Homo sapiens GN=ATP8A1
Peptidyl-prolyl cis-trans isomerase-like 1 OS=Homo sapiens GN=PPIL1 PE=1 SV=1
Sorting nexin-5 OS=Homo sapiens GN=SNX5 PE=1 SV=1
Laminin subunit gamma-3 OS=Homo sapiens GN=LAMC3 PE=1 SV=3
Amphiphysin OS=Homo sapiens GN=AMPH PE=4 SV=1
Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=3 SV=1
Vesicle-associated membrane protein-associated protein B/C OS=Homo sapiens GN=VAP
Collagen alpha-1(XVIII) chain OS=Homo sapiens GN=COL18A1 PE=1 SV=5
Dystrobrevin alpha OS=Homo sapiens GN=DTNA PE=2 SV=1
Guanine nucleotide-binding protein subunit gamma OS=Homo sapiens GN=GNG2 PE=3 SV=3
Membrane-associated progesterone receptor component 1 OS=Homo sapiens GN=PGRMC1 P
Sialic acid synthase OS=Homo sapiens GN=NANS PE=1 SV=2
40S ribosomal protein S6 OS=Homo sapiens GN=RPS6 PE=1 SV=1
Cluster of Serine/arginine-rich splicing factor 4 OS=Homo sapiens GN=SRSF4 PE=1 S
Heterogeneous nuclear ribonucleoprotein H3 OS=Homo sapiens GN=HNRNPH3 PE=1 SV=2
Translin OS=Homo sapiens GN=TSN PE=4 SV=1
Calumenin OS=Homo sapiens GN=CALU PE=1 SV=2
Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1
S-adenosylmethionine synthase isoform type-2 OS=Homo sapiens GN=MAT2A PE=1 SV=1
Glycine--tRNA ligase OS=Homo sapiens GN=GARS PE=1 SV=3
Protein phosphatase methylesterase 1 OS=Homo sapiens GN=PPME1 PE=3 SV=1
Neutral amino acid transporter A OS=Homo sapiens GN=SLC1A4 PE=1 SV=1
Oxysterol-binding protein 1 OS=Homo sapiens GN=OSBP PE=1 SV=1
Rho-related GTP-binding protein RhoB OS=Homo sapiens GN=RHOB PE=1 SV=1
Hydroxymethylglutaryl-CoA synthase, cytoplasmic OS=Homo sapiens GN=HMGCS1 PE=1 SV=1
Microsomal glutathione S-transferase 1 OS=Homo sapiens GN=MGST1 PE=1 SV=1
Malectin (Fragment) OS=Homo sapiens GN=MLEC PE=4 SV=1
Peptidyl-prolyl cis-trans isomerase FKBP1A OS=Homo sapiens GN=FKBP1A PE=1 SV=2
Rho GDP-dissociation inhibitor 2 (Fragment) OS=Homo sapiens GN=ARHGDI2 PE=4 SV=1
Protein transport protein Sec24C OS=Homo sapiens GN=SEC24C PE=4 SV=2

Guanine nucleotide-binding protein G(z) subunit alpha OS=Homo sapiens GN=GNAZ PE=15 kDa selenoprotein OS=Homo sapiens GN=SEP15 PE=1 SV=3
SH3 domain-binding glutamic acid-rich-like protein OS=Homo sapiens GN=SH3BGRL PE=Microtubule-associated protein 6 OS=Homo sapiens GN=MAP6 PE=1 SV=2
Histidine triad nucleotide-binding protein 2, mitochondrial OS=Homo sapiens GN=HI
Estradiol 17-beta-dehydrogenase 12 OS=Homo sapiens GN=HSD17B12 PE=1 SV=2
Ubiquitin-conjugating enzyme E2 L3 OS=Homo sapiens GN=UBE2L3 PE=1 SV=1
Integrin alpha-M OS=Homo sapiens GN=ITGAM PE=1 SV=2
Synaptojanin-2-binding protein OS=Homo sapiens GN=SYNJ2BP PE=1 SV=2
Transmembrane emp24 domain-containing protein 9 OS=Homo sapiens GN=TMED9 PE=1 SV=40S ribosomal protein S27 OS=Homo sapiens GN=RPS27L PE=3 SV=1
Cluster of Cyclin-dependent kinase 5 OS=Homo sapiens GN=CDK5 PE=1 SV=3 (Q00535)
Vigilin OS=Homo sapiens GN=HDLBP PE=1 SV=2
Beta-2-microglobulin form pI 5.3 OS=Homo sapiens GN=B2M PE=3 SV=1
TRAF2 and NCK-interacting protein kinase OS=Homo sapiens GN=TNIK PE=4 SV=1
Acidic leucine-rich nuclear phosphoprotein 32 family member B OS=Homo sapiens GN=RELA protein OS=Homo sapiens GN=RELA PE=2 SV=1
Cyclin-Y OS=Homo sapiens GN=CCNY PE=2 SV=1
Zinc-alpha-2-glycoprotein OS=Homo sapiens GN=AZGP1 PE=3 SV=1
Breast carcinoma-amplified sequence 1 OS=Homo sapiens GN=BCAS1 PE=2 SV=1
2-oxoisovalerate dehydrogenase subunit alpha, mitochondrial OS=Homo sapiens GN=BC
Cold-inducible RNA-binding protein OS=Homo sapiens GN=CIRBP PE=2 SV=1
Glutathione synthetase OS=Homo sapiens GN=GSS PE=2 SV=1
60S ribosomal protein L23 (Fragment) OS=Homo sapiens GN=RPL23 PE=3 SV=1
Synaptojanin-1 OS=Homo sapiens GN=SYNJ1 PE=4 SV=1
Thymidine phosphorylase (Fragment) OS=Homo sapiens GN=TYMP PE=4 SV=1
Annexin OS=Homo sapiens GN=ANXA3 PE=3 SV=1
Tubulin-specific chaperone A OS=Homo sapiens GN=TBCA PE=4 SV=1
MAGUK p55 subfamily member 2 OS=Homo sapiens GN=MPP2 PE=4 SV=1
Interferon-induced, double-stranded RNA-activated protein kinase OS=Homo sapiens
CD82 antigen (Fragment) OS=Homo sapiens GN=CD82 PE=4 SV=1
Tight junction protein ZO-2 OS=Homo sapiens GN=TJP2 PE=4 SV=1
Limbic system-associated membrane protein OS=Homo sapiens GN=LSAMP PE=4 SV=1
Active breakpoint cluster region-related protein OS=Homo sapiens GN=ABR PE=4 SV=1
Dynamin-1-like protein OS=Homo sapiens GN=DNM1L PE=3 SV=1
E3 ubiquitin-protein ligase HUWE1 (Fragment) OS=Homo sapiens GN=HUWE1 PE=4 SV=1
NSFL1 cofactor p47 OS=Homo sapiens GN=NSFL1C PE=4 SV=1
Synaptosomal-associated protein 23 OS=Homo sapiens GN=SNAP23 PE=1 SV=1
Exportin-T OS=Homo sapiens GN=XPOT PE=1 SV=2
ATPase ASNA1 OS=Homo sapiens GN=ASNA1 PE=1 SV=2
PC4 and SFRS1-interacting protein OS=Homo sapiens GN=PSIP1 PE=1 SV=1
Heat shock factor-binding protein 1 OS=Homo sapiens GN=HSBP1 PE=1 SV=1
Interferon-inducible double stranded RNA-dependent protein kinase activator A OS=
Kelch repeat and BTB domain-containing protein 11 OS=Homo sapiens GN=KBTBD11 PE=1
Complement C1q subcomponent subunit C OS=Homo sapiens GN=C1QC PE=1 SV=3
Carnitine O-palmitoyltransferase 2, mitochondrial OS=Homo sapiens GN=CPT2 PE=1 SV
Adenylate kinase isoenzyme 4, mitochondrial OS=Homo sapiens GN=AK4 PE=1 SV=1

3-hydroxyisobutyrate dehydrogenase, mitochondrial OS=Homo sapiens GN=HIBADH PE=1
Serine hydroxymethyltransferase, mitochondrial OS=Homo sapiens GN=SHMT2 PE=1 SV=3
Cluster of Ran GTPase-activating protein 1 OS=Homo sapiens GN=RANGAP1 PE=1 SV=1 (Glutamine--tRNA ligase OS=Homo sapiens GN=QARS PE=1 SV=1
Eukaryotic translation initiation factor 5 OS=Homo sapiens GN=EIF5 PE=1 SV=2
40S ribosomal protein S8 OS=Homo sapiens GN=RPS8 PE=1 SV=2
60S ribosomal protein L31 OS=Homo sapiens GN=RPL31 PE=1 SV=1
Nuclease-sensitive element-binding protein 1 OS=Homo sapiens GN=YBX1 PE=1 SV=3
Twinfilin-1 OS=Homo sapiens GN=TWF1 PE=1 SV=3
Aminoacyl tRNA synthase complex-interacting multifunctional protein 1 OS=Homo sap
Peptidyl-prolyl cis-trans isomerase NIMA-interacting 1 OS=Homo sapiens GN=PIN1 PE
Thiosulfate sulfurtransferase OS=Homo sapiens GN=TST PE=1 SV=4
NAD kinase domain-containing protein 1 OS=Homo sapiens GN=NADKD1 PE=1 SV=2
Ectonucleotide pyrophosphatase/phosphodiesterase family member 6 OS=Homo sapiens
Leucine-rich repeat-containing protein 47 OS=Homo sapiens GN=LRRC47 PE=1 SV=1
Phosphatidylinositol 3,4,5-trisphosphate-dependent Rac exchanger 1 protein OS=Hom
Glucosamine-6-phosphate isomerase 2 OS=Homo sapiens GN=GNPDA2 PE=1 SV=1
Choline transporter-like protein 1 OS=Homo sapiens GN=SLC44A1 PE=1 SV=1
Optineurin OS=Homo sapiens GN=OPTN PE=1 SV=2
PDZ and LIM domain protein 5 OS=Homo sapiens GN=PDLIM5 PE=1 SV=5
COP9 signalosome complex subunit 4 OS=Homo sapiens GN=COPS4 PE=1 SV=1
Toll-interacting protein OS=Homo sapiens GN=TOLLIP PE=1 SV=1
Protein FAM49B OS=Homo sapiens GN=FAM49B PE=1 SV=1
26S proteasome non-ATPase regulatory subunit 13 OS=Homo sapiens GN=PSMD13 PE=1 SV
Cluster of Serine/threonine-protein kinase MRCK beta OS=Homo sapiens GN=CDC42BPB
Importin subunit alpha-1 (Fragment) OS=Homo sapiens GN=KPNA1 PE=4 SV=1
Protein KIAA0368 OS=Homo sapiens GN=KIAA0368 PE=4 SV=1
Tumor protein D54 OS=Homo sapiens GN=TPD52L2 PE=1 SV=2
Lysosomal alpha-glucosidase OS=Homo sapiens GN=GAA PE=1 SV=4
Nucleoprotein TPR OS=Homo sapiens GN=TPR PE=1 SV=3
Ran-specific GTPase-activating protein OS=Homo sapiens GN=RANBP1 PE=1 SV=1
40S ribosomal protein S10 OS=Homo sapiens GN=RPS10 PE=1 SV=1
Serine/threonine-protein kinase PAK 1 OS=Homo sapiens GN=PAK1 PE=1 SV=2
Metalloproteinase inhibitor 3 OS=Homo sapiens GN=TIMP3 PE=4 SV=1
C-terminal-binding protein 1 (Fragment) OS=Homo sapiens GN=CTBP1 PE=3 SV=1
Thioredoxin-like protein 1 OS=Homo sapiens GN=TXNL1 PE=1 SV=3
Proteasome subunit alpha type-2 OS=Homo sapiens GN=PSMA2 PE=1 SV=2
Phosphatidylinositol transfer protein alpha isoform OS=Homo sapiens GN=PITPNA PE=
Isobutyryl-CoA dehydrogenase, mitochondrial OS=Homo sapiens GN=ACAD8 PE=1 SV=1
UMP-CMP kinase OS=Homo sapiens GN=CMPK1 PE=3 SV=1
Glia maturation factor beta OS=Homo sapiens GN=GMFB PE=1 SV=2
Crk-like protein OS=Homo sapiens GN=CRKL PE=1 SV=1
NADPH:adrenodoxin oxidoreductase, mitochondrial OS=Homo sapiens GN=FDXR PE=2 SV=1
Rho-related GTP-binding protein RhoG OS=Homo sapiens GN=RHOG PE=1 SV=1
L-asparaginase OS=Homo sapiens GN=ASRGL1 PE=1 SV=2
Protein quaking (Fragment) OS=Homo sapiens GN=QKI PE=4 SV=1
Prostaglandin-H2 D-isomerase OS=Homo sapiens GN=PTGDS PE=3 SV=1

Carboxymethylenebutenolidase homolog OS=Homo sapiens GN=CMBL PE=1 SV=1
Phosphoribosyl pyrophosphate synthase-associated protein 2 OS=Homo sapiens GN=PRP
Protein XRP2 OS=Homo sapiens GN=RP2 PE=1 SV=4
IST1 homolog OS=Homo sapiens GN=IST1 PE=2 SV=1
Cluster of HCG2044781 OS=Homo sapiens GN=TMEM189 PE=4 SV=1 (G3V2F7)
Integrin beta-2 OS=Homo sapiens GN=ITGB2 PE=1 SV=2
SRC kinase-signaling inhibitor 1 (Fragment) OS=Homo sapiens GN=SRCIN1 PE=4 SV=2
Charged multivesicular body protein 5 OS=Homo sapiens GN=CHMP5 PE=1 SV=1
Isoform 3 of Neurochondrin OS=Homo sapiens GN=NCDN
Ribosome maturation protein SBDS OS=Homo sapiens GN=SBDS PE=1 SV=4
Cluster of Calretinin (Fragment) OS=Homo sapiens GN=CALB2 PE=4 SV=2 (H3BN14)
COP9 signalosome complex subunit 6 OS=Homo sapiens GN=COPS6 PE=4 SV=1
Protein S100-A11 OS=Homo sapiens GN=S100A11 PE=1 SV=2
Isoform 2 of Sorting nexin-18 OS=Homo sapiens GN=SNX18
40S ribosomal protein S4, X isoform OS=Homo sapiens GN=RPS4X PE=1 SV=2
GTP-binding protein SAR1a OS=Homo sapiens GN=SAR1A PE=1 SV=1
Peflin OS=Homo sapiens GN=PEF1 PE=1 SV=1
HCG37564, isoform CRA_c OS=Homo sapiens GN=TIRAP3 PE=2 SV=1
Visinin-like protein 1 (Fragment) OS=Homo sapiens GN=VSNL1 PE=4 SV=1
Cluster of Eukaryotic translation initiation factor 2 subunit 1 OS=Homo sapiens G
von Willebrand factor A domain-containing protein 5A OS=Homo sapiens GN=VWA5A PE=
Phospholysine phosphohistidine inorganic pyrophosphate phosphatase OS=Homo sapien
Peptidyl-prolyl cis-trans isomerase F, mitochondrial OS=Homo sapiens GN=PPIF PE=1
Histidine-rich glycoprotein OS=Homo sapiens GN=HRG PE=1 SV=1
Decorin OS=Homo sapiens GN=DCN PE=1 SV=1
Ras-related protein Rap-2b OS=Homo sapiens GN=RAP2B PE=1 SV=1
Cadherin-13 OS=Homo sapiens GN=CDH13 PE=2 SV=1
Prostate leucine zipper variant 2 OS=Homo sapiens GN=TPD52 PE=2 SV=2
Proline synthase co-transcribed bacterial homolog protein (Fragment) OS=Homo sapi
Dickkopf-related protein 3 OS=Homo sapiens GN=DKK3 PE=4 SV=2
Glycolipid transfer protein OS=Homo sapiens GN=GLTP PE=4 SV=1
Enoyl-CoA delta isomerase 1, mitochondrial (Fragment) OS=Homo sapiens GN=ECI1 PE=
Proteasome subunit beta type-10 (Fragment) OS=Homo sapiens GN=PSMB10 PE=4 SV=1
Glia maturation factor gamma OS=Homo sapiens GN=GMFG PE=1 SV=1
Protein ETHE1, mitochondrial OS=Homo sapiens GN=ETHE1 PE=1 SV=2
Basic leucine zipper and W2 domain-containing protein 1 OS=Homo sapiens GN=BZW1 P
Retinal dehydrogenase 1 OS=Homo sapiens GN=ALDH1A1 PE=1 SV=2
26S proteasome non-ATPase regulatory subunit 12 OS=Homo sapiens GN=PSMD12 PE=1 SV
DnaJ homolog subfamily B member 1 OS=Homo sapiens GN=DNAJB1 PE=2 SV=1
Alpha-2-macroglobulin receptor-associated protein OS=Homo sapiens GN=LRPAP1 PE=1
Syntaxin-1A OS=Homo sapiens GN=STX1A PE=4 SV=1
Constitutive coactivator of PPAR-gamma-like protein 1 OS=Homo sapiens GN=FAM120A
CD59 glycoprotein OS=Homo sapiens GN=CD59 PE=4 SV=1
Neuronal membrane glycoprotein M6-b OS=Homo sapiens GN=GPM6B PE=1 SV=2
Acidic leucine-rich nuclear phosphoprotein 32 family member A OS=Homo sapiens GN=
Phosphatidylinositol 4-phosphate 5-kinase type-1 gamma OS=Homo sapiens GN=PIP5K1C
Actin-related protein 2/3 complex subunit 3 OS=Homo sapiens GN=ARPC3 PE=1 SV=3

Vesicular glutamate transporter 1 OS=Homo sapiens GN=SLC17A7 PE=2 SV=1
WD40 repeat-containing protein SMU1 OS=Homo sapiens GN=SMU1 PE=1 SV=2
Hsp90 co-chaperone Cdc37 OS=Homo sapiens GN=CDC37 PE=1 SV=1
Protein FAM162A OS=Homo sapiens GN=FAM162A PE=4 SV=1
Protein PRRC2A OS=Homo sapiens GN=PRRC2A PE=4 SV=1
Sterol-4-alpha-carboxylate 3-dehydrogenase, decarboxylating (Fragment) OS=Homo sa
Cytochrome c oxidase subunit 7A2, mitochondrial (Fragment) OS=Homo sapiens GN=COX
Malonyl-CoA decarboxylase, mitochondrial OS=Homo sapiens GN=MLYCD PE=1 SV=3
Ubiquitin carboxyl-terminal hydrolase OS=Homo sapiens GN=USP14 PE=3 SV=2
Copine I OS=Homo sapiens GN=CPNE1 PE=4 SV=1
Eukaryotic translation initiation factor 3 subunit H OS=Homo sapiens GN=EIF3H PE=
Sorting nexin 6, isoform CRA_e OS=Homo sapiens GN=SNX6 PE=2 SV=1
Protein transport protein Sec31A OS=Homo sapiens GN=SEC31A PE=2 SV=1
RNA-binding protein 39 OS=Homo sapiens GN=RBM39 PE=4 SV=1
Farnesyl pyrophosphate synthase OS=Homo sapiens GN=FDPS PE=3 SV=1
Acidic leucine-rich nuclear phosphoprotein 32 family member E (Fragment) OS=Homo
COP9 signalosome complex subunit 7a (Fragment) OS=Homo sapiens GN=COPS7A PE=4 SV=
CADM1 protein OS=Homo sapiens GN=CADM1 PE=2 SV=1
Protein canopy homolog 2 (Fragment) OS=Homo sapiens GN=CNPY2 PE=4 SV=1
DNA-(apurinic or apyrimidinic site) lyase (Fragment) OS=Homo sapiens GN=APEX1 PE=
Lipid phosphate phosphohydrolase 3 OS=Homo sapiens GN=PPAP2B PE=1 SV=1
Serine/threonine-protein kinase DCLK1 OS=Homo sapiens GN=DCLK1 PE=1 SV=2
Small glutamine-rich tetratricopeptide repeat-containing protein alpha OS=Homo sa
N-acetylserotonin O-methyltransferase-like protein OS=Homo sapiens GN=ASMTL PE=1
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=Homo sapiens GN=N
Cluster of HLA class II histocompatibility antigen, DRB1-11 beta chain OS=Homo sa
Methylmalonyl-CoA mutase, mitochondrial OS=Homo sapiens GN=MUT PE=1 SV=4
Splicing factor U2AF 65 kDa subunit OS=Homo sapiens GN=U2AF2 PE=1 SV=4
Adenylosuccinate synthetase isozyme 2 OS=Homo sapiens GN=ADSS PE=1 SV=3
NADP-dependent malic enzyme OS=Homo sapiens GN=ME1 PE=1 SV=1
Hepatoma-derived growth factor OS=Homo sapiens GN=HDGF PE=1 SV=1
Coatomer subunit alpha OS=Homo sapiens GN=COPA PE=1 SV=2
26S protease regulatory subunit 10B OS=Homo sapiens GN=PSMC6 PE=1 SV=1
Tyrosine-protein phosphatase non-receptor type 11 OS=Homo sapiens GN=PTPN11 PE=1
Ras-related protein Rab-31 OS=Homo sapiens GN=RAB31 PE=1 SV=1
Protein prune homolog OS=Homo sapiens GN=PRUNE PE=1 SV=2
NEDD8-activating enzyme E1 catalytic subunit OS=Homo sapiens GN=UBA3 PE=1 SV=2
5' (3')-deoxyribonucleotidase, cytosolic type OS=Homo sapiens GN=NT5C PE=1 SV=2
COP9 signalosome complex subunit 5 OS=Homo sapiens GN=COPS5 PE=1 SV=4
Fructosamine-3-kinase OS=Homo sapiens GN=FN3K PE=1 SV=1
Histidine triad nucleotide-binding protein 3 OS=Homo sapiens GN=HINT3 PE=1 SV=1
Actin-binding protein anillin OS=Homo sapiens GN=ANLN PE=1 SV=2
Adseverin OS=Homo sapiens GN=SCIN PE=1 SV=4
60S ribosomal protein L3 (Fragment) OS=Homo sapiens GN=RPL3 PE=3 SV=1
COP9 signalosome complex subunit 1 OS=Homo sapiens GN=GPS1 PE=4 SV=2
Cytoplasmic dynein 1 light intermediate chain 1 OS=Homo sapiens GN=DYNC1LI1 PE=4
60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4

Adapter molecule crk OS=Homo sapiens GN=CRK PE=1 SV=2
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 6 OS=Homo sapiens GN=N
40S ribosomal protein S7 OS=Homo sapiens GN=RPS7 PE=1 SV=1
FACT complex subunit SSRP1 OS=Homo sapiens GN=SSRP1 PE=1 SV=1
Paraspeckle component 1 OS=Homo sapiens GN=PSPC1 PE=1 SV=1
Far upstream element-binding protein 2 OS=Homo sapiens GN=KHSRP PE=1 SV=4
1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase beta-1 OS=Homo sapiens
Phenylalanine--tRNA ligase alpha subunit OS=Homo sapiens GN=FARSA PE=1 SV=3
Ras-related protein R-Ras2 OS=Homo sapiens GN=RRAS2 PE=2 SV=1
Protein 4.1 OS=Homo sapiens GN=EPB41 PE=4 SV=1
Protein transport protein Sec23A OS=Homo sapiens GN=SEC23A PE=4 SV=1
Carnitine O-acetyltransferase OS=Homo sapiens GN=CRAT PE=1 SV=5
Signal transducer and activator of transcription 3 OS=Homo sapiens GN=STAT3 PE=4
Probable ATP-dependent RNA helicase DDX6 OS=Homo sapiens GN=DDX6 PE=1 SV=2
Methionine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=MARS PE=1 SV=2
60S ribosomal protein L7a OS=Homo sapiens GN=RPL7A PE=1 SV=2
3-hydroxyisobutyryl-CoA hydrolase, mitochondrial OS=Homo sapiens GN=HIBCH PE=1 SV
Cluster of Abl interactor 1 OS=Homo sapiens GN=ABI1 PE=4 SV=2 (A6NFN2)
AP-1 complex subunit gamma-1 OS=Homo sapiens GN=AP1G1 PE=2 SV=1
Neural cell adhesion molecule L1-like protein OS=Homo sapiens GN=CHL1 PE=1 SV=4
Peptidyl-prolyl cis-trans isomerase FKBP8 OS=Homo sapiens GN=FKBP8 PE=4 SV=1
Sorting and assembly machinery component 50 homolog OS=Homo sapiens GN=SAMM50 PE=
Thioredoxin reductase 1, cytoplasmic OS=Homo sapiens GN=TXNRD1 PE=2 SV=1
Phosphatidylinositol 5-phosphate 4-kinase type-2 alpha OS=Homo sapiens GN=PIP4K2A
Uncharacterized protein OS=Homo sapiens PE=4 SV=1
Gamma-glutamylcyclotransferase OS=Homo sapiens GN=GGCT PE=1 SV=1
Phosphatidylinositol transfer protein beta isoform OS=Homo sapiens GN=PITPNB PE=2
Copine-2 OS=Homo sapiens GN=CPNE2 PE=1 SV=3
PITH domain-containing protein 1 OS=Homo sapiens GN=PITHD1 PE=1 SV=1
40S ribosomal protein S19 OS=Homo sapiens GN=RPS19 PE=1 SV=2
Protein scribble homolog OS=Homo sapiens GN=SCRIB PE=1 SV=4
Isochorismatase domain-containing protein 1 OS=Homo sapiens GN=ISOC1 PE=1 SV=3
Phosphoacetylglucosamine mutase OS=Homo sapiens GN=PGM3 PE=3 SV=1
Interferon-induced GTP-binding protein Mx1 OS=Homo sapiens GN=MX1 PE=3 SV=1
Acylphosphatase-2 OS=Homo sapiens GN=ACYP2 PE=1 SV=2
Sorting nexin-27 OS=Homo sapiens GN=SNX27 PE=1 SV=2
ProSAAS OS=Homo sapiens GN=PCSK1N PE=1 SV=1
Exocyst complex component 7 OS=Homo sapiens GN=EXOC7 PE=1 SV=3
Glutathione S-transferase Mu 1 OS=Homo sapiens GN=GSTM1 PE=1 SV=3
ADP-ribosylation factor-like protein 8B OS=Homo sapiens GN=ARL8B PE=1 SV=1
Acyl-coenzyme A thioesterase 9, mitochondrial (Fragment) OS=Homo sapiens GN=ACOT9
Branched-chain-amino-acid aminotransferase, cytosolic (Fragment) OS=Homo sapiens
CD151 antigen OS=Homo sapiens GN=CD151 PE=4 SV=1
UV excision repair protein RAD23 homolog B OS=Homo sapiens GN=RAD23B PE=1 SV=1
Serine-threonine kinase receptor-associated protein OS=Homo sapiens GN=STRAP PE=2
40S ribosomal protein S17 (Fragment) OS=Homo sapiens GN=RPS17 PE=3 SV=1
Double-stranded RNA-specific adenosine deaminase OS=Homo sapiens GN=ADAR PE=4 SV=

Proliferating cell nuclear antigen OS=Homo sapiens GN=PCNA PE=1 SV=1
Exocyst complex component 1 OS=Homo sapiens GN=EXOC1 PE=1 SV=4
Protein TFG (Fragment) OS=Homo sapiens GN=TFG PE=4 SV=1
Polymerase I and transcript release factor OS=Homo sapiens GN=PTRF PE=1 SV=1
Protein tweety homolog 1 OS=Homo sapiens GN=TTYH1 PE=2 SV=1
Reticulocalbin-2 OS=Homo sapiens GN=RCN2 PE=4 SV=1
60S ribosomal protein L14 OS=Homo sapiens GN=RPL14 PE=4 SV=1
Adenosine kinase OS=Homo sapiens GN=ADK PE=1 SV=2
DnaJ homolog subfamily B member 11 OS=Homo sapiens GN=DNAJB11 PE=1 SV=1
Splicing factor 3B subunit 3 OS=Homo sapiens GN=SF3B3 PE=1 SV=4
Protein ERGIC-53 OS=Homo sapiens GN=LMAN1 PE=1 SV=2
Cystathionine beta-synthase OS=Homo sapiens GN=CBS PE=4 SV=1
Protein PML OS=Homo sapiens GN=PML PE=1 SV=3
Nucleobindin-1 OS=Homo sapiens GN=NUCB1 PE=1 SV=4
Heat shock protein beta-8 OS=Homo sapiens GN=HSPB8 PE=1 SV=1
EPS15L1 protein OS=Homo sapiens GN=EPS15L1 PE=2 SV=1
Importin-7 OS=Homo sapiens GN=IPO7 PE=1 SV=1
Omega-amidase NIT2 OS=Homo sapiens GN=NIT2 PE=1 SV=1
NHP2 non-histone chromosome protein 2-like 1 (*S. cerevisiae*) OS=Homo sapiens GN=N
40S ribosomal protein S3a OS=Homo sapiens GN=RPS3A PE=3 SV=1
Glutathione S-transferase kappa 1 OS=Homo sapiens GN=GSTK1 PE=4 SV=1
Glycine cleavage system H protein, mitochondrial OS=Homo sapiens GN=GCSH PE=1 SV=
Translation initiation factor eIF-2B subunit beta OS=Homo sapiens GN=EIF2B2 PE=1
Ubiquitin-fold modifier 1 OS=Homo sapiens GN=UFM1 PE=1 SV=1
40S ribosomal protein S28 OS=Homo sapiens GN=RPS28 PE=1 SV=1
EPM2A-interacting protein 1 OS=Homo sapiens GN=EPM2AIP1 PE=1 SV=1
Calcium/calmodulin-dependent protein kinase type 1D OS=Homo sapiens GN=CAMK1D PE=
Small VCP/p97-interacting protein OS=Homo sapiens GN=SVIP PE=2 SV=1
Calcineurin-like phosphoesterase domain-containing protein 1 OS=Homo sapiens GN=C
Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-12 OS=Homo sapien
Tropomodulin-2 OS=Homo sapiens GN=TMOD2 PE=1 SV=1
60S ribosomal protein L9 (Fragment) OS=Homo sapiens GN=RPL9 PE=4 SV=1
Septin-4 OS=Homo sapiens GN=SEPT4 PE=3 SV=1
RNA-binding protein FUS OS=Homo sapiens GN=FUS PE=4 SV=1
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 10, mitochondrial OS=H
ADP-ribosylation factor 6 OS=Homo sapiens GN=ARF6 PE=1 SV=2
Metalloreductase STEAP3 OS=Homo sapiens GN=STEAP3 PE=1 SV=2
Dynein light chain 2, cytoplasmic OS=Homo sapiens GN=DYNLL2 PE=1 SV=1
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 9, mitochondrial OS=Ho
Thioredoxin-related transmembrane protein 1 OS=Homo sapiens GN=TMX1 PE=1 SV=1
AP-2 complex subunit sigma OS=Homo sapiens GN=AP2S1 PE=1 SV=2
Myoglobin (Fragment) OS=Homo sapiens GN=MB PE=3 SV=1
MARCKS-related protein OS=Homo sapiens GN=MARCKSL1 PE=1 SV=2
Exocyst complex component 4 OS=Homo sapiens GN=EXOC4 PE=1 SV=1
Cysteine and histidine-rich domain-containing protein 1 OS=Homo sapiens GN=CHORDC
Mitochondrial carrier homolog 2 OS=Homo sapiens GN=MTCH2 PE=1 SV=1
Proteasome activator complex subunit 2 OS=Homo sapiens GN=PSME2 PE=4 SV=1

Replication protein A 70 kDa DNA-binding subunit OS=Homo sapiens GN=RPA1 PE=1 SV=1
Mitochondrial import inner membrane translocase subunit TIM44 OS=Homo sapiens GN=
Cytochrome b-245 heavy chain OS=Homo sapiens GN=CYBB PE=1 SV=2
Oxidation resistance protein 1 OS=Homo sapiens GN=OXR1 PE=1 SV=2
Uncharacterized protein OS=Homo sapiens GN=YJEFN3 PE=4 SV=1
Ragulator complex protein LAMTOR3 OS=Homo sapiens GN=LAMTOR3 PE=4 SV=1
Vacuolar protein sorting-associated protein VTA1 homolog OS=Homo sapiens GN=VTA1
Raft-linking protein, isoform CRA_c OS=Homo sapiens GN=RFTN1 PE=4 SV=1
Acyl-CoA synthetase family member 2, mitochondrial OS=Homo sapiens GN=ACSF2 PE=2
Elongation factor 1-beta OS=Homo sapiens GN=EEF1B2 PE=1 SV=3
Nuclear pore complex protein Nup93 OS=Homo sapiens GN=NUP93 PE=4 SV=1
Ketosamine-3-kinase OS=Homo sapiens GN=FN3KRP PE=4 SV=1
Microtubule-associated protein 1S OS=Homo sapiens GN=MAP1S PE=2 SV=1
Suppressor of G2 allele of SKP1 homolog OS=Homo sapiens GN=SUGT1 PE=1 SV=3
Ornithine aminotransferase, mitochondrial OS=Homo sapiens GN=OAT PE=1 SV=1
Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-5 OS=Homo sapiens
PRA1 family protein 3 OS=Homo sapiens GN=ARL6IP5 PE=1 SV=1
Myelin expression factor 2 OS=Homo sapiens GN=MYEF2 PE=1 SV=3
ATP synthase subunit epsilon, mitochondrial OS=Homo sapiens GN=ATP5E PE=1 SV=2
Gamma-butyrobetaine dioxygenase OS=Homo sapiens GN=BBOX1 PE=1 SV=1
Solute carrier family 15 member 2 OS=Homo sapiens GN=SLC15A2 PE=2 SV=2
RNA-binding protein Musashi homolog 2 OS=Homo sapiens GN=MSI2 PE=2 SV=1
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 3 (Fragment) OS=Homo sa
Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3
Ig heavy chain V-I region V35 OS=Homo sapiens PE=1 SV=1
Microtubule-associated proteins 1A/1B light chain 3A OS=Homo sapiens GN=MAP1LC3A
Calcium-binding protein 39 OS=Homo sapiens GN=CAB39 PE=1 SV=1
Zinc finger CCCH-type antiviral protein 1 OS=Homo sapiens GN=ZC3HAV1 PE=4 SV=1
40S ribosomal protein S13 OS=Homo sapiens GN=RPS13 PE=1 SV=2
Myotubularin OS=Homo sapiens GN=MTM1 PE=2 SV=1
D-dopachrome decarboxylase-like protein OS=Homo sapiens GN=DDTL PE=2 SV=1
KH domain-containing, RNA-binding, signal transduction-associated protein 1 OS=Ho
Cytochrome c oxidase subunit 6C OS=Homo sapiens GN=COX6C PE=1 SV=2
26S protease regulatory subunit 6B OS=Homo sapiens GN=PSMC4 PE=1 SV=2
Rho guanine nucleotide exchange factor 1 OS=Homo sapiens GN=ARHGEF1 PE=1 SV=2
Serine palmitoyltransferase 2 OS=Homo sapiens GN=SPTLC2 PE=1 SV=1
Remodeling and spacing factor 1 (Fragment) OS=Homo sapiens GN=RSF1 PE=4 SV=1
Thy-1 membrane glycoprotein (Fragment) OS=Homo sapiens GN=THY1 PE=4 SV=1
60S ribosomal protein L10 OS=Homo sapiens GN=RPL10 PE=1 SV=4
Atlantin-3 OS=Homo sapiens GN=ATL3 PE=4 SV=1
Immunoglobulin superfamily member 1 OS=Homo sapiens GN=IGSF1 PE=4 SV=1
Leucine-rich repeat-containing protein 7 OS=Homo sapiens GN=LRRC7 PE=1 SV=1
Myelin P2 protein OS=Homo sapiens GN=PMP2 PE=1 SV=3
OCIA domain-containing protein 1 OS=Homo sapiens GN=OCIAD1 PE=4 SV=1
DCN1-like protein 1 (Fragment) OS=Homo sapiens GN=DCUN1D1 PE=4 SV=1
Short-chain-specific acyl-CoA dehydrogenase, mitochondrial OS=Homo sapiens GN=ACA
Ubiquitin-like-conjugating enzyme ATG3 OS=Homo sapiens GN=ATG3 PE=4 SV=1

Apolipoprotein L2 OS=Homo sapiens GN=APOL2 PE=4 SV=1
Sepiapterin reductase OS=Homo sapiens GN=SPR PE=1 SV=1
Acyl-CoA-binding domain-containing protein 7 OS=Homo sapiens GN=ACBD7 PE=1 SV=1
Ras and Rab interactor 1 OS=Homo sapiens GN=RIN1 PE=4 SV=1
Protein NEDD8-MDP1 (Fragment) OS=Homo sapiens GN=NEDD8-MDP1 PE=4 SV=1
Metalloproteinase inhibitor 1 OS=Homo sapiens GN=TIMP1 PE=1 SV=1
Palmitoyl-protein thioesterase 1 OS=Homo sapiens GN=PPT1 PE=1 SV=1
28 kDa heat- and acid-stable phosphoprotein OS=Homo sapiens GN=PDAP1 PE=1 SV=1
Peroxisomal acyl-coenzyme A oxidase 1 OS=Homo sapiens GN=ACOX1 PE=1 SV=3
Transmembrane protein 65 OS=Homo sapiens GN=TMEM65 PE=1 SV=2
CD109 antigen OS=Homo sapiens GN=CD109 PE=1 SV=2
Pyridoxal phosphate phosphatase OS=Homo sapiens GN=PDXP PE=1 SV=2
COP9 signalosome complex subunit 3 OS=Homo sapiens GN=COPS3 PE=1 SV=3
Cluster of EF-hand domain-containing protein D1 OS=Homo sapiens GN=EFHD1 PE=4 SV=
Beta-arrestin-1 (Fragment) OS=Homo sapiens GN=ARRB1 PE=4 SV=1
Proteasome subunit beta type-4 OS=Homo sapiens GN=PSMB4 PE=1 SV=4
40S ribosomal protein S14 OS=Homo sapiens GN=RPS14 PE=1 SV=3
GLI pathogenesis-related 2 OS=Homo sapiens GN=GLIPR2 PE=2 SV=1
AP-3 complex subunit beta-1 OS=Homo sapiens GN=AP3B1 PE=4 SV=1
L-xylulose reductase (Fragment) OS=Homo sapiens GN=DCXR PE=3 SV=1
Phospholipid transfer protein OS=Homo sapiens GN=PLTP PE=2 SV=1
Myosin regulatory light chain 12A OS=Homo sapiens GN=MYL12A PE=4 SV=1
Prolyl endopeptidase OS=Homo sapiens GN=PREP PE=1 SV=2
Eukaryotic translation initiation factor 3 subunit B OS=Homo sapiens GN=EIF3B PE=
HCG1770615, isoform CRA_b OS=Homo sapiens GN=TSC22D4 PE=2 SV=1
Apoptosis inhibitor 5 OS=Homo sapiens GN=API5 PE=4 SV=1
Mitochondrial import receptor subunit TOM40 homolog OS=Homo sapiens GN=TOMM40 PE=
UDP-glucose:glycoprotein glucosyltransferase 1 OS=Homo sapiens GN=UGGT1 PE=1 SV=3
Glycerol-3-phosphate dehydrogenase [NAD(+)], cytoplasmic (Fragment) OS=Homo sapie
SUMO-activating enzyme subunit 2 OS=Homo sapiens GN=UBA2 PE=1 SV=2
DCC-interacting protein 13-alpha OS=Homo sapiens GN=APPL1 PE=1 SV=1
Apolipoprotein B-100 OS=Homo sapiens GN=APOB PE=1 SV=2
Actin-binding LIM protein 1 OS=Homo sapiens GN=ABLIM1 PE=4 SV=1
Tumor suppressor p53-binding protein 1 OS=Homo sapiens GN=TP53BP1 PE=4 SV=2
Lysophospholipase-like protein 1 OS=Homo sapiens GN=LYPLAL1 PE=1 SV=3
Protein enabled homolog OS=Homo sapiens GN=ENAH PE=1 SV=2
Apolipoprotein 0 OS=Homo sapiens GN=APO0 PE=4 SV=1
Nuclear receptor-binding protein OS=Homo sapiens GN=NRBP1 PE=4 SV=1
Galectin-3 OS=Homo sapiens GN=LGALS3 PE=1 SV=5
E3 ubiquitin/ISG15 ligase TRIM25 OS=Homo sapiens GN=TRIM25 PE=1 SV=2
Phosphomevalonate kinase OS=Homo sapiens GN=PMVK PE=1 SV=3
Phosphoglycolate phosphatase OS=Homo sapiens GN=PGP PE=1 SV=1
Cluster of Fragile X mental retardation syndrome-related protein 1 OS=Homo sapien
Insulin-like growth factor II OS=Homo sapiens GN=IGF2 PE=3 SV=2
Complement component C6 OS=Homo sapiens GN=C6 PE=1 SV=3
Opioid binding protein/cell adhesion molecule-like isoform c OS=Homo sapiens GN=0
ATP-binding cassette sub-family F member 1 (Fragment) OS=Homo sapiens GN=ABCF1 PE

Collagen alpha-2(IV) chain OS=Homo sapiens GN=COL4A2 PE=1 SV=4
Signal recognition particle 9 kDa protein OS=Homo sapiens GN=SRP9 PE=1 SV=2
Eukaryotic translation initiation factor 2 subunit 3 OS=Homo sapiens GN=EIF2S3 PE=1 SV=2
Ferritin light chain OS=Homo sapiens GN=FTL PE=1 SV=2
Leucine-rich repeat-containing protein 4B OS=Homo sapiens GN=LRRC4B PE=1 SV=3
Ig kappa chain V-III region SIE OS=Homo sapiens PE=1 SV=1
Nicastrin OS=Homo sapiens GN=NCSTN PE=4 SV=2
ADP-ribosylation factor-like protein 8A OS=Homo sapiens GN=ARL8A PE=1 SV=1
Cluster of 3-hydroxymethyl-3-methylglutaryl-Coenzyme A lyase OS=Homo sapiens GN=HMGCL
Wolframin OS=Homo sapiens GN=WFS1 PE=1 SV=2
60S ribosomal protein L27a OS=Homo sapiens GN=RPL27A PE=3 SV=1
GTPase NRas OS=Homo sapiens GN=NRAS PE=1 SV=1
Aquaporin-1 OS=Homo sapiens GN=AQP1 PE=2 SV=1
Laminin subunit alpha-4 OS=Homo sapiens GN=LAMA4 PE=1 SV=4
Citrate lyase subunit beta-like protein, mitochondrial OS=Homo sapiens GN=CLYBL P
C-Jun-amino-terminal kinase-interacting protein 4 OS=Homo sapiens GN=SPAG9 PE=1 SV=1
Leucine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=LARS PE=2 SV=1
26S proteasome non-ATPase regulatory subunit 7 OS=Homo sapiens GN=PSMD7 PE=2 SV=1
Membrane protein MLC1 OS=Homo sapiens GN=MLC1 PE=2 SV=1
Thioredoxin reductase 2, mitochondrial OS=Homo sapiens GN=TXNRD2 PE=3 SV=1
NudC domain-containing protein 2 OS=Homo sapiens GN=NUDCD2 PE=4 SV=1
Ras GTPase-activating protein-binding protein 1 (Fragment) OS=Homo sapiens GN=G3BP1
Serine/threonine-protein phosphatase 2A 56 kDa regulatory subunit delta isoform 0
Rab GTPase-binding effector protein 1 OS=Homo sapiens GN=RABEP1 PE=4 SV=1
Adaptin ear-binding coat-associated protein 2 OS=Homo sapiens GN=NECAP2 PE=4 SV=1
60S ribosomal protein L18 (Fragment) OS=Homo sapiens GN=RPL18 PE=4 SV=1
Tight junction protein 1 (Zona occludens 1), isoform CRA_a OS=Homo sapiens GN=TJP1
Regulator of microtubule dynamics protein 3 (Fragment) OS=Homo sapiens GN=FAM82A2
Cleavage and polyadenylation-specificity factor subunit 5 (Fragment) OS=Homo sapiens GN=CPSF5
Hydroxyacylglutathione hydrolase, mitochondrial (Fragment) OS=Homo sapiens GN=HAGHL
E3 ubiquitin-protein ligase CHIP (Fragment) OS=Homo sapiens GN=STUB1 PE=4 SV=1
Microsomal glutathione S-transferase 3 OS=Homo sapiens GN=MGST3 PE=1 SV=1
Hepatocyte growth factor-regulated tyrosine kinase substrate OS=Homo sapiens GN=HGFR
Non-syndromic hearing impairment protein 5 OS=Homo sapiens GN=DFNA5 PE=1 SV=2
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 7 OS=Homo sapiens GN=ND7
Serine/threonine-protein kinase OSR1 OS=Homo sapiens GN=OXSR1 PE=1 SV=1
Cytochrome b5 OS=Homo sapiens GN=CYB5A PE=1 SV=2
Hypoxanthine-guanine phosphoribosyltransferase OS=Homo sapiens GN=HPRT1 PE=1 SV=2
Cathepsin G OS=Homo sapiens GN=CTSG PE=1 SV=2
Mesencephalic astrocyte-derived neurotrophic factor OS=Homo sapiens GN=MANF PE=1 SV=1
40S ribosomal protein S26 OS=Homo sapiens GN=RPS26 PE=1 SV=3
Guanidinoacetate N-methyltransferase OS=Homo sapiens GN=GAMT PE=1 SV=1
Ras suppressor protein 1 OS=Homo sapiens GN=RSU1 PE=1 SV=3
Splicing factor 3A subunit 1 OS=Homo sapiens GN=SF3A1 PE=1 SV=1
Putative ATP-dependent Clp protease proteolytic subunit, mitochondrial OS=Homo sapiens GN=CLPP
ER membrane protein complex subunit 1 OS=Homo sapiens GN=EMC1 PE=1 SV=1
Trans-L-3-hydroxyproline dehydratase OS=Homo sapiens GN=L3HYPDH PE=1 SV=2

Chloride intracellular channel protein 6 OS=Homo sapiens GN=CLIC6 PE=2 SV=3
Charged multivesicular body protein 4b OS=Homo sapiens GN=CHMP4B PE=1 SV=1
Calcium-regulated heat stable protein 1 OS=Homo sapiens GN=CARHSP1 PE=1 SV=2
Rap guanine nucleotide exchange factor 2 OS=Homo sapiens GN=RAPGEF2 PE=1 SV=1
Lysosome-associated membrane glycoprotein 2 OS=Homo sapiens GN=LAMP2 PE=2 SV=1
Basigin OS=Homo sapiens GN=BSG PE=4 SV=1
Regulator of nonsense transcripts 1 OS=Homo sapiens GN=UPF1 PE=1 SV=2
Translocon-associated protein subunit alpha OS=Homo sapiens GN=SSR1 PE=4 SV=1
Adenine phosphoribosyltransferase OS=Homo sapiens GN=APRT PE=4 SV=1
ATP-binding cassette sub-family B member 7, mitochondrial OS=Homo sapiens GN=ABCB
Fatty acid-binding protein, heart OS=Homo sapiens GN=FABP3 PE=1 SV=4
Dipeptidyl aminopeptidase-like protein 6 OS=Homo sapiens GN=DPP6 PE=4 SV=1
Eukaryotic peptide chain release factor subunit 1 OS=Homo sapiens GN=ETF1 PE=2 SV=1
Unconventional myosin-XVIIIa OS=Homo sapiens GN=MYO18A PE=4 SV=1
Pirin OS=Homo sapiens GN=PIR PE=1 SV=1
Casein kinase II subunit beta OS=Homo sapiens GN=CSNK2B PE=1 SV=1
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 4 OS=Homo sapiens GN=ND
Keratin, type II cytoskeletal 5 OS=Homo sapiens GN=KRT5 PE=1 SV=3
Glypican-1 (Fragment) OS=Homo sapiens GN=GPC1 PE=3 SV=1
Casein kinase II subunit alpha OS=Homo sapiens GN=CSNK2A1 PE=4 SV=1
Protein phosphatase 2A activator, regulatory subunit 4 OS=Homo sapiens GN=PPP2R4
Inorganic pyrophosphatase 2, mitochondrial OS=Homo sapiens GN=PPA2 PE=4 SV=1
Eukaryotic translation initiation factor 3 subunit J OS=Homo sapiens GN=EIF3J PE=1
Synaptobrevin homolog YKT6 OS=Homo sapiens GN=YKT6 PE=1 SV=1
Reticulocalbin-1 OS=Homo sapiens GN=RCN1 PE=1 SV=1
Cluster of Protein phosphatase 1A OS=Homo sapiens GN=PPM1A PE=3 SV=1 (J3KNM0)
AP-1 complex subunit mu-1 OS=Homo sapiens GN=AP1M1 PE=4 SV=1
Plexin domain-containing protein 2 OS=Homo sapiens GN=PLXDC2 PE=4 SV=1
Tyrosine-protein phosphatase non-receptor type 23 OS=Homo sapiens GN=PTPN23 PE=2
Bis(5'-nucleosyl)-tetrphosphatase [asymmetrical] OS=Homo sapiens GN=NUDT2 PE=1 S
GTPase KRas OS=Homo sapiens GN=KRAS PE=1 SV=1
Myeloid cell nuclear differentiation antigen OS=Homo sapiens GN=MNDA PE=1 SV=1
Caprin-1 OS=Homo sapiens GN=CAPRIN1 PE=4 SV=1
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 11 OS=Homo sapiens GN=ND
Calcyclin binding protein OS=Homo sapiens GN=CACYBP PE=2 SV=1
Zyxin (Fragment) OS=Homo sapiens GN=ZYX PE=4 SV=1
Epididymal secretory protein E1 OS=Homo sapiens GN=NPC2 PE=2 SV=1
Ras GTPase-activating-like protein IQGAP2 OS=Homo sapiens GN=IQGAP2 PE=1 SV=4
60S ribosomal protein L36 OS=Homo sapiens GN=RPL36 PE=1 SV=3
Protein C15orf38-AP3S2 OS=Homo sapiens GN=C15orf38-AP3S2 PE=4 SV=1
RNA binding motif protein, X-linked-like-1 OS=Homo sapiens GN=RBMXL1 PE=1 SV=1
CD99 antigen OS=Homo sapiens GN=CD99 PE=4 SV=1
von Willebrand factor A domain-containing protein 1 OS=Homo sapiens GN=VWA1 PE=2
Transformer-2 protein homolog beta (Fragment) OS=Homo sapiens GN=TRA2B PE=4 SV=1
Prefoldin subunit 5 (Fragment) OS=Homo sapiens GN=PFDN5 PE=4 SV=1
Cluster of Ig kappa chain V-III region VG (Fragment) OS=Homo sapiens PE=1 SV=1 (P
Calcineurin subunit B type 1 OS=Homo sapiens GN=PPP3R1 PE=4 SV=1

Fibulin-5 OS=Homo sapiens GN=FBLN5 PE=4 SV=1

Protein arginine N-methyltransferase 5, N-terminally processed (Fragment) OS=Homo sapiens PE=4 SV=2
Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=2
Putative pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15 OS=Homo sapiens PE=1 SV=6
Translational activator GCN1 OS=Homo sapiens GN=GCN1L1 PE=1 SV=6
Integrin-linked protein kinase OS=Homo sapiens GN=ILK PE=2 SV=1
Serine/arginine repetitive matrix protein 2 OS=Homo sapiens GN=SRRM2 PE=1 SV=2
Uncharacterized protein C9orf142 OS=Homo sapiens GN=C9orf142 PE=1 SV=2
Methylcrotonoyl-CoA carboxylase subunit alpha, mitochondrial (Fragment) OS=Homo sapiens PE=1 SV=1
Hepatitis B virus X-interacting protein OS=Homo sapiens GN=HBXIP PE=4 SV=1
Centrin-2 OS=Homo sapiens GN=CETN2 PE=1 SV=1
Alkyldihydroxyacetonephosphate synthase, peroxisomal OS=Homo sapiens GN=AGPS PE=1 SV=1
Syntaxin-binding protein 6 OS=Homo sapiens GN=STXBP6 PE=4 SV=1
Ubiquitin carboxyl-terminal hydrolase L5 OS=Homo sapiens GN=UCHL5 PE=2 SV=1
Quinone oxidoreductase-like protein 1 OS=Homo sapiens GN=CRYZL1 PE=4 SV=1
Lysosome-associated membrane glycoprotein 1 OS=Homo sapiens GN=LAMP1 PE=1 SV=3
Sulfurtransferase OS=Homo sapiens GN=MPST PE=4 SV=1
CDK5 regulatory subunit-associated protein 3 OS=Homo sapiens GN=CDK5RAP3 PE=4 SV=1
Neurabin-2 OS=Homo sapiens GN=PPP1R9B PE=1 SV=2
Inosine triphosphate pyrophosphatase OS=Homo sapiens GN=ITPA PE=1 SV=2
Diphosphoinositol polyphosphate phosphohydrolase 1 OS=Homo sapiens GN=NUDT3 PE=1 SV=1
Dipeptidyl peptidase 3 OS=Homo sapiens GN=DPP3 PE=4 SV=1
Alpha-parvin OS=Homo sapiens GN=PARVA PE=2 SV=1
AMP deaminase 2 OS=Homo sapiens GN=AMPD2 PE=4 SV=1
Ubiquitin-conjugating enzyme E2 D3 OS=Homo sapiens GN=UBE2D3 PE=1 SV=1
Synaptic vesicle membrane protein VAT-1 homolog-like OS=Homo sapiens GN=VAT1L PE=1 SV=4
40S ribosomal protein S5 OS=Homo sapiens GN=RPS5 PE=1 SV=4
Eukaryotic translation initiation factor 3 subunit L OS=Homo sapiens GN=EIF3EIP PE=1 SV=1
HEAT repeat-containing protein 2 OS=Homo sapiens GN=HEATR2 PE=1 SV=4

Fibulin-2 OS=Homo sapiens GN=FBLN2 PE=4 SV=1

Protein mago nashi homolog 2 OS=Homo sapiens GN=MAGOHB PE=4 SV=1
Importin subunit alpha-4 OS=Homo sapiens GN=KPNA4 PE=1 SV=1
Apolipoprotein C-I OS=Homo sapiens GN=APOC1 PE=1 SV=1
Microfibril-associated glycoprotein 4 OS=Homo sapiens GN=MFAP4 PE=4 SV=1
KIF1-binding protein OS=Homo sapiens GN=KIAA1279 PE=1 SV=1
Protein phosphatase 1G OS=Homo sapiens GN=PPM1G PE=1 SV=1
Dual specificity protein phosphatase 3 OS=Homo sapiens GN=DUSP3 PE=1 SV=1
Aminoacylase-1 OS=Homo sapiens GN=ACY1 PE=2 SV=1
Ras-related protein Rab-21 OS=Homo sapiens GN=RAB21 PE=1 SV=3
Acylpyruvase FAHD1, mitochondrial OS=Homo sapiens GN=FAHD1 PE=4 SV=1
ELAV (Embryonic lethal, abnormal vision, Drosophila)-like 4 (Hu antigen D) OS=Homo sapiens PE=1 SV=1
CTP synthase 2 OS=Homo sapiens GN=CTPS2 PE=1 SV=1
Neutrophil defensin 1 OS=Homo sapiens GN=DEFA1 PE=1 SV=1
Ubiquitin-conjugating enzyme E2 K OS=Homo sapiens GN=UBE2K PE=2 SV=1
AH receptor-interacting protein OS=Homo sapiens GN=AIP PE=1 SV=2
Complement factor H-related 5 OS=Homo sapiens GN=CFHR5 PE=2 SV=1
Alpha-1-acid glycoprotein 2 OS=Homo sapiens GN=ORM2 PE=1 SV=2

Synaptic vesicle glycoprotein 2A OS=Homo sapiens GN=SV2A PE=1 SV=1
Stearoyl-CoA desaturase 5 OS=Homo sapiens GN=SCD5 PE=2 SV=2
Exocyst complex component 8 OS=Homo sapiens GN=EXOC8 PE=1 SV=2
Protein BRICK1 OS=Homo sapiens GN=BRK1 PE=4 SV=1
Acyl carrier protein (Fragment) OS=Homo sapiens GN=NDUFAB1 PE=3 SV=1
Sodium/potassium-transporting ATPase subunit beta-3 OS=Homo sapiens GN=ATP1B3 PE=
Cytochrome b-c1 complex subunit 6, mitochondrial OS=Homo sapiens GN=UQCRH PE=1 SV
Alpha-N-acetylgalactosaminidase OS=Homo sapiens GN=NAGA PE=1 SV=2
Beta-1,3-glucosyltransferase OS=Homo sapiens GN=B3GALTL PE=1 SV=2
Mitotic checkpoint protein BUB3 OS=Homo sapiens GN=BUB3 PE=2 SV=1
Neurocalcin-delta OS=Homo sapiens GN=NCALD PE=2 SV=2
Mimecan OS=Homo sapiens GN=OGN PE=1 SV=1
Mitochondria-eating protein OS=Homo sapiens GN=SPATA18 PE=4 SV=1
Histone-binding protein RBBP7 OS=Homo sapiens GN=RBBP7 PE=4 SV=1
Protein flightless-1 homolog OS=Homo sapiens GN=FLII PE=4 SV=1
Signal recognition particle 14 kDa protein OS=Homo sapiens GN=SRP14 PE=1 SV=2
Clathrin interactor 1 OS=Homo sapiens GN=CLINT1 PE=1 SV=1
Protein FAM186B OS=Homo sapiens GN=FAM186B PE=2 SV=1
Kinesin-like protein KIF13B OS=Homo sapiens GN=KIF13B PE=3 SV=1
Small ubiquitin-related modifier 3 OS=Homo sapiens GN=SUMO3 PE=4 SV=1
Apolipoprotein C-III OS=Homo sapiens GN=APOC3 PE=4 SV=1
Nucleoporin p54 OS=Homo sapiens GN=NUP54 PE=2 SV=1
60S ribosomal protein L24 OS=Homo sapiens GN=RPL24 PE=4 SV=1
Nascent polypeptide-associated complex subunit alpha OS=Homo sapiens GN=NACA PE=4
Protein FAM184A OS=Homo sapiens GN=FAM184A PE=4 SV=1
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4 OS=Homo sapiens GN=N
Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma-4 OS=Homo sapiens
ATP synthase subunit e, mitochondrial OS=Homo sapiens GN=ATP5I PE=1 SV=2
Peptidase inhibitor 16 OS=Homo sapiens GN=PI16 PE=1 SV=1
Neuronal growth regulator 1 OS=Homo sapiens GN=NEGR1 PE=1 SV=3
Lysophosphatidylcholine acyltransferase 1 OS=Homo sapiens GN=LPCAT1 PE=1 SV=2
DnaJ homolog subfamily C member 5 OS=Homo sapiens GN=DNAJC5 PE=1 SV=1
Ig lambda chain V-III region LOI OS=Homo sapiens PE=1 SV=1
14 kDa phosphohistidine phosphatase OS=Homo sapiens GN=PHPT1 PE=1 SV=1
UV excision repair protein RAD23 homolog A OS=Homo sapiens GN=RAD23A PE=1 SV=1
Integrin beta OS=Homo sapiens GN=ITGB8 PE=2 SV=1
Tropomodulin-3 (Fragment) OS=Homo sapiens GN=TMOD3 PE=4 SV=1
HCG2027369, isoform CRA_a OS=Homo sapiens GN=KIF21B PE=2 SV=1
Vacuolar protein sorting-associated protein 13C OS=Homo sapiens GN=VPS13C PE=1 SV
Regulation of nuclear pre-mRNA domain containing 1B (Fragment) OS=Homo sapiens GN
ADP-sugar pyrophosphatase OS=Homo sapiens GN=NUDT5 PE=3 SV=1
Syntaxin 4A (Placental), isoform CRA_c OS=Homo sapiens GN=STX4 PE=2 SV=1
Target of Myb protein 1 (Fragment) OS=Homo sapiens GN=TOM1 PE=4 SV=1
Guanine nucleotide binding protein-like 1 (Fragment) OS=Homo sapiens GN=GNL1 PE=4
Dynein light chain roadblock-type 1 OS=Homo sapiens GN=DYNLRB1 PE=4 SV=1
NADH dehydrogenase [ubiquinone] iron-sulfur protein 7, mitochondrial OS=Homo sapi
Retinol dehydrogenase 11 OS=Homo sapiens GN=RDH11 PE=2 SV=1

Eukaryotic translation initiation factor 3 subunit D OS=Homo sapiens GN=EIF3D PE=Galactokinase OS=Homo sapiens GN=GALK1 PE=2 SV=1
Sec1 family domain containing 1, isoform CRA_b OS=Homo sapiens GN=SCFD1 PE=2 SV=1
Eukaryotic translation initiation factor 3 subunit K OS=Homo sapiens GN=EIF3K PE=Peroxisomal carnitine 0-octanoyltransferase OS=Homo sapiens GN=CROT PE=3 SV=1
60S ribosomal protein L28 OS=Homo sapiens GN=RPL28 PE=4 SV=1
Myoferlin OS=Homo sapiens GN=MYOF PE=4 SV=2
Acyl-protein thioesterase 1 (Fragment) OS=Homo sapiens GN=LYPLA1 PE=4 SV=1
Pyruvate dehydrogenase protein X component, mitochondrial OS=Homo sapiens GN=PDHX
Protein Clorf27 OS=Homo sapiens GN=Clorf27 PE=4 SV=1
N-alpha-acetyltransferase 38, NatC auxiliary subunit OS=Homo sapiens GN=NAA38 PE=Neuroplastin OS=Homo sapiens GN=NPTN PE=4 SV=1
Neural cell adhesion molecule L1 OS=Homo sapiens GN=L1CAM PE=4 SV=1
Neutral cholesterol ester hydrolase 1 OS=Homo sapiens GN=NCEH1 PE=4 SV=1
LIM domain and actin-binding protein 1 OS=Homo sapiens GN=LIMA1 PE=4 SV=1
UPF0160 protein MYG1, mitochondrial OS=Homo sapiens GN=C12orf10 PE=4 SV=1
V-type proton ATPase subunit D OS=Homo sapiens GN=ATP6V1D PE=4 SV=1
Leucine-rich repeat-containing protein 57 (Fragment) OS=Homo sapiens GN=LRRC57 PE=Zinc finger protein ZPR1 (Fragment) OS=Homo sapiens GN=ZNF259 PE=4 SV=1
Nodal modulator 3 OS=Homo sapiens GN=NOMO3 PE=4 SV=1
ARF GTPase-activating protein GIT1 OS=Homo sapiens GN=GIT1 PE=4 SV=1
Interferon-induced protein with tetratricopeptide repeats 3 OS=Homo sapiens GN=IF
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 2 OS=Homo sapiens GN=N
Pentraxin-related protein PTX3 OS=Homo sapiens GN=PTX3 PE=1 SV=3
Peptidyl-prolyl cis-trans isomerase FKBP2 OS=Homo sapiens GN=FKBP2 PE=1 SV=2
OX-2 membrane glycoprotein OS=Homo sapiens GN=CD200 PE=1 SV=4
COP9 signalosome complex subunit 2 OS=Homo sapiens GN=COPS2 PE=1 SV=1
Complement C1q tumor necrosis factor-related protein 5 OS=Homo sapiens GN=C1QTNF5
LanC-like protein 2 OS=Homo sapiens GN=LANCL2 PE=1 SV=1
Prostaglandin F2 receptor negative regulator OS=Homo sapiens GN=PTGFRN PE=1 SV=2
C-type lectin domain family 11 member A OS=Homo sapiens GN=CLEC11A PE=1 SV=1
Mitochondrial fission 1 protein OS=Homo sapiens GN=FIS1 PE=1 SV=2
Exportin-7 OS=Homo sapiens GN=XP07 PE=4 SV=1
Gamma-glutamyltransferase 5 OS=Homo sapiens GN=GGT5 PE=4 SV=1
Translin-associated protein X OS=Homo sapiens GN=TSNAX PE=1 SV=1
Membrane-associated progesterone receptor component 2 OS=Homo sapiens GN=PGRMC2 P
Sulfide:quinone oxidoreductase, mitochondrial OS=Homo sapiens GN=SQRDL PE=1 SV=1
Chromobox homolog 5 (HP1 alpha homolog, Drosophila), isoform CRA_a OS=Homo sapien
COP9 signalosome complex subunit 8 OS=Homo sapiens GN=COPS8 PE=4 SV=1
Mitochondrial-processing peptidase subunit alpha OS=Homo sapiens GN=PMPCA PE=1 SV
LIM and senescent cell antigen-like-containing domain protein 1 OS=Homo sapiens G
Low molecular weight phosphotyrosine protein phosphatase OS=Homo sapiens GN=ACP1
BTB/POZ domain-containing protein KCTD21 (Fragment) OS=Homo sapiens GN=KCTD21 PE=SPARC-like protein 1 OS=Homo sapiens GN=SPARCL1 PE=2 SV=1
B-cell receptor-associated protein 29 OS=Homo sapiens GN=BCAP29 PE=4 SV=1
Proteasome subunit beta type-6 OS=Homo sapiens GN=PSMB6 PE=1 SV=4
ADP-ribosylation factor-like protein 2 OS=Homo sapiens GN=ARL2 PE=1 SV=4

Golgi apparatus protein 1 OS=Homo sapiens GN=GLG1 PE=1 SV=2
Protein NDRG4 OS=Homo sapiens GN=NDRG4 PE=2 SV=1
L-aminoadipate-semialdehyde dehydrogenase-phosphopantetheinyl transferase (Fragme
TBC1 domain family member 17 OS=Homo sapiens GN=TBC1D17 PE=4 SV=1
Signal recognition particle 68 kDa protein OS=Homo sapiens GN=SRP68 PE=4 SV=1
60 kDa SS-A/Ro ribonucleoprotein OS=Homo sapiens GN=TROVE2 PE=1 SV=2
Peptidyl-prolyl cis-trans isomerase FKBP3 OS=Homo sapiens GN=FKBP3 PE=1 SV=1
NADPH--cytochrome P450 reductase OS=Homo sapiens GN=POR PE=4 SV=1
Isoform 2 of Formin-like protein 2 OS=Homo sapiens GN=FMNL2
Ethanolamine-phosphate cytidyltransferase OS=Homo sapiens GN=PCYT2 PE=2 SV=1
Disco-interacting protein 2 homolog B OS=Homo sapiens GN=DIP2B PE=1 SV=3
Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERBP1 P
Spermidine synthase OS=Homo sapiens GN=SRM PE=1 SV=1
Pre-mRNA branch site protein p14 OS=Homo sapiens GN=SF3B14 PE=1 SV=1
Ganglioside-induced differentiation-associated protein 1 OS=Homo sapiens GN=GDAP1
Arf-GAP domain and FG repeat-containing protein 1 OS=Homo sapiens GN=AGFG1 PE=4 S
Protein VAC14 homolog OS=Homo sapiens GN=VAC14 PE=4 SV=1
26S proteasome non-ATPase regulatory subunit 9 OS=Homo sapiens GN=PSMD9 PE=4 SV=1
Endoplasmic reticulum lectin 1 OS=Homo sapiens GN=ERLEC1 PE=4 SV=1
Secreted glypican-4 OS=Homo sapiens GN=GPC4 PE=2 SV=1
Nuclear receptor coactivator 5 OS=Homo sapiens GN=NCOA5 PE=1 SV=2
Protein kinase C and casein kinase substrate in neurons 2 (Fragment) OS=Homo sapi
Epidermal growth factor receptor OS=Homo sapiens GN=EGFR PE=3 SV=2
UPF0553 protein C9orf64 OS=Homo sapiens GN=C9orf64 PE=1 SV=1
Large neutral amino acids transporter small subunit 1 OS=Homo sapiens GN=SLC7A5 P
Splicing factor U2AF 35 kDa subunit OS=Homo sapiens GN=U2AF1 PE=1 SV=3
60S ribosomal protein L17 (Fragment) OS=Homo sapiens GN=RPL17 PE=3 SV=1
Sorting nexin-12 OS=Homo sapiens GN=SNX12 PE=1 SV=3
Lysophospholipid acyltransferase 7 OS=Homo sapiens GN=MBOAT7 PE=1 SV=2
Anoctamin-10 OS=Homo sapiens GN=ANO10 PE=1 SV=2
ER membrane protein complex subunit 2 OS=Homo sapiens GN=EMC2 PE=1 SV=1
Adenylate kinase isoenzyme 5 OS=Homo sapiens GN=AK5 PE=1 SV=2
Kynurenine--oxoglutarate transaminase 3 OS=Homo sapiens GN=CCBL2 PE=1 SV=1
Ras-related protein Rab-8B (Fragment) OS=Homo sapiens GN=RAB8B PE=3 SV=1
Probable ubiquitin carboxyl-terminal hydrolase FAF-Y OS=Homo sapiens GN=USP9Y PE=
Unconventional myosin-IXb OS=Homo sapiens GN=MYO9B PE=1 SV=3
Synaptopodin-2 (Fragment) OS=Homo sapiens GN=SYNP02 PE=4 SV=1
Nuclear ubiquitous casein and cyclin-dependent kinase substrate 1 OS=Homo sapiens
Choline dehydrogenase, mitochondrial OS=Homo sapiens GN=CHDH PE=2 SV=2
Aldehyde dehydrogenase family 16 member A1 OS=Homo sapiens GN=ALDH16A1 PE=4 SV=1
Protein OSCP1 OS=Homo sapiens GN=OSCP1 PE=4 SV=2
Deoxyuridine 5'-triphosphate nucleotidohydrolase, mitochondrial (Fragment) OS=Hom
Signal recognition particle 72 kDa protein OS=Homo sapiens GN=SRP72 PE=4 SV=1
Polyadenylate-binding protein-interacting protein 1 OS=Homo sapiens GN=PAIP1 PE=4
Katanin p80 WD40-containing subunit B1 OS=Homo sapiens GN=KATNB1 PE=1 SV=1
Estradiol 17-beta-dehydrogenase 11 OS=Homo sapiens GN=HSD17B11 PE=4 SV=1
Monofunctional C1-tetrahydrofolate synthase, mitochondrial OS=Homo sapiens GN=MTH

Cadherin-20 OS=Homo sapiens GN=CDH20 PE=2 SV=2
Prolargin OS=Homo sapiens GN=PRELP PE=1 SV=1
Mothers against decapentaplegic homolog 5 OS=Homo sapiens GN=SMAD5 PE=4 SV=1
Prefoldin subunit 3 OS=Homo sapiens GN=VBP1 PE=4 SV=1
U1 small nuclear ribonucleoprotein A OS=Homo sapiens GN=SNRPA PE=1 SV=3
WASH complex subunit 7 OS=Homo sapiens GN=KIAA1033 PE=1 SV=2
Glycerol-3-phosphate dehydrogenase 1-like protein OS=Homo sapiens GN=GPD1L PE=1 S
Gamma-glutamyl hydrolase OS=Homo sapiens GN=GGH PE=1 SV=2
Ataxin-10 OS=Homo sapiens GN=ATXN10 PE=1 SV=1
Splicing factor 3B subunit 1 OS=Homo sapiens GN=SF3B1 PE=1 SV=3
Diphosphoinositol polyphosphate phosphohydrolase 2 OS=Homo sapiens GN=NUDT4 PE=1
PRA1 family protein 2 OS=Homo sapiens GN=PRAF2 PE=1 SV=1
Calponin-2 OS=Homo sapiens GN=CNN2 PE=2 SV=1
Phosphatidylinositide phosphatase SAC1 OS=Homo sapiens GN=SACM1L PE=2 SV=1
Platelet-activating factor acetylhydrolase IB subunit beta OS=Homo sapiens GN=PAF
Procollagen-lysine,2-oxoglutarate 5-dioxygenase 2 OS=Homo sapiens GN=PLOD2 PE=4 S
Integrin alpha-6 (Fragment) OS=Homo sapiens GN=ITGA6 PE=3 SV=1
Lysosomal protective protein OS=Homo sapiens GN=CTSA PE=1 SV=2
1-acyl-sn-glycerol-3-phosphate acyltransferase gamma OS=Homo sapiens GN=AGPAT3 PE
RNA-binding protein Nova-2 OS=Homo sapiens GN=NOVA2 PE=1 SV=1
Afamin OS=Homo sapiens GN=AFM PE=1 SV=1
Soluble CD163 OS=Homo sapiens GN=CD163 PE=4 SV=1
Nuclear pore complex protein Nup205 OS=Homo sapiens GN=NUP205 PE=1 SV=3
Cathepsin Z OS=Homo sapiens GN=CTSZ PE=1 SV=1
Histone H1.0 OS=Homo sapiens GN=H1F0 PE=1 SV=3
Spectrin beta chain, non-erythrocytic 4 OS=Homo sapiens GN=SPTBN4 PE=4 SV=1
Armadillo repeat protein deleted in velo-cardio-facial syndrome OS=Homo sapiens G
Dedicator of cytokinesis protein 4 (Fragment) OS=Homo sapiens GN=DOCK4 PE=4 SV=1
Protein arginine N-methyltransferase 3 OS=Homo sapiens GN=PRMT3 PE=1 SV=3
Cytochrome b-c1 complex subunit 8 OS=Homo sapiens GN=UQCRQ PE=1 SV=4
Putative hydroxypyruvate isomerase (Fragment) OS=Homo sapiens GN=HYI PE=4 SV=1
Elongator complex protein 4 OS=Homo sapiens GN=ELP4 PE=2 SV=1
High mobility group protein B2 OS=Homo sapiens GN=HMGB2 PE=1 SV=2
Isoform 2 of PDZ and LIM domain protein 3 OS=Homo sapiens GN=PDLIM3
Serine dehydratase-like OS=Homo sapiens GN=SDSL PE=1 SV=1
Protein RCC2 OS=Homo sapiens GN=RCC2 PE=1 SV=2
Na(+)/H(+) exchange regulatory cofactor NHE-RF2 (Fragment) OS=Homo sapiens GN=SLC
60S ribosomal protein L30 OS=Homo sapiens GN=RPL30 PE=1 SV=2
Spermine synthase OS=Homo sapiens GN=SMS PE=2 SV=1
Atlantin-1 OS=Homo sapiens GN=ATL1 PE=4 SV=1
Trans-2,3-enoyl-CoA reductase OS=Homo sapiens GN=TECR PE=2 SV=1
Ankyrin repeat and MYND domain containing 2, isoform CRA_c OS=Homo sapiens GN=ANK
Bifunctional coenzyme A synthase OS=Homo sapiens GN=COASY PE=1 SV=4
Thymopietin OS=Homo sapiens GN=TMPO PE=2 SV=1
Isoform 2 of Mitochondrial import inner membrane translocase subunit TIM50 OS=Hom
Prostaglandin E synthase 2 OS=Homo sapiens GN=PTGES2 PE=4 SV=2
LUC7-like (S. cerevisiae) OS=Homo sapiens GN=LUC7L PE=4 SV=1

Phospholipid scramblase 1 OS=Homo sapiens GN=PLSCR1 PE=2 SV=1
Uncharacterized protein Clorf198 OS=Homo sapiens GN=Clorf198 PE=4 SV=1
GMP reductase 2 OS=Homo sapiens GN=GMPR2 PE=3 SV=1
SUMO-conjugating enzyme UBC9 OS=Homo sapiens GN=UBE2I PE=3 SV=1
Complement C5 OS=Homo sapiens GN=C5 PE=1 SV=4
Receptor-type tyrosine-protein phosphatase F OS=Homo sapiens GN=PTPRF PE=1 SV=2
Enhancer of mRNA-decapping protein 4 OS=Homo sapiens GN=EDC4 PE=1 SV=1
60S ribosomal protein L34 OS=Homo sapiens GN=RPL34 PE=1 SV=3
Endophilin-B1 OS=Homo sapiens GN=SH3GLB1 PE=1 SV=1
mRNA export factor OS=Homo sapiens GN=RAE1 PE=4 SV=1
Transcription elongation factor B (SIII), polypeptide 2 (18kDa, elongin B), isofo
Protein SCO1 homolog, mitochondrial OS=Homo sapiens GN=SCO1 PE=4 SV=1
Isoform 2 of Cytoplasmic dynein 1 intermediate chain 1 OS=Homo sapiens GN=DYNC1I1
Receptor-type tyrosine-protein phosphatase delta OS=Homo sapiens GN=PTPRD PE=4 SV
CD63 antigen OS=Homo sapiens GN=CD63 PE=4 SV=1
Serine/threonine-protein phosphatase 2A 56 kDa regulatory subunit gamma isoform 0
Coiled-coil domain-containing protein 124 OS=Homo sapiens GN=CCDC124 PE=1 SV=1
Junctional adhesion molecule C OS=Homo sapiens GN=JAM3 PE=1 SV=1
Synaptopodin OS=Homo sapiens GN=SYNPO PE=1 SV=2
MAP/microtubule affinity-regulating kinase 4 OS=Homo sapiens GN=MARK4 PE=1 SV=1
Transmembrane protein 205 OS=Homo sapiens GN=TMEM205 PE=1 SV=1
Sideroflexin-5 OS=Homo sapiens GN=SFXN5 PE=2 SV=1
Protein FAM49A OS=Homo sapiens GN=FAM49A PE=1 SV=1
Importin subunit alpha OS=Homo sapiens GN=KPNA6 PE=3 SV=1
60S ribosomal protein L21 OS=Homo sapiens GN=RPL21 PE=1 SV=2
E3 UFM1-protein ligase 1 OS=Homo sapiens GN=UFL1 PE=1 SV=2
Long-chain-fatty-acid--CoA ligase 6 OS=Homo sapiens GN=ACSL6 PE=4 SV=1
Prolyl 4-hydroxylase subunit alpha-1 OS=Homo sapiens GN=P4HA1 PE=4 SV=1
Lipoamide acyltransferase component of branched-chain alpha-keto acid dehydrogena
ATP-dependent Clp protease ATP-binding subunit clpX-like, mitochondrial OS=Homo s
ATPase inhibitor, mitochondrial OS=Homo sapiens GN=ATPIF1 PE=1 SV=1
Kinesin-like protein KIF2A OS=Homo sapiens GN=KIF2A PE=3 SV=1
Prefoldin subunit 2 OS=Homo sapiens GN=PFDN2 PE=1 SV=1
Copper transport protein ATOX1 OS=Homo sapiens GN=ATOX1 PE=4 SV=1
Signal recognition particle 54 kDa protein OS=Homo sapiens GN=SRP54 PE=1 SV=1
Optic atrophy 3 protein OS=Homo sapiens GN=OPA3 PE=1 SV=1
HLA class I histocompatibility antigen, B-56 alpha chain (Fragment) OS=Homo sapie
Serine/threonine-protein phosphatase (Fragment) OS=Homo sapiens GN=PPP5C PE=3 SV=
Seizure 6-like protein 2 OS=Homo sapiens GN=SEZ6L2 PE=4 SV=1
Azurocidin OS=Homo sapiens GN=AZU1 PE=1 SV=3
60S ribosomal protein L11 (Fragment) OS=Homo sapiens GN=RPL11 PE=2 SV=1
Hepatoma-derived growth factor-related protein 3 (Fragment) OS=Homo sapiens GN=HD
Ig heavy chain V-III region BRO OS=Homo sapiens PE=1 SV=1
Engulfment and cell motility protein 1 OS=Homo sapiens GN=ELM01 PE=1 SV=2
Rap1 GTPase-activating protein 1 OS=Homo sapiens GN=RAP1GAP PE=4 SV=1
DNA repair protein RAD50 (Fragment) OS=Homo sapiens GN=RAD50 PE=4 SV=1
Isoform 2 of Receptor-type tyrosine-protein phosphatase epsilon OS=Homo sapiens G

Leucine-rich repeat-containing protein 59 OS=Homo sapiens GN=LRR59 PE=1 SV=1
PLEKHA5 protein OS=Homo sapiens GN=PLEKHA5 PE=2 SV=1
Junctional adhesion molecule B OS=Homo sapiens GN=JAM2 PE=4 SV=1
Protein TSSC1 OS=Homo sapiens GN=TSSC1 PE=4 SV=2
Protein SEC13 homolog OS=Homo sapiens GN=SEC13 PE=4 SV=1
Leucine-rich repeat flightless-interacting protein 2 OS=Homo sapiens GN=LRRFIP2 P
Guanylate kinase OS=Homo sapiens GN=GUK1 PE=4 SV=1
Small nuclear ribonucleoprotein-associated protein OS=Homo sapiens GN=SNRPN PE=2
Engulfment and cell motility protein 2 OS=Homo sapiens GN=ELM2 PE=2 SV=1
Transforming acidic coiled-coil-containing protein 1 OS=Homo sapiens GN=TACC1 PE=
Replication protein A 14 kDa subunit OS=Homo sapiens GN=RPA3 PE=4 SV=1
Calpastatin OS=Homo sapiens GN=CAST PE=2 SV=1
BRO1 domain-containing protein BROX OS=Homo sapiens GN=Clorf58 PE=2 SV=1
Glycogen [starch] synthase, muscle (Fragment) OS=Homo sapiens GN=GYS1 PE=4 SV=1
Selenocysteine-specific elongation factor OS=Homo sapiens GN=EEFSEC PE=4 SV=1
Protein BUD31 homolog OS=Homo sapiens GN=BUD31 PE=4 SV=1
Protein FAM54B (Fragment) OS=Homo sapiens GN=FAM54B PE=4 SV=1
Echinoderm microtubule-associated protein-like 2 OS=Homo sapiens GN=EML2 PE=4 SV=
Synaptophysin-like protein 1 OS=Homo sapiens GN=SYPL1 PE=4 SV=1
Complement C1q subcomponent subunit B OS=Homo sapiens GN=C1QB PE=4 SV=1
Cytochrome b5 type B OS=Homo sapiens GN=CYB5B PE=3 SV=1
Carboxypeptidase Q (Fragment) OS=Homo sapiens GN=CPQ PE=4 SV=1
Aggrecan core protein OS=Homo sapiens GN=ACAN PE=4 SV=2
ATP-binding cassette sub-family D member 3 OS=Homo sapiens GN=ABCD3 PE=3 SV=1
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 5, mitochondrial OS=Hom
Ubiquitin-like modifier-activating enzyme ATG7 OS=Homo sapiens GN=ATG7 PE=4 SV=1
Neurolysin, mitochondrial OS=Homo sapiens GN=NLN PE=3 SV=1
D-beta-hydroxybutyrate dehydrogenase, mitochondrial OS=Homo sapiens GN=BDH1 PE=3
Protein O-GlcNAcase OS=Homo sapiens GN=MGEA5 PE=4 SV=1
60S ribosomal protein L8 OS=Homo sapiens GN=RPL8 PE=4 SV=1
Vacuolar protein sorting-associated protein 51 homolog (Fragment) OS=Homo sapiens
Methylosome subunit pICln OS=Homo sapiens GN=CLNS1A PE=4 SV=1
Parkinson disease 7 domain-containing protein 1 OS=Homo sapiens GN=PDDC1 PE=4 SV=
tRNA methyltransferase 112 homolog OS=Homo sapiens GN=TRMT112 PE=4 SV=1
Acyl-coenzyme A thioesterase 13 OS=Homo sapiens GN=ACOT13 PE=4 SV=1
Tumor susceptibility gene 101 protein OS=Homo sapiens GN=TSG101 PE=4 SV=1
Serum paraoxonase/arylesterase 1 OS=Homo sapiens GN=PON1 PE=4 SV=1
NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 12 OS=Homo sapiens GN=
60S ribosomal protein L6 (Fragment) OS=Homo sapiens GN=RPL6 PE=3 SV=1
Torsin-1A-interacting protein 1 (Fragment) OS=Homo sapiens GN=TOR1AIP1 PE=4 SV=1
Diphosphomevalonate decarboxylase (Fragment) OS=Homo sapiens GN=MVD PE=4 SV=1
Glutathione S-transferase theta-1 (Fragment) OS=Homo sapiens GN=GSTT1 PE=4 SV=1
Serrate RNA effector molecule homolog (Fragment) OS=Homo sapiens GN=SRRT PE=4 SV=
Thioredoxin domain-containing protein 17 OS=Homo sapiens GN=TXNDC17 PE=4 SV=1
Phosphatidylinositol 4-kinase alpha OS=Homo sapiens GN=PI4KA PE=4 SV=1
60S ribosomal protein L38 (Fragment) OS=Homo sapiens GN=RPL38 PE=3 SV=1
Unconventional myosin-If OS=Homo sapiens GN=MYO1F PE=1 SV=3

Nucleolar protein 56 OS=Homo sapiens GN=NOP56 PE=1 SV=4
Lipid phosphate phosphohydrolase 1 OS=Homo sapiens GN=PPAP2A PE=1 SV=1
Regulator of G-protein signaling 10 OS=Homo sapiens GN=RGS10 PE=1 SV=2
DnaJ homolog subfamily C member 13 OS=Homo sapiens GN=DNAJC13 PE=1 SV=5
Phosphatidate cytidyltransferase 2 OS=Homo sapiens GN=CDS2 PE=1 SV=1
Ig kappa chain V-I region CAR OS=Homo sapiens PE=1 SV=1
Ig kappa chain V-IV region Len OS=Homo sapiens PE=1 SV=2
Replication protein A 32 kDa subunit OS=Homo sapiens GN=RPA2 PE=1 SV=1
DNA-directed RNA polymerases I, II, and III subunit RPABC1 OS=Homo sapiens GN=POL
Serum amyloid A-4 protein OS=Homo sapiens GN=SAA4 PE=1 SV=2
60S ribosomal protein L29 OS=Homo sapiens GN=RPL29 PE=1 SV=2
Glutamate--cysteine ligase regulatory subunit OS=Homo sapiens GN=GCLM PE=1 SV=1
Proteolipid protein 2 OS=Homo sapiens GN=PLP2 PE=1 SV=1
Cullin-3 OS=Homo sapiens GN=CUL3 PE=1 SV=2
Eukaryotic translation initiation factor 4H OS=Homo sapiens GN=EIF4H PE=1 SV=5
RNA binding protein, autoantigenic (HnRNP-associated with lethal yellow homolog (Cell adhesion molecule 3 OS=Homo sapiens GN=CADM3 PE=1 SV=1
Protein canopy homolog 4 OS=Homo sapiens GN=CNPY4 PE=2 SV=1
Osteoclast-stimulating factor 1 OS=Homo sapiens GN=OSTF1 PE=1 SV=2
Sortilin OS=Homo sapiens GN=SORT1 PE=1 SV=3
Coronin-1B OS=Homo sapiens GN=CORO1B PE=1 SV=1
Synaptotagmin-11 OS=Homo sapiens GN=SYT11 PE=1 SV=2
Fibronectin type III and SPRY domain-containing protein 1 OS=Homo sapiens GN=FSD1
GrpE protein homolog 1, mitochondrial OS=Homo sapiens GN=GRPEL1 PE=1 SV=2
Translation initiation factor eIF-2B subunit gamma OS=Homo sapiens GN=EIF2B3 PE=1
Poly(ADP-ribose) glycohydrolase ARH3 OS=Homo sapiens GN=ADPRHL2 PE=1 SV=1
C-type mannose receptor 2 OS=Homo sapiens GN=MRC2 PE=1 SV=2
Protein NipSnap homolog 3A OS=Homo sapiens GN=NIPSNAP3A PE=1 SV=2
Neudesin OS=Homo sapiens GN=NENF PE=1 SV=1
UPF0568 protein C14orf166 OS=Homo sapiens GN=C14orf166 PE=1 SV=1
Coiled-coil domain-containing protein 22 OS=Homo sapiens GN=CCDC22 PE=1 SV=1
Mitochondrial import inner membrane translocase subunit Tim13 OS=Homo sapiens GN=
Endothelin B receptor-like protein 2 OS=Homo sapiens GN=GPR37L1 PE=1 SV=2
Ig lambda chain V region 4A OS=Homo sapiens PE=4 SV=1
COMM domain containing 6 OS=Homo sapiens GN=COMMD6 PE=4 SV=1
2-aminoethanethiol dioxygenase OS=Homo sapiens GN=ADO PE=1 SV=2
Ig kappa chain V-I region AG OS=Homo sapiens PE=1 SV=1
Leucine-rich repeat and calponin homology domain-containing protein 3 OS=Homo sap
Spartin OS=Homo sapiens GN=SPG20 PE=1 SV=1
Ewing sarcoma breakpoint region 1 OS=Homo sapiens GN=EWSR1 PE=4 SV=1
Sickle tail protein homolog OS=Homo sapiens GN=KIAA1217 PE=4 SV=2
Aspartyl/asparaginyl beta-hydroxylase OS=Homo sapiens GN=ASPH PE=4 SV=1
Dolichol-phosphate mannosyltransferase subunit 3 OS=Homo sapiens GN=DPM3 PE=1 SV=
Small nuclear ribonucleoprotein E OS=Homo sapiens GN=SNRPE PE=4 SV=1
Tubulin tyrosine ligase-like family, member 12 OS=Homo sapiens GN=TTLL12 PE=4 SV=
Parathymosin OS=Homo sapiens GN=PTMS PE=4 SV=1
Ig lambda chain V-I region HA OS=Homo sapiens PE=1 SV=1

Trafficking protein particle complex 3, isoform CRA_a OS=Homo sapiens GN=TRAPPC3
A kinase (PRKA) anchor protein 13, isoform CRA_a OS=Homo sapiens GN=AKAP13 PE=4 S
Golgi reassembly-stacking protein 2 OS=Homo sapiens GN=GORASP2 PE=2 SV=1
UPF0452 protein C7orf41 (Fragment) OS=Homo sapiens GN=C7orf41 PE=4 SV=1
Serum amyloid P-component OS=Homo sapiens GN=APCS PE=1 SV=2
Cytochrome b-c1 complex subunit 9 OS=Homo sapiens GN=UQCR10 PE=1 SV=3
Coiled-coil domain-containing protein 58 OS=Homo sapiens GN=CCDC58 PE=4 SV=1
G-protein-coupled receptor family C group 5 member B OS=Homo sapiens GN=GPRC5B PE
NEDD8-activating enzyme E1 regulatory subunit OS=Homo sapiens GN=NAE1 PE=4 SV=2
Allograft inflammatory factor 1 OS=Homo sapiens GN=AIF1 PE=1 SV=1
Urea transporter 1 OS=Homo sapiens GN=SLC14A1 PE=2 SV=2
Caveolin (Fragment) OS=Homo sapiens GN=CAV1 PE=3 SV=1
Inositol 1,4,5-trisphosphate receptor type 1 OS=Homo sapiens GN=ITPR1 PE=4 SV=2
Cleavage and polyadenylation specificity factor subunit 1 OS=Homo sapiens GN=CPSF
Synaptogyrin 1 OS=Homo sapiens GN=SYNGR1 PE=4 SV=1
Synaptogyrin-3 OS=Homo sapiens GN=SYNGR3 PE=4 SV=1
EH domain-containing protein 4 OS=Homo sapiens GN=EHD4 PE=1 SV=1
Hippocalcin-like protein 1 OS=Homo sapiens GN=HPCAL1 PE=1 SV=3
Phosphatidylinositol 5-phosphate 4-kinase type-2 beta OS=Homo sapiens GN=PIP4K2B
Serine/arginine-rich-splicing factor 5 OS=Homo sapiens GN=SRSF5 PE=2 SV=1
Pterin-4-alpha-carbinolamine dehydratase 2 (Fragment) OS=Homo sapiens GN=PCBD2 PE
Heterogeneous nuclear ribonucleoprotein L-like OS=Homo sapiens GN=HNRPLL PE=4 SV=
AFG3-like protein 2 OS=Homo sapiens GN=AFG3L2 PE=1 SV=2
Ras-related protein Rab-35 (Fragment) OS=Homo sapiens GN=RAB35 PE=3 SV=1
DmX-like protein 2 OS=Homo sapiens GN=DMXL2 PE=4 SV=1
Protein NDRG3 OS=Homo sapiens GN=NDRG3 PE=2 SV=1
Beta-2-syntrophin OS=Homo sapiens GN=SNTB2 PE=1 SV=1
Probable Xaa-Pro aminopeptidase 3 OS=Homo sapiens GN=XPNPEP3 PE=1 SV=1
Ig heavy chain V-III region GAL OS=Homo sapiens PE=1 SV=1
UDP-N-acetylhexosamine pyrophosphorylase OS=Homo sapiens GN=UAP1 PE=1 SV=3
Serine/threonine-protein kinase 3 OS=Homo sapiens GN=STK3 PE=2 SV=1
Protein diaphanous homolog 1 OS=Homo sapiens GN=DIAPH1 PE=2 SV=1
Gamma-interferon-inducible lysosomal thiol reductase OS=Homo sapiens GN=IFI30 PE=
Integrin alpha-2 OS=Homo sapiens GN=ITGA2 PE=3 SV=1
1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase gamma-1 OS=Homo sapiens
2',5'-phosphodiesterase 12 OS=Homo sapiens GN=PDE12 PE=4 SV=1
Chymase OS=Homo sapiens GN=CMA1 PE=1 SV=1
La-related protein 1 OS=Homo sapiens GN=LARP1 PE=1 SV=2
Growth arrest-specific protein 7 OS=Homo sapiens GN=GAS7 PE=2 SV=1
Proteasome (Prosome, macropain) subunit, beta type, 9 (Large multifunctional pept
RAB3GAP1 protein OS=Homo sapiens GN=RAB3GAP1 PE=2 SV=1
Coatomer protein complex, subunit epsilon, isoform CRA_e OS=Homo sapiens GN=COPE
Protein SET OS=Homo sapiens GN=SET PE=2 SV=2
Putative Golgi pH regulator C OS=Homo sapiens GN=GPR89C PE=5 SV=2
Adaptin ear-binding coat-associated protein 1 OS=Homo sapiens GN=NECAP1 PE=2 SV=1
Protein FAM98B OS=Homo sapiens GN=FAM98B PE=4 SV=1
Cat eye syndrome critical region protein 5 OS=Homo sapiens GN=CECR5 PE=4 SV=1

Serine/arginine repetitive matrix 1 OS=Homo sapiens GN=SRRM1 PE=4 SV=1
Periostin OS=Homo sapiens GN=POSTN PE=4 SV=1
Rho guanine nucleotide exchange factor (GEF) 7 OS=Homo sapiens GN=ARHGEF7 PE=4 SV=1
Epidermal growth factor receptor pathway substrate 15 OS=Homo sapiens GN=EPS15 PE=4 SV=1
ICSU OS=Homo sapiens GN=ISCU PE=2 SV=1
Ran-binding protein 9 OS=Homo sapiens GN=RANBP9 PE=2 SV=1
SUMO-1 activating enzyme subunit 1, isoform CRA_a OS=Homo sapiens GN=SAE1 PE=2 SV=1
Eukaryotic translation initiation factor 3 subunit F OS=Homo sapiens GN=EIF3F PE=4 SV=1
Coronin OS=Homo sapiens GN=CORO7 PE=2 SV=1
CUGBP Elav-like family member 2 OS=Homo sapiens GN=CELF2 PE=2 SV=1
F-box only protein 17 OS=Homo sapiens GN=FBXO17 PE=2 SV=1
Chromosome 2 open reading frame 18, isoform CRA_c OS=Homo sapiens GN=C2orf18 PE=2 SV=1
Mitochondrial dicarboxylate carrier OS=Homo sapiens GN=SLC25A10 PE=2 SV=1
Methylosome protein 50 OS=Homo sapiens GN=WDR77 PE=2 SV=1
Far upstream element-binding protein 1 OS=Homo sapiens GN=FUBP1 PE=2 SV=1
Serine/arginine-rich-splicing factor 11 OS=Homo sapiens GN=SRSF11 PE=2 SV=1
Methionine aminopeptidase OS=Homo sapiens GN=METAP2 PE=2 SV=1
26S proteasome non-ATPase regulatory subunit 5 OS=Homo sapiens GN=PSMD5 PE=2 SV=1
Ribulose-phosphate 3-epimerase OS=Homo sapiens GN=RPE PE=2 SV=1
Soluble Alc-alpha OS=Homo sapiens GN=CLSTN1 PE=2 SV=1
Glutathione S-transferase theta-2 OS=Homo sapiens GN=GSTT2 PE=4 SV=2
HCG2043423, isoform CRA_a OS=Homo sapiens GN=ATP6V1G2 PE=4 SV=1
Tryptophan--tRNA ligase, mitochondrial OS=Homo sapiens GN=WARS2 PE=2 SV=1
Cytosolic Fe-S cluster assembly factor NUBP2 OS=Homo sapiens GN=NUBP2 PE=2 SV=1
Glutamate receptor delta-1 subunit OS=Homo sapiens GN=GRID1 PE=2 SV=1
Monoglyceride lipase OS=Homo sapiens GN=MGLL PE=2 SV=1
Hyccin OS=Homo sapiens GN=FAM126A PE=4 SV=1
Integral membrane protein 2C OS=Homo sapiens GN=ITM2C PE=4 SV=2
Elongation factor G, mitochondrial OS=Homo sapiens GN=GFM1 PE=4 SV=1
Rho GTPase-activating protein 17 (Fragment) OS=Homo sapiens GN=ARHGAP17 PE=4 SV=3
Lanosterol synthase (Fragment) OS=Homo sapiens GN=LSS PE=4 SV=1
Beta-galactosidase (Fragment) OS=Homo sapiens GN=GLB1 PE=4 SV=1
Ubiquitin-like modifier-activating enzyme 5 (Fragment) OS=Homo sapiens GN=UBA5 PE=4 SV=1
Synaptosomal-associated protein (Fragment) OS=Homo sapiens GN=SNAP29 PE=3 SV=1
Phosphoserine phosphatase (Fragment) OS=Homo sapiens GN=PSPH PE=4 SV=1
Secernin-3 (Fragment) OS=Homo sapiens GN=SCRN3 PE=4 SV=1
Acylamino-acid-releasing enzyme OS=Homo sapiens GN=APEH PE=4 SV=1
STAM-binding protein (Fragment) OS=Homo sapiens GN=STAMPB PE=4 SV=1
L-2-hydroxyglutarate dehydrogenase, mitochondrial OS=Homo sapiens GN=L2HGDH PE=4 SV=1
Neurologin 1 OS=Homo sapiens GN=nlgn1 PE=2 SV=1
40S ribosomal protein S23 OS=Homo sapiens GN=RPS23 PE=3 SV=1
Dynactin subunit 6 OS=Homo sapiens GN=DCTN6 PE=4 SV=1
Eukaryotic translation initiation factor 3 subunit E (Fragment) OS=Homo sapiens GN=EIF3E PE=4 SV=1
Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B alpha isoform
Protein LYRIC OS=Homo sapiens GN=MTDH PE=4 SV=1
Apoptotic chromatin condensation inducer in the nucleus OS=Homo sapiens GN=ACIN1 PE=4 SV=1
Mitochondrial-processing peptidase subunit beta OS=Homo sapiens GN=PMPCB PE=3 SV=1

Nucleolysin TIAR (Fragment) OS=Homo sapiens GN=TIAL1 PE=4 SV=1
Stromal membrane-associated protein 1 OS=Homo sapiens GN=SMAP1 PE=4 SV=1
Anoctamin OS=Homo sapiens GN=ANO6 PE=3 SV=1
THO complex subunit 4 OS=Homo sapiens GN=ALYREF PE=4 SV=1
Suprabasin OS=Homo sapiens GN=SBSN PE=4 SV=1
DNA topoisomerase 2 (Fragment) OS=Homo sapiens GN=TOP2B PE=3 SV=1
Apolipoprotein L1 OS=Homo sapiens GN=APOL1 PE=4 SV=1
7-dehydrocholesterol reductase (Fragment) OS=Homo sapiens GN=DHCR7 PE=4 SV=1
Acyl-coenzyme A thioesterase 8 OS=Homo sapiens GN=ACOT8 PE=4 SV=1
Signal recognition particle receptor subunit alpha OS=Homo sapiens GN=SRPR PE=4 SV=1
GDP-L-fucose synthase (Fragment) OS=Homo sapiens GN=TSTA3 PE=4 SV=1
Phosphatidylinositol-binding clathrin assembly protein OS=Homo sapiens GN=PICALM
Eukaryotic translation initiation factor 3 subunit M (Fragment) OS=Homo sapiens GN=EIF3M
Stromal interaction molecule 1 OS=Homo sapiens GN=STIM1 PE=4 SV=1
Poly(U)-binding-splicing factor PUF60 (Fragment) OS=Homo sapiens GN=PUF60 PE=4 SV=1
Probable asparagine--tRNA ligase, mitochondrial (Fragment) OS=Homo sapiens GN=NARL
Vacuolar protein sorting-associated protein 26B OS=Homo sapiens GN=VPS26B PE=4 SV=1
Oligoribonuclease, mitochondrial OS=Homo sapiens GN=REXO2 PE=4 SV=1
Diablo homolog, mitochondrial (Fragment) OS=Homo sapiens GN=DIABLO PE=4 SV=1
Complement component C8 beta chain OS=Homo sapiens GN=C8B PE=4 SV=1
Vacuolar protein sorting-associated protein 45 OS=Homo sapiens GN=VPS45 PE=4 SV=1
Ubiquitin thioesterase OTUB1 OS=Homo sapiens GN=OTUB1 PE=4 SV=1
N-acylneuraminate cytidyltransferase (Fragment) OS=Homo sapiens GN=CMAS PE=4 SV=1
ELKS/Rab6-interacting/CAST family member 1 OS=Homo sapiens GN=ERC1 PE=4 SV=1
Kinase suppressor of Ras 1 OS=Homo sapiens GN=KSR1 PE=4 SV=1
Oligophrenin-1 OS=Homo sapiens GN=OPHN1 PE=4 SV=1
N-sulphoglucosamine sulphohydrolase OS=Homo sapiens GN=SGSH PE=4 SV=1
Epimerase family protein SDR39U1 OS=Homo sapiens GN=SDR39U1 PE=4 SV=1
2-methoxy-6-polyprenyl-1,4-benzoquinol methylase, mitochondrial (Fragment) OS=Homo sapiens GN=HVCN1 PE=4 SV=1
Voltage-gated hydrogen channel 1 OS=Homo sapiens GN=HVCN1 PE=4 SV=1
Lysozyme C OS=Homo sapiens GN=LYZ PE=3 SV=1
Iron-sulfur protein NUBPL (Fragment) OS=Homo sapiens GN=NUBPL PE=4 SV=1
GMP synthase [glutamine-hydrolyzing] OS=Homo sapiens GN=GMPS PE=4 SV=1
Golgi integral membrane protein 4 OS=Homo sapiens GN=GOLIM4 PE=4 SV=1
Exocyst complex component 5 OS=Homo sapiens GN=EXOC5 PE=4 SV=1
Abhydrolase domain-containing protein 14B OS=Homo sapiens GN=ABHD14B PE=4 SV=1
Polyribonucleotide nucleotidyltransferase 1, mitochondrial OS=Homo sapiens GN=PNP2
Developmentally-regulated GTP-binding protein 1 OS=Homo sapiens GN=DRG1 PE=4 SV=1
Integrin alpha-7 light chain (Fragment) OS=Homo sapiens GN=ITGA7 PE=4 SV=1
SRA stem-loop-interacting RNA-binding protein, mitochondrial OS=Homo sapiens GN=SRP
Myotubularin-related protein 5 OS=Homo sapiens GN=SBF1 PE=4 SV=1
Immunoglobulin superfamily member 21 (Fragment) OS=Homo sapiens GN=IGSF21 PE=4 SV=1
Mitochondrial carrier homolog 1 (Fragment) OS=Homo sapiens GN=MTCH1 PE=3 SV=1
Protein C6orf108 (Fragment) OS=Homo sapiens GN=C6orf108 PE=4 SV=1
Ribonucleoside-diphosphate reductase subunit M2 B (Fragment) OS=Homo sapiens GN=RDR
Protein DDI1 homolog 2 (Fragment) OS=Homo sapiens GN=DDI2 PE=4 SV=1
Dehydrogenase/reductase SDR family member 7 (Fragment) OS=Homo sapiens GN=DHRS7 P

Pro-interleukin-16 (Fragment) OS=Homo sapiens GN=IL16 PE=4 SV=2
U2 small nuclear ribonucleoprotein A' (Fragment) OS=Homo sapiens GN=SNRPA1 PE=4 SV=1
WD repeat-containing protein 61 OS=Homo sapiens GN=WDR61 PE=4 SV=1
Dynactin subunit 5 OS=Homo sapiens GN=DCTN5 PE=4 SV=1
E3 ubiquitin-protein ligase ARIH1 (Fragment) OS=Homo sapiens GN=ARIH1 PE=4 SV=1
Nucleoside diphosphate kinase OS=Homo sapiens GN=NME3 PE=3 SV=1
Uncharacterized protein OS=Homo sapiens PE=4 SV=1
Melanoma-associated antigen D2 OS=Homo sapiens GN=MAGED2 PE=4 SV=1
Methionine adenosyltransferase 2 subunit beta (Fragment) OS=Homo sapiens GN=MAT2B
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 2, mitochondrial (Fragment)
Uncharacterized protein OS=Homo sapiens PE=4 SV=1
Leucine carboxyl methyltransferase 1 (Fragment) OS=Homo sapiens GN=LCMT1 PE=4 SV=1
Antibacterial protein LL-37 OS=Homo sapiens GN=CAMP PE=4 SV=1
Lymphocyte antigen 6H OS=Homo sapiens GN=LY6H PE=4 SV=1
Dehydrogenase/reductase SDR family member 7B (Fragment) OS=Homo sapiens GN=DHRS7B
Ribosomal protein L19 OS=Homo sapiens GN=RPL19 PE=3 SV=1
Syntaxin-binding protein 3 OS=Homo sapiens GN=STXBP3 PE=1 SV=2
Agrin OS=Homo sapiens GN=AGRN PE=1 SV=4
Programmed cell death protein 5 OS=Homo sapiens GN=PDCD5 PE=1 SV=3
Tax1-binding protein 3 OS=Homo sapiens GN=TAX1BP3 PE=1 SV=2
Secretory carrier-associated membrane protein 1 OS=Homo sapiens GN=SCAMP1 PE=1 SV=1
Neurosecretory protein VGF OS=Homo sapiens GN=VGF PE=1 SV=2
ER membrane protein complex subunit 8 OS=Homo sapiens GN=EMC8 PE=1 SV=1
Striatin OS=Homo sapiens GN=STRN PE=1 SV=4
Retinal rod rhodopsin-sensitive cGMP 3',5'-cyclic phosphodiesterase subunit delta
Mitochondrial import inner membrane translocase subunit Tim8 A OS=Homo sapiens GN=TIM8A
28S ribosomal protein S14, mitochondrial OS=Homo sapiens GN=MRPS14 PE=1 SV=1
NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 1 OS=Homo sapiens GN=ND1
Cytokine receptor-like factor 1 OS=Homo sapiens GN=CRLF1 PE=1 SV=1
Vesicle-associated membrane protein 5 OS=Homo sapiens GN=VAMP5 PE=1 SV=1
Myelin protein zero-like protein 1 OS=Homo sapiens GN=MPZL1 PE=1 SV=1
Ig delta chain C region OS=Homo sapiens GN=IGHD PE=1 SV=2
Apolipoprotein C-II OS=Homo sapiens GN=APOC2 PE=1 SV=1
Complement C1q subcomponent subunit A OS=Homo sapiens GN=C1QA PE=1 SV=2
Metallothionein-2 OS=Homo sapiens GN=MT2A PE=1 SV=1
Alkaline phosphatase, tissue-nonspecific isozyme OS=Homo sapiens GN=ALPL PE=1 SV=1
Intercellular adhesion molecule 1 OS=Homo sapiens GN=ICAM1 PE=1 SV=2
Complement component C8 alpha chain OS=Homo sapiens GN=C8A PE=1 SV=2
Complement component C8 gamma chain OS=Homo sapiens GN=C8G PE=1 SV=3
U1 small nuclear ribonucleoprotein 70 kDa OS=Homo sapiens GN=SNRNP70 PE=1 SV=2
Secretogranin-2 OS=Homo sapiens GN=SCG2 PE=1 SV=2
Integrin beta-4 OS=Homo sapiens GN=ITGB4 PE=1 SV=5
Tubulin gamma-1 chain OS=Homo sapiens GN=TUBG1 PE=1 SV=2
Protein S100-A4 OS=Homo sapiens GN=S100A4 PE=1 SV=1
Glutaredoxin-1 OS=Homo sapiens GN=GLRX PE=1 SV=2
Alpha-taxilin OS=Homo sapiens GN=TXLNA PE=1 SV=3
Tyrosine-protein kinase CSK OS=Homo sapiens GN=CSK PE=1 SV=1

Cell surface glycoprotein MUC18 OS=Homo sapiens GN=MCAM PE=1 SV=2
Heat shock 70 kDa protein 13 OS=Homo sapiens GN=HSPA13 PE=1 SV=1
Emerin OS=Homo sapiens GN=EMD PE=1 SV=1
Serpin B9 OS=Homo sapiens GN=SERPINB9 PE=1 SV=1
Sulfite oxidase, mitochondrial OS=Homo sapiens GN=SUOX PE=1 SV=2
Vesicle-associated membrane protein 7 OS=Homo sapiens GN=VAMP7 PE=1 SV=3
Delta-1-pyrroline-5-carboxylate synthase OS=Homo sapiens GN=ALDH18A1 PE=1 SV=2
Nucleoside diphosphate kinase homolog 5 OS=Homo sapiens GN=NME5 PE=1 SV=1
Hormone-sensitive lipase OS=Homo sapiens GN=LIPE PE=1 SV=4
3-ketodihydrosphingosine reductase OS=Homo sapiens GN=KDSR PE=1 SV=1
Amidophosphoribosyltransferase OS=Homo sapiens GN=PPAT PE=1 SV=1
Acetyl-CoA carboxylase 1 OS=Homo sapiens GN=ACACA PE=1 SV=2
Methylglutaconyl-CoA hydratase, mitochondrial OS=Homo sapiens GN=AUH PE=1 SV=1
Zinc transporter ZIP14 OS=Homo sapiens GN=SLC39A14 PE=1 SV=3
Surfeit locus protein 1 OS=Homo sapiens GN=SURF1 PE=1 SV=1
TGF-beta-activated kinase 1 and MAP3K7-binding protein 1 OS=Homo sapiens GN=TAB1
AP2-associated protein kinase 1 OS=Homo sapiens GN=AAK1 PE=1 SV=3
Vacuolar ATPase assembly integral membrane protein VMA21 OS=Homo sapiens GN=VMA21
Quinone oxidoreductase PIG3 OS=Homo sapiens GN=TP53I3 PE=1 SV=2
PDZ domain-containing protein 11 OS=Homo sapiens GN=PDZD11 PE=1 SV=2
Cytochrome c oxidase protein 20 homolog OS=Homo sapiens GN=COX20 PE=1 SV=2
Cartilage acidic protein 1 (Fragment) OS=Homo sapiens GN=CRTAC1 PE=2 SV=1
E3 ubiquitin-protein ligase UBR4 OS=Homo sapiens GN=UBR4 PE=1 SV=1
CD276 antigen OS=Homo sapiens GN=CD276 PE=1 SV=1
Calcium-binding mitochondrial carrier protein SCaMC-2 OS=Homo sapiens GN=SLC25A25
Tetratricopeptide repeat protein 37 OS=Homo sapiens GN=TTC37 PE=1 SV=1
Aspartate--tRNA ligase, mitochondrial OS=Homo sapiens GN=DARS2 PE=1 SV=1
Zinc transporter 9 OS=Homo sapiens GN=SLC30A9 PE=1 SV=1
3-hydroxyacyl-CoA dehydratase 2 OS=Homo sapiens GN=PTPLB PE=1 SV=1
Protein phosphatase 1 regulatory subunit 21 OS=Homo sapiens GN=PPP1R21 PE=1 SV=1
ATP-binding cassette sub-family F member 2 OS=Homo sapiens GN=ABCF2 PE=2 SV=1
Amyloid beta A4 precursor protein-binding family B member 1-interacting protein 0
Fermitin family homolog 3 OS=Homo sapiens GN=FERMT3 PE=1 SV=1
Potassium intermediate/small conductance calcium-activated channel, subfamily N,
WAS protein family, member 3, isoform CRA_a OS=Homo sapiens GN=WASF3 PE=2 SV=1
Endophilin-A3 OS=Homo sapiens GN=SH3GL3 PE=2 SV=1
WD repeat-containing protein 16 OS=Homo sapiens GN=WDR16 PE=1 SV=3
1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase delta-3 OS=Homo sapiens
Minor histocompatibility antigen H13 OS=Homo sapiens GN=HM13 PE=1 SV=1
D-tyrosyl-tRNA(Tyr) deacylase 1 OS=Homo sapiens GN=DTD1 PE=1 SV=2
Palladin OS=Homo sapiens GN=PALLD PE=1 SV=3
Protein phosphatase 1E OS=Homo sapiens GN=PPM1E PE=1 SV=2
Immunity-related GTPase family Q protein OS=Homo sapiens GN=IRGQ PE=1 SV=1
Estradiol 17-beta-dehydrogenase 8 OS=Homo sapiens GN=HSD17B8 PE=1 SV=2
Tetratricopeptide repeat protein 9A OS=Homo sapiens GN=TTC9 PE=2 SV=3
UPF0556 protein C19orf10 OS=Homo sapiens GN=C19orf10 PE=1 SV=1
Nicalin OS=Homo sapiens GN=NCLN PE=1 SV=2

Zinc finger protein 428 OS=Homo sapiens GN=ZNF428 PE=1 SV=2
Nucleoporin SEH1 OS=Homo sapiens GN=SEH1L PE=1 SV=3
Mannose-1-phosphate guanyltransferase alpha OS=Homo sapiens GN=GMPPA PE=1 SV=1
Exocyst complex component 2 OS=Homo sapiens GN=EXOC2 PE=1 SV=1
Dehydrogenase/reductase SDR family member 1 OS=Homo sapiens GN=DHRS1 PE=1 SV=1
Actin-related protein 2/3 complex subunit 5-like protein OS=Homo sapiens GN=ARPC5
O-acetyl-ADP-ribose deacetylase MACROD1 OS=Homo sapiens GN=MACROD1 PE=1 SV=2
KxDL motif-containing protein 1 OS=Homo sapiens GN=KXD1 PE=1 SV=2
Protein syndesmos OS=Homo sapiens GN=NUDT16L1 PE=1 SV=1
Uncharacterized protein C19orf52 OS=Homo sapiens GN=C19orf52 PE=1 SV=2
UPF0368 protein Cxorf26 OS=Homo sapiens GN=CXorf26 PE=1 SV=1
Transmembrane protein 120A OS=Homo sapiens GN=TMEM120A PE=2 SV=1
Ubiquitin-conjugating enzyme E2 0 OS=Homo sapiens GN=UBE20 PE=1 SV=3
Tubulointerstitial nephritis antigen-like OS=Homo sapiens GN=TINAGL1 PE=1 SV=1
UPF0687 protein C20orf27 OS=Homo sapiens GN=C20orf27 PE=1 SV=3
Rabenosyn-5 OS=Homo sapiens GN=ZFYVE20 PE=1 SV=2
MIP18 family protein FAM96A OS=Homo sapiens GN=FAM96A PE=1 SV=1
Plasminogen receptor (KT) OS=Homo sapiens GN=PLGRKT PE=1 SV=1
Mitochondrial import receptor subunit TOM22 homolog OS=Homo sapiens GN=TOMM22 PE=
DnaJ homolog subfamily B member 4 OS=Homo sapiens GN=DNAJB4 PE=1 SV=1
Ras-related protein Rab-23 OS=Homo sapiens GN=RAB23 PE=1 SV=1
Intercellular adhesion molecule 5 OS=Homo sapiens GN=ICAM5 PE=1 SV=3
Regulator complex protein LAMTOR2 OS=Homo sapiens GN=LAMTOR2 PE=1 SV=1
U6 snRNA-associated Sm-like protein LSM2 OS=Homo sapiens GN=LSM2 PE=1 SV=1
Coatmer subunit gamma-1 OS=Homo sapiens GN=COPG1 PE=1 SV=1
Coiled-coil-helix-coiled-coil-helix domain-containing protein 2, mitochondrial OS
Kinesin-like protein KIF1A OS=Homo sapiens GN=KIF1A PE=3 SV=1
Pleckstrin homology-like domain family B member 1 OS=Homo sapiens GN=PHLDB1 PE=1
Serine/threonine-protein kinase TBK1 (Fragment) OS=Homo sapiens GN=TBK1 PE=4 SV=1
Protein RER1 OS=Homo sapiens GN=RER1 PE=1 SV=1
Krev interaction trapped protein 1 (Fragment) OS=Homo sapiens GN=KRIT1 PE=4 SV=1
TBC1 domain family member 15 OS=Homo sapiens GN=TBC1D15 PE=2 SV=1
High mobility group protein HMG-I/HMG-Y OS=Homo sapiens GN=HMGA1 PE=4 SV=1
Importin subunit alpha-3 OS=Homo sapiens GN=KPNA3 PE=1 SV=2
Cullin-5 OS=Homo sapiens GN=CUL5 PE=1 SV=4
Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=1
Dynein light chain Tctex-type 1 OS=Homo sapiens GN=DYNLT1 PE=1 SV=1
Translocation protein SEC63 homolog OS=Homo sapiens GN=SEC63 PE=1 SV=2
5'-AMP-activated protein kinase catalytic subunit alpha-2 OS=Homo sapiens GN=PRKA
2-oxoisovalerate dehydrogenase subunit beta, mitochondrial OS=Homo sapiens GN=BCK
Arfaptin-1 OS=Homo sapiens GN=ARFIP1 PE=2 SV=1
Nitrilase 1 OS=Homo sapiens GN=NIT1 PE=4 SV=1
Matrix metalloproteinase-9 OS=Homo sapiens GN=MMP9 PE=1 SV=3
CDGSH iron-sulfur domain-containing protein 2 OS=Homo sapiens GN=CISD2 PE=4 SV=1
Pterin-4-alpha-carbinolamine dehydratase OS=Homo sapiens GN=PCBD1 PE=1 SV=2
Glycylpeptide N-tetradecanoyltransferase 1 OS=Homo sapiens GN=NMT1 PE=1 SV=2
Synembryn-A (Fragment) OS=Homo sapiens GN=RIC8A PE=4 SV=1

Neutrophil gelatinase-associated lipocalin OS=Homo sapiens GN=LCN2 PE=3 SV=1
CCR4-NOT transcription complex subunit 1 OS=Homo sapiens GN=CNOT1 PE=1 SV=2
NudC domain-containing protein 3 OS=Homo sapiens GN=NUDC3 PE=4 SV=1
Epsin-1 OS=Homo sapiens GN=EPN1 PE=1 SV=2
GDH/6PGL endoplasmic bifunctional protein OS=Homo sapiens GN=H6PD PE=1 SV=2
GRB2-associated-binding protein 1 (Fragment) OS=Homo sapiens GN=GAB1 PE=4 SV=1
Protein NOXP20 OS=Homo sapiens GN=DKFZp686F20250 PE=2 SV=1
Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=1
Kinesin light chain 4 OS=Homo sapiens GN=KLC4 PE=2 SV=1
Proline-rich transmembrane protein 2 OS=Homo sapiens GN=PRRT2 PE=1 SV=1
Protein C11orf31 (Fragment) OS=Homo sapiens GN=C11orf31 PE=4 SV=1
REST corepressor 3 (Fragment) OS=Homo sapiens GN=RCOR3 PE=4 SV=1
Argininosuccinate lyase OS=Homo sapiens GN=ASL PE=3 SV=1
Dihydropyrimidine dehydrogenase [NADP(+)] OS=Homo sapiens GN=DPYD PE=1 SV=2
Inositol-3-phosphate synthase 1 OS=Homo sapiens GN=ISYNA1 PE=2 SV=1
CDC42 effector protein (Rho GTPase binding) 1 (Fragment) OS=Homo sapiens GN=CDC42
Basic leucine zipper and W2 domain-containing protein 2 OS=Homo sapiens GN=BZW2 P
Pituitary adenylate cyclase-activating polypeptide type I receptor (Fragment) OS=
NADH dehydrogenase [ubiquinone] 1 subunit C2 OS=Homo sapiens GN=NDUFC2 PE=3 SV=1
Collagen alpha-1(XI) chain OS=Homo sapiens GN=COL21A1 PE=4 SV=1
Neural Wiskott-Aldrich syndrome protein OS=Homo sapiens GN=WASL PE=1 SV=2
DnaJ homolog subfamily C member 8 OS=Homo sapiens GN=DNAJC8 PE=1 SV=2
Dual specificity mitogen-activated protein kinase kinase 3 OS=Homo sapiens GN=MAP
40S ribosomal protein S11 OS=Homo sapiens GN=RPS11 PE=1 SV=3
Aminoacyl tRNA synthase complex-interacting multifunctional protein 2 OS=Homo sap
Mitochondrial inner membrane organizing system protein 1 OS=Homo sapiens GN=MINOS
WD repeat-containing protein 11 OS=Homo sapiens GN=WDR11 PE=1 SV=1
Apoptosis-associated speck-like protein containing a CARD OS=Homo sapiens GN=PYCA
Complement C1s subcomponent heavy chain OS=Homo sapiens GN=C1S PE=3 SV=1
Protein farnesyltransferase/geranylgeranyltransferase type-1 subunit alpha (Fragm
CDP-diacylglycerol--inositol 3-phosphatidyltransferase OS=Homo sapiens GN=CDIPT P
Myotubularin-related protein 10 OS=Homo sapiens GN=MTMR10 PE=4 SV=1
Eukaryotic translation initiation factor 6 (Fragment) OS=Homo sapiens GN=EIF6 PE=
Ubiquitin-like protein 3 OS=Homo sapiens GN=UBL3 PE=1 SV=1
CKLF-like MARVEL transmembrane domain-containing protein 4 (Fragment) OS=Homo sap
Glucosamine--fructose-6-phosphate aminotransferase [isomerizing] 1 OS=Homo sapien
ATP-dependent (S)-NAD(P)H-hydrate dehydratase OS=Homo sapiens GN=CARKD PE=2 SV=1
Protein FADD OS=Homo sapiens GN=FADD PE=1 SV=1
Acidic fibroblast growth factor intracellular-binding protein (Fragment) OS=Homo
Sphingosine-1-phosphate lyase 1 OS=Homo sapiens GN=SGPL1 PE=1 SV=3
UPF0670 protein THEM6 OS=Homo sapiens GN=THEM6 PE=1 SV=2
IQ and ubiquitin-like domain-containing protein OS=Homo sapiens GN=IQUB PE=1 SV=2
Casein kinase II subunit alpha' (Fragment) OS=Homo sapiens GN=CSNK2A2 PE=4 SV=1
Pantothenate kinase 4 OS=Homo sapiens GN=PANK4 PE=4 SV=1
Protein transport protein Sec23B OS=Homo sapiens GN=SEC23B PE=1 SV=2
Unconventional myosin-Ic OS=Homo sapiens GN=MYO1C PE=4 SV=1
ADP-ribosylation factor-like protein 3 OS=Homo sapiens GN=ARL3 PE=1 SV=2

Diacylglycerol kinase iota OS=Homo sapiens GN=DGKI PE=4 SV=1
Bisphosphoglycerate mutase OS=Homo sapiens GN=BPGM PE=1 SV=2
Protein-tyrosine sulfotransferase 1 (Fragment) OS=Homo sapiens GN=TPST1 PE=4 SV=1
Phosphatidylinositol 3-kinase regulatory subunit beta OS=Homo sapiens GN=PIK3R2 P
Signal transducer and activator of transcription 5A OS=Homo sapiens GN=STAT5A PE=
HCG1811539, isoform CRA_b OS=Homo sapiens GN=TXNDC5 PE=2 SV=1
Extracellular matrix protein 1 OS=Homo sapiens GN=ECM1 PE=1 SV=2
40S ribosomal protein S24 OS=Homo sapiens GN=RPS24 PE=3 SV=1
Protein YIPF5 (Fragment) OS=Homo sapiens GN=YIPF5 PE=4 SV=1
Lipoma-preferred partner OS=Homo sapiens GN=LPP PE=1 SV=1
Protein RBM14-RBM4 OS=Homo sapiens GN=RBM14/RBM4 fusion PE=2 SV=1
ATP-binding cassette sub-family E member 1 OS=Homo sapiens GN=ABCE1 PE=4 SV=1
Zinc finger protein GLI2 OS=Homo sapiens GN=GLI2 PE=1 SV=4
Complement factor I light chain OS=Homo sapiens GN=CFI PE=3 SV=1
Protein phosphatase 1F OS=Homo sapiens GN=PPM1F PE=3 SV=1
Multidrug and toxin extrusion protein 1 (Fragment) OS=Homo sapiens GN=SLC47A1 PE=

Access Number	Grade I	Grade II	Grade III	Grade IV
P14136	1	2	1	0.67
P02768	1	0.55	1.73	0.91
Q15149	1	1.9	1.8	0.9
P68871	1	1	0.89	1.11
P08670	1	0.58	1.25	0.83
Q13509	1	0.89	0.67	1
P60709	1	1.11	1.33	1.56
P68363	1	0.9	0.8	0.6
Q13813	1	1.09	0.73	0.64
P69905	1	1	0.89	1.22
Q01082	1	1.4	1	0.8
P14618	1	0.7	0.8	1.2
P04406	1	0.7	0.7	2.3
Q00610	1	0.89	0.89	0.89
P21333	1	0.5	0.8	0.9
Q14204	1	1.3	1	1
B1AKY9	1	1.4	1	0.8
P02675	1	1	1	12.5
P02671	1	0.8	0.9	9.2
Q16555	1	0.8	0.8	0.7
P12814	1	0.7	1.5	0.8
P10809	1	1.2	0.8	1.1
P25705	1	0.9	0.7	0.9
P02647	1	0.8	2.4	4.9
P09543	1	0.9	0.5	0.6
P12277	1	0.7	0.5	0.4
P06733	1	0.9	0.7	1.3
P01024	1	0.82	1.73	1.64
P00367	1	2.78	0.89	1.11
P68104	1	0.8	1.2	1.7
P02787	1	0.7	1.8	1.9
P02686	1	1.9	0.1	0.5
P07196	1	0.9	0.5	0.7
P13611	1	7.25	1.25	1.33
P07900	1	0.82	0.73	0.73
A6NN80	1	1.56	1.22	1.33
P02679	1	0.8	0.8	8.3
P04083	1	0.45	0.64	0.91
Q09666	1	1.1	1.4	1
P62258	1	1.5	1	0.7
P48681	1	1	3.55	1.82
P11142	1	0.9	1	1.1
E7ESP9	1	1.33	0.56	0.89
P24821	1	3.55	3.09	1.45
P46821	1	0.9	1	1.6
F8VPV9	1	0.9	0.7	0.8

P01023	1	1.1	1.7	1.7
A2BHY4	1	1.4	2.6	2.7
P06396	1	0.9	0.5	1.4
P00558	1	0.7	0.7	2.2
P01857	1	0.64	1.55	1.18
P08758	1	1.3	1.4	1.3
P23527	1	2	3	2.6
P35579	1	1	1.6	2.1
A2A274	1	1.1	1	0.8
P04075	1	0.73	0.73	1
P40926	1	1	0.78	1
Q05193	1	0.6	0.4	0.3
P02511	1	3.5	1.1	1.1
P09936	1	0.56	0.67	0.56
P62937	1	0.7	0.7	2.1
Q14315	1	0.4	0.5	0.6
Q96GW7	1	2.5	1.2	1.2
E7EPA7	1	0.9	0.7	1
G5EA52	1	0.9	1.4	1.7
P09104	1	0.9	0.7	0.6
P16152	1	0.6	0.5	0.6
P60201	1	2.89	0.67	1
Q16352	1	0.91	0.45	0.73
P07355	1	0.5	0.9	1.6
P08238	1	1.33	1.11	1.11
P09972	1	1.5	0.8	1
P08107	1	0.83	0.83	1.17
E7ENR4	1	0.6	0.5	0.7
P14625	1	1	1.45	1.18
P62805	1	1	1.9	2.4
P01009	1	0.64	1.27	1.64
P49411	1	1.2	0.9	1
P62873	1	0.82	0.64	0.45
Q06830	1	1	0.7	1.1
Q16181	1	1.5	1	0.7
B7Z242	1	0.5	1.2	0.6
P09382	1	0.8	1.6	1.4
P11021	1	0.9	1.5	2.6
E7ER44	1	1.1	2	9
P31150	1	1	0.8	0.5
P07195	1	1.7	1.1	0.9
POCG05	1	0.7	1.7	3.4
Q92752	1	4.22	1.22	1.22
P13591	1	2.6	1.1	1.1
P04792	1	0.45	0.73	1.18
Q01484	1	1	0.8	0.9
Q9Y490	1	1	1.4	1.5

P30086	1	0.9	0.7	0.7
P63104	1	0.89	1	0.78
Q9Y617	1	1.8	1.1	0.6
B4DGP8	1	1.7	2.1	1.4
P18669	1	0.9	0.7	0.7
P23471	1	3.22	2	1.11
P23634	1	1.4	1.8	0.9
P60174	1	1	0.89	2.33
O94760	1	1	0.64	0.55
P13639	1	0.9	1	1.7
P46459	1	0.7	0.6	0.7
Q14195	1	1.3	1.4	1
E7EVA0	1	1	2.8	1.2
P61764	1	0.6	0.4	0.6
E9PK25	1	1	0.9	1.6
P38646	1	1.4	0.9	1
P48735	1	2	1.22	0.89
P26038	1	0.7	1.2	1.3
P55072	1	1.1	1.6	1
Q13423	1	1.4	1	1.2
B4DV12	1	1.11	1.89	1.22
E9PHY5	1	1.4	1.1	0.7
P15311	1	1.64	0.91	0.73
B5MCX3	1	1.89	1.22	1.11
P27348	1	1.2	1	0.7
P28482	1	1.2	0.8	0.9
O75083	1	0.8	0.8	1.3
P01834	1	0.73	1.91	2.09
P49189	1	1	0.7	0.9
O15061	1	0.6	0.9	0.7
Q9NQC3	1	1	1	1
F5H098	1	0.9	0.7	0.6
O75369	1	0.8	0.6	0.6
P40939	1	1.5	1.1	1.1
P07237	1	1.1	2	4.2
POCOS5	1	0.8	1.1	5.5
P11216	1	1.3	0.6	0.6
P49419	1	1.4	0.7	0.7
Q96KP4	1	0.8	0.6	0.7
P11137	1	0.6	0.7	0.7
P32119	1	0.9	0.6	1.2
Q14764	1	0.6	1.1	1.2
E9PCV6	1	0.5	0.4	2.1
P02649	1	2.6	1.1	1.7
P21796	1	1.1	0.8	1.1
P63010	1	0.8	1	0.8
B7Z1R5	1	0.7	0.6	0.7

043175	1	1.8	1.1	1
Q12860	1	2.1	1	0.9
G5E9B2	1	1	0.8	1.3
J3KNU8	1	3.2	1.2	1.2
P00738	1	0.9	3.4	5.3
P02545	1	0.7	0.9	0.8
P09211	1	0.8	0.5	1
P09471	1	1.3	0.7	0.7
P30041	1	0.6	0.5	0.7
P51178	1	0.4	0.3	0.3
P55268	1	0.55	0.45	0.55
E9PE77	1	1	1	8.78
P06744	1	0.64	0.64	1.09
P22314	1	1	1.1	0.9
Q14697	1	0.7	0.9	0.8
E9PLK3	1	1.2	0.8	0.9
H3BRN4	1	2.9	1.5	1.1
P07737	1	0.73	0.73	2.91
P49327	1	1.1	1.3	1.1
F8VXI5	1	3	0.9	0.6
043865	1	2.5	0.9	1.1
P04899	1	1.4	1.6	1.1
P30044	1	0.64	0.36	0.55
P54652	1	0.64	0.27	0.45
G3V5I3	1	1	2.8	3.6
P00568	1	0.45	0.27	0.27
P12036	1	1	0.6	1
P13667	1	1.1	1.1	2.2
P21281	1	0.6	0.5	0.7
P27797	1	1	2	1.3
Q99832	1	1	0.9	1.1
Q9BPU6	1	0.82	0.82	0.73
Q9Y2J2	1	1.2	1.3	1.1
B4DGU4	1	1.4	1.3	1
P17600	1	0.7	0.4	0.7
P37802	1	0.6	2.2	1
P60880	1	0.5	0.3	0.4
Q01518	1	1.2	1.5	1.2
Q16658	1	1.1	1.5	1.5
B4E2W0	1	1.6	1.1	1.3
B7ZAR1	1	1.1	1	1.1
F5GWF6	1	1.1	1	1
P06702	1	0.9	0.7	23.5
P22626	1	1.2	2	1.2
B4DQJ8	1	0.8	0.9	1.2
P00747	1	0.67	1	3.56
P04264	1	1.73	1.82	1.45

P12235	1	1.2	1	1
Q13642	1	1.4	1.9	0.8
Q99497	1	0.89	0.67	1.22
Q9BW30	1	0.5	1.4	0.4
F8VTL3	1	1.4	1.1	1.1
P10909	1	0.82	1.09	1.64
P17252	1	1.5	1.1	1
P43490	1	1.1	1.2	10.1
Q07954	1	1.2	1.5	1.4
Q86VP6	1	1.1	1.2	0.9
E9PDA4	1	1.7	0.8	0.7
E9PL22	1	1.2	1.3	1.4
075947	1	0.9	0.6	0.6
P00441	1	1.09	0.64	0.91
P17987	1	1	0.9	1
P34932	1	0.9	0.9	0.9
P45880	1	1.2	0.8	1
Q16891	1	1	0.8	1.1
B3KUK2	1	1.4	2.6	3.1
E7EMB3	1	1.69	1.92	0.62
G5E949	1	2.6	0.9	1.2
P00338	1	0.9	0.8	2.1
P00918	1	0.6	0.6	1.3
P02774	1	0.8	1.5	4.2
P49588	1	1.2	1.1	1.4
Q00839	1	0.9	1.7	2.2
Q14974	1	1.1	1.2	1.2
F5HOT1	1	1	1.1	1
P00505	1	0.7	0.6	0.6
P18206	1	1.2	1.4	0.8
P24752	1	1.7	0.8	1.1
Q9UEY8	1	1.67	1.56	0.89
B4DJA5	1	1.4	1.5	1.9
P04843	1	1.4	1.7	1.7
P09960	1	1	0.9	1.2
P19823	1	0.9	2.4	3.6
P28331	1	0.9	0.7	0.8
P28838	1	0.9	1.3	2.1
P52272	1	1.2	1.4	1.5
P55809	1	0.9	0.5	0.6
Q96AQ6	1	1	1.5	0.7
B4DJV2	1	1.22	1.11	1.11
E9PD68	1	0.9	1.1	0.8
E9PH29	1	0.9	0.7	0.7
F5H2A9	1	0.9	0.6	1
095782	1	0.8	0.9	0.8
P01876	1	1	2.2	3.5

P07099	1	1.3	0.8	0.7
P07954	1	1.27	0.82	0.91
P11047	1	0.5	0.5	0.9
P13804	1	2	1.4	0.8
P22695	1	1.4	0.9	1
P33176	1	1.1	1.1	1.2
P49368	1	0.8	0.8	1.2
Q9Y2J8	1	0.9	0.6	0.7
B4DZP4	1	0.9	0.9	1
E5RJR5	1	1.44	1.22	0.67
H7BY58	1	0.8	0.7	0.8
P00387	1	1.7	2.7	0.8
P00915	1	0.64	0.55	0.91
P11177	1	0.91	0.64	0.73
P18085	1	1.2	1.2	0.9
P20700	1	2	2.45	1.36
P23246	1	1.2	1.7	1.5
P31146	1	1	1.1	2.2
P55087	1	1.9	1.2	0.6
P61981	1	1	1	1.38
Q15907	1	1.67	1.33	1.89
Q7KZF4	1	1.1	1.4	1.7
Q9Y230	1	1	1	0.8
Q9Y696	1	1.3	1.6	1.2
B4DY08	1	0.9	1.2	1.3
E9PGC8	1	0.6	0.6	0.7
F5H3X9	1	1	0.9	1.1
000499	1	0.9	0.6	0.6
P05164	1	1	1	9.2
P13489	1	0.8	0.7	1
P22392	1	1.18	0.82	1.82
P31930	1	1	0.7	0.8
P51659	1	1.5	1.6	1.5
Q02218	1	0.9	0.6	1.2
Q15365	1	0.6	0.8	1.7
Q6UVK1	1	0.6	0.8	0.6
Q9UQM7	1	0.55	0.36	0.36
B0UZR7	1	1	0.67	0.89
E5RHW4	1	0.9	1.2	0.7
E7EU23	1	1.18	1.09	0.82
F8W791	1	0.9	0.6	0.6
G3V192	1	3.2	0.9	1.6
J3KPX7	1	1.5	1	1.3
P00488	1	0.8	1	4.9
P02730	1	1.11	0.78	0.78
P05026	1	0.67	0.67	0.44
P10412	1	0.7	1.7	3

P16949	1	0.82	0.91	0.64
P21266	1	1.33	0.67	0.67
P23284	1	1.2	1.8	2.2
P23526	1	1	1.1	1.2
P30038	1	2.1	1.4	1.1
P67775	1	1.1	1.1	1.3
Q05707	1	0.5	0.4	0.5
Q7L576	1	1.7	1.5	0.8
Q99962	1	0.9	0.7	0.9
B4DEB1	1	0.4	0.6	12.9
D6RER5	1	0.8	1	1.1
E9PBS1	1	1.7	1.6	1.1
E9PBU3	1	1.1	1	0.8
F5H2S7	1	1	1	0.8
O14594	1	2	1.1	1.2
O95336	1	0.9	0.8	1
P12532	1	0.7	0.5	0.5
P16615	1	1.36	1.45	0.91
P17174	1	0.78	0.56	0.56
P19827	1	0.9	4.1	2.5
P23396	1	0.8	0.9	2
P36871	1	1	0.8	0.8
P46940	1	0.7	1.6	1.7
P61604	1	1.8	1.3	1
P98160	1	0.55	0.55	1.09
Q08209	1	0.9	0.7	0.7
Q08211	1	1.5	1.6	1.4
Q8IXJ6	1	0.73	0.45	0.55
A2A3N8	1	0.9	1.1	0.9
A8MTH6	1	1.5	1	1.1
H7BYY1	1	0.55	0.36	0.36
O95865	1	0.9	1.7	1.1
P02790	1	1	1.7	1.2
P04233	1	1.6	5.9	1.8
P07339	1	0.8	1	0.9
P09417	1	1	0.6	0.8
P13010	1	1.09	1.27	1
Q86UP2	1	1	1	1.1
B1AHC7	1	1	1.2	1.2
B1AHR1	1	1.18	0.91	0.36
C9JMU5	1	1.1	1.4	1.6
C9JW96	1	1.4	1.1	1.3
E7EMK3	1	1.5	1.5	1.7
E7EPV7	1	0.4	0.3	0.4
F5H7H8	1	0.4	0.3	0.3
F8VVM2	1	1.67	1.33	0.89
P00450	1	1	1.7	2.1

P15104	1	0.9	0.5	1.4
P19338	1	1.4	1.7	1.4
P31946	1	0.9	1	0.7
P37837	1	0.91	0.73	1.09
P40227	1	0.91	0.73	0.91
P43304	1	1.5	1.2	1.2
P45974	1	0.9	1.2	1.3
P50502	1	1.1	1.5	0.8
P50991	1	1.1	1	1
P51991	1	0.8	1.1	0.9
P54920	1	1	1.1	0.9
Q01813	1	1.1	0.5	0.9
Q13938	1	0.5	0.4	0.2
Q99536	1	1.3	1.2	1.5
Q9HDC9	1	1.2	1.2	2
B5MDF5	1	1	1.1	1.5
E9PFU1	1	0.8	1	1.4
E9PH06	1	0.9	0.9	0.9
E9PHI5	1	2.56	1.33	1.11
000154	1	0.5	0.4	0.7
043301	1	0.82	0.45	0.64
094769	1	0.4	0.3	0.5
P04271	1	1.22	1	0.33
P10636	1	0.6	0.5	0.9
P10644	1	1.18	1.09	0.73
P61158	1	1	1.27	1.09
P61978	1	1	1.4	1.2
P67936	1	1.7	2.2	1.4
Q07065	1	1.4	1.9	7.1
Q9UDT6	1	1.1	1.4	1
A8MY36	1	1.3	1.1	1.1
B1AK87	1	1.4	1.6	1.5
B5MCQ5	1	1.3	1.8	1.6
F5GXC8	1	0.9	0.6	0.7
F5GZS6	1	1.6	1.1	0.8
F5H740	1	0.8	0.5	0.6
G3V2B8	1	1.3	1.1	1.9
060506	1	1.2	1.6	1
060664	1	0.9	1	2.3
P00390	1	1	0.8	1.1
P01859	1	0.5	1.6	3.4
P01871	1	0.7	1.2	1.6
P12109	1	0.6	0.6	1.2
P13645	1	2.1	1.9	1.4
P13746	1	1.9	3.2	2
P28907	1	6.8	1.3	1.4
P35998	1	1	1	1

P36543	1	0.6	0.5	0.8
P78417	1	1	0.6	1.2
Q07021	1	1.27	0.73	0.45
Q15334	1	1.6	1.1	1
Q9Y3I0	1	1.2	1.1	0.9
B3KSI9	1	0.6	0.5	0.8
B4DTU0	1	3.64	1.73	0.82
E7EMM4	1	1.22	1	1
E9PBG7	1	0.7	0.9	1.4
E9PIT3	1	0.8	1.5	3.4
O94811	1	0.67	0.44	0.44
O94973	1	0.9	0.8	0.9
P02549	1	1.1	0.8	0.9
P04040	1	0.8	0.7	1.8
P09455	1	0.5	0.8	0.7
P09525	1	0.9	0.9	1
P09874	1	1.27	1.73	1.09
P11766	1	0.91	0.82	0.64
P14314	1	1.18	1.64	1.09
P17858	1	1.33	1.44	1.78
P29966	1	8.89	2.78	2.44
P30042	1	1.9	1.2	1
P38117	1	1.82	1.36	1
P50148	1	1.44	1.33	1
P51149	1	1.1	1.1	1.7
Q14CZ8	1	3.3	1.1	0.8
Q15691	1	1	1.27	0.82
Q5VU66	1	1.2	1.6	1.1
Q92499	1	1.2	1.1	0.9
Q92598	1	0.6	0.6	0.7
Q9BY11	1	0.5	0.3	0.4
Q9P2R3	1	1.3	0.9	1
A8MV58	1	1.56	2.78	1.44
B1AVU8	1	1.7	1.9	1.1
B4DT69	1	1.6	1	1.1
B4E1Z4	1	0.8	1.7	1.7
C9JPG0	1	1.2	1.1	1.2
E7EPC6	1	1.1	2.4	2.1
HOYL72	1	1	0.7	0.7
H3BPE1	1	1.3	1.3	1.4
I3L397	1	1	0.89	0.89
J3KQA0	1	0.25	0.25	0.17
J9JID7	1	1.5	1.6	1.3
O95817	1	1.5	1.6	1.1
P04080	1	0.78	0.67	0.44
P15121	1	1	0.7	1
P24539	1	1.5	1	1

P42765	1	0.9	1.2	1
P60981	1	0.8	0.8	1
Q00796	1	1.8	0.7	0.9
Q04828	1	0.5	0.4	0.4
Q04917	1	0.8	0.8	0.7
Q14980	1	1.6	1.5	1.6
Q15075	1	1.3	2.5	1.2
Q96I99	1	2.27	1.45	1.09
Q9BX66	1	1.1	1.2	1.2
Q9UHD8	1	1.3	0.9	0.9
A6NN60	1	1	0.8	0.9
B4DE86	1	1.7	0.8	0.6
B4DLZ5	1	1	1.1	1
B7Z6B8	1	2.4	1.6	1.3
C9J9K3	1	0.91	1.09	1.27
E9PM69	1	1	0.9	1.2
F8VRQ1	1	1.55	1.73	1.18
F8VW21	1	1.1	1.4	1.6
O15144	1	1.1	1.5	1.5
P05109	1	1.09	1.09	14.18
P08603	1	0.82	1.64	2.64
P14866	1	1	1.82	1.09
P14868	1	1.9	1.2	1.5
P24043	1	0.56	0.44	0.56
P28161	1	1.55	0.91	0.73
P28289	1	0.82	1.18	0.64
P32455	1	0.6	0.9	0.9
P36776	1	1.36	1	1
P43004	1	1.27	0.64	0.91
P47755	1	0.9	1.1	0.7
P53396	1	0.8	0.8	1.2
P54577	1	1.2	1.1	1.3
P55795	1	1.2	1.5	1.6
P60842	1	1.2	1.1	1.4
P63000	1	1.55	1.64	1.55
P78324	1	0.9	0.6	0.5
Q15417	1	0.9	1.2	1.1
Q9H4M9	1	0.9	0.7	0.7
Q9NSD9	1	1.1	0.9	0.8
Q9Y2T3	1	0.7	0.8	0.5
P14174	1	1.1	0.9	1.4
A6NFQ9	1	0.9	0.9	0.9
B1AKZ3	1	3.1	2	0.7
B1ALW1	1	1.2	1.5	1.1
E7EUC7	1	0.9	1	1.3
E7EUY0	1	1	1.3	1.8
E9PBJ5	1	1.22	1	1.44

E9PND2	1	1	1.2	1.1
F5GYQ1	1	0.8	0.7	0.8
F8WCA0	1	0.8	0.6	0.7
G5E9M5	1	1.1	1	1.8
H0YL69	1	0.9	1.1	0.8
014980	1	0.8	0.9	0.9
075131	1	1.2	1.4	1.1
075781	1	0.7	0.6	0.6
P00966	1	0.7	1.9	0.8
P06748	1	1	1.2	1.6
P07203	1	1.2	1.2	2.3
P07384	1	1.4	1.2	1.1
P07814	1	0.73	0.91	1.18
P10606	1	1.2	1.1	1.4
P14550	1	1.1	0.8	1.3
P25398	1	1.2	1.4	1.7
P26232	1	1	0.8	0.7
P31689	1	1	1.1	1.2
P36957	1	0.9	0.7	1.2
P54136	1	1	1.2	1
P60953	1	1.73	1.73	1.09
P61160	1	0.9	1	2.7
Q02952	1	0.8	1.7	1.1
Q13228	1	0.8	0.6	0.6
Q14240	1	0.9	0.9	0.8
Q15233	1	0.7	0.9	1.9
Q16799	1	1.1	1.3	0.9
Q92597	1	1	0.91	1.82
Q9BUT1	1	0.7	0.6	0.8
Q9UQ03	1	1.1	1.4	0.9
F8W0P2	1	0.9	4.4	1.1
B4DGB4	1	0.8	1	1.3
B4DL14	1	1	0.9	0.8
B4DTG2	1	1	0.9	1.1
B4E241	1	1	0.8	1.4
B7Z6Z4	1	0.6	1	4.7
B7Z8M7	1	0.7	0.7	2.5
C9JF17	1	1.27	1.55	1.73
E7EW34	1	1	0.9	1.3
E9PK01	1	1.1	1.1	1
F2Z2U8	1	1.18	1.27	0.91
F8VQ10	1	1.1	1.2	1
F8W1A4	1	1.18	1.27	1.36
G3V0I5	1	1.09	1	0.64
G3V1K3	1	2.89	2.33	1.44
H7BYD0	1	1.11	1	0.78
J3KNX3	1	2	1.3	1.1

043776	1	1.27	1.18	0.55
076070	1	0.8	0.6	0.8
094925	1	0.5	0.3	0.4
P04004	1	0.8	0.8	4.8
P11233	1	2.7	1.9	1.8
P17677	1	1.7	2.2	1.5
P22102	1	1.1	1	1.1
P26599	1	0.64	1.09	1
P35221	1	1.8	1.6	1.3
P35237	1	0.9	0.7	0.6
P48047	1	1.1	0.9	0.9
P58546	1	0.9	0.8	1.1
P60033	1	1.8	1.3	0.9
P61970	1	1.27	1.27	0.64
P80723	1	2.67	2.11	0.67
Q15029	1	1.3	1.5	1.4
Q99714	1	1.4	1.4	1.4
Q9BS26	1	0.9	1.1	1.6
Q9H4G0	1	0.6	0.7	0.5
Q9NZ45	1	1	0.7	0.8
Q9Y2A7	1	1.2	1.2	1.1
Q9Y4G6	1	1.2	1	1
Q13185	1	1	1.7	2.1
A6NIZ1	1	1.9	2.1	1.2
A6NLG9	1	0.4	0.4	0.6
B0V043	1	1.2	1.1	1.9
B3KR64	1	1	0.8	0.7
B3KV44	1	1.18	1.64	0.73
B4DT77	1	1.5	1.3	1.3
B4DVE7	1	0.8	0.7	0.9
B7ZKJ8	1	0.7	1.4	1.2
C9J712	1	0.78	0.78	1.33
D3DP09	1	0.8	0.6	0.6
E7EPT4	1	0.8	0.8	1.1
E7EQI5	1	0.7	0.8	0.7
F5GYC0	1	0.8	0.5	0.8
H0Y4R1	1	1.1	1.3	1.5
J3KQ17	1	1.4	0.9	1
O14818	1	1.1	1.1	1.2
043390	1	1.1	1.6	1
P01008	1	0.7	1.4	1.6
P05165	1	1.1	0.9	0.9
P06703	1	0.75	3.08	8.42
P11277	1	1.1	0.8	0.9
P11498	1	2.3	1	0.6
P14415	1	1.3	0.7	1.2
P17655	1	1.2	1.5	0.8

P19022	1	1.3	1.1	1.1
P23381	1	1	1.7	1.4
P31947	1	0.33	0.33	0.33
P35612	1	0.9	0.9	0.9
P43034	1	1.2	1	0.9
P60891	1	1.1	1	0.8
P61026	1	2.6	1.7	1.5
P61266	1	0.36	0.18	0.45
Q02790	1	0.73	1.27	0.91
Q14247	1	1.4	6	1.6
Q969P0	1	0.9	0.6	0.7
Q9H4A4	1	0.89	0.56	0.56
Q9H9B4	1	1.4	0.8	1.2
Q9P035	1	1.18	1.27	0.64
Q9UBQ7	1	1.2	0.9	0.8
Q9Y266	1	0.7	0.6	0.5
Q9Y281	1	0.8	0.6	1
Q04760	1	0.9	0.8	1.3
P07108	1	6.3	4.3	0.8
F5H774	1	1	0.8	1
A7MAP0	1	1.1	1.1	2.7
B4DF14	1	2.82	0.82	0.55
B4DUS9	1	1.2	1	1
B7Z9I1	1	2	1.8	1.3
C9JME2	1	2.3	1.2	1
C9JWM7	1	0.82	0.82	1.73
C9JZR2	1	1.1	1.3	1.4
E7ERU0	1	1.1	1.3	1.2
F5H816	1	0.7	0.8	3
F8VSD4	1	0.7	0.6	0.9
G3V126	1	0.7	0.5	0.8
J3KQ83	1	1.17	0.92	0.92
J3QRD1	1	1.45	2.18	1.18
014745	1	3.3	2.1	1.2
015230	1	0.73	0.64	0.64
015394	1	3.3	1.7	0.8
015540	1	0.8	1.7	0.7
075874	1	0.91	1.09	2
095831	1	2.38	1.13	1.63
P02749	1	0.7	2	2.9
P04217	1	0.9	1.5	2.3
P04844	1	1.3	1.9	0.7
P11413	1	0.73	0.82	1.09
P13073	1	1.27	1	1.18
P20674	1	0.55	0.45	1.45
P25786	1	1.2	1.1	1.5
P30043	1	0.78	0.56	1.11

P30049	1	1.13	0.63	1
P47985	1	1.6	1.1	1.1
P49591	1	1	1.1	1.3
P50440	1	0.7	0.8	0.6
P61163	1	1	1	1.11
Q12905	1	1	1.3	0.7
Q13011	1	1.4	0.9	1.2
Q16531	1	0.8	0.7	1.1
Q8N3J6	1	1.9	1.1	0.6
Q99747	1	1	0.78	1
Q9UIJ7	1	2.3	1.2	0.7
Q9Y265	1	0.8	1	0.9
Q9Y3Z3	1	0.8	1.3	1.4
P10768	1	0.9	0.5	0.8
C9JV77	1	0.64	1.82	1.18
Q13424	1	1.64	1.27	0.36
A8K3Z3	1	1	0.9	0.9
B4DJ62	1	0.4	0.4	0.3
B4DTC3	1	1.44	1.67	1.22
B4DZI8	1	1.1	1.2	1
C9JFR7	1	0.8	1	0.9
E7EQV3	1	0.9	1.2	2.3
E9PCE8	1	1.4	1.5	1.2
F5H7V7	1	1.67	1.44	1
F8VQX6	1	1.7	0.9	1.6
F8VY02	1	0.7	1.2	1.5
F8W998	1	2.5	1.2	1.2
H0YCY6	1	0.7	0.6	0.9
J3KPM7	1	1.2	1	0.8
J3KQ32	1	0.7	0.6	0.9
J3QRU1	1	2.1	1.4	1.2
075323	1	1.8	0.8	1
075380	1	0.6	0.5	0.9
075955	1	2	1.3	1.7
P01019	1	1.6	1.1	1
P04003	1	0.7	0.6	3.1
P05387	1	1.4	2.2	2.5
P06727	1	0.8	1.2	1.6
P09496	1	1	0.8	1.6
P14543	1	0.7	0.6	1.2
P20618	1	1	0.9	1.8
P42224	1	1	1.3	1.4
P51884	1	0.4	0.3	0.5
P53618	1	1	1.1	1
P55060	1	0.82	0.91	0.55
P69891	1	0.7	0.4	0.4
Q06323	1	1.1	0.9	0.9

Q13177	1	1.11	1.11	1.44
Q13409	1	1.2	0.9	1.1
Q14019	1	1	0.8	1
Q14344	1	3	1.6	1.1
Q15435	1	0.9	0.8	1.2
Q5JWF2	1	1.67	1.56	1.33
Q6YN16	1	2.22	1.22	1.11
Q8IVF2	1	0.5	0.6	0.5
Q8TAM6	1	0.8	0.5	0.6
Q92973	1	1.3	1.4	0.8
Q96QK1	1	1	0.9	1
Q9H115	1	0.6	0.4	0.5
Q9HC38	1	1.1	0.6	0.8
Q9Y6R1	1	5.6	1.4	1.4
Q14914	1	0.88	0.75	0.63
I3L3P7	1	1.1	1.3	1.9
A8MXP9	1	1.3	1.3	1
B1AHM9	1	1.4	0.7	2.1
B4DLR8	1	0.5	0.5	0.5
E5KLJ5	1	1	1	0.6
E5RHG8	1	1	1	1.1
E7EMS6	1	1	1.27	0.64
E9PDN5	1	1.2	0.9	0.5
E9PHR3	1	0.5	0.5	1.1
F5GXN1	1	0.9	1	0.6
F5H569	1	0.6	0.6	0.9
F5H897	1	1.7	1.1	1.3
F6SBX2	1	1	0.8	0.8
G3V295	1	1	1	1.8
G3XAI2	1	0.7	0.6	1.1
G8JLA8	1	1	0.78	3.89
HOYBL1	1	1.8	1.3	1.1
J3KR54	1	1.1	1.4	1.1
000571	1	1	1.3	1.7
014531	1	0.9	0.8	0.8
014979	1	1.56	2.78	0.89
043157	1	3.5	1.6	1
043488	1	1.1	0.5	0.3
075367	1	1	1.2	1.8
075891	1	2.27	1.73	1.45
P02652	1	0.7	2.8	2.3
P05556	1	1.3	1.7	2.7
P08247	1	0.5	0.4	0.4
P08559	1	1.3	0.9	0.7
P12110	1	0.78	0.67	1.11
P21283	1	1	0.8	0.9
P21399	1	0.8	0.7	1

P23368	1	1.7	1.3	1
P25391	1	0.7	0.6	0.6
P35573	1	1.2	1	0.8
P36578	1	0.91	2	1.09
P38919	1	1.36	1.64	1
P42704	1	1.22	1.11	1
P46777	1	0.9	1.5	2.1
P50454	1	0.82	1.45	2.27
P51553	1	0.8	0.7	1.2
P51572	1	1.2	1.5	1.2
P52758	1	3.3	1.6	0.8
P53597	1	1.3	0.8	0.9
P54289	1	0.9	1.5	1.3
P61106	1	1.1	1.2	1.2
P62140	1	1.1	1.2	1.8
Q04837	1	3.4	2.4	1.4
Q14008	1	1	1	1.1
Q15181	1	1.2	1.2	1.1
Q9BPW8	1	1.4	1.1	0.9
Q9GZV7	1	1.1	0.3	0.6
Q9NTX5	1	1.09	1.09	1.27
Q9NZ08	1	0.91	1.18	1.09
Q9UPY8	1	0.6	0.5	0.5
E7EV01	1	1.2	2.1	1
H3BPB8	1	1	0.7	1
P00403	1	0.6	0.6	0.8
P13796	1	1	1.3	3.9
P23515	1	0.75	0.33	0.58
Q00765	1	0.6	1.2	1.7
G3V1D4	1	2.9	1.4	1.4
P09429	1	1.4	1.2	1.9
P13693	1	0.8	0.6	1.3
Q9H2M9	1	0.6	0.6	0.9
P60866	1	0.7	1	2.4
Q6P2I3	1	1.2	0.8	0.8
Q14108	1	1.7	1.7	1.3
P53985	1	1.09	1	2.36
B8ZZ77	1	0.6	0.5	0.8
A0AVT1	1	0.9	1.2	1.4
B4DDP6	1	1.1	1.3	1.6
B4DU58	1	0.45	0.64	1.45
B4DUC8	1	1.36	1.09	0.82
B4E2V5	1	1.6	0.6	4.4
B7Z233	1	1.1	1.1	0.9
C9JMC5	1	0.44	0.33	0.33
D6RAC2	1	0.9	1.1	2.4
E9PEJ4	1	1.2	0.8	0.8

F5GYR8	1	0.9	1.1	1.6
F5H668	1	1.2	1	1.2
F8VTY8	1	1.2	1.2	1.6
F8W6H6	1	1	0.9	0.8
G5E9Y2	1	1.11	1.11	1
H0Y8C6	1	1.4	1.4	1.2
I6L8B7	1	0.45	0.91	1.09
043242	1	1.33	1.17	1.42
060884	1	1.1	1.1	0.9
075489	1	1.4	1.1	0.6
076041	1	0.82	0.73	0.45
P00491	1	1.4	0.8	1.1
P01034	1	1.2	1.1	1
P02763	1	1.36	6.64	1.73
P08574	1	0.9	0.8	1.3
P13861	1	1.1	1.1	1.4
P17302	1	2.8	0.7	1
P21589	1	1.5	1	2.8
P30084	1	1.2	0.9	0.9
P35527	1	1.56	1.56	1.56
P36222	1	1.3	2.7	14.2
P49753	1	1.18	0.82	0.82
P52597	1	0.73	0.82	0.82
P61081	1	0.8	1	0.8
P61586	1	1	0.6	2.8
Q02750	1	1	0.89	1
Q07960	1	0.7	1	1.3
Q08722	1	1.7	1.6	1
Q12805	1	0.8	0.6	1.7
Q13263	1	1.1	2.2	1.4
Q13554	1	0.64	0.36	0.45
Q14152	1	1.1	1	1.3
Q5W0J0	1	1.64	1.27	1.45
Q8IWA5	1	2.8	1.7	1
Q96I24	1	0.91	1.45	1.64
Q9BSJ8	1	0.6	0.7	0.9
Q9HAV0	1	1.3	1.7	0.7
Q9NRV9	1	1.3	0.9	0.9
Q9P2E9	1	0.9	1.2	2.4
Q9UDR5	1	2.91	1.55	0.91
Q9UHD9	1	1.1	1.3	1.1
Q9UMS4	1	1	1.3	1.1
Q9Y6R7	1	1.2	5.8	2
P09497	1	0.55	0.55	0.64
B5ME97	1	1.82	1.73	0.73
J3KND9	1	1.6	1	1.4
A0MZ66	1	1.3	0.8	0.8

P02748	1	1.11	1	5.78
P07858	1	1	1.64	2.27
H3BQZ7	1	1.08	1.08	0.5
000231	1	0.91	0.82	1
B4DP11	1	0.9	0.9	0.8
A6NNI4	1	3.2	2.4	0.7
P60903	1	0.3	0.9	0.3
A8MVQ3	1	1.11	1.44	1.11
B1AJQ6	1	1.1	1.6	1.2
B1APG4	1	0.9	0.8	0.9
B4DHP6	1	1.6	2.5	1.7
B4DRZ7	1	1.3	1.3	1
B4DUR9	1	0.7	0.7	0.7
B5ME19	1	1	1.1	1.5
B7Z3I9	1	1.3	1.1	1.2
B7Z6C0	1	0.9	0.9	0.7
C9JDE9	1	2	1.78	1.89
C9JRZ6	1	1.82	1.36	1.27
D3DNT2	1	0.9	1	0.9
D3DQV9	1	0.8	0.8	0.9
E3W994	1	1.1	1	0.7
E7EVH7	1	0.9	0.9	1.2
E7EWZ6	1	1.7	1.5	1.2
E9PKG1	1	1	1.5	1
F5GXH5	1	0.9	1	0.9
F5GY03	1	1.9	1.4	2.3
F8VW96	1	1.09	1	1.64
G8JLP2	1	1.5	0.9	1.2
H0Y2M5	1	1	1.1	1.6
H9KVC2	1	0.7	0.4	0.7
J3KQG6	1	1.1	1.1	1.2
000151	1	0.56	0.89	0.44
043813	1	1.3	0.8	0.9
060493	1	0.8	0.7	1.5
075396	1	1.2	1.2	1.6
094826	1	1	0.8	1.1
095202	1	1.2	1.1	0.8
P08697	1	0.9	1.2	4
P10915	1	4	1	1.3
P15880	1	0.82	1.36	1.36
P16401	1	1.2	4.2	5.3
P20336	1	0.6	0.7	0.8
P21912	1	1	1	1.3
P25788	1	0.7	0.8	1.6
P26639	1	0.91	1	0.82
P28066	1	1	1.1	1.3
P30050	1	0.9	1.3	3.6

P30740	1	0.64	0.55	2.55
P40123	1	0.7	0.7	0.9
P50416	1	1.7	4.4	1.6
P53004	1	1	1.7	1.6
P62191	1	0.91	1.27	1.18
P62993	1	1.1	1.1	2.3
P78357	1	1.2	1.2	1
Q00577	1	1.1	0.8	0.7
Q01995	1	0.5	0.4	0.9
Q08752	1	0.7	0.6	1
Q13151	1	1	1.2	2.4
Q13464	1	1.2	1.2	1.1
Q13555	1	0.9	0.6	0.5
Q13884	1	1.4	0.9	1.8
Q15102	1	1.1	0.9	0.8
Q16143	1	0.6	0.5	0.7
Q5SSJ5	1	0.9	1	1.5
Q6IBS0	1	0.64	1.09	1.09
Q8N111	1	1.56	1.78	1.22
Q8NBX0	1	1	0.7	0.9
Q8NCW5	1	1	0.8	0.7
Q8NFZ8	1	1.5	1.3	0.9
Q96CX2	1	1.67	1.75	0.92
Q96F85	1	0.82	0.55	0.55
Q96FC7	1	1.5	0.9	1.1
Q96G03	1	1.4	1.3	1.2
Q99426	1	1	1.09	1
Q99436	1	0.6	0.7	2.2
Q9BRX8	1	1.9	1.4	0.8
Q9BTV4	1	0.9	1.7	0.9
Q9H936	1	0.7	0.5	0.8
Q9HCC0	1	1.6	1	1.5
Q9NVI7	1	1.89	1.33	1.44
Q9NZU5	1	0.6	0.5	2.1
Q9POL0	1	1.5	0.9	1
Q9UHQ9	1	1	0.7	1
Q9Y2S2	1	1.3	0.6	1.3
B7Z2R2	1	0.8	0.5	1
E9PFH8	1	1	0.82	0.82
E9PPW7	1	1.6	1.1	0.9
P02766	1	1.2	1.9	2.7
B0FLL2	1	1.1	0.6	1.1
095881	1	1.2	1.7	2.8
P46976	1	2.1	1.9	1.9
P50135	1	1.3	0.8	1.1
H0YJ34	1	1.1	1.2	1.9
P49773	1	0.3	0.4	1.9

014773	1	1.3	1	1.4
P05161	1	0.6	1.2	0.4
A8MUS3	1	0.82	1.64	1.55
D6R9P3	1	1.4	1.9	1
075340	1	1.3	1.1	0.9
000487	1	0.7	0.6	0.8
P63167	1	0.3	0.3	2
P04632	1	0.9	0.8	1.4
060262	1	1.7	1	0.4
Q5T123	1	0.7	0.8	2
E7EMM2	1	0.8	0.9	1.2
A8MXH2	1	1.2	0.8	0.7
BOYIW6	1	0.9	1	1.4
B3KWE1	1	1.3	1.3	1.4
B4DEK4	1	1	1	1.7
B4DP62	1	3.1	2.5	1.4
B4DVB8	1	0.8	1.5	1.3
B4E1F0	1	0.91	1.64	1.09
B4E1K7	1	1.3	1.2	1.2
B7Z815	1	0.9	0.9	1.1
C9JQS9	1	1.3	1.1	0.8
E7EPC8	1	1.5	0.9	1.4
E7ES43	1	0.6	0.8	0.9
E9PC15	1	1.4	1	1.4
E9PFW3	1	0.8	1	1.7
F2Z3F8	1	0.9	0.8	0.6
G5E9L9	1	1.4	1.5	0.7
I3L430	1	1.1	0.8	1.2
J3KPM8	1	0.73	0.55	0.55
J3KTL2	1	1.8	2.4	1.8
000299	1	0.3	0.3	2.3
015031	1	1.2	1.3	0.9
015400	1	1.1	1.6	1.1
015511	1	1.2	1.7	2.3
043837	1	1.1	0.7	1
060437	1	0.4	0.3	0.4
075521	1	2.1	1.4	1.7
075643	1	1.1	0.9	1.7
075964	1	0.9	0.7	1.3
076054	1	1.3	1.1	2
094919	1	0.91	0.82	1.73
P02462	1	0.64	0.55	2.45
P18124	1	0.6	1.6	1.3
P35908	1	2	1.8	1.6
P36955	1	0.5	0.9	1.4
P48506	1	1.6	1	0.7
P53999	1	1.8	1.1	1

P58107	1	0.4	0.3	0.3
Q13136	1	1	2.1	1.2
Q14112	1	0.4	0.4	0.9
Q14118	1	1.3	1.6	0.6
Q14847	1	1	0.9	1.6
Q15046	1	1	1.1	1.2
Q16204	1	0.8	0.8	1
Q4V328	1	0.9	1.1	0.9
Q6PCE3	1	0.8	0.6	1.1
Q8N163	1	0.9	1.1	2.5
Q92747	1	1.33	1.67	1.11
Q99460	1	1.1	1.3	2.4
Q9BZV1	1	0.8	0.7	0.6
Q9UHG3	1	1.3	1	1.3
Q9UHY7	1	0.8	0.6	0.7
Q9UI15	1	0.89	0.67	0.33
Q9UK22	1	1.27	1.09	0.91
Q9Y2Q0	1	0.6	0.4	0.4
Q9Y3C6	1	1	1.1	1.3
Q9Y5X3	1	1	1.2	1.2
Q9Y6N6	1	0.4	0.3	0.4
F5H533	1	0.82	0.55	0.64
H0YFH3	1	0.8	0.7	1.2
095292	1	0.9	1.2	1.1
P39060	1	0.4	0.4	0.7
B4DIR0	1	1.2	1	1
G3V2N0	1	1	1.1	0.9
000264	1	1	1	0.9
Q9NR45	1	0.7	0.5	2.6
P62753	1	0.9	1.1	1.7
Q08170	1	1.4	1.6	2
P31942	1	1.5	1.7	2.2
E9PGT1	1	1.9	1.3	2.2
043852	1	1.6	4.5	1.6
Q08380	1	1.2	1.9	1.3
P31153	1	0.9	0.7	1.6
P41250	1	0.8	0.7	0.9
J3QT22	1	1.3	0.8	1.1
P43007	1	5.8	1.6	1.5
P22059	1	1.1	0.9	1.2
P62745	1	0.6	0.4	0.8
Q01581	1	0.89	1.56	1
P10620	1	1.4	1.3	0.9
F5H1S8	1	0.91	1	1.18
P62942	1	0.9	1.1	1.6
H0YGX7	1	0.7	0.9	2.4
E7EP00	1	0.9	0.8	2.2

P19086	1	1.2	1.1	0.7
060613	1	0.9	1.5	1.8
075368	1	1.6	1.9	0.3
Q96JE9	1	0.56	0.44	0.33
Q9BX68	1	1	0.8	0.7
Q53GQ0	1	1.5	1.1	1.8
P68036	1	0.7	0.9	1
P11215	1	1.4	1.4	5.1
P57105	1	1.3	1.1	1.1
Q9BVK6	1	1.5	2.2	1.4
HOYMV8	1	1.18	1.27	1.73
Q00535	1	0.8	0.7	0.9
Q00341	1	0.6	1	2.9
F5H6I0	1	2.9	3.7	1.4
F5H5M9	1	1.6	1.4	1.3
Q92688	1	1	1.5	1.4
Q2TAM5	1	1.7	1.1	2.2
B7Z8E4	1	0.8	0.9	0.6
C9JEV0	1	0.8	2.4	1
B4DFL4	1	1	0.45	0.55
B4DP47	1	2	1.1	1
B4E2X2	1	0.8	0.9	0.5
B7Z514	1	0.6	0.7	1.3
C9JD32	1	0.9	1.3	1.8
C9JFZ1	1	0.7	0.6	0.6
C9JGI3	1	0.7	1.3	3.8
D6RA82	1	0.6	0.8	5.1
E5RIW3	1	1.3	1.1	1.4
E7EV80	1	1.1	1.4	0.8
E9PC80	1	1.2	1.1	1
E9PJC7	1	2.3	0.9	1.2
F5H301	1	2.18	1	0.64
F5H5G1	1	1.9	1.8	0.7
F5H8B3	1	0.8	0.5	0.9
G8JLD5	1	0.78	0.89	1.56
HOY5W0	1	1.2	1.2	1.3
J3QK90	1	0.9	1.3	1.6
000161	1	1.36	1.45	1.45
043592	1	1	1	0.9
043681	1	1.1	1.2	1.1
075475	1	1.6	2	1.1
075506	1	1	1.1	0.8
075569	1	0.8	0.8	1.1
094819	1	1.8	0.8	1
P02747	1	1.2	1.7	1.9
P23786	1	1.9	1.4	0.7
P27144	1	1.1	0.8	1.5

P31937	1	2.1	1.2	1.4
P34897	1	1.7	1.5	1.7
P46060	1	0.9	0.7	1.5
P47897	1	1.1	1	1.2
P55010	1	0.7	0.6	2.5
P62241	1	0.9	1.5	1.4
P62899	1	1	1.3	1.7
P67809	1	1.56	3.33	2.11
Q12792	1	0.9	1.2	0.8
Q12904	1	0.8	1	1.8
Q13526	1	0.6	0.6	0.8
Q16762	1	2.5	1.2	1.2
Q4G0N4	1	1.5	0.9	1.2
Q6UWR7	1	2.2	1	1.8
Q8N1G4	1	1.7	1.3	1.4
Q8TCU6	1	1.3	1.9	1.1
Q8TDQ7	1	1.3	0.8	1
Q8WWI5	1	1.9	1	0.8
Q96CV9	1	0.55	1	0.82
Q96HC4	1	1.1	0.9	0.8
Q9BT78	1	1	0.9	1
Q9H0E2	1	0.8	0.9	1
Q9NUQ9	1	1.1	1.1	1.7
Q9UNM6	1	1.1	1.1	1
Q9Y5S2	1	1.4	1.5	2.1
C9JYI4	1	1	1.2	1.2
J3KN16	1	0.9	1.1	1.8
043399	1	1.4	2	1.4
P10253	1	0.9	1	0.8
P12270	1	1.2	1.3	1
P43487	1	1.11	1.11	1
P46783	1	1.3	1.2	1.8
Q13153	1	0.5	0.5	0.7
B1AJV7	1	0.4	0.4	0.5
D6RAX2	1	1	1.1	0.8
043396	1	0.9	0.8	0.5
P25787	1	0.7	0.5	4.2
Q00169	1	1.1	0.8	1.1
Q9UKU7	1	2.6	1.1	3.1
E9PGI8	1	0.6	0.6	1.8
P60983	1	1	0.8	1.4
P46109	1	1.4	1.3	0.8
B4DHX5	1	0.8	0.6	0.4
P84095	1	0.8	0.9	4.6
Q7L266	1	1.5	1.2	1.1
F5GXS8	1	0.8	1	1.6
F8W7I2	1	3.3	0.6	0.5

Q96DG6	1	1.1	0.8	0.7
B4E329	1	1.8	1.1	1
075695	1	1.1	1.6	1.3
A8KAH5	1	1.4	1.3	2.1
G3V2F7	1	1.3	1.1	1.3
P05107	1	1.7	1.3	6
F5H376	1	1.2	0.6	0.7
Q9NZZ3	1	1.2	1.6	0.9
Q9UBB6	1	0.8	0.7	0.6
Q9Y3A5	1	1.2	0.9	1.1
H3BN14	1	1.2	1.1	1
E7EM64	1	1	0.8	0.7
P31949	1	0.2	1	1.3
Q96RF0	1	1.7	1	1.4
P62701	1	0.6	1.7	0.6
Q9NR31	1	1.3	0.9	1.9
Q9UBV8	1	1	1	0.9
Q6JUT2	1	2	1.44	2
E7ER47	1	2.22	0.89	0.56
P05198	1	0.7	1	1.5
000534	1	0.9	1	0.8
Q5T1Z0	1	0.9	0.7	0.8
P30405	1	3.1	1.2	1.6
P04196	1	1.09	2	3.91
P07585	1	0.3	0.2	0.7
P61225	1	1.2	0.9	2.3
B7Z590	1	1	0.7	0.5
C9J502	1	0.5	1.1	1.1
E5RFX7	1	1	0.9	0.9
E7EUD0	1	1.6	0.7	0.4
F5H0U5	1	1.3	1	1.3
H3BP91	1	2.3	1.7	1.3
J3QQN1	1	1.18	1.18	1.82
060234	1	1.18	1.09	3
095571	1	1.7	1	1.6
Q7L1Q6	1	1.1	1.1	1.6
P00352	1	1.7	1.1	0.7
000232	1	1	0.9	1.1
B4DX52	1	0.9	1.6	1.3
P30533	1	1	1.9	1.4
A8MZ54	1	0.73	0.55	0.45
Q9NZB2	1	1.1	1.1	1.6
E9PNW4	1	1.2	1	1.6
Q13491	1	2.5	1.4	0.5
H0YN26	1	1.1	1.7	2.1
060331	1	1.2	1.4	0.7
015145	1	1.4	1.3	0.9

B4DG46	1	1.4	0.4	2.5
Q2TAY7	1	0.8	0.8	0.4
Q16543	1	0.8	1	0.9
F8W7Q4	1	1	0.8	2.5
C9J1F6	1	1.5	1.3	1
C9JDR0	1	1.3	2	0.9
D6RGV5	1	1.5	1.1	0.5
095822	1	0.9	0.7	0.7
A6NJA2	1	0.9	0.9	0.7
B0QZ18	1	1.3	1	1.8
B3KS98	1	1.1	1.2	1.9
B4DJS7	1	0.9	1	1.5
B7ZL00	1	0.8	1	1.9
E1P5S2	1	0.9	1.2	1.7
E9PCI9	1	1.09	0.82	0.91
E9PPH5	1	0.9	0.9	2.7
F5GYF7	1	0.8	0.7	0.9
F5H0J4	1	1.5	1.1	1
F8VXJ7	1	1	1.3	2.3
G3V3C7	1	1.1	0.9	0.9
014495	1	4.7	2.3	1.2
015075	1	0.6	0.7	0.7
043765	1	1	1.5	1.9
095671	1	1.4	1	1.5
096000	1	1.4	1.1	1.1
P20039	1	1.1	3.8	2.6
P22033	1	1.1	1.3	1.4
P26368	1	1	1.4	1
P30520	1	0.89	0.67	1.33
P48163	1	1	1.8	0.6
P51858	1	0.9	1.4	1.4
P53621	1	1.1	1.2	1.8
P62333	1	1	1	1
Q06124	1	1.2	1	1
Q13636	1	1.6	1.6	3.4
Q86TP1	1	0.7	0.7	0.6
Q8TBC4	1	0.9	0.9	0.8
Q8TCD5	1	1.3	0.8	1.3
Q92905	1	1	0.8	0.8
Q9H479	1	0.9	0.6	0.8
Q9NQE9	1	1.3	1.3	1.3
Q9NQW6	1	1.5	0.6	1
Q9Y6U3	1	0.9	1.6	2.3
B5MCW2	1	0.82	1.27	1.45
C9JFE4	1	0.9	0.7	1
E9PHI6	1	1.2	1.1	1.6
P26373	1	0.6	1.1	4.2

P46108	1	1.89	1	0.89
P56556	1	1.2	0.8	1
P62081	1	1.1	1.3	1.2
Q08945	1	1.4	1.7	1.2
Q8WXF1	1	1.2	1.2	1.1
Q92945	1	1	1.9	2.4
Q9NQ66	1	1.3	0.9	0.8
Q9Y285	1	0.9	0.8	0.7
B7Z5Z2	1	1.3	0.8	1
E9PEW9	1	1.3	1	1.2
F5H365	1	1	0.8	1.1
P43155	1	2.2	2	1.8
G8JLH9	1	0.9	0.9	1.5
P26196	1	1.4	1.3	1
P56192	1	0.9	0.9	1.3
P62424	1	0.9	1.7	3.6
Q6NVY1	1	1.4	0.7	1.1
A6NFN2	1	1.3	1.3	1.3
B3KXW5	1	0.8	0.8	0.6
000533	1	1.6	1.1	0.9
J3KQ73	1	1.3	1	1.4
Q9Y512	1	1.3	1.1	1.4
B2R5P6	1	0.8	0.7	1.8
B4DGX2	1	1.18	1	0.82
G3V3Y1	1	0.73	0.45	0.73
075223	1	1	0.8	1.9
B7Z7Q0	1	1.8	1.5	0.6
Q96FN4	1	0.9	0.9	1.1
Q9GZP4	1	0.9	0.6	0.5
P39019	1	0.9	1.8	2.8
Q14160	1	1.3	1.5	1.2
Q96CN7	1	1.3	1.2	1.4
D6RF12	1	1.6	1.2	1.1
F8W8T1	1	0.6	1.3	1
P14621	1	1.7	0.9	1
Q96L92	1	1.2	1.2	2.1
Q9UHG2	1	0.4	0.4	0.7
Q9UPT5	1	1.2	1.1	1.1
P09488	1	0.7	0.7	2.2
Q9NVJ2	1	1.6	1.6	1.2
H7C5Q2	1	1.9	1.5	0.7
F5H2F2	1	1.3	2.6	6.3
E9PL82	1	0.9	1.6	0.5
P54727	1	1.1	1.5	0.7
B4DNJ6	1	1.09	1.36	0.55
H0YK46	1	1.1	1.3	3.1
E7ENU4	1	1.1	1.5	2.5

P12004	1	0.6	0.8	1.8
Q9NV70	1	0.9	1.1	1.6
C9JJP5	1	1.2	1.4	1.3
Q6NZI2	1	0.9	1.1	1.2
Q9H313	1	2.1	1	0.8
A8MXP8	1	1.33	1.22	1.33
E7EPB3	1	0.4	0.9	3.3
P55263	1	1.2	1	0.4
Q9UBS4	1	1	1.5	0.83
Q15393	1	1	1.3	0.9
P49257	1	1.36	1.36	1.45
F5H2U1	1	1.7	1.2	1.5
P29590	1	1.2	1.5	2.1
Q02818	1	1.1	1.6	1.6
Q9UJY1	1	1.5	1	1
A2RRF3	1	1.1	1.1	0.6
095373	1	1.7	1.6	0.7
Q9NQR4	1	1.1	0.5	0.8
B1AHD1	1	1.5	1.5	2.2
D6RAT0	1	0.9	1.4	1.8
E9PFN5	1	1.3	2.2	2.2
P23434	1	2.6	1.5	0.9
P49770	1	1.08	1.17	1.17
P61960	1	0.5	0.6	1
P62857	1	1.55	1.91	1.45
Q7L775	1	0.8	0.8	0.8
Q8IU85	1	1.6	0.9	0.8
Q8NHG7	1	1.9	0.8	1.6
Q9BRF8	1	0.8	1.3	0.7
Q9UBI6	1	1.7	3.8	0.8
Q9NZR1	1	2.09	1.45	0.45
D6RAN4	1	1	1.3	1.6
J3KSZ7	1	0.6	0.8	1.1
H3BPE7	1	2.3	2.4	1.3
E7ESZ7	1	1.1	1.1	2.3
P62330	1	1.5	1	1.6
Q658P3	1	0.6	0.9	1.2
Q96FJ2	1	1.1	1.4	0.3
F5H0J3	1	1.2	0.8	0.9
Q9H3N1	1	1.27	1.27	1.82
P53680	1	1.3	1.2	0.9
B0QYF8	1	0.9	0.9	2.4
P49006	1	4.8	2.8	2.4
Q96A65	1	1.2	1	0.9
Q9UHD1	1	0.6	0.8	1
Q9Y6C9	1	0.9	0.8	0.7
HOYM70	1	0.9	0.9	1.3

P27694	1	0.9	0.7	0.6
043615	1	1	1.1	0.8
P04839	1	0.7	2.2	8.4
Q8N573	1	1.2	0.5	0.9
E7ENQ6	1	2.4	1.6	0.9
J3KMX4	1	1.1	1	1.9
Q9NP79	1	0.73	0.82	1.09
G3XAJ6	1	1	1.9	2.1
B4DFQ6	1	1.5	1.3	1.6
P24534	1	0.9	0.6	1.9
H3BVG0	1	1.2	1.3	0.9
F5H4E4	1	1.3	1	0.6
B4DH53	1	0.8	1.4	1.6
Q9Y2Z0	1	1.11	0.78	1.11
P04181	1	2	0.9	1.5
P63218	1	3.6	2.7	1.4
075915	1	1.1	1.8	1.1
Q9P2K5	1	1.3	1.4	1.3
P56381	1	2.1	1.6	1.1
075936	1	0.8	0.5	0.8
Q16348	1	4.5	0.7	1.3
B4DHE8	1	1.2	1.1	0.5
C9JKQ2	1	1.7	1.4	0.8
P05546	1	0.3	1	1.1
P23083	1	0.8	2.1	1.9
Q9H492	1	0.8	0.8	0.5
Q9Y376	1	1.1	1	0.6
C9J6P4	1	1.2	1.3	4.3
P62277	1	0.9	1.8	1.2
B7Z491	1	1.4	1	0.9
A6NHG4	1	1.5	1.6	1
Q07666	1	1.2	1.6	0.8
P09669	1	0.9	0.8	0.9
P43686	1	0.9	0.9	2.3
Q92888	1	1.1	1.9	1.9
015270	1	1.7	2.1	0.8
H0YCN2	1	1.4	1.2	1.3
E9PIM6	1	1	1	1.2
P27635	1	1	1.7	2.9
C9JU1	1	1.2	1.6	2
H9KV64	1	1.7	0.9	1.3
Q96NW7	1	1.5	1.3	1
P02689	1	2.1	1.6	1.2
D6RBN5	1	3.5	1.5	1.6
C9JOB2	1	0.9	1	0.5
E9PE82	1	1.5	1.1	1.1
F8WDI0	1	0.9	1.3	1.6

J3KQL8	1	1.1	3.3	3
P35270	1	1.2	0.8	1.1
Q8N6N7	1	1.2	0.9	0.8
E9PMB8	1	1.8	1	1.6
E9PL57	1	0.3	0.2	2.6
P01033	1	1.5	1.2	16
P50897	1	1.2	1.4	1.5
Q13442	1	2	2.7	1.7
Q15067	1	2.1	1.6	1.3
Q6PI78	1	2.1	1.4	1.2
Q6YHK3	1	0.7	1	0.5
Q96GD0	1	1	0.7	0.8
Q9UNS2	1	1.2	1	1.2
E9PFH3	1	1.1	1.2	1.6
E9PM35	1	1.6	0.7	1
P28070	1	0.7	0.8	1.1
P62263	1	1	1.4	1.6
Q5VZR0	1	0.8	1.1	0.9
E5RJ68	1	1	1.3	1.2
J3KS22	1	2.2	0.9	1.1
B3KUE5	1	0.8	0.8	0.8
J3QRS3	1	1.5	1.5	2.4
P48147	1	0.8	0.7	1.2
P55884	1	1.6	1.6	1.4
A8MWR6	1	1.5	1.3	1.2
G3V1C3	1	1.2	1.4	1.5
096008	1	1.4	1.5	1
Q9NYU2	1	0.9	1	1.4
F8VVR9	1	0.7	0.5	0.6
Q9UBT2	1	1.1	1.1	2.2
Q9UKG1	1	1.2	1.2	1.3
P04114	1	0.8	0.7	1.1
F8W8M4	1	1	1	0.7
A6NNK5	1	0.9	0.9	0.9
Q5VWZ2	1	1.2	0.9	0.8
Q8N8S7	1	1.3	1.9	1.1
C9J574	1	1.3	1.4	0.8
F8W6G1	1	1.1	1.1	0.5
P17931	1	1	1.1	2.9
Q14258	1	1.1	1.8	0.9
Q15126	1	1.1	1	1.9
A6NDG6	1	1.4	0.9	1
B4DXZ6	1	1.3	1	1.3
C9JAF2	1	0.4	0.4	0.7
P13671	1	0.7	0.9	2.8
B2CZX2	1	1.2	0.8	0.6
HOYGW7	1	2.4	1.6	1.5

P08572	1	0.3	0.3	3.4
P49458	1	1.6	0.9	2
P41091	1	0.9	1.2	1.6
P02792	1	1.6	1.1	1.8
Q9NT99	1	1.8	1.3	1
P01620	1	0.6	1.5	1.7
E7ENA9	1	1.5	1.4	1.3
Q96BM9	1	1.3	1.3	1
B1AK13	1	1.3	1.7	1.1
O76024	1	0.8	0.9	1.3
E9PJD9	1	0.9	2.1	1.3
P01111	1	1.1	1	1.2
B4E220	1	5.7	1.3	1.6
Q16363	1	0.9	0.9	4.4
B4DU60	1	1.1	0.6	1.2
O60271	1	1.1	1.3	0.9
B4DER1	1	1.2	1.2	2.2
B4DXI8	1	0.5	0.7	2.4
B7Z2T9	1	3.9	1.2	1
D3YTF8	1	1.5	1.1	2.1
E5RFP0	1	0.6	0.5	1.4
E5RIZ6	1	0.6	1	1.6
E9PFR3	1	1.1	1.3	1.5
F5H355	1	1.6	1	0.5
F6SKB8	1	0.8	0.9	1.2
F8VUA6	1	0.9	1.7	3.4
G3V1L9	1	2.2	1.6	1.1
H0YLG5	1	1.6	1.3	1.4
H3BND3	1	1.2	1.5	0.9
H3BPK3	1	1.44	1	1
H3BS86	1	0.9	1	1.1
O14880	1	1	0.8	1.4
O14964	1	1.1	0.9	1.1
O60443	1	1.2	1.1	1.2
O95182	1	0.6	0.6	1.1
O95747	1	1	1.11	1.33
P00167	1	1.2	1.2	1.4
P00492	1	0.6	0.5	0.7
P08311	1	0.7	0.8	14
P55145	1	1.36	2.73	1.18
P62854	1	1	1.8	2
Q14353	1	0.64	0.64	0.73
Q15404	1	0.9	1	1.1
Q15459	1	0.9	1.4	1.6
Q16740	1	1.64	1.36	1.55
Q8N766	1	1.2	1.3	1.3
Q96EM0	1	0.9	0.6	0.8

Q96NY7	1	0.6	0.4	0.9
Q9H444	1	1.3	1.5	2.7
Q9Y2V2	1	0.5	0.6	0.6
Q9Y4G8	1	0.6	0.5	0.7
B4E2S7	1	1.7	2.6	1.1
I3NI00	1	0.64	0.73	2.91
Q92900	1	0.9	0.9	1.9
C9IZQ1	1	1.4	2	1.4
H3BQZ9	1	1.1	1.8	0.6
G3XAC4	1	1.2	1.4	0.8
P05413	1	0.7	0.8	1.5
E9PDL2	1	2	1.33	0.33
B7Z7P8	1	0.8	0.8	1.1
J3KNX9	1	1.7	1.5	0.7
000625	1	0.6	0.5	0.8
P67870	1	0.9	1	1.2
095168	1	1.3	0.7	1
P13647	1	1.3	1.2	1
H7C410	1	1.7	5.2	2.8
E7EU96	1	1.1	1.2	1.4
A6PVN5	1	1.11	0.89	1.56
E2QRM6	1	1.1	0.7	0.8
F5H425	1	1.1	1.1	1.1
015498	1	1.3	1.4	1
Q15293	1	2	2	2.2
J3KNM0	1	1.2	0.9	1.5
E7ENJ6	1	1.1	1.3	1.1
B7WP84	1	1	0.9	0.7
B4DST5	1	1.1	1.3	1
P50583	1	0.9	0.8	0.9
P01116	1	0.82	0.82	1.36
P41218	1	0.9	1.2	3.9
G3V153	1	1	1.2	1.2
C9JT23	1	1.3	0.9	0.4
Q5R370	1	1.1	1.1	1.8
H0Y2Y8	1	0.7	2.5	11.1
B4DQV7	1	0.7	1.1	2
Q13576	1	0.7	0.7	0.9
Q9Y3U8	1	1	1.8	1.5
E2QRD5	1	0.9	1.1	1.9
Q96E39	1	1.2	2	1.4
A6NIW1	1	3.3	2.2	1
Q6PCB0	1	0.3	0.4	0.6
H7BXF3	1	1.2	1.6	1.2
H3BPF6	1	1.4	0.9	2.5
P04433	1	0.7	2.3	3.5
D3YTA9	1	2	1.7	0.9

G3V4U0	1	0.5	0.4	0.6
G3V2L6	1	0.7	0.6	1
I3L4J1	1	0.8	0.9	1.1
043143	1	1.4	1.3	1.3
Q92616	1	1.3	1.6	1.1
B7Z1I0	1	0.6	0.7	0.8
Q9UQ35	1	0.8	1.2	1.8
Q9BUH6	1	1.3	1.3	2.5
E9PG35	1	1.6	1.4	1.4
E9PLX3	1	1.2	1.2	2.1
P41208	1	1	0.8	0.8
000116	1	1.4	1.5	2.2
F8VZZ2	1	1.3	1	1.1
Q5LJA5	1	1.44	1.33	0.78
A6NMA8	1	1.4	1.1	1
P11279	1	1.2	1.2	1.4
B1AH49	1	1.7	1.1	1.4
F5H3I5	1	1	1.7	1.3
Q96SB3	1	1.4	2	1.1
Q9BY32	1	1	1.8	1.3
095989	1	1.1	0.9	0.4
F5H8L6	1	1.3	0.9	0.8
B7Z952	1	0.6	0.7	0.4
E9PNG0	1	1.5	1.5	1.6
P61077	1	0.7	0.7	1.3
Q9HCJ6	1	1.2	2.1	0.8
P46782	1	0.9	1.5	3.7
B0QY89	1	0.9	1.3	1
Q86Y56	1	1	1.2	0.6
F5H1F3	1	0.4	0.1	0.2
F5H6P7	1	1	1.6	0.9
000629	1	0.8	0.9	0.6
P02654	1	0.6	1.3	0.8
A8MVM2	1	0.3	0.2	0.2
Q96EK5	1	0.9	0.8	0.7
015355	1	1.1	1	1
P51452	1	0.7	0.5	2.7
B4DNW0	1	1	0.7	1
Q9UL25	1	1.6	1.6	0.9
B1AK40	1	2.4	1.7	0.8
B1APY8	1	1.1	1.3	1.2
Q9NRF8	1	0.8	0.8	1
P59665	1	1.6	1.7	21.1
B4DIZ2	1	1	0.7	1.1
000170	1	0.8	0.7	1.1
Q5VYL6	1	0.4	1.5	3
P19652	1	1	1.5	1.7

Q7L0J3	1	0.4	1.1	1.9
Q86SK9	1	1.44	1.11	1.78
Q8IYI6	1	1.7	0.9	1.5
F5H577	1	2.2	2.1	0.7
I3L505	1	1.9	1.1	0.3
B7Z1N7	1	1.8	3	2.3
P07919	1	0.9	1	0.6
P17050	1	0.7	0.6	1
Q6Y288	1	0.8	1.3	0.9
B4DDM6	1	1	1	2.1
P61601	1	1.4	2.3	0.4
P20774	1	1	1.1	1.6
E7EVV3	1	0.5	0.5	0.8
E9PC52	1	6.6	1.5	0.8
E7EPM0	1	0.8	0.9	3.4
P37108	1	1	1.3	1.7
Q14677	1	1.3	1.2	1
B4DZ15	1	0.9	0.5	0.5
F8VPJ2	1	3.2	1.3	0.3
A8MU27	1	0.9	0.9	1.7
BOYIW2	1	0.6	1.4	1.8
B4DT35	1	0.9	0.9	0.9
C9JNW5	1	1	0.9	2.2
E9PAV3	1	1.1	2	1.7
F8W8D6	1	2.9	1	0.5
000483	1	2.1	1.5	0.5
P50150	1	1.2	1.3	0.7
P56385	1	1.5	1.2	0.7
Q6UXB8	1	2.4	1.2	0.2
Q7Z3B1	1	1.2	1.1	0.4
Q8NF37	1	0.8	1.9	2.1
Q9H3Z4	1	3.1	3.2	1.2
P80748	1	0.5	1.3	2.9
Q9NRX4	1	0.6	0.6	2.5
P54725	1	1	1.3	1
B4DHD4	1	0.8	1.4	1.7
HOYKU1	1	0.7	0.9	2
B2RP62	1	1.3	0.9	1.5
Q709C8	1	1.3	1.1	0.6
A2A2M0	1	0.9	1.1	1.3
A6NCQ0	1	2.3	1.8	0.9
A8MXY0	1	2.7	2	2
BOQY01	1	1.2	1	0.8
BOVOL0	1	1.1	1	1.7
B1AKR6	1	2	1.6	0.8
B3KRI2	1	0.9	0.9	0.8
B4DDW0	1	1.3	1.1	1.7

B4DVY1	1	1	1.1	0.9
B4E1G6	1	1.2	1	2.1
B7Z4U7	1	0.7	0.8	1.3
B7ZAM9	1	1.2	1.1	1.1
C9J3D7	1	1.3	1.3	0.8
C9JB50	1	1.4	1.6	2.1
C9JCN0	1	0.6	0.7	1.5
E5RGR0	1	0.9	1	2
E9PB14	1	1.7	0.9	1.4
E9PFR7	1	1.1	1.1	0.7
F2Z2Y6	1	1.4	1.6	1.9
F5GZD0	1	0.7	0.7	0.7
F5H025	1	0.6	0.5	0.8
F5H7K4	1	0.7	0.8	2
F8VQE1	1	0.9	2.1	2.3
F8VQQ3	1	1	0.9	1.1
G3V2S6	1	0.7	0.7	1.4
H3BSW0	1	1	0.8	1.3
H7BZM7	1	1.2	1.1	1.5
J3KN36	1	1.2	1.8	1.8
J3QRU8	1	0.8	0.9	0.9
014879	1	1.6	1.1	0.9
043678	1	1.2	1.1	1.2
P26022	1	1.1	3.4	8.7
P26885	1	1.2	1.2	1.6
P41217	1	1.4	0.9	0.8
P61201	1	1	0.8	1.3
Q9BXJ0	1	0.5	0.3	0.7
Q9NS86	1	1.1	0.8	1.2
Q9P2B2	1	1.3	2.7	1.8
Q9Y240	1	0.2	0.1	0.2
Q9Y3D6	1	1.7	1.2	1.5
E7ESC6	1	1	1.2	1.2
E7EUG3	1	1.2	1	1.6
Q99598	1	1.1	0.9	0.9
015173	1	1.4	1.7	1.3
Q9Y6N5	1	0.8	1.7	1.4
G3V1X9	1	1.3	2.6	0.9
E9PGT6	1	1	0.8	1.8
Q10713	1	1.1	1.2	2.5
P48059	1	1	1	0.9
B5MCC7	1	1	0.8	1.5
E9PJJ5	1	1.2	1.9	1.3
B4E2Z0	1	3.9	2.1	1.5
G5E9L4	1	1.5	1.4	1.7
P28072	1	0.9	0.7	1.8
P36404	1	1.3	0.9	0.7

Q92896	1	1.3	1.2	1.7
B4DK66	1	0.7	1.3	0.4
E9PLW6	1	0.9	0.7	0.7
F5H1W7	1	1.6	1.4	1.5
G3V1U4	1	1.1	1.2	2.1
P10155	1	1.5	1.8	1.1
Q00688	1	1.4	1.1	1.9
E7EMD0	1	0.9	1	1.2
Q96PY5	1	1.4	0.9	1.1
B7Z7A5	1	1	1.2	0.6
Q9P265	1	1.2	1.2	1.6
Q8NC51	1	0.9	2.7	4.1
P19623	1	0.9	1	1.1
Q9Y3B4	1	1	1.1	1.2
Q8TB36	1	1.5	1	0.8
B8ZZY2	1	0.9	0.8	1
H3BUU8	1	1.6	1.2	1
F5GX23	1	0.9	1.1	1.8
B5MC72	1	0.9	0.6	1.2
B4E2C0	1	1.4	1.1	0.9
Q9HCD5	1	1.8	1.8	1.5
B0QYG7	1	0.9	1.1	0.9
E9PFD7	1	3.7	2.2	2.5
Q5T6V5	1	0.6	0.7	0.8
Q01650	1	1.5	1	1.7
Q01081	1	1	1.4	1.5
J3KRX5	1	0.8	1.4	1.7
Q9UMY4	1	1.1	0.8	1.8
Q96N66	1	0.6	0.9	1.3
Q9NW15	1	1.5	1.2	0.5
Q15006	1	1.2	1	1.5
Q9Y6K8	1	0.7	0.3	0.4
Q6YP21	1	2.5	2	1
HOYNE9	1	1.9	1.6	2.9
000507	1	1.2	1.1	1
Q13459	1	1.4	1.6	1.4
HOY9Y3	1	0.4	0.4	0.2
Q9H1E3	1	3.5	5.3	1.5
Q8NE62	1	2.8	1.4	1
F5H4B6	1	1.3	2.1	1.2
C9JF58	1	0.5	0.4	0.3
HOYKC5	1	1.8	1.1	1.7
G5E9Z8	1	0.9	1.1	1.7
A6NKV8	1	1.3	3.1	0.9
Q9BVA0	1	0.8	0.7	1.4
D6RCDO	1	1.3	1.4	1.6
Q6UB35	1	1.6	1.5	0.9

Q9HBT6	1	4.3	1	0.8
P51888	1	0.8	0.5	1
F5GWU7	1	0.9	0.9	1.6
F5H2A7	1	1	0.9	1.4
P09012	1	0.9	1.4	1.5
Q2M389	1	1.2	1.3	1.3
Q8N335	1	1.6	1	1.4
Q92820	1	0.9	1.7	2.1
Q9UBB4	1	1.6	1.3	1
075533	1	1.2	1.8	1.5
Q9NZJ9	1	1.5	0.9	0.6
060831	1	0.9	2	1.2
B4DDF4	1	0.7	0.6	3
B4DK71	1	1.4	1.7	1.2
P68402	1	0.9	0.5	0.9
E7ETU9	1	5.2	1.5	14
C9JK10	1	5.1	1.3	1.4
P10619	1	0.8	1.6	1
Q9NRZ7	1	1.8	1.2	1
Q9UNW9	1	1.2	1.3	1.7
P43652	1	0.7	1.5	2.9
C9JHR8	1	1.3	1.4	3
Q92621	1	1.2	1.3	3
Q9UBR2	1	1.2	1.3	2.8
P07305	1	1.1	1.1	0.9
C9JRP8	1	0.6	0.6	0.9
C9JJX6	1	0.8	0.5	0.5
H0Y599	1	2	1	1.1
060678	1	1.1	1.3	1.3
014949	1	1	1	1.6
J9JIE9	1	1.1	0.7	0.4
B4E3W0	1	1.5	1.4	1.3
P26583	1	1.8	3.6	2.9
Q53GG5	1	0.7	0.9	2
Q96GA7	1	1.2	0.7	0.9
Q9P258	1	1.5	1	1
H3BUQ9	1	1.5	0.8	2.1
P62888	1	0.7	1.5	1.8
B4DE40	1	1.2	1.7	3.9
F8WD04	1	1.3	0.8	0.4
B3KSQ1	1	1.4	1.4	0.8
G3V0G5	1	0.7	1	1.2
Q13057	1	1.6	0.8	0.7
A2T926	1	1.4	1.8	1.7
Q3ZCQ8	1	2	2.6	0.5
A6NHH0	1	1.3	1.2	2.2
A8MYV2	1	1.2	1.6	1.6

B4DTE8	1	1.4	3.1	3.1
E5RFY9	1	0.7	1.3	0.4
H0YLV5	1	1.4	0.8	1.7
B0QYN7	1	0.7	0.9	2.2
P01031	1	0.6	1.3	3.9
P10586	1	1.5	1.2	0.6
Q6P2E9	1	1.2	1.7	2.1
P49207	1	0.6	1.9	3.1
Q9Y371	1	1.2	1.7	1.2
E9PQ57	1	1	1	1.6
B7WPD3	1	1.1	1.2	0.9
J3QL56	1	1	0.8	1.6
014576	1	0.7	0.7	0.8
F5GWR7	1	1.5	1.2	0.8
C9JV86	1	1.2	1.6	1.8
F5GWP3	1	0.4	0.8	0.3
Q96CT7	1	1.2	1.2	1.2
Q9BX67	1	1.1	0.7	1.4
Q8N3V7	1	0.5	0.9	0.8
Q96L34	1	0.9	0.7	1.3
Q6UW68	1	1	0.9	1
Q8TD22	1	1.5	0.9	0.7
Q9H0Q0	1	0.5	0.6	1.2
F5GYL8	1	0.6	0.9	1
P46778	1	0.8	1.1	1.5
094874	1	0.9	1.1	1.6
E7ERD7	1	1.9	1.4	2.8
C9JL12	1	0.6	0.5	1.1
P11182	1	1.6	0.9	1.4
076031	1	0.9	0.7	1.6
Q9UII2	1	0.7	0.5	0.5
E9PB70	1	0.8	0.9	3.2
Q9UHV9	1	3.2	0.9	2.1
E5RGN3	1	0.9	0.5	1.7
P61011	1	1.3	1	1.5
Q9H6K4	1	1	0.9	1.6
H0Y4I7	1	1.7	2.5	4
H0YDU8	1	0.6	0.5	1.5
F5H293	1	0.7	0.9	0.7
P20160	1	0.6	0.5	4.4
Q5VVC8	1	0.9	1.2	2.2
H3BTV7	1	2.3	2.8	1.1
P01766	1	0.7	1.3	1.9
Q92556	1	1	0.9	1.4
F2Z357	1	0.9	1	0.6
E7EN38	1	0.9	0.9	1
P23469	1	1.3	1.6	0.3

Q96AG4	1	1.2	1.3	1
A0JP02	1	1.4	1.3	0.6
A8MQ45	1	2.8	1.5	1
A8MUM1	1	1.4	0.9	1
A8MV37	1	0.7	0.8	1.6
A8MXR0	1	0.8	0.9	0.6
B1ANG9	1	0.7	0.7	0.8
B3KVR1	1	1.7	2.4	0.6
B4DRL5	1	1.5	0.7	0.7
B4E3H6	1	1	1.4	0.9
B5MC59	1	0.9	0.7	1
B7Z468	1	1.2	1.9	0.7
B7Z9G5	1	0.7	0.8	0.9
C9IYM4	1	4.1	4.5	2.2
C9J8T0	1	1.1	1.1	1.7
C9JCD9	1	1	0.9	1.3
C9JF50	1	0.7	0.7	1
C9JRL6	1	0.6	0.6	0.8
C9JYN0	1	3.5	1.4	1
D6R934	1	1.6	2.3	0.6
D6RFH4	1	1.6	1.8	3.5
E5RH35	1	0.9	1	1.5
E7ENV9	1	0.9	1	0.8
E7EUE1	1	2.7	2.7	2.4
E7EWP0	1	1.7	1.4	0.9
E9PB95	1	1	1.2	1.7
E9PCB6	1	2.2	2.3	1.3
E9PCG9	1	0.7	0.9	0.4
E9PGF9	1	0.7	1.2	0.8
E9PIZ3	1	0.5	1.3	1.6
E9PJ36	1	1.2	0.9	2.6
E9PJF4	1	1.4	1.2	1.4
E9PLI5	1	1	1	0.9
F5GX77	1	0.8	0.6	1.7
F5H2L4	1	1.6	1.2	0.7
F5H442	1	0.8	0.7	0.7
F5H4W9	1	0.6	1.3	1.1
F8VQS7	1	1.9	1.6	1
F8W181	1	1.7	1.9	3.4
H0YD16	1	1.1	1.2	2
H3BP35	1	0.9	1	1.9
H7BZG9	1	1	1.2	0.2
H7C159	1	1	1.4	1.7
I3L2R6	1	0.9	0.7	1.4
J3KN10	1	0.7	1	1
J3KSP2	1	0.7	1.4	4
000160	1	0.9	0.8	4.2

000567	1	1.5	1.2	1.4
014494	1	0.8	0.7	0.3
043665	1	0.5	0.7	0.7
075165	1	1	1.3	2
095674	1	1.7	1.2	1.3
P01596	1	0.5	2.9	3
P01625	1	0.8	2.2	3.3
P15927	1	0.9	1	0.9
P19388	1	1	1	1
P35542	1	0.9	1.6	2.8
P47914	1	0.7	3.8	5.5
P48507	1	0.6	0.6	0.8
Q04941	1	0.9	0.9	0.7
Q13618	1	0.9	0.7	3.2
Q15056	1	1.2	1.3	1.1
Q5QPL9	1	1.6	1.8	1.2
Q8N126	1	0.6	1.1	0.7
Q8N129	1	1.1	1.5	1
Q92882	1	0.9	1	2
Q99523	1	1.2	1.6	1
Q9BR76	1	1.2	0.9	3
Q9BT88	1	0.6	0.7	1.1
Q9BTV5	1	1	0.9	0.8
Q9HAV7	1	1.4	1.1	1.5
Q9NR50	1	1.1	1	0.9
Q9NX46	1	0.9	0.9	1
Q9UBG0	1	0.9	1.7	0.9
Q9UFN0	1	1.5	1.6	0.4
Q9UMX5	1	1.4	2.4	0.8
Q9Y224	1	1.2	0.9	0.6
060826	1	0.9	1	1.4
Q9Y5L4	1	1.9	1.3	1.5
060883	1	1.2	0.3	1.4
P04211	1	0.8	1	1.7
B0QZ40	1	1.7	1	1.5
Q96SZ5	1	1.6	1.4	1.8
P01593	1	0.67	3.67	3.67
B4E0T7	1	1.3	1.1	1.7
Q8NOX7	1	0.5	0.6	0.7
B0QYK0	1	2.2	3.6	0.9
A6NLF3	1	1.2	1.1	0.7
F5H667	1	1.1	0.9	1.2
Q9P2X0	1	0.8	1	1.5
A6NHK2	1	0.4	0.6	8.2
B1AH89	1	1.3	1.1	0.9
F5GXR3	1	1.1	1.6	1.4
P01700	1	1.6	1.5	8.4

A6NDN0	1	0.9	0.8	1.1
A8MYJ1	1	1.1	1.1	2
B4DKT0	1	1	1.1	1.8
C9JBT1	1	0.9	1	1
P02743	1	0.7	1.2	1
Q9UDW1	1	1.7	1.3	1
C9JQ41	1	1.6	1.5	1.6
B7Z831	1	1.3	0.8	2.2
A6NCK0	1	1	0.9	0.9
P55008	1	1.4	1.9	2.5
Q13336	1	1.6	0.6	0.8
C9JKI3	1	0.8	1.9	2.6
E7EPX7	1	0.9	0.6	0.9
Q10570	1	0.8	0.9	1.5
A2A283	1	0.5	0.3	0.3
H3BP41	1	0.8	0.8	0.4
Q9H223	1	0.5	0.9	4.6
P37235	1	1.3	1.4	0.7
P78356	1	1.2	0.8	1.3
B4DJK0	1	1	1.9	0.9
H0YA52	1	1.8	1.3	1
B7WPG3	1	0.9	0.8	1.8
Q9Y4W6	1	1.3	0.8	1
F5H157	1	1.5	1.3	1.2
F5GWF1	1	0.8	0.7	2.8
B7Z341	1	1.2	1	0.7
Q13425	1	1	1	2.1
Q9NQH7	1	1.1	1.5	1.7
P01781	1	0.5	1.3	1
Q16222	1	1.5	1.2	2.8
B3KYA7	1	1.1	1	4.4
B4E2I7	1	0.5	0.9	1.1
P13284	1	1.7	1	3.8
E7ESP4	1	1.9	1.3	1.3
P19174	1	0.9	1.2	1.7
F6T1Q0	1	0.8	0.9	1.5
P23946	1	1.1	0.8	1.3
Q6PKG0	1	0.9	1.1	3.1
B7Z1Q2	1	1.4	1	0.9
A2ACR0	1	1.2	1.3	3.3
A6H8Z3	1	0.5	0.8	2.4
A6NE29	1	0.8	1.1	0.6
A6NGV1	1	2.9	3.7	0.4
A6NKF9	1	1.3	1.9	1.2
A8K3C2	1	0.7	0.5	0.9
A8MUW5	1	1	0.9	0.5
A8MYZ9	1	7.2	3.3	0.8

A9Z1X7	1	0.8	1.6	0.6
B1ALD8	1	1.4	0.9	15.8
B1ALK7	1	1.2	1.1	1
B1AUU8	1	1.3	1.9	1.2
B1P7G3	1	1.3	1.3	0.8
B2R8E1	1	1.1	1.2	1.4
B3KNJ4	1	1.1	1.4	1.2
B3KSH1	1	1.5	1.3	1.6
B3KSY4	1	0.8	1.1	2.7
B4DIB6	1	0.9	1.1	1.5
B4DJM9	1	0.7	0.6	0.6
B4DLH2	1	1.9	1.7	2.9
B4DLN1	1	1.2	1.7	1.4
B4DP38	1	1.1	0.9	2
B4DT31	1	1	1.2	2.1
B4DTC1	1	1	0.9	1.2
B4DUX5	1	2.1	1.9	0.6
B4DZM8	1	0.9	1.4	1
B4E016	1	0.7	0.7	1.6
B4E3Q1	1	1.2	1	1.2
B5MCL2	1	10.2	1.9	0.7
B5MEF0	1	1	0.4	0.5
B7Z5X7	1	2.2	0.9	0.8
B7Z6P0	1	0.7	1.1	0.9
B7Z7L0	1	1.3	1.5	0.6
B7Z9D1	1	0.9	1	1.1
B8ZZJ1	1	0.8	0.8	1.1
B8ZZM6	1	1.1	1.7	0.7
C9IZ01	1	2.4	1.9	1.5
C9IZD3	1	1	2.1	0.9
C9J315	1	1.5	2.1	1.1
C9J4G9	1	0.9	1.3	1
C9J5W5	1	1	0.8	0.6
C9JAF7	1	0.8	0.9	2
C9JBI3	1	0.8	0.6	1.4
C9JID8	1	0.9	0.5	0.6
C9JIF9	1	0.8	1.1	0.8
C9JK83	1	1	1	0.9
C9JVN9	1	1.4	0.9	1.1
D2X2H5	1	3.2	2.7	1
D6RD47	1	0.9	1.4	1.6
E5RFU3	1	1.4	1.3	1.3
E5RIT4	1	1	1	2.1
E5RIY1	1	1.1	1.2	1.1
E5RJU9	1	1.3	1.7	2.2
E7EQT4	1	2.6	1.5	0.6
E7ERZ4	1	1.2	1	1.3

E7ETJ9	1	1.2	1.3	1.8
E7ETU6	1	1	1.3	0.8
E9PB30	1	1.6	2.3	1
E9PB61	1	2	3.9	1.6
E9PBV3	1	3	1.3	2.6
E9PCY5	1	0.8	1.1	1.1
E9PF24	1	1.2	1.3	2.1
E9PJ54	1	1.5	2	0.9
E9PJN0	1	1.4	1.8	1.2
E9PJS4	1	1.1	1.2	2.1
E9PKL9	1	1.2	0.9	1
E9PN05	1	1.2	1.6	0.9
E9PN86	1	0.8	0.7	1.4
E9PNJ4	1	1	0.7	0.9
E9PQ56	1	0.6	0.7	0.7
E9PRK2	1	2.4	7.6	1.3
E9PRT4	1	0.9	0.8	1.4
F5GX07	1	1	1.1	1.4
F5GXT8	1	2.4	1.8	1.2
F5GY80	1	0.7	1.3	14.5
F5GYC3	1	1.4	1.6	0.4
F5GYJ8	1	0.9	0.5	0.2
F5GYM0	1	1.1	1.1	1.5
F5GZU8	1	1.2	3	0.5
F5HOK8	1	6.2	1.5	1.8
F5H2E3	1	1.9	1.3	0.8
F5H6A3	1	1	1.1	1.3
F6RWC2	1	1.1	0.8	0.8
F8VP53	1	1.2	0.9	1
F8VPF7	1	1.2	1.3	1.6
F8VV32	1	1.5	0.9	4.2
F8W061	1	1.4	1	0.9
F8W720	1	1	0.8	1.1
F8W785	1	1.4	1.5	1.2
F8W9B8	1	1.1	0.9	0.9
F8W9U3	1	0.6	0.6	1.2
F8WBI3	1	1.5	1.2	1.9
F8WEE0	1	0.9	0.9	1.9
G3V2C6	1	1.1	1.4	1.5
G3V2S9	1	1.5	1.6	2.6
G5E933	1	1.6	1.1	0.6
H0Y415	1	1.1	0.7	0.8
H0Y8C3	1	1.6	1.1	2
H0Y8X4	1	1.7	1.1	0.9
H0YAV1	1	1.1	0.8	1.2
H0YI90	1	0.9	0.7	1.3
H0YJ66	1	1.1	1.3	1.1

HOYLH9	1	1.2	1.2	1.6
HOYLR3	1	1.1	1.9	1.6
HOYM76	1	1	1.3	1.1
H3BN51	1	1.1	0.9	0.9
H3BNB9	1	0.8	1.1	1.1
H3BPR2	1	1.6	0.8	1.9
H3BQ06	1	1.3	1.1	1.5
H7BXW3	1	0.6	1.6	0.7
H7COX7	1	1.3	1	1.3
H7C5B8	1	2.9	1.3	2.2
I3L097	1	1.6	1	1.5
I3L2E3	1	0.5	0.6	1.1
J3KNB4	1	0.7	0.6	14.2
J3KQI0	1	0.3	0.2	0.2
J3KRS1	1	1.1	1.3	0.5
J3KTE4	1	0.6	2.4	2.4
000186	1	1.7	1.3	1.2
000468	1	0.6	1.2	2.5
014737	1	1.3	2.1	0.9
014907	1	0.9	0.8	0.6
015126	1	0.8	1	1
015240	1	0.8	0.5	1.1
043402	1	1.3	1.3	1.1
043815	1	0.9	1	0.7
043924	1	1.1	1	0.9
060220	1	1.2	1.4	1
060783	1	1.7	1.3	1.2
075438	1	1.1	0.9	1.4
075462	1	0.6	0.3	0.4
095183	1	2.3	1.4	2.7
095297	1	3.3	2.9	1.2
P01880	1	1	4.7	2.1
P02655	1	0.7	1.4	2.9
P02745	1	1.2	2.2	2.2
P02795	1	0.6	0.6	4.4
P05186	1	1.7	2.9	1.1
P05362	1	1.3	1.4	1.7
P07357	1	1.4	2	7
P07360	1	0.4	0.7	4.9
P08621	1	1.2	3	3.7
P13521	1	0.4	0.3	0.7
P16144	1	1.2	2.1	0.4
P23258	1	1.5	1.3	0.7
P26447	1	0.5	0.6	2.3
P35754	1	1.3	1.5	2.7
P40222	1	1.1	1.1	6.6
P41240	1	1	1.1	2.3

P43121	1	1.1	1.2	1.7
P48723	1	1.5	1.3	0.3
P50402	1	1.2	1	1.1
P50453	1	1.1	1.2	1.3
P51687	1	1.3	0.7	0.9
P51809	1	0.8	1.2	1.3
P54886	1	0.8	0.8	1.2
P56597	1	0.4	0.3	0.6
Q05469	1	0.9	0.7	0.8
Q06136	1	1.8	1.4	1.6
Q06203	1	2.1	2.5	1
Q13085	1	1.3	1.5	0.8
Q13825	1	1.2	0.9	1.3
Q15043	1	1	0.9	1
Q15526	1	1.5	1.1	0.9
Q15750	1	1.2	1.2	0.8
Q2M2I8	1	0.5	0.4	0.1
Q3ZAQ7	1	0.9	2	1.6
Q53FA7	1	0.6	1.2	0.8
Q5EBL8	1	0.8	0.9	0.8
Q5RI15	1	1.8	1.5	1.7
Q5T4F6	1	4	0.8	0.7
Q5T4S7	1	1.1	1.1	1.2
Q5ZPR3	1	1.1	1.2	1.2
Q6KCM7	1	1.9	1.5	1.1
Q6PGP7	1	1.2	1.5	0.9
Q6PI48	1	1.7	1.1	1.4
Q6PML9	1	2.3	1.6	0.6
Q6Y1H2	1	1.1	0.9	1.2
Q6ZMI0	1	0.6	0.6	0.7
Q75MJ1	1	2.5	1.6	2.2
Q7Z5R6	1	1.2	1.3	2.3
Q86UX7	1	0.7	1	9.2
Q86VF9	1	5.4	1.5	1.1
Q86VQ2	1	1.9	0.8	0.5
Q8IVP1	1	0.5	0.4	0.5
Q8N1V2	1	0.4	0.4	0.4
Q8N3E9	1	4.7	1.5	0.4
Q8TCT9	1	1.4	1.9	1.7
Q8TEA8	1	1	1.4	3.1
Q8WX93	1	0.7	1.1	3
Q8WY54	1	1	0.6	0.8
Q8WZA9	1	2.4	1.6	0.6
Q92506	1	1.4	0.9	1.2
Q92623	1	0.9	0.9	1
Q969H8	1	1.1	1.4	1.8
Q969V3	1	1	1	1.3

Q96B54	1	1.6	1.9	4.6
Q96EE3	1	1.2	1	1.3
Q96IJ6	1	0.8	0.7	5.3
Q96KP1	1	1.4	1.3	1.5
Q96LJ7	1	1.3	0.9	3.3
Q9BPX5	1	1.4	1.4	0.8
Q9BQ69	1	1.7	1.5	1.2
Q9BQD3	1	1.1	0.9	1
Q9BRJ7	1	1.5	1.1	1.3
Q9BSF4	1	1.1	0.7	1.2
Q9BVG4	1	0.9	1.1	2
Q9BXJ8	1	1.5	1.3	0.6
Q9C0C9	1	1.5	1.1	1
Q9GZM7	1	0.8	0.6	0.9
Q9GZN8	1	0.9	0.9	1.5
Q9H1K0	1	0.8	1.1	0.6
Q9H5X1	1	1.1	0.8	1.1
Q9HBL7	1	3	2.6	2
Q9NS69	1	2.4	2.1	0.8
Q9UDY4	1	1	1.2	1.1
Q9ULC3	1	0.8	0.7	0.5
Q9UMF0	1	0.4	0.2	0.2
Q9Y2Q5	1	1.2	1.6	1.7
Q9Y333	1	1	1	1.3
Q9Y678	1	0.8	1.2	1.5
Q9Y6H1	1	2.1	2.7	1.6
F5H045	1	1.3	1.2	1.1
Q86UU1	1	1.3	0.9	1.8
F5GZI4	1	0.9	0.9	0.9
015258	1	1.3	1.9	0.6
C9JIY2	1	0.9	1.1	1.9
A8K8E1	1	1.5	1.2	1.1
H7BYM6	1	0.7	0.6	7.3
000505	1	0.9	0.9	0.6
Q93034	1	0.9	0.9	0.9
HOYGE4	1	0.8	0.7	1
P63172	1	1.4	1.1	1.1
Q9UGP8	1	1.7	0.9	1.5
P54646	1	1.3	1	1
P21953	1	1.5	1.2	3
B4DS69	1	0.7	1.2	1
B1AQP4	1	1.4	0.9	0.8
P14780	1	1.5	1.5	8.3
D6RCF4	1	1.5	2.1	1.7
P61457	1	1	1	0.9
P30419	1	0.8	1	2.9
HOYENO	1	0.8	0.6	1.3

H9KV70	1	1.2	1.2	6.3
A5YKK6	1	1.3	1.4	1.1
C9IYD9	1	0.6	1	2.5
Q9Y6I3	1	0.6	0.7	0.8
095479	1	2.1	1.6	2.4
D6RIF8	1	1.1	1.1	0.7
Q6MZV4	1	1	1	1.2
H3BQ15	1	1	0.9	1.3
B4DME9	1	0.7	0.6	2.1
Q7Z6L0	1	1.9	0.6	0.4
HOYE28	1	1.3	1	1.5
HOYDR8	1	3.4	1.5	0.8
E7EMIO	1	0.8	1	2
Q12882	1	1.3	1.3	0.9
B7Z3K3	1	0.6	0.8	1.2
B0QYC6	1	1.8	1.1	0.7
B5MCE7	1	0.8	0.9	0.6
C9JVH3	1	4.9	2.1	1.3
E9PRB2	1	2	1.6	1
F5GZK2	1	1.2	0.4	0.2
000401	1	1.1	1.3	1.8
075937	1	1.3	1.4	1.4
P46734	1	1	0.9	2.5
P62280	1	0.8	1.8	1
Q13155	1	1.1	1.1	1.3
Q5TGZ0	1	1.5	1.1	0.8
Q9BZH6	1	1.8	1.7	0.7
Q9ULZ3	1	1.9	0.8	2.6
F8WCZ6	1	1	0.9	1.9
E9PQP6	1	1.1	1.2	0.7
B3KY94	1	1.3	1.4	0.9
H3BUS9	1	1	0.9	2.1
B7ZBH1	1	0.7	0.7	2.5
095164	1	3.2	1.1	1.3
J3QRP2	1	3	1.5	1.3
Q06210	1	0.8	0.8	0.8
B7Z3Q0	1	0.9	0.5	1.1
Q13158	1	0.9	1.2	1.3
HOYCE7	1	1.1	1.2	0.9
095470	1	1.7	1.8	1.5
Q8WUY1	1	1.1	1	0.9
Q8NA54	1	4	2.5	0.7
H3BNI9	1	1.2	1.1	0.8
E9PHT6	1	1.2	1.2	1.3
Q15437	1	1	1.3	1.2
F5H6E2	1	0.8	1.1	1.9
P36405	1	0.6	0.6	0.9

E7EM72	1	0.8	0.5	1.2
P07738	1	1	0.9	1
C9J3I4	1	0.8	1.3	1.6
000459	1	2	1.6	1
P42229	1	0.8	0.8	3
B2RDM2	1	1.3	2.1	2.1
Q16610	1	1.1	1.3	3.4
E7EPK6	1	0.8	0.9	1.7
E5RGR9	1	1	1.1	1.7
Q93052	1	1.1	1	2.4
BOLM41	1	0.8	1.1	0.9
D6R9I9	1	1.3	1.6	2.8
P10070	1	2.6	1	0.8
E7ETH0	1	0.6	2	2.8
B5MCT7	1	1.1	1.1	1.3
J3KRZ3	1	0.9	2.4	0.8