

Gene Name	Name of Proteins (UniProtID_SPECES, protein name, origin species, gene name)	Peptide Count (FP)	Peptide Count (iPS)	Fold Enrichment
LUTP6	Q9NYH9LUTP6_HUMAN U3 small nucleolar RNA-associated protein 6 homolog OS=Homo sapiens GN=LUTP6 PE=2 SV=2	9	0	Not in IgG control
HYDN	Q4G0P3HYDN_HUMAN Hydrocephalus-inducing protein homolog OS=Homo sapiens GN=HYDN PE=1 SV=3	5	0	Not in IgG control
CHSY3	Q70JA7CHSY3_HUMAN Chondroitin sulfate synthase 3 OS=Homo sapiens GN=CHSY3 PE=2 SV=3	5	0	Not in IgG control
No_GeneID	MOR2C6MOR2C6_HUMAN Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=1	4	0	Not in IgG control
DDX19B	Q9UMR2DDX19B_HUMAN ATP-dependent RNA helicase DDX19B OS=Homo sapiens GN=DDX19B PE=1 SV=1	3	0	Not in IgG control
CCDC148	Q8NFR7CC148_HUMAN Colled-coil domain-containing protein 148 OS=Homo sapiens GN=CCDC148 PE=1 SV=2	3	0	Not in IgG control
TBC1D8	O95759TBCD8_HUMAN TBC1 domain family member 8 OS=Homo sapiens GN=TBC1D8 PE=1 SV=3	3	0	Not in IgG control
ANKRD36	A6QL64IAN36A_HUMAN Ankyrin repeat domain-containing protein 36A OS=Homo sapiens GN=ANKRD36 PE=2 SV=3	3	0	Not in IgG control
MUC5AC	P98088MUC5A_HUMAN Mucin-5AC OS=Homo sapiens GN=MUC5AC PE=1 SV=4	3	0	Not in IgG control
AZGP1	P25311AZA2G_HUMAN Zinc-alpha-2-glycoprotein OS=Homo sapiens GN=AZGP1 PE=1 SV=2	2	0	Not in IgG control
LILRB4	Q8NHJ6LILRB4_HUMAN Leukocyte immunoglobulin-like receptor subfamily B member 4 OS=Homo sapiens GN=LILRB4 PE=1 SV=3	2	0	Not in IgG control
CECR2	Q98XF3CECR2_HUMAN Cat eye syndrome critical region protein 2 OS=Homo sapiens GN=CECR2 PE=1 SV=2	2	0	Not in IgG control
ETFDH	Q16134IETFH_HUMAN Electron transfer flavoprotein-ubiquinone oxidoreductase, mitochondrial OS=Homo sapiens GN=ETFDH PE=1 SV=2	2	0	Not in IgG control
CCDC88A	Q3V6T2GRDN_HUMAN Girdin OS=Homo sapiens GN=CCDC88A PE=1 SV=2	2	0	Not in IgG control
CLSTN2	Q9H4DDICSTN2_HUMAN Calsyntenin-2 OS=Homo sapiens GN=CLSTN2 PE=1 SV=2	2	0	Not in IgG control
HSP90AA1	P07900HSP90A_HUMAN Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5	2	0	Not in IgG control
ASAP2	O43150IASAP2_HUMAN Arf-GAP with SH3 domain, ANK repeat and PH domain-containing protein 2 OS=Homo sapiens GN=ASAP2 PE=1 SV=3	2	0	Not in IgG control
SYNRG	Q9UM22SYNRG_HUMAN Synergin gamma OS=Homo sapiens GN=SYNRG PE=1 SV=2	2	0	Not in IgG control
ACTBL2	Q562R1ACTBL_HUMAN Beta-actin-like protein 2 OS=Homo sapiens GN=ACTBL2 PE=1 SV=2	2	0	Not in IgG control
ZNF671	Q8TAW3ZNF671_HUMAN Zinc finger protein 671 OS=Homo sapiens GN=ZNF671 PE=2 SV=2	2	0	Not in IgG control
TRIM13	O60858TRIM13_HUMAN E3 ubiquitin-protein ligase TRIM13 OS=Homo sapiens GN=TRIM13 PE=1 SV=2	2	0	Not in IgG control
STK25	O00506STK25_HUMAN Serine/threonine-protein kinase 25 OS=Homo sapiens GN=STK25 PE=1 SV=1	2	0	Not in IgG control
TEX11	Q8YF3ITEX11_HUMAN Testis-expressed sequence 11 protein OS=Homo sapiens GN=TEX11 PE=1 SV=3	2	0	Not in IgG control
CRH	P06850ICRF_HUMAN Corticotropin-releasing hormone receptor type 1 OS=Homo sapiens GN=CRH PE=1 SV=1	2	0	Not in IgG control
RAVER1	Q8Y67IRAVR1_HUMAN Ribonucleoprotein PTB-binding 1 OS=Homo sapiens GN=RAVER1 PE=1 SV=1	2	0	Not in IgG control
DDX58	O95786DDX58_HUMAN Probable ATP-dependent RNA helicase DDX58 OS=Homo sapiens GN=DDX58 PE=1 SV=2	2	0	Not in IgG control
F5	P12259FAS_HUMAN Coagulation factor V OS=Homo sapiens GN=F5 PE=1 SV=4	1	0	Not in IgG control
GINS3	Q9BRX5GINS3_HUMAN DNA replication complex GINS protein PSF3 OS=Homo sapiens GN=GINS3 PE=1 SV=1	1	0	Not in IgG control
ANXA1	P04083IANXA1_HUMAN Annexin A1 OS=Homo sapiens GN=ANXA1 PE=1 SV=2	1	0	Not in IgG control
USF3	Q680E3IUSF3_HUMAN Basic helix-loop-helix domain-containing protein USF3 OS=Homo sapiens GN=USF3 PE=1 SV=3	1	0	Not in IgG control
CLCF1	Q9UBD9CLCF1_HUMAN Cardiotrophin-like cytokine factor 1 OS=Homo sapiens GN=CLCF1 PE=1 SV=1	1	0	Not in IgG control
LENG9	Q96B70LENG9_HUMAN Leukocyte receptor cluster member 9 OS=Homo sapiens GN=LENG9 PE=2 SV=2	1	0	Not in IgG control
WDR60	Q8WVS4WDR60_HUMAN WD repeat-containing protein 60 OS=Homo sapiens GN=WDR60 PE=1 SV=3	1	0	Not in IgG control
C9	P02749C09_HUMAN Complement component C9 OS=Homo sapiens GN=C9 PE=1 SV=2	1	0	Not in IgG control
GTF2H1	P32780ITF2H1_HUMAN General transcription factor IIH subunit 1 OS=Homo sapiens GN=GTF2H1 PE=1 SV=1	1	0	Not in IgG control
FKBP2	P26885FKBP2_HUMAN Peptidyl-prolyl cis-trans isomerase FKBP2 OS=Homo sapiens GN=FKBP2 PE=1 SV=2	1	0	Not in IgG control
PDZK1P1	ABMUH7PDZ1P_HUMAN Putative PDZ domain-containing protein PDZK1P1 OS=Homo sapiens GN=PDZK1P1 PE=5 SV=2	1	0	Not in IgG control
ANKRD20A12P	Q8NF67IA2012_HUMAN Putative ankyrin repeat domain-containing protein 20A12 pseudogene OS=Homo sapiens GN=ANKRD20A12P PE=5 SV=3	1	0	Not in IgG control
FBXO18	Q8NF20FBH1_HUMAN F-box DNA helicase 1 OS=Homo sapiens GN=FBXO18 PE=1 SV=2	1	0	Not in IgG control
RBM34	P42696RBM34_HUMAN RNA-binding protein 34 OS=Homo sapiens GN=RBM34 PE=1 SV=2	1	0	Not in IgG control
KLK14	Q9P0G3KLK14_HUMAN Kallikrein-14 OS=Homo sapiens GN=KLK14 PE=1 SV=2	1	0	Not in IgG control
HNRNPD	Q14103HNRNP_HUMAN Heterogeneous nuclear ribonucleoprotein D0 OS=Homo sapiens GN=HNRNPD PE=1 SV=1	1	0	Not in IgG control
NLE1	Q9NVX2NLE1_HUMAN Notchless protein homolog 1 OS=Homo sapiens GN=NLE1 PE=1 SV=4	1	0	Not in IgG control
HSPA7	P48741HSP77_HUMAN Putative heat shock 70 kDa protein 7 OS=Homo sapiens GN=HSPA7 PE=5 SV=2	1	0	Not in IgG control
CASP14	P31944ICASP14_HUMAN Caspase-14 OS=Homo sapiens GN=CASP14 PE=1 SV=2	1	0	Not in IgG control
ZSWIM8	A7E2V4IZSWIM8_HUMAN Zinc finger SWIM domain-containing protein 8 OS=Homo sapiens GN=ZSWIM8 PE=1 SV=1	1	0	Not in IgG control
RASSF10	A6NK89RASSF10_HUMAN Ras association domain-containing protein 10 OS=Homo sapiens GN=RASSF10 PE=2 SV=3	1	0	Not in IgG control
IGHV3-72	AOA0B4J1Y9IA0B4J1Y9_HUMAN Protein IGHV3-72 (Fragment) OS=Homo sapiens GN=IGHV3-72 PE=1 SV=1	1	0	Not in IgG control
HRASLS2	Q9NWW9HRSL2_HUMAN HRAS-like suppressor 2 OS=Homo sapiens GN=HRASLS2 PE=1 SV=1	1	0	Not in IgG control
PGBD2	Q6P38IPGBD2_HUMAN PiggyBac transposable element-derived protein 2 OS=Homo sapiens GN=PGBD2 PE=2 SV=1	1	0	Not in IgG control
INO80	Q9ULG1INO80_HUMAN DNA helicase INO80 OS=Homo sapiens GN=INO80 PE=1 SV=2	1	0	Not in IgG control
CREBRF	Q8URIGCREBRF_HUMAN CREB3 regulatory factor OS=Homo sapiens GN=CREBRF PE=1 SV=2	1	0	Not in IgG control
RPS25	P62851RPS25_HUMAN 40S ribosomal protein S25 OS=Homo sapiens GN=RPS25 PE=1 SV=1	1	0	Not in IgG control
MGAT1	P26572MGAT1_HUMAN Alpha-1,3-mannosyl-glycoprotein 2-beta-N-acetylglucosaminyltransferase OS=Homo sapiens GN=MGAT1 PE=1 SV=2	1	0	Not in IgG control
STAU2	Q9NUL3STAU2_HUMAN Double-stranded RNA-binding protein Staufen homolog 2 OS=Homo sapiens GN=STAU2 PE=1 SV=1	14	3	4.67
C3	P01024IC03_HUMAN Complement C3 OS=Homo sapiens GN=C3 PE=1 SV=2	22	7	3.14
RAI1	Q7254IRAI1_HUMAN Retinoic acid-induced protein 1 OS=Homo sapiens GN=RAI1 PE=1 SV=2	5	2	2.50
PLEC	Q15149PLEC_HUMAN Plectin OS=Homo sapiens GN=PLEC PE=1 SV=3	9	4	2.25
VIPAS39	Q9H9C1IPSE39_HUMAN Spermatogenesis-defective protein 39 homolog OS=Homo sapiens GN=VIPAS39 PE=1 SV=1	2	1	2.00
POLDIP3	Q9BY77POLDIP3_HUMAN Polymerase delta-interacting protein 3 OS=Homo sapiens GN=POLDIP3 PE=1 SV=2	2	1	2.00
TUBB2A	Q13885ITB2A_HUMAN Tubulin beta-2A chain OS=Homo sapiens GN=TUBB2A PE=1 SV=1	2	1	2.00
TFG	Q92734ITFG_HUMAN Protein TFG OS=Homo sapiens GN=TFG PE=1 SV=2	5	3	1.67
LPCAT1	Q8NF37IPCAT1_HUMAN Lysophosphatidylcholine acyltransferase 1 OS=Homo sapiens GN=LPCAT1 PE=1 SV=2	5	3	1.67
DDX28	Q9NUL7IDDX28_HUMAN Probable ATP-dependent RNA helicase DDX28 OS=Homo sapiens GN=DDX28 PE=1 SV=2	3	2	1.50
SURF4	O15260ISURF4_HUMAN Surflet locus protein 4 OS=Homo sapiens GN=SURF4 PE=1 SV=3	3	2	1.50
ACSF3	Q4G176ACSF3_HUMAN Acyl-CoA synthetase family member 3, mitochondrial OS=Homo sapiens GN=ACSF3 PE=1 SV=3	3	2	1.50
PSMD5	Q16401IPSD5_HUMAN 26S proteasome non-ATPase regulatory subunit 5 OS=Homo sapiens GN=PSMD5 PE=1 SV=3	3	2	1.50
CAP2	P40123ICAP2_HUMAN Adenylyl cyclase-associated protein 2 OS=Homo sapiens GN=CAP2 PE=1 SV=1	4	3	1.33
RUVBL2	Q9230RUVB2_HUMAN RuvB-like 2 OS=Homo sapiens GN=RUVBL2 PE=1 SV=3	10	8	1.25
ABCD3	P28288IABCD3_HUMAN ATP-binding cassette sub-family D member 3 OS=Homo sapiens GN=ABCD3 PE=1 SV=1	7	6	1.17
HNRNPH1	P31943HNRH1_HUMAN Heterogeneous nuclear ribonucleoprotein H OS=Homo sapiens GN=HNRNPH1 PE=1 SV=4	10	9	1.11
DCD	P81605IDCD_HUMAN Dermicin OS=Homo sapiens GN=DCD PE=1 SV=2	2	2	1.00
KPNA1	P52294IMAS_HUMAN Importin subunit alpha-5 OS=Homo sapiens GN=KPNA1 PE=1 SV=3	1	1	1.00
TMEM266	Q2M3C6ITM266_HUMAN Transmembrane protein 266 OS=Homo sapiens GN=TMEM266 PE=2 SV=2	1	1	1.00
ZC3H15	Q8WU90I0ZC3H15_HUMAN Zinc finger CCH domain-containing protein 15 OS=Homo sapiens GN=ZC3H15 PE=1 SV=1	2	2	1.00
SLC16A1	P53985IMOT1_HUMAN Monocarboxylate transporter 1 OS=Homo sapiens GN=SLC16A1 PE=1 SV=3	1	1	1.00
SHC1	P29353ISHC1_HUMAN SHC-transforming protein 1 OS=Homo sapiens GN=SHC1 PE=1 SV=4	1	1	1.00
PIP	P12273PIP_HUMAN Prolectin-inducible protein OS=Homo sapiens GN=PIP PE=1 SV=1	1	1	1.00
GAPDH	P04406IG3P_HUMAN Glycerinaldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	3	3	1.00
SERPIND1	P05546IHP2_HUMAN Heparin cofactor 2 OS=Homo sapiens GN=SERPIND1 PE=1 SV=3	1	1	1.00
PSMD12	O00232IPSD12_HUMAN 26S proteasome non-ATPase regulatory subunit 12 OS=Homo sapiens GN=PSMD12 PE=1 SV=3	2	2	1.00
S100A8	P05109S10A8_HUMAN Protein S100-A8 OS=Homo sapiens GN=S100A8 PE=1 SV=1	1	1	1.00
REXO4	Q9G2R2REXO4_HUMAN RNA exonuclease 4 OS=Homo sapiens GN=REXO4 PE=1 SV=2	2	2	1.00
LAMB1	P07942ILAMB1_HUMAN Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	1	1	1.00
DDX49	Q9Y6V7DDX49_HUMAN Probable ATP-dependent RNA helicase DDX49 OS=Homo sapiens GN=DDX49 PE=1 SV=1	1	1	1.00
Cys-GrB-Fc-RSP01	ADA024R160IA0A024R160_HUMAN Cys-GrB-Fc-RSP01 OS=Homo sapiens GN=Cys-GrB-Fc-RSP01 PE=3 SV=1	3	3	1.00
NEFM	P07197INFM_HUMAN Neurofilament medium polypeptide OS=Homo sapiens GN=NEFM PE=1 SV=3	4	4	1.00

RETSAT	Q6NUM9RETST_HUMAN All-trans-retinol 13,14-reductase OS=Homo sapiens GN=RETSAT PE=1 SV=2	4	4	1.00
IGLL5	B9A064IGLL5_HUMAN Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2	1	1	1.00
STT3B	Q8TCJ2ISTT3B_HUMAN Dolchyl-diphosphooligosaccharide--protein glycosyltransferase subunit STT3B OS=Homo sapiens GN=STT3B PE=1 SV=1	2	2	1.00
CDK17	Q00537CDK17_HUMAN Cyclin-dependent kinase 17 OS=Homo sapiens GN=CDK17 PE=1 SV=2	2	2	1.00
PPP2R2A	P63151I2ABA_HUMAN Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B alpha isoform OS=Homo sapiens GN=PPP2R2A PE=1 SV=1	1	1	1.00
GD2	P50395IGDIB_HUMAN Rab GDP dissociation inhibitor beta OS=Homo sapiens GN=GD2 PE=1 SV=2	1	1	1.00
SLC27A2	O14975IS27A2_HUMAN Very long-chain acyl-CoA synthetase OS=Homo sapiens GN=SLC27A2 PE=1 SV=2	4	4	1.00
ENO1	P06733ENO1_HUMAN Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	8	8	1.00
UBA52	P62987RL40_HUMAN Ubiquitin-60S ribosomal protein L40 OS=Homo sapiens GN=UBA52 PE=1 SV=2	2	2	1.00
TUBB3	Q13509ITB83_HUMAN Tubulin beta-3 chain OS=Homo sapiens GN=TUBB3 PE=1 SV=2	6	6	1.00
TUBA1B	P68363ITBA1B_HUMAN Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1	20	21	0.95
TUBB	P07437ITB85_HUMAN Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2	24	26	0.92
UGDH	O60701IUGDH_HUMAN UDP-glucose 6-dehydrogenase OS=Homo sapiens GN=UGDH PE=1 SV=1	22	24	0.92
HNRNPK	P61978HNRNPK_HUMAN Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK PE=1 SV=1	19	21	0.90
YME1L1	Q96T2IYME1L1_HUMAN ATP-dependent zinc metalloprotease YME1L1 OS=Homo sapiens GN=YME1L1 PE=1 SV=2	5	6	0.83
SQLE	Q14534IERG1_HUMAN Squalene monooxygenase OS=Homo sapiens GN=SQLE PE=1 SV=3	5	6	0.83
CCT4	P50991ITCPD_HUMAN T-complex protein 1 subunit delta OS=Homo sapiens GN=CCT4 PE=1 SV=4	19	23	0.83
SNTB1	Q13884ISNTB1_HUMAN Beta-1-syntrophin OS=Homo sapiens GN=SNTB1 PE=1 SV=3	9	11	0.82
TIN	Q8W242ITITN_HUMAN Titin OS=Homo sapiens GN=TIN PE=1 SV=4	35	47	0.81
GLUD1	P00367IDHE3_HUMAN Glutamate dehydrogenase 1, mitochondrial OS=Homo sapiens GN=GLUD1 PE=1 SV=2	30	33	0.81
MARCKS	P29966IMARCS_HUMAN Myristoylated alanine-rich C-kinase substrate OS=Homo sapiens GN=MARCKS PE=1 SV=4	4	5	0.80
DHCR24	Q15392IDHC24_HUMAN Delta(24)-sterol reductase OS=Homo sapiens GN=DHCR24 PE=1 SV=2	4	5	0.80
NOP58	Q9Y2X3INOP58_HUMAN Nucleolar protein 58 OS=Homo sapiens GN=NOP58 PE=1 SV=1	12	15	0.80
ABCB10	Q9NRK6IACB10_HUMAN ATP-binding cassette sub-family B member 10, mitochondrial OS=Homo sapiens GN=ABCB10 PE=1 SV=2	4	5	0.80
CYP27A1	Q02318ICP27A1_HUMAN Sterol 26-hydroxylase, mitochondrial OS=Homo sapiens GN=CYP27A1 PE=1 SV=1	8	10	0.80
ATP5B	P06576IATP5B_HUMAN ATP synthase subunit beta, mitochondrial OS=Homo sapiens GN=ATP5B PE=1 SV=3	28	35	0.80
TUBA1C	Q9BQ83IAT1C_HUMAN Tubulin alpha-1C chain OS=Homo sapiens GN=TUBA1C PE=1 SV=1	4	5	0.80
RSL1D1	O76021IRL1D1_HUMAN Ribosomal L1 domain-containing protein 1 OS=Homo sapiens GN=RSL1D1 PE=1 SV=3	7	9	0.78
UGP2	Q16851IUGPA_HUMAN UTP--glucose-1-phosphate uridylyltransferase OS=Homo sapiens GN=UGP2 PE=1 SV=5	10	13	0.77
CYP51A1	Q16850CYP51A1_HUMAN Lanosterol 14-alpha demethylase OS=Homo sapiens GN=CYP51A1 PE=1 SV=3	10	13	0.77
ATP5A1	P25705IATPA_HUMAN ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 PE=1 SV=1	20	26	0.77
NONO	Q15233INONO_HUMAN Non-POU domain-containing octamer-binding protein OS=Homo sapiens GN=NONO PE=1 SV=4	15	20	0.75
TUBB4B	P68371ITB4B_HUMAN Tubulin beta-4B chain OS=Homo sapiens GN=TUBB4B PE=1 SV=1	6	8	0.75
PTBP1	P26599PTBP1_HUMAN Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1 PE=1 SV=1	9	12	0.75
ATP6V1B2	P21281IVATB2_HUMAN V-type proton ATPase subunit B, brain isoform OS=Homo sapiens GN=ATP6V1B2 PE=1 SV=3	3	4	0.75
KPNA6	O60684IIMA7_HUMAN Importin subunit alpha-7 OS=Homo sapiens GN=KPNA6 PE=1 SV=1	3	4	0.75
LBR	Q14739ILBR_HUMAN Lamin-B receptor OS=Homo sapiens GN=LBR PE=1 SV=2	3	4	0.75
GTF3C5	Q9Y5Q8ITF3C5_HUMAN General transcription factor 3C polypeptide 5 OS=Homo sapiens GN=GTF3C5 PE=1 SV=2	3	4	0.75
DDX5	P17844IDDX5_HUMAN Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 PE=1 SV=1	14	19	0.74
PKM	P14618IKPYM_HUMAN Pyruvate kinase PKM OS=Homo sapiens GN=PKM PE=1 SV=4	25	34	0.74
ALDH1B1	P30837AL1B1_HUMAN Aldehyde dehydrogenase X, mitochondrial OS=Homo sapiens GN=ALDH1B1 PE=1 SV=3	14	20	0.70
STT3A	P46977ISTT3A_HUMAN Dolchyl-diphosphooligosaccharide--protein glycosyltransferase subunit STT3A OS=Homo sapiens GN=STT3A PE=1 SV=2	7	10	0.70
GTPBP6	O43824IGTPB6_HUMAN Putative GTP-binding protein 6 OS=Homo sapiens GN=GTPBP6 PE=2 SV=3	2	3	0.67
DDX56	Q9N9Y3IDDX56_HUMAN Probable ATP-dependent RNA helicase DDX56 OS=Homo sapiens GN=DDX56 PE=1 SV=1	2	3	0.67
UMPS	P11172IUMPS_HUMAN Uridine 5'-monophosphate synthase OS=Homo sapiens GN=UMPS PE=1 SV=1	2	3	0.67
ZNF207	O43670IZNF207_HUMAN BUB3-interacting and GLEBS motif-containing protein ZNF207 OS=Homo sapiens GN=ZNF207 PE=1 SV=1	2	3	0.67
ACAD9	Q9H845IACAD9_HUMAN Acyl-CoA dehydrogenase family member 9, mitochondrial OS=Homo sapiens GN=ACAD9 PE=1 SV=1	4	6	0.67
SLC2A1	P11166GTR1_HUMAN Solute carrier family 2, facilitated glucose transporter member 1 OS=Homo sapiens GN=SLC2A1 PE=1 SV=2	4	6	0.67
ALDH3A2	P51648IAL3A2_HUMAN Fatty aldehyde dehydrogenase OS=Homo sapiens GN=ALDH3A2 PE=1 SV=1	2	3	0.67
DLST	P36957IOD02_HUMAN Dihydrodipolysine-residue succinyltransferase component of 2-oxoglutarate dehydrogenase complex, mitochondrial OS=Homo sapiens GN=DLST PE=1 SV=4	6	9	0.67
SLC39A14	Q15043IS39AE_HUMAN Zinc transporter ZIP14 OS=Homo sapiens GN=SLC39A14 PE=1 SV=3	2	3	0.67
RNMT	O43148IMCES_HUMAN mRNA cap guanine-N7 methyltransferase OS=Homo sapiens GN=RNMT PE=1 SV=1	4	6	0.67
ZC3H12C	Q9C00DIZC12C_HUMAN Probable ribonuclease ZC3H12C OS=Homo sapiens GN=ZC3H12C PE=1 SV=2	2	3	0.67
EEF1A1	P68104IEF1A1_HUMAN Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1	4	6	0.67
SQSTM1	Q13501ISQSTM_HUMAN Sequestosome-1 OS=Homo sapiens GN=SQSTM1 PE=1 SV=1	2	3	0.67
KNOP1	Q1ED39KNOP1_HUMAN Lysine-rich nucleolar protein 1 OS=Homo sapiens GN=KNOP1 PE=1 SV=1	2	3	0.67
CCT6A	P40227ITCP2_HUMAN T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3	13	20	0.65
P4HB	P07237PDA1_HUMAN Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3	20	31	0.65
PDIA3	P30101IPDIA3_HUMAN Protein disulfide-isomerase A3 OS=Homo sapiens GN=PDIA3 PE=1 SV=4	19	30	0.63
GORASP2	Q9H8Y8IGORS2_HUMAN Golgi reassembly-stacking protein 2 OS=Homo sapiens GN=GORASP2 PE=1 SV=3	5	8	0.63
ATP1B1	P05026IAT1B1_HUMAN Sodium/potassium-transporting ATPase subunit beta-1 OS=Homo sapiens GN=ATP1B1 PE=1 SV=1	5	8	0.63
PHGDH	O43175ISERA_HUMAN D-3-phosphoglycerate dehydrogenase OS=Homo sapiens GN=PHGDH PE=1 SV=4	10	16	0.63
EIF253	P41091IF2G_HUMAN Eukaryotic translation initiation factor 2 subunit 3 OS=Homo sapiens GN=EIF253 PE=1 SV=3	5	8	0.63
PDIA5	Q14554IPDIA5_HUMAN Protein disulfide-isomerase A5 OS=Homo sapiens GN=PDIA5 PE=1 SV=1	8	13	0.62
ANXA11	P50995IANX11_HUMAN Annexin A11 OS=Homo sapiens GN=ANXA11 PE=1 SV=1	3	5	0.60
BSG	P35613IBAS1_HUMAN Basigin OS=Homo sapiens GN=BSG PE=1 SV=2	6	10	0.60
PDIA6	Q15084IPDIA6_HUMAN Protein disulfide-isomerase A6 OS=Homo sapiens GN=PDIA6 PE=1 SV=1	3	5	0.60
NOP56	O00567NOP56_HUMAN Nucleolar protein 56 OS=Homo sapiens GN=NOP56 PE=1 SV=4	3	5	0.60
CCT8	P50990ITCP3_HUMAN T-complex protein 1 subunit theta OS=Homo sapiens GN=CCT8 PE=1 SV=4	16	27	0.59
P4HA2	O15460IP4HA2_HUMAN Prolyl 4-hydroxylase subunit alpha-2 OS=Homo sapiens GN=P4HA2 PE=1 SV=1	7	12	0.58
CCT2	P78371ITCPB_HUMAN T-complex protein 1 subunit beta OS=Homo sapiens GN=CCT2 PE=1 SV=4	14	24	0.58
HNRNPL	P14866HNRPL_HUMAN Heterogeneous nuclear ribonucleoprotein L OS=Homo sapiens GN=HNRNPL PE=1 SV=2	8	14	0.57
CCT7	Q99832ITCPH_HUMAN T-complex protein 1 subunit eta OS=Homo sapiens GN=CCT7 PE=1 SV=2	13	23	0.57
MCCC2	Q9HCC0MCCB_HUMAN Methylcrotonyl-CoA carboxylase beta chain, mitochondrial OS=Homo sapiens GN=MCCC2 PE=1 SV=1	5	9	0.56
DARS	P14868ISYDC_HUMAN Aspartate--tRNA ligase, cytoplasmic OS=Homo sapiens GN=DARS PE=1 SV=2	13	24	0.54
ALDH7A1	P49419IAL7A1_HUMAN Alpha-aminoacidic semialdehyde dehydrogenase OS=Homo sapiens GN=ALDH7A1 PE=1 SV=5	7	13	0.54
PSMC1	P62191IPRS4_HUMAN 26S protease regulatory subunit 4 OS=Homo sapiens GN=PSMC1 PE=1 SV=1	7	13	0.54
STAU1	O95793ISTAU1_HUMAN Double-stranded RNA-binding protein Staufen homolog 1 OS=Homo sapiens GN=STAU1 PE=1 SV=2	7	13	0.54
P4HA1	P13674IP4HA1_HUMAN Prolyl 4-hydroxylase subunit alpha-1 OS=Homo sapiens GN=P4HA1 PE=1 SV=2	10	19	0.53
SHMT2	P34897ILYLM_HUMAN Serine hydroxymethyltransferase, mitochondrial OS=Homo sapiens GN=SHMT2 PE=1 SV=3	12	23	0.52
TCP1	P17987ITCPA_HUMAN T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1	14	27	0.52
HSPD1	P10809ICH60_HUMAN 60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2	28	54	0.52
CCT3	P49368ITCPG_HUMAN T-complex protein 1 subunit gamma OS=Homo sapiens GN=CCT3 PE=1 SV=4	12	24	0.50
SMARCD1	Q96GM5ISMRD1_HUMAN SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily D member 1 OS=Homo sapiens GN=SMARCD1 PE=1 SV=2	1	2	0.50
YES1	P07947YES_HUMAN Tyrosine-protein kinase Yes OS=Homo sapiens GN=YES1 PE=1 SV=3	1	2	0.50
ALDH1A2	O94788IAL1A2_HUMAN Retinal dehydrogenase 2 OS=Homo sapiens GN=ALDH1A2 PE=1 SV=3	1	2	0.50
ATP6V1H	Q9U12IVATH_HUMAN V-type proton ATPase subunit H OS=Homo sapiens GN=ATP6V1H PE=1 SV=1	1	2	0.50
UAP1	Q16222IUAP1_HUMAN UDP-N-acetylhexosamine pyrophosphorylase OS=Homo sapiens GN=UAP1 PE=1 SV=3	2	4	0.50

FAAH2	Q6MR7FAAH2_HUMAN Fatty-acid amide hydrolase 2 OS=Homo sapiens GN=FAAH2 PE=2 SV=1	1	2	0.50
ABAT	P80404IGABT_HUMAN 4-aminobutyrate aminotransferase, mitochondrial OS=Homo sapiens GN=ABAT PE=1 SV=3	3	6	0.50
HACD3	Q8P035HACD3_HUMAN Very-long-chain (3R)-3-hydroxyacyl-CoA dehydratase 3 OS=Homo sapiens GN=HACD3 PE=1 SV=2	1	2	0.50
TXNRD1	Q16881TXNRD1_HUMAN Thioredoxin reductase 1, cytoplasmic OS=Homo sapiens GN=TXNRD1 PE=1 SV=3	1	2	0.50
FGB	P02675FIBB_HUMAN Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1 SV=2	3	6	0.50
TUBAL3	A6NHL2TUBAL3_HUMAN Tubulin alpha chain-like 3 OS=Homo sapiens GN=TUBAL3 PE=1 SV=2	1	2	0.50
TMPO	P42166LAP2A_HUMAN Lamina-associated polypeptide 2, isoform alpha OS=Homo sapiens GN=TMPO PE=1 SV=2	4	8	0.50
ADSL	P30566PUR8_HUMAN Adenylosuccinate lyase OS=Homo sapiens GN=ADSL PE=1 SV=2	2	4	0.50
PFKFB4	Q16877IF264_HUMAN 6-phosphofructo-2-kinase/fructose-2,6-bisphosphatase 4 OS=Homo sapiens GN=PFKFB4 PE=2 SV=6	1	2	0.50
GLS	O94925GLSK_HUMAN Glutaminase kidney isoform, mitochondrial OS=Homo sapiens GN=GLS PE=1 SV=1	6	12	0.50
OXSR1	O95747OXSR1_HUMAN Serine/threonine-protein kinase OSR1 OS=Homo sapiens GN=OXSR1 PE=1 SV=1	2	4	0.50
RPN2	P04844RPN2_HUMAN Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2 OS=Homo sapiens GN=RPN2 PE=1 SV=3	10	20	0.50
AGFG2	O95081AGFG2_HUMAN Arf-GAP domain and FG repeat-containing protein 2 OS=Homo sapiens GN=AGFG2 PE=1 SV=2	1	2	0.50
TBC1D24	Q9ULP9ITBC24_HUMAN TBC1 domain family member 24 OS=Homo sapiens GN=TBC1D24 PE=1 SV=2	1	2	0.50
CLPB	Q9H078CLPB_HUMAN Caseolytic peptidase B protein homolog OS=Homo sapiens GN=CLPB PE=1 SV=1	3	6	0.50
CDS2	O95674CDS2_HUMAN Phosphatidate cytidyltransferase 2 OS=Homo sapiens GN=CDS2 PE=1 SV=1	1	2	0.50
TRAP1	Q12931TRAP1_HUMAN Heat shock protein 75 kDa, mitochondrial OS=Homo sapiens GN=TRAP1 PE=1 SV=3	2	4	0.50
DYNC1L1	Q9Y6G9IDC1L1_HUMAN Cytoplasmic dynein 1 light intermediate chain 1 OS=Homo sapiens GN=DYNC1L1 PE=1 SV=3	2	4	0.50
CES1	P23141EST1_HUMAN Liver carboxylesterase 1 OS=Homo sapiens GN=CES1 PE=1 SV=2	14	29	0.48
ALDH2	P05091ALDH2_HUMAN Aldehyde dehydrogenase, mitochondrial OS=Homo sapiens GN=ALDH2 PE=1 SV=2	11	23	0.48
ACSL5	Q9ULCSACLS5_HUMAN Long-chain-fatty-acid--CoA ligase 5 OS=Homo sapiens GN=ACSL5 PE=1 SV=1	7	15	0.47
RRBP1	Q9P2E9IRBP1_HUMAN Ribosome-binding protein 1 OS=Homo sapiens GN=RRBP1 PE=1 SV=4	6	13	0.46
HARS	P12081ISYHC_HUMAN Histidine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=HARS PE=1 SV=2	5	11	0.45
FARSA	Q9Y285ISYFA_HUMAN Phenylalanine--tRNA ligase alpha subunit OS=Homo sapiens GN=FARSA PE=1 SV=3	4	9	0.44
CAT	P04040ICATA_HUMAN Catalase OS=Homo sapiens GN=CAT PE=1 SV=3	8	18	0.44
SRC	P12931SRC_HUMAN Proto-oncogene tyrosine-protein kinase Src OS=Homo sapiens GN=SRC PE=1 SV=3	4	9	0.44
TKFC	Q3LXA3TKFC_HUMAN Triokinase/FMN cyclase OS=Homo sapiens GN=TKFC PE=1 SV=2	7	16	0.44
CCTS	P48643ITCPE_HUMAN T-complex protein 1 subunit epsilon OS=Homo sapiens GN=CCTS PE=1 SV=1	10	23	0.43
EIF3D	O15371EIF3D_HUMAN Eukaryotic translation initiation factor 3 subunit D OS=Homo sapiens GN=EIF3D PE=1 SV=1	3	7	0.43
AGT	P01019IANGT_HUMAN Angiotensinogen OS=Homo sapiens GN=AGT PE=1 SV=1	3	7	0.43
EIF3L	Q9Y262EIF3L_HUMAN Eukaryotic translation initiation factor 3 subunit L OS=Homo sapiens GN=EIF3L PE=1 SV=1	6	14	0.43
CPNE3	O75131CPNE3_HUMAN Copine-3 OS=Homo sapiens GN=CPNE3 PE=1 SV=1	5	12	0.42
ALB	P02768IALBU_HUMAN Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	12	29	0.41
DNPEP	Q9ULADINPEP_HUMAN Aspartyl aminopeptidase OS=Homo sapiens GN=DNPEP PE=1 SV=1	2	5	0.40
PKC2	Q16822IPCKGM_HUMAN Phosphoenolpyruvate carboxykinase [GTP], mitochondrial OS=Homo sapiens GN=PKC2 PE=1 SV=3	10	25	0.40
ABCE1	P61221ABCE1_HUMAN ATP-binding cassette sub-family E member 1 OS=Homo sapiens GN=ABCE1 PE=1 SV=1	2	5	0.40
PRPF19	Q9UM45IPRP19_HUMAN Pre-mRNA-processing factor 19 OS=Homo sapiens GN=PRPF19 PE=1 SV=1	6	15	0.40
PRPF4	O43172IPRP4_HUMAN U4/U6 small nuclear ribonucleoprotein Ptp4 OS=Homo sapiens GN=PRPF4 PE=1 SV=2	2	5	0.40
TBRG4	Q96920TBRG4_HUMAN Protein TBRG4 OS=Homo sapiens GN=TBRG4 PE=1 SV=1	2	5	0.40
ASPSR1	Q9BZE9IASPC1_HUMAN Tether containing UBX domain for GLUT4 OS=Homo sapiens GN=ASPSR1 PE=1 SV=1	2	5	0.40
SYNE1	Q8NF91ISYNE1_HUMAN Nesprin-1 OS=Homo sapiens GN=SYNE1 PE=1 SV=4	4	10	0.40
PCYOX1	Q9UH33PCYOX_HUMAN Prenylcysteine oxidase 1 OS=Homo sapiens GN=PCYOX1 PE=1 SV=3	6	15	0.40
ATIC	P31939IPUR9_HUMAN Bifunctional purine biosynthesis protein PURH OS=Homo sapiens GN=ATIC PE=1 SV=3	13	33	0.39
PSMD3	O43242PSMD3_HUMAN 26S proteasome non-ATPase regulatory subunit 3 OS=Homo sapiens GN=PSMD3 PE=1 SV=2	7	18	0.39
HNRNPR	O43390HNRNPR_HUMAN Heterogeneous nuclear ribonucleoprotein R OS=Homo sapiens GN=HNRNPR PE=1 SV=1	5	13	0.38
CALR	P27797CALR_HUMAN Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1	10	26	0.38
GP2	Q8TD30ALAT2_HUMAN Alanine aminotransferase 2 OS=Homo sapiens GN=GP2 PE=1 SV=1	8	21	0.38
PDHX	O00330IDOPK_HUMAN Pyruvate dehydrogenase protein X component, mitochondrial OS=Homo sapiens GN=PDHX PE=1 SV=3	3	8	0.38
SF3A3	Q12874SF3A3_HUMAN Splicing factor 3A subunit 3 OS=Homo sapiens GN=SF3A3 PE=1 SV=1	3	8	0.38
WDR1	O75083WDR1_HUMAN WD repeat-containing protein 1 OS=Homo sapiens GN=WDR1 PE=1 SV=4	6	16	0.38
TOR1AIP1	Q5JTV8ITOP1_HUMAN Torsin-1A-interacting protein 1 OS=Homo sapiens GN=TOR1AIP1 PE=1 SV=2	3	8	0.38
CES2	O00748EST2_HUMAN Cocaine esterase OS=Homo sapiens GN=CES2 PE=1 SV=1	3	8	0.38
PAK2	Q13177IPAK2_HUMAN Serine/threonine-protein kinase PAK 2 OS=Homo sapiens GN=PAK2 PE=1 SV=3	3	8	0.38
EROT1A	Q96HE7ERO1A_HUMAN ERO1-like protein alpha OS=Homo sapiens GN=ERO1A PE=1 SV=2	7	19	0.37
PCLR	P30613KPYR_HUMAN Pyruvate kinase PCLR OS=Homo sapiens GN=PCLR PE=1 SV=2	5	14	0.36
GK	P32189GLPK_HUMAN Glycerol kinase OS=Homo sapiens GN=GK PE=1 SV=3	5	14	0.36
ALDH4A1	P30038IAL4A1_HUMAN Delta-1-pyrroline-5-carboxylate dehydrogenase, mitochondrial OS=Homo sapiens GN=ALDH4A1 PE=1 SV=3	6	17	0.35
LMAN1	P49257LMAN1_HUMAN Protein ERGIC-53 OS=Homo sapiens GN=LMAN1 PE=1 SV=2	7	20	0.35
HNF4A	P41235HNF4A_HUMAN Hepatocyte nuclear factor 4-alpha OS=Homo sapiens GN=HNF4A PE=1 SV=3	1	3	0.33
DLD	P09622DLDL_HUMAN Dihydrodipolyl dehydrogenase, mitochondrial OS=Homo sapiens GN=DLD PE=1 SV=2	3	9	0.33
GPI	P06744IGPL_HUMAN Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI PE=1 SV=4	7	21	0.33
HSPA5	P11021GRP78_HUMAN 78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2	4	12	0.33
PLIN2	Q99541PLIN2_HUMAN Perilipin-2 OS=Homo sapiens GN=PLIN2 PE=1 SV=2	1	3	0.33
KPNA3	O00505IMA4_HUMAN Importin subunit alpha-4 OS=Homo sapiens GN=KPNA3 PE=1 SV=2	1	3	0.33
VTN	P04004VTNC_HUMAN Vitronectin OS=Homo sapiens GN=VTN PE=1 SV=1	1	3	0.33
UTP18	Q9Y5J1UTP18_HUMAN U3 small nucleolar RNA-associated protein 18 homolog OS=Homo sapiens GN=UTP18 PE=1 SV=3	1	3	0.33
KPNA4	O00629IMA3_HUMAN Importin subunit alpha-3 OS=Homo sapiens GN=KPNA4 PE=1 SV=1	2	6	0.33
AGF1	P52594IAGF1_HUMAN Arf-GAP domain and FG repeat-containing protein 1 OS=Homo sapiens GN=AGF1 PE=1 SV=2	1	3	0.33
USP39	Q53G59SUNT2_HUMAN U4/U6.U5 tri-snRNP-associated protein 2 OS=Homo sapiens GN=USP39 PE=1 SV=2	1	3	0.33
SRP72	O76094SRP72_HUMAN Signal recognition particle subunit SRP72 OS=Homo sapiens GN=SRP72 PE=1 SV=3	1	3	0.33
ILVBL	A110L0LILVBL_HUMAN Acetolactate synthase-like protein OS=Homo sapiens GN=ILVBL PE=1 SV=2	2	6	0.33
LMNA	P02545LMNA_HUMAN Prelamin-A/C OS=Homo sapiens GN=LMNA PE=1 SV=1	9	29	0.31
KHDRBS1	Q07666KHDR1_HUMAN KH domain-containing, RNA-binding, signal transduction-associated protein 1 OS=Homo sapiens GN=KHDRBS1 PE=1 SV=1	4	13	0.31
GGPD	P11413IG6PD_HUMAN Glucose-6-phosphate 1-dehydrogenase OS=Homo sapiens GN=GGPD PE=1 SV=4	4	13	0.31
DDX6	P26196DDX6_HUMAN Probable ATP-dependent RNA helicase DDX6 OS=Homo sapiens GN=DDX6 PE=1 SV=2	3	10	0.30
PUF60	Q9UHX1PUF60_HUMAN Poly(U)-binding-splicing factor PUF60 OS=Homo sapiens GN=PUF60 PE=1 SV=1	3	10	0.30
UZAF2	P26368UZAF2_HUMAN Splicing factor UZAF 65 kDa subunit OS=Homo sapiens GN=UZAF2 PE=1 SV=4	2	7	0.29
ATCF	Q9NQ94A1CF_HUMAN APOBEC1 complementation factor OS=Homo sapiens GN=ATCF PE=1 SV=1	4	14	0.29
GPC3	P51654GPC3_HUMAN Glypican-3 OS=Homo sapiens GN=GPC3 PE=1 SV=1	2	7	0.29
SERPINA1	P01009A1AT_HUMAN Alpha-1-antitrypsin OS=Homo sapiens GN=SERPINA1 PE=1 SV=3	2	7	0.29
ALDH1A1	P00352IAL1A1_HUMAN Retinal dehydrogenase 1 OS=Homo sapiens GN=ALDH1A1 PE=1 SV=2	6	22	0.27
LAP3	P28838IAMPL_HUMAN Cytosol aminopeptidase OS=Homo sapiens GN=LAP3 PE=1 SV=3	6	22	0.27
NCLN	Q969V3INCLN_HUMAN Nicotin OS=Homo sapiens GN=NCLN PE=1 SV=2	3	11	0.27
AFP	P02771IFETA_HUMAN Alpha-fetoprotein OS=Homo sapiens GN=AFP PE=1 SV=1	6	23	0.26
TM9SF4	Q92544TM9SF4_HUMAN Transmembrane 9 superfamily member 4 OS=Homo sapiens GN=TM9SF4 PE=1 SV=2	1	4	0.25
ALDH6A1	Q02252IMMSA_HUMAN Methylmalonate-semialdehyde dehydrogenase [acylating], mitochondrial OS=Homo sapiens GN=ALDH6A1 PE=1 SV=2	4	16	0.25
RRP9	O43818IUP2_HUMAN U3 small nucleolar RNA-interacting protein 2 OS=Homo sapiens GN=RRP9 PE=1 SV=1	2	8	0.25
ETFDH	Q16134IETFD_HUMAN Electron transfer flavoprotein-ubiquinone oxidoreductase, mitochondrial OS=Homo sapiens GN=ETFDH PE=1 SV=2	1	4	0.25

DPYSL2	Q16555DPYSL2_HUMAN Dihydropyrimidinase-related protein 2 OS=Homo sapiens GN=DPYSL2 PE=1 SV=1	5	20	0.25
SAMM50	Q9Y512SAM50_HUMAN Sorting and assembly machinery component 50 homolog OS=Homo sapiens GN=SAMM50 PE=1 SV=3	3	12	0.25
RUVBL1	Q9Y265RUVBL1_HUMAN RuvB-like 1 OS=Homo sapiens GN=RUVBL1 PE=1 SV=1	4	16	0.25
DNAJC3	Q13217DNAJC3_HUMAN DnaJ homolog subfamily C member 3 OS=Homo sapiens GN=DNAJC3 PE=1 SV=1	2	8	0.25
EARS2	Q5JPH6EARS2_HUMAN Probable glutamate-tRNA ligase, mitochondrial OS=Homo sapiens GN=EARS2 PE=1 SV=2	1	4	0.25
CAP1	Q01518CAP1_HUMAN Adenylyl cyclase-associated protein 1 OS=Homo sapiens GN=CAP1 PE=1 SV=5	3	13	0.23
PGM1	P36871PGM1_HUMAN Phosphoglucomutase-1 OS=Homo sapiens GN=PGM1 PE=1 SV=3	4	18	0.22
YARS	P54577YARS_HUMAN Tyrosine-tRNA ligase, cytoplasmic OS=Homo sapiens GN=YARS PE=1 SV=4	4	18	0.22
FSCN1	Q16658FSCN1_HUMAN Fascin OS=Homo sapiens GN=FSCN1 PE=1 SV=3	2	9	0.22
PMPCA	Q10713PMPCA_HUMAN Mitochondrial-processing peptidase subunit alpha OS=Homo sapiens GN=PMPCA PE=1 SV=2	2	9	0.22
NARS	O43776NARS_HUMAN Asparagine-tRNA ligase, cytoplasmic OS=Homo sapiens GN=NARS PE=1 SV=1	4	18	0.22
MAOB	P27338MAOB_HUMAN Amine oxidase (flavin-containing) B OS=Homo sapiens GN=MAOB PE=1 SV=3	2	9	0.22
ACSF2	Q96CM8ACSF2_HUMAN Acyl-CoA synthetase family member 2, mitochondrial OS=Homo sapiens GN=ACSF2 PE=1 SV=2	3	14	0.21
GRWD1	Q98Q67GRWD1_HUMAN Glutamate-rich WD repeat-containing protein 1 OS=Homo sapiens GN=GRWD1 PE=1 SV=1	1	5	0.20
RCC2	Q9P25BRCC2_HUMAN Protein RCC2 OS=Homo sapiens GN=RCC2 PE=1 SV=2	1	5	0.20
ARCN1	P48444ARCN1_HUMAN Costumer subunit delta OS=Homo sapiens GN=ARCN1 PE=1 SV=1	2	10	0.20
ADCK3	Q8N60IADCK3_HUMAN Atypical kinase ADCK3, mitochondrial OS=Homo sapiens GN=ADCK3 PE=1 SV=1	1	5	0.20
CPNE1	Q98293CPNE1_HUMAN Copine-1 OS=Homo sapiens GN=CPNE1 PE=1 SV=1	2	10	0.20
RTCB	Q9Y301RTCB_HUMAN tRNA-splicing ligase RtcB homolog OS=Homo sapiens GN=RTCB PE=1 SV=1	3	15	0.20
CNDP2	Q96K4PCNDP2_HUMAN Cytosolic non-specific dipeptidase OS=Homo sapiens GN=CNDP2 PE=1 SV=2	1	5	0.20
NTSDC2	Q9H857NTSDC2_HUMAN 5'-nucleotidase domain-containing protein 2 OS=Homo sapiens GN=NTSDC2 PE=1 SV=1	1	5	0.20
SPATS2L	Q9NUQ6SPATS2L_HUMAN SPATS2-like protein OS=Homo sapiens GN=SPATS2L PE=1 SV=2	1	5	0.20
PSPC1	Q8WXF1PSPC1_HUMAN Paraspeckle component 1 OS=Homo sapiens GN=PSPC1 PE=1 SV=1	2	11	0.18
GLB1	P16278GLB1_HUMAN Beta-galactosidase OS=Homo sapiens GN=GLB1 PE=1 SV=2	2	11	0.18
ALDH5A1	P51649ALDH5A1_HUMAN Succinate-semialdehyde dehydrogenase, mitochondrial OS=Homo sapiens GN=ALDH5A1 PE=1 SV=2	2	11	0.18
ATAD3A	Q9NV7IATAD3A_HUMAN ATPase family AAA domain-containing protein 3A OS=Homo sapiens GN=ATAD3A PE=1 SV=2	3	17	0.18
GSR	P00390GSR_HUMAN Glutathione reductase, mitochondrial OS=Homo sapiens GN=GSR PE=1 SV=2	2	12	0.17
SMARCE1	Q969G3SMARCE1_HUMAN SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily E member 1 OS=Homo sapiens GN=SMARCE1 PE=1 SV=2	1	6	0.17
LEMED2	Q8NC56LEMED2_HUMAN LEM domain-containing protein 2 OS=Homo sapiens GN=LEMED2 PE=1 SV=1	1	6	0.17
FTCD	O95954FTCD_HUMAN Formimidoyltransferase-cyclodeaminase OS=Homo sapiens GN=FTCD PE=1 SV=2	2	12	0.17
SARS	P49591SARS_HUMAN Serine-tRNA ligase, cytoplasmic OS=Homo sapiens GN=SARS PE=1 SV=3	2	13	0.15
AIFM1	O95831AIFM1_HUMAN Apoptosis-inducing factor 1, mitochondrial OS=Homo sapiens GN=AIFM1 PE=1 SV=1	3	20	0.15
RPA1	P27694RPA1_HUMAN Replication protein A 70 kDa DNA-binding subunit OS=Homo sapiens GN=RPA1 PE=1 SV=2	1	7	0.14
HMGC51	Q01581HMGC51_HUMAN Hydroxymethylglutaryl-CoA synthase, cytoplasmic OS=Homo sapiens GN=HMGC51 PE=1 SV=2	2	14	0.14
PGM3	O95294PGM3_HUMAN Phosphoglucoamylase OS=Homo sapiens GN=PGM3 PE=1 SV=1	1	7	0.14
HDAC1	Q13547HDAC1_HUMAN Histone deacetylase 1 OS=Homo sapiens GN=HDAC1 PE=1 SV=1	1	7	0.14
ACADVL	P49748ACADVL_HUMAN Very long-chain specific acyl-CoA dehydrogenase, mitochondrial OS=Homo sapiens GN=ACADVL PE=1 SV=1	2	15	0.13
PPP2R1A	P301532PPP2R1A_HUMAN Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha isoform OS=Homo sapiens GN=PPP2R1A PE=1 SV=4	2	15	0.13
FKBP4	Q02790FKBP4_HUMAN Peptidyl-prolyl cis-trans isomerase FKBP4 OS=Homo sapiens GN=FKBP4 PE=1 SV=3	2	16	0.13
KPNA2	P52292KPNA2_HUMAN Importin subunit alpha-1 OS=Homo sapiens GN=KPNA2 PE=1 SV=1	1	8	0.13
MAOA	P21397MAOA_HUMAN Amine oxidase (flavin-containing) A OS=Homo sapiens GN=MAOA PE=1 SV=1	1	8	0.13
CHDH	Q8NE62CHDH_HUMAN Choline dehydrogenase, mitochondrial OS=Homo sapiens GN=CHDH PE=1 SV=2	1	8	0.13
PCK1	P35558PCK1_HUMAN Phosphoenolpyruvate carboxylase, cytosolic [GTP] OS=Homo sapiens GN=PCK1 PE=1 SV=3	1	9	0.11
IGF2BP2	Q9Y6M1IGF2BP2_HUMAN Insulin-like growth factor 2 mRNA-binding protein 2 OS=Homo sapiens GN=IGF2BP2 PE=1 SV=2	1	9	0.11
GALNT2	Q10471GALNT2_HUMAN Polypeptide N-acetylgalactosaminyltransferase 2 OS=Homo sapiens GN=GALNT2 PE=1 SV=1	1	9	0.11
UTP15	Q8TED0UTP15_HUMAN U3 small nucleolar RNA-associated protein 15 homolog OS=Homo sapiens GN=UTP15 PE=1 SV=3	1	9	0.11
HSP90B1	P14625HSP90B1_HUMAN Endoplasmic reticulum protein OS=Homo sapiens GN=HSP90B1 PE=1 SV=1	1	9	0.11
STIP1	P31948STIP1_HUMAN Stress-induced-phosphoprotein 1 OS=Homo sapiens GN=STIP1 PE=1 SV=1	2	20	0.10
USP14	P54578USP14_HUMAN Ubiquitin carboxyl-terminal hydrolase 14 OS=Homo sapiens GN=USP14 PE=1 SV=3	1	11	0.09
SGPL1	O95470SGPL1_HUMAN Sphingosine-1-phosphate lyase 1 OS=Homo sapiens GN=SGPL1 PE=1 SV=3	0	5	0.00
DAPK3	O43293DAPK3_HUMAN Death-associated protein kinase 3 OS=Homo sapiens GN=DAPK3 PE=1 SV=1	0	2	0.00
ENTPD6	O75345ENTPD6_HUMAN Ectonucleoside triphosphate diphosphohydrolase 6 OS=Homo sapiens GN=ENTPD6 PE=1 SV=3	0	2	0.00
SERINC1	Q9NRX5SERINC1_HUMAN Serine incorporator 1 OS=Homo sapiens GN=SERINC1 PE=1 SV=1	0	1	0.00
ITGB1	P05556ITGB1_HUMAN Integrin beta-1 OS=Homo sapiens GN=ITGB1 PE=1 SV=2	0	2	0.00
ZNF512	Q96ME7ZNF512_HUMAN Zinc finger protein 512 OS=Homo sapiens GN=ZNF512 PE=1 SV=2	0	6	0.00
CPT2	P23786CPT2_HUMAN Carnitine O-palmitoyltransferase 2, mitochondrial OS=Homo sapiens GN=CPT2 PE=1 SV=2	0	5	0.00
NRBP1	Q9UHY1NRBP1_HUMAN Nuclear receptor-binding protein OS=Homo sapiens GN=NRBP1 PE=1 SV=1	0	3	0.00
YTHDF1	Q98Y30YTHDF1_HUMAN YTH domain-containing family protein 1 OS=Homo sapiens GN=YTHDF1 PE=1 SV=1	0	3	0.00
PTPN11	Q06124PTPN11_HUMAN Tyrosine-protein phosphatase non-receptor type 11 OS=Homo sapiens GN=PTPN11 PE=1 SV=2	0	2	0.00
HSPA13	P48723HSPA13_HUMAN Heat shock 70 kDa protein 13 OS=Homo sapiens GN=HSPA13 PE=1 SV=1	0	1	0.00
CTSA	P10619PP2R1A_HUMAN Lysosomal protective protein OS=Homo sapiens GN=CTSA PE=1 SV=2	0	1	0.00
FER	P16591FER_HUMAN Tyrosine-protein kinase Fer OS=Homo sapiens GN=FER PE=1 SV=2	0	2	0.00
SMU1	Q27AY7SMU1_HUMAN WD40 repeat-containing protein SMU1 OS=Homo sapiens GN=SMU1 PE=1 SV=2	0	3	0.00
PCCB	P05166PCCB_HUMAN Propionyl-CoA carboxylase beta chain, mitochondrial OS=Homo sapiens GN=PCCB PE=1 SV=3	0	9	0.00
RANGAP1	P46060RANGAP1_HUMAN Ran GTPase-activating protein 1 OS=Homo sapiens GN=RANGAP1 PE=1 SV=1	0	3	0.00
CDC47	Q96A33CDC47_HUMAN Colicoid domain-containing protein 47 OS=Homo sapiens GN=CDC47 PE=1 SV=1	0	11	0.00
G3BP1	Q13283G3BP1_HUMAN Ras GTPase-activating protein-binding protein 1 OS=Homo sapiens GN=G3BP1 PE=1 SV=1	0	9	0.00
CPVL	Q9H3G5CPVL_HUMAN Probable serine carboxypeptidase CPVL OS=Homo sapiens GN=CPVL PE=1 SV=2	0	1	0.00
LACTB	P83111LACTB_HUMAN Serine beta-lactamase-like protein LACTB, mitochondrial OS=Homo sapiens GN=LACTB PE=1 SV=2	0	3	0.00
ACSM2A	Q08A83ACSM2A_HUMAN Acyl-coenzyme A synthetase ACSM2A, mitochondrial OS=Homo sapiens GN=ACSM2A PE=1 SV=2	0	1	0.00
FAM134C	Q86VR2F134C_HUMAN Protein FAM134C OS=Homo sapiens GN=FAM134C PE=1 SV=1	0	4	0.00
AHSG	P02765AHSG_HUMAN Alpha-2-HS-glycoprotein OS=Homo sapiens GN=AHSG PE=1 SV=1	0	4	0.00
RTN4	Q9NQC3RTN4_HUMAN Reticulon-4 OS=Homo sapiens GN=RTN4 PE=1 SV=2	0	1	0.00
SLC1A4	P43007SLC1A4_HUMAN Neutral amino acid transporter A OS=Homo sapiens GN=SLC1A4 PE=1 SV=1	0	1	0.00
SELENBP1	Q13228SELENBP1_HUMAN Selenium-binding protein 1 OS=Homo sapiens GN=SELENBP1 PE=1 SV=2	0	6	0.00
PNKP	Q96T60PNKP_HUMAN Bifunctional polynucleotide phosphatase/kinase OS=Homo sapiens GN=PNKP PE=1 SV=1	0	2	0.00
DDX19A	Q9NUJ7DD19A_HUMAN ATP-dependent RNA helicase DDX19A OS=Homo sapiens GN=DDX19A PE=1 SV=1	0	6	0.00
EPHX2	P34913EPHX2_HUMAN Bifunctional epoxide hydrolase 2 OS=Homo sapiens GN=EPHX2 PE=1 SV=2	0	4	0.00
ZNHIT6	Q9NWK9ZNHIT6_HUMAN Box C/D snoRNA protein 1 OS=Homo sapiens GN=ZNHIT6 PE=1 SV=1	0	2	0.00
DDX17	Q92841DDX17_HUMAN Probable ATP-dependent RNA helicase DDX17 OS=Homo sapiens GN=DDX17 PE=1 SV=2	0	1	0.00
BCAM	P50895BCAM_HUMAN Basal cell adhesion molecule OS=Homo sapiens GN=BCAM PE=1 SV=2	0	2	0.00
CDC73	Q6P1J9CDC73_HUMAN Parafibromin OS=Homo sapiens GN=CDC73 PE=1 SV=1	0	4	0.00
MARS2	Q96G91MARS2_HUMAN Methionine-tRNA ligase, mitochondrial OS=Homo sapiens GN=MARS2 PE=1 SV=2	0	4	0.00
PELP1	Q8IZL8PELP1_HUMAN Proline-, glutamic acid- and leucine-rich protein 1 OS=Homo sapiens GN=PELP1 PE=1 SV=2	0	3	0.00
TOE1	Q96GM8TOE1_HUMAN Target of EGR1 protein 1 OS=Homo sapiens GN=TOE1 PE=1 SV=1	0	3	0.00
NEK2	P51955NEK2_HUMAN Serine/threonine-protein kinase Nek2 OS=Homo sapiens GN=NEK2 PE=1 SV=1	0	2	0.00
NUDCD1	Q96RS6NUDCD1_HUMAN NudC domain-containing protein 1 OS=Homo sapiens GN=NUDCD1 PE=1 SV=2	0	2	0.00
PPP4R2	Q9NY27PPP4R2_HUMAN Serine/threonine-protein phosphatase 4 regulatory subunit 2 OS=Homo sapiens GN=PPP4R2 PE=1 SV=3	0	3	0.00

KIAA0391	O15091MRP3_HUMAN Mitochondrial ribonuclease P protein 3 OS=Homo sapiens GN=KIAA0391 PE=1 SV=2	0	2	0.00
HDAC2	Q92769HDAC2_HUMAN Histone deacetylase 2 OS=Homo sapiens GN=HDAC2 PE=1 SV=2	0	2	0.00
STXB3	O00186STXB3_HUMAN Syntaxin-binding protein 3 OS=Homo sapiens GN=STXB3 PE=1 SV=2	0	3	0.00
APH	P02749APH_HUMAN Beta-2-glycoprotein 1 OS=Homo sapiens GN=APH PE=1 SV=3	0	3	0.00
No_GeneID	AOA1BOGU03AOA1BOGU03_HUMAN Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0	1	0.00
SEPT11	Q9NVA2SEPT11_HUMAN Septin-11 OS=Homo sapiens GN=SEPT11 PE=1 SV=3	0	4	0.00
KMT2A	Q03164KMT2A_HUMAN Histone-lysine N-methyltransferase 2A OS=Homo sapiens GN=KMT2A PE=1 SV=5	0	7	0.00
NUCB1	Q02818NUCB1_HUMAN Nucleobindin-1 OS=Homo sapiens GN=NUCB1 PE=1 SV=4	0	9	0.00
C1QB	Q07021C1QB_HUMAN Complement component 1 Q subcomponent-binding protein, mitochondrial OS=Homo sapiens GN=C1QB PE=1 SV=1	0	1	0.00
PICALM	Q13492PICALM_HUMAN Phosphatidylinositol-binding clathrin assembly protein OS=Homo sapiens GN=PICALM PE=1 SV=2	0	2	0.00
GNL3	Q9BVP2GNL3_HUMAN Guanine nucleotide-binding protein-like 3 OS=Homo sapiens GN=GNL3 PE=1 SV=2	0	6	0.00
ADAM10	O14672ADAM10_HUMAN Disintegrin and metalloprotease domain-containing protein 10 OS=Homo sapiens GN=ADAM10 PE=1 SV=1	0	2	0.00
SNW1	Q13573SNW1_HUMAN SNW domain-containing protein 1 OS=Homo sapiens GN=SNW1 PE=1 SV=1	0	4	0.00
PEX14	O75381PEX14_HUMAN Peroxisomal membrane protein PEX14 OS=Homo sapiens GN=PEX14 PE=1 SV=1	0	2	0.00
ALDOC	P09972ALDOC_HUMAN Fructose-bisphosphate aldolase C OS=Homo sapiens GN=ALDOC PE=1 SV=2	0	1	0.00
CDYL	Q9Y232CDYL1_HUMAN Chromodomain Y-like protein OS=Homo sapiens GN=CDYL PE=1 SV=2	0	2	0.00
RELA	Q04206ITF65_HUMAN Transcription factor p65 OS=Homo sapiens GN=RELA PE=1 SV=2	0	3	0.00
NOP9	Q8U38INOP9_HUMAN Nucleolar protein 9 OS=Homo sapiens GN=NOP9 PE=1 SV=1	0	8	0.00
CA9	Q16790CAH9_HUMAN Carbonic anhydrase 9 OS=Homo sapiens GN=CA9 PE=1 SV=2	0	1	0.00
HNRNP	Q00839HNRNP_HUMAN Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNP PE=1 SV=6	0	3	0.00
WR37	Q9Y281WR37_HUMAN WD repeat-containing protein 37 OS=Homo sapiens GN=WR37 PE=1 SV=2	0	1	0.00
MFAP1	P55081MFAP1_HUMAN Microfibrillar-associated protein 1 OS=Homo sapiens GN=MFAP1 PE=1 SV=2	0	2	0.00
IMPDH2	P12268IMPDH2_HUMAN Inosine-5'-monophosphate dehydrogenase 2 OS=Homo sapiens GN=IMPDH2 PE=1 SV=2	0	11	0.00
CWF19L1	Q9BYN2C19L1_HUMAN CWF19-like protein 1 OS=Homo sapiens GN=CWF19L1 PE=1 SV=2	0	1	0.00
EHD1	Q9HAM9EHD1_HUMAN EH domain-containing protein 1 OS=Homo sapiens GN=EHD1 PE=1 SV=2	0	4	0.00
UTP3	Q9NQZ2IAS10_HUMAN Something about silencing protein 10 OS=Homo sapiens GN=UTP3 PE=1 SV=1	0	1	0.00
SRP54	P61011SRP54_HUMAN Signal recognition particle 54 kDa protein OS=Homo sapiens GN=SRP54 PE=1 SV=1	0	7	0.00
NAE1	Q13564IULA1_HUMAN NEDD8-activating enzyme E1 regulatory subunit OS=Homo sapiens GN=NAE1 PE=1 SV=1	0	4	0.00
PPAT	Q06203PPAT_HUMAN Amidophosphoribosyltransferase OS=Homo sapiens GN=PPAT PE=1 SV=1	0	5	0.00
LPAT2	Q7L5N7PCAT2_HUMAN Lysophosphatidylcholine acyltransferase 2 OS=Homo sapiens GN=LPAT2 PE=1 SV=1	0	3	0.00
APMAP	Q9HDC9APMAP_HUMAN Adipocyte plasma membrane-associated protein OS=Homo sapiens GN=APMAP PE=1 SV=2	0	2	0.00
DPYSL4	O14531DPYSL4_HUMAN Dihydropyrimidinase-related protein 4 OS=Homo sapiens GN=DPYSL4 PE=1 SV=2	0	2	0.00
PGK1	P00558PGK1_HUMAN Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3	0	2	0.00
SLC7A5	Q01650LAT1_HUMAN Large neutral amino acids transporter small subunit 1 OS=Homo sapiens GN=SLC7A5 PE=1 SV=2	0	1	0.00
PEPD	P12955PEPD_HUMAN Xaa-Pro dipeptidase OS=Homo sapiens GN=PEPD PE=1 SV=3	0	8	0.00
NPM1	P06748NPM_HUMAN Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2	0	4	0.00
ZC3HC1	Q86W0NIP4_HUMAN Nuclear-interacting partner of ALK OS=Homo sapiens GN=ZC3HC1 PE=1 SV=1	0	2	0.00
GDI1	P31150GIDIA_HUMAN Rab GDP dissociation inhibitor alpha OS=Homo sapiens GN=GDI1 PE=1 SV=2	0	3	0.00
PHK2A	Q9BTU6PHK2A_HUMAN Phosphatidylinositol 4-kinase type 2-alpha OS=Homo sapiens GN=PHK2A PE=1 SV=1	0	2	0.00
PDIA4	P13667PDIA4_HUMAN Protein disulfide-isomerase A4 OS=Homo sapiens GN=PDIA4 PE=1 SV=2	0	4	0.00
TAOK2	Q9UL54ITAO2_HUMAN Serine/threonine-protein kinase TAO2 OS=Homo sapiens GN=TAOK2 PE=1 SV=2	0	3	0.00
PPP3CA	Q08209PPP2B8_HUMAN Serine/threonine-protein phosphatase 2B catalytic subunit alpha isoform OS=Homo sapiens GN=PPP3CA PE=1 SV=1	0	4	0.00
UTP15	Q8TED0UTP15_HUMAN U3 small nucleolar RNA-associated protein 15 homolog OS=Homo sapiens GN=UTP15 PE=1 SV=3	0	1	0.00
AHCYL1	O43865SAHH2_HUMAN Adenosylhomocysteinase 2 OS=Homo sapiens GN=AHCYL1 PE=1 SV=2	0	3	0.00
ACTB	P60709IACB_HUMAN Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	0	9	0.00
GPS1	Q13098ICSN1_HUMAN COP9 signalosome complex subunit 1 OS=Homo sapiens GN=GPS1 PE=1 SV=4	0	2	0.00
STIM1	Q13586STIM1_HUMAN Stromal interaction molecule 1 OS=Homo sapiens GN=STIM1 PE=1 SV=3	0	1	0.00
PEG10	Q86T67PEG10_HUMAN Retrotransposon-derived protein PEG10 OS=Homo sapiens GN=PEG10 PE=1 SV=2	0	1	0.00
PRPF31	Q8WY3PRPF31_HUMAN U4/U6 small nuclear ribonucleoprotein Pp31 OS=Homo sapiens GN=PRPF31 PE=1 SV=2	0	2	0.00
FKBP8	Q14318FKBP8_HUMAN Peptidyl-prolyl cis-trans isomerase FKBP8 OS=Homo sapiens GN=FKBP8 PE=1 SV=2	0	4	0.00
ODR4	Q5SW8IODR4_HUMAN Protein odr-4 homolog OS=Homo sapiens GN=ODR4 PE=1 SV=1	0	2	0.00
FAM69A	Q5T7M9IFA69A_HUMAN Protein FAM69A OS=Homo sapiens GN=FAM69A PE=2 SV=1	0	1	0.00
STXB2	Q15833STXB2_HUMAN Syntaxin-binding protein 2 OS=Homo sapiens GN=STXB2 PE=1 SV=2	0	5	0.00
ARSE	P51690ARSE_HUMAN Arylsulfatase E OS=Homo sapiens GN=ARSE PE=1 SV=2	0	6	0.00
CCDC88B	A6NC98CC88B_HUMAN Coiled-coil domain-containing protein 88 OS=Homo sapiens GN=CCDC88B PE=1 SV=1	0	4	0.00
CDKN2AIP	Q9NXV6ICARF_HUMAN CDKN2A-interacting protein OS=Homo sapiens GN=CDKN2AIP PE=1 SV=3	0	3	0.00
POCD4	Q53EL6POCD4_HUMAN Programmed cell death protein 4 OS=Homo sapiens GN=POCD4 PE=1 SV=2	0	2	0.00
WHHD1	O757171WHHD1_HUMAN WD repeat and HMG-box DNA-binding protein 1 OS=Homo sapiens GN=WHHD1 PE=1 SV=1	0	2	0.00
TMEM209	Q96SK2TM209_HUMAN Transmembrane protein 209 OS=Homo sapiens GN=TMEM209 PE=1 SV=2	0	2	0.00
PPP5C	P53041PPP5_HUMAN Serine/threonine-protein phosphatase 5 OS=Homo sapiens GN=PPP5C PE=1 SV=1	0	5	0.00
PIGT	Q969N2PIGT_HUMAN GPI transamidase component PIG-T OS=Homo sapiens GN=PIGT PE=1 SV=1	0	3	0.00
GK2	Q14410GLPK2_HUMAN Glycerol kinase 2 OS=Homo sapiens GN=GK2 PE=2 SV=2	0	2	0.00
EIF2B4	Q9U10E2B4_HUMAN Translation initiation factor eIF-2B subunit delta OS=Homo sapiens GN=EIF2B4 PE=1 SV=2	0	4	0.00
EIF2B3	Q9NR50E2B3_HUMAN Translation initiation factor eIF-2B subunit gamma OS=Homo sapiens GN=EIF2B3 PE=1 SV=1	0	2	0.00
NAP1L4	Q9Y733INP1L4_HUMAN Nucleosome assembly protein 1-like 4 OS=Homo sapiens GN=NAP1L4 PE=1 SV=1	0	7	0.00
HSP90A1	P08238HSP90A_HUMAN Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90A1 PE=1 SV=4	0	7	0.00
NELFCD	Q8IXH7NELFD_HUMAN Negative elongation factor C/D OS=Homo sapiens GN=NELFCD PE=1 SV=2	0	3	0.00
RAP1GDS1	P52306GDS1_HUMAN Rap1 GTPase-GDP dissociation stimulator 1 OS=Homo sapiens GN=RAP1GDS1 PE=1 SV=3	0	3	0.00
HSPA6	P17066HSP76_HUMAN Heat shock 70 kDa protein 6 OS=Homo sapiens GN=HSPA6 PE=1 SV=2	0	1	0.00
TOR1AIP2	Q8NFQ8TOIP2_HUMAN Torsin-1A-interacting protein 2 OS=Homo sapiens GN=TOR1AIP2 PE=1 SV=1	0	1	0.00
FN1	P02751FN1_HUMAN Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4	0	5	0.00
MTA3	Q9BTC8MTA3_HUMAN Metastasis-associated protein MTA3 OS=Homo sapiens GN=MTA3 PE=1 SV=2	0	2	0.00
SLC27A1	Q9PCB7IS27A1_HUMAN Long-chain fatty acid transport protein 1 OS=Homo sapiens GN=SLC27A1 PE=2 SV=1	0	2	0.00
PACSN2	Q9UNF0PACN2_HUMAN Protein kinase C and casein kinase substrate in neurons protein 2 OS=Homo sapiens GN=PACSN2 PE=1 SV=2	0	7	0.00
FARS5	Q9NSD9ISYF8_HUMAN Phenylalanine--tRNA ligase beta subunit OS=Homo sapiens GN=FARS5 PE=1 SV=3	0	8	0.00
NARS2	Q96I59IYNM_HUMAN Probable asparagine--tRNA ligase, mitochondrial OS=Homo sapiens GN=NARS2 PE=1 SV=3	0	3	0.00
ACSM3	Q53F22ACSM3_HUMAN Acyl-coenzyme A synthetase ACSM3, mitochondrial OS=Homo sapiens GN=ACSM3 PE=1 SV=2	0	3	0.00
RBM39	Q14498RBM39_HUMAN RNA-binding protein 39 OS=Homo sapiens GN=RBM39 PE=1 SV=2	0	1	0.00
PIG5	Q96522PIG5_HUMAN GPI transamidase component PIG-5 OS=Homo sapiens GN=PIG5 PE=1 SV=3	0	4	0.00
CLPTM1L	Q96KA5CLPTM1L_HUMAN Clef, lip and palate transmembrane protein 1-like protein OS=Homo sapiens GN=CLPTM1L PE=1 SV=1	0	3	0.00
SERPINA7	P05543THBC_HUMAN Throxine-binding globulin OS=Homo sapiens GN=SERPINA7 PE=1 SV=2	0	3	0.00
MGST2	Q99735MGST2_HUMAN Microsomal glutathione S-transferase 2 OS=Homo sapiens GN=MGST2 PE=1 SV=1	0	1	0.00
NAP1L1	P55209INP1L1_HUMAN Nucleosome assembly protein 1-like 1 OS=Homo sapiens GN=NAP1L1 PE=1 SV=1	0	3	0.00
FKBP5	Q13451FKBP5_HUMAN Peptidyl-prolyl cis-trans isomerase FKBP5 OS=Homo sapiens GN=FKBP5 PE=1 SV=2	0	6	0.00
CORO1B	Q9BR76COR1B_HUMAN Coronin-1B OS=Homo sapiens GN=CORO1B PE=1 SV=1	0	6	0.00
SLC16A3	O15427MOT4_HUMAN Monocarboxylate transporter 4 OS=Homo sapiens GN=SLC16A3 PE=1 SV=1	0	1	0.00
XPNPE3	Q9NQH7XPP3_HUMAN Probable Xaa-Pro aminopeptidase 3 OS=Homo sapiens GN=XPNPE3 PE=1 SV=1	0	1	0.00
EXOSC9	Q06265EXOSC9_HUMAN Exosome complex component RRP45 OS=Homo sapiens GN=EXOSC9 PE=1 SV=3	0	2	0.00

MTAP	Q9NVV4IPAD1_HUMAN Poly(A) RNA polymerase, mitochondrial OS=Homo sapiens GN=MTAP PE=1 SV=1	0	3	0.00
SLC44A1	Q8WVISL1CT1_HUMAN Choline transporter-like protein 1 OS=Homo sapiens GN=SLC44A1 PE=1 SV=1	0	2	0.00
SUOX	P51687SUOX_HUMAN Sulfite oxidase, mitochondrial OS=Homo sapiens GN=SUOX PE=1 SV=2	0	3	0.00
EWSR1	Q01844EWS_HUMAN RNA-binding protein EWS OS=Homo sapiens GN=EWSR1 PE=1 SV=1	0	1	0.00
M6PR	P20645IMPRD_HUMAN Cation-dependent mannose-6-phosphate receptor OS=Homo sapiens GN=M6PR PE=1 SV=1	0	1	0.00
PIPSK1A	Q99755PIPS1A_HUMAN Phosphatidylinositol 4-phosphate 5-kinase type-1 alpha OS=Homo sapiens GN=PIPSK1A PE=1 SV=1	0	4	0.00
SEC62	Q99442SEC62_HUMAN Translocation protein SEC62 OS=Homo sapiens GN=SEC62 PE=1 SV=1	0	3	0.00
FUBP3	Q96124FUBP3_HUMAN Far upstream element-binding protein 3 OS=Homo sapiens GN=FUBP3 PE=1 SV=2	0	12	0.00
TMX3	Q96171TMX3_HUMAN Protein disulfide-isomerase TMX3 OS=Homo sapiens GN=TMX3 PE=1 SV=2	0	7	0.00
ATE1	O95260ATE1_HUMAN Arginyl-tRNA--protein transferase 1 OS=Homo sapiens GN=ATE1 PE=1 SV=2	0	2	0.00
NTSC2	P49902ISNTC_HUMAN Cytosolic purine 5'-nucleotidase OS=Homo sapiens GN=NTSC2 PE=1 SV=1	0	1	0.00
CTPS2	Q9NFB8PYRG2_HUMAN CTP synthase 2 OS=Homo sapiens GN=CTPS2 PE=1 SV=1	0	1	0.00
P3H4	Q82791ISC65_HUMAN Synaptonemal complex protein SC65 OS=Homo sapiens GN=P3H4 PE=1 SV=1	0	6	0.00
CPQ	Q9Y646CPQ_HUMAN Carboxypeptidase Q OS=Homo sapiens GN=CPQ PE=1 SV=1	0	4	0.00
NCOA5	Q9HCD5NCOA5_HUMAN Nuclear receptor coactivator 5 OS=Homo sapiens GN=NCOA5 PE=1 SV=2	0	5	0.00
SACM1L	Q9NTJ5ISAC1_HUMAN Phosphatidylinositol phosphatase SAC1 OS=Homo sapiens GN=SACM1L PE=1 SV=2	0	3	0.00
DYNC1H1	Q14204IDYHC1_HUMAN Cytoplasmic dynein 1 heavy chain 1 OS=Homo sapiens GN=DYNC1H1 PE=1 SV=5	0	3	0.00
F10	P00742F10_HUMAN Coagulation factor X OS=Homo sapiens GN=F10 PE=1 SV=2	0	4	0.00
LZTS3	O60299LZTS3_HUMAN Leucine zipper putative tumor suppressor 3 OS=Homo sapiens GN=LZTS3 PE=2 SV=1	0	3	0.00
HYOU1	Q9Y41HYOU1_HUMAN Hypoxia up-regulated protein 1 OS=Homo sapiens GN=HYOU1 PE=1 SV=1	0	4	0.00
NUP50	Q9UKX7NUP50_HUMAN Nuclear pore complex protein Nup50 OS=Homo sapiens GN=NUP50 PE=1 SV=2	0	3	0.00
PRRC1	Q96M27IPRC1_HUMAN Protein PRRC1 OS=Homo sapiens GN=PRRC1 PE=1 SV=1	0	2	0.00
AJUBA	Q96F1AJUBA_HUMAN LIM domain-containing protein ajuba OS=Homo sapiens GN=AJUBA PE=1 SV=1	0	6	0.00
CKAP4	Q07065CKAP4_HUMAN Cytoskeleton-associated protein 4 OS=Homo sapiens GN=CKAP4 PE=1 SV=2	0	20	0.00
SGSH	P51688SGSH_HUMAN N-sulphoglucosamine sulphohydrolase OS=Homo sapiens GN=SGSH PE=1 SV=1	0	2	0.00
TECR	Q9NZ01ITECR_HUMAN Very-long-chain enoyl-CoA reductase OS=Homo sapiens GN=TECR PE=1 SV=1	0	1	0.00
B4GAT1	O43505B4GAT1_HUMAN Beta-1,4-glucuronyltransferase 1 OS=Homo sapiens GN=B4GAT1 PE=1 SV=1	0	2	0.00
METTL7A	Q9H8H3MET7A_HUMAN Methyltransferase-like protein 7A OS=Homo sapiens GN=METTL7A PE=1 SV=1	0	1	0.00
SEPT8	Q92599SEPT8_HUMAN Septin-8 OS=Homo sapiens GN=SEPT8 PE=1 SV=4	0	2	0.00
HSPA1L	P34931H517L_HUMAN Heat shock 70 kDa protein 1-like OS=Homo sapiens GN=HSPA1L PE=1 SV=2	0	3	0.00
DST	Q03001IDYST_HUMAN Dystonin OS=Homo sapiens GN=DST PE=1 SV=4	0	6	0.00
ME2	P23368MAOM_HUMAN NAD-dependent malic enzyme, mitochondrial OS=Homo sapiens GN=ME2 PE=1 SV=1	0	4	0.00
RBBP4	Q09028RBBP4_HUMAN Histone-binding protein RBBP4 OS=Homo sapiens GN=RBBP4 PE=1 SV=3	0	1	0.00
DPY19L3	Q62PD9D19L3_HUMAN Probable C-mannosyltransferase DPY19L3 OS=Homo sapiens GN=DPY19L3 PE=2 SV=1	0	1	0.00
POZK1	Q5T2W1NHRF3_HUMAN Nsf(+)/VH(+) exchange regulatory cofactor NHE-RF3 OS=Homo sapiens GN=POZK1 PE=1 SV=2	0	2	0.00
CD55	P08174IDAF_HUMAN Complement decay-accelerating factor OS=Homo sapiens GN=CD55 PE=1 SV=4	0	1	0.00
RCC1	P18754IRCC1_HUMAN Regulator of chromosome condensation OS=Homo sapiens GN=RCC1 PE=1 SV=1	0	3	0.00
CDK5RAP3	Q96JB5ICK5P3_HUMAN CDK5 regulatory subunit-associated protein 3 OS=Homo sapiens GN=CDK5RAP3 PE=1 SV=2	0	6	0.00
SNX27	Q96L92SNX27_HUMAN Sorting nexin-27 OS=Homo sapiens GN=SNX27 PE=1 SV=2	0	3	0.00
PP1L4	Q8WUA2PP1L4_HUMAN Peptidyl-prolyl cis-trans isomerase-like 4 OS=Homo sapiens GN=PP1L4 PE=1 SV=1	0	2	0.00
PLD3	Q8V08IPLD3_HUMAN Phospholipase D3 OS=Homo sapiens GN=PLD3 PE=1 SV=1	0	3	0.00
TF	P02787ITRFE_HUMAN Serotransferrin OS=Homo sapiens GN=TF PE=1 SV=3	0	3	0.00
AKAP1	Q92667IAKAP1_HUMAN A-kinase anchor protein 1, mitochondrial OS=Homo sapiens GN=AKAP1 PE=1 SV=1	0	2	0.00
LSM14A	Q8ND56LSM14A_HUMAN Protein LSM14 homolog A OS=Homo sapiens GN=LSM14A PE=1 SV=3	0	3	0.00
TOM1	O60784TOM1_HUMAN Target of Myb protein 1 OS=Homo sapiens GN=TOM1 PE=1 SV=2	0	4	0.00
ZGPAT	Q8NSA5ZGPAT_HUMAN Zinc finger CCH-type with G patch domain-containing protein OS=Homo sapiens GN=ZGPAT PE=1 SV=3	0	2	0.00
GSDMD	P57764IGSDMD_HUMAN Gasdermin-D OS=Homo sapiens GN=GSDMD PE=1 SV=1	0	2	0.00
PRICKLE2	Q7Z3G6PRIC2_HUMAN Prickle-like protein 2 OS=Homo sapiens GN=PRICKLE2 PE=1 SV=2	0	3	0.00
RPN1	P04843RPN1_HUMAN Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPN1 PE=1 SV=1	0	9	0.00
TMEI23	Q8N131IPORM_HUMAN Porin OS=Homo sapiens GN=TMEI23 PE=1 SV=1	0	1	0.00
YBX1	P67809YBOX1_HUMAN Nuclease-sensitive element-binding protein 1 OS=Homo sapiens GN=YBX1 PE=1 SV=3	0	1	0.00
GLTSCR2	Q9NZM5GSCR2_HUMAN Glioma tumor suppressor candidate region gene 2 protein OS=Homo sapiens GN=GLTSCR2 PE=1 SV=2	0	2	0.00
RAD23B	P54727IRD23B_HUMAN UV excision repair protein RAD23 homolog B OS=Homo sapiens GN=RAD23B PE=1 SV=1	0	6	0.00
SYNCRP	O60506HNRPQ_HUMAN Heterogeneous nuclear ribonucleoprotein Q OS=Homo sapiens GN=SYNCRP PE=1 SV=2	0	16	0.00
FTO	Q9C081FTO_HUMAN Alpha-ketoglutarate-dependent dioxygenase FTO OS=Homo sapiens GN=FTO PE=1 SV=3	0	4	0.00
CDC86	Q9H65FICDC86_HUMAN Coiled-coil domain-containing protein 86 OS=Homo sapiens GN=CDC86 PE=1 SV=1	0	2	0.00
DKC1	O60832DKC1_HUMAN H/ACA ribonucleoprotein complex subunit 4 OS=Homo sapiens GN=DKC1 PE=1 SV=3	0	4	0.00
SNRP70	P08621RUI7_HUMAN U1 small nuclear ribonucleoprotein 70 kDa OS=Homo sapiens GN=SNRP70 PE=1 SV=2	0	3	0.00
RPL8	P62917RPL8_HUMAN 60S ribosomal protein L8 OS=Homo sapiens GN=RPL8 PE=1 SV=2	0	1	0.00
KMT5B	Q4FZB7KMT5B_HUMAN Histone-lysine N-methyltransferase KMT5B OS=Homo sapiens GN=KMT5B PE=1 SV=4	0	2	0.00
SERPINA3	P01011IAACT_HUMAN Alpha-1-antichymotrypsin OS=Homo sapiens GN=SERPINA3 PE=1 SV=2	0	7	0.00
IQGAP1	P46940IQGAP1_HUMAN Ras GTPase-activating-like protein IQGAP1 OS=Homo sapiens GN=IQGAP1 PE=1 SV=1	0	3	0.00
ABCB7	O75027IACB7_HUMAN ATP-binding cassette sub-family B member 7, mitochondrial OS=Homo sapiens GN=ABCB7 PE=1 SV=2	0	2	0.00
TKT	P29401ITKT_HUMAN Transketolase OS=Homo sapiens GN=TKT PE=1 SV=3	0	6	0.00
UBP1	Q9NZI7IUBP1_HUMAN Upstream-binding protein 1 OS=Homo sapiens GN=UBP1 PE=1 SV=1	0	1	0.00
HEXA	P06865HEXA_HUMAN Beta-hexosaminidase subunit alpha OS=Homo sapiens GN=HEXA PE=1 SV=2	0	11	0.00
SIL1	Q9H173SIL1_HUMAN Nucleotide exchange factor SIL1 OS=Homo sapiens GN=SIL1 PE=1 SV=1	0	2	0.00
CARM1	Q86X5SICARM1_HUMAN Histone-arginine methyltransferase CARM1 OS=Homo sapiens GN=CARM1 PE=1 SV=3	0	2	0.00
HSPA9	P38646GRP75_HUMAN Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2	0	6	0.00
LNP	Q9C081LNP_HUMAN Protein lunapark OS=Homo sapiens GN=LNP PE=1 SV=2	0	2	0.00
TMSF3	Q9HD45ITMS3_HUMAN Transmembrane 9 superfamily member 3 OS=Homo sapiens GN=TMSF3 PE=1 SV=2	0	2	0.00
LTA4H	P09960LKH44_HUMAN Leukotriene A-4 hydrolase OS=Homo sapiens GN=LTA4H PE=1 SV=2	0	14	0.00
SPTBN1	Q01082ISPTB2_HUMAN Spectrin beta chain, non-erythrocytic 1 OS=Homo sapiens GN=SPTBN1 PE=1 SV=2	0	10	0.00
LRRC1	Q9BTT6LRRC1_HUMAN Leucine-rich repeat-containing protein 1 OS=Homo sapiens GN=LRRC1 PE=1 SV=1	0	2	0.00
SMARCD2	Q92925SMRD2_HUMAN SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily D member 2 OS=Homo sapiens GN=SMARCD2 PE=1 SV=3	0	3	0.00
VASN	Q9EMK4VASN_HUMAN Vasorin OS=Homo sapiens GN=VASN PE=1 SV=1	0	1	0.00
TUBB2B	Q9BVA1ITBB2B_HUMAN Tubulin beta-2B chain OS=Homo sapiens GN=TUBB2B PE=1 SV=1	0	4	0.00
ANAPC7	Q9UJX3IAPC7_HUMAN Anaphase-promoting complex subunit 7 OS=Homo sapiens GN=ANAPC7 PE=1 SV=4	0	2	0.00
TRIM27	P14373TRIM27_HUMAN Zinc finger protein RFP OS=Homo sapiens GN=TRIM27 PE=1 SV=1	0	5	0.00
FLNA	P21333FLNA_HUMAN Filamin-A OS=Homo sapiens GN=FLNA PE=1 SV=4	0	5	0.00
No_GeneID	AOA087WZ62IAOA087WZ62_HUMAN Mannosyltransferase OS=Homo sapiens GN=No_GeneID PE=3 SV=1	0	3	0.00
ATAD3B	Q5T9A4IATD3B_HUMAN ATPase family AAA domain-containing protein 3B OS=Homo sapiens GN=ATAD3B PE=1 SV=1	0	3	0.00
SYPL1	Q16563SYPL1_HUMAN Synaptophysin-like protein 1 OS=Homo sapiens GN=SYPL1 PE=1 SV=1	0	2	0.00
GLYR1	Q49A26GLYR1_HUMAN Putative oxidoreductase GLYR1 OS=Homo sapiens GN=GLYR1 PE=1 SV=3	0	1	0.00
RABGGTA	Q92696PGTA_HUMAN Geranylgeranyl transferase type-2 subunit alpha OS=Homo sapiens GN=RABGGTA PE=1 SV=2	0	7	0.00
RANBP3	Q9HGZ4RANB3_HUMAN Ran-binding protein 3 OS=Homo sapiens GN=RANBP3 PE=1 SV=1	0	8	0.00
TNIP3	Q96KP6ITNIP3_HUMAN TNFAIP3-interacting protein 3 OS=Homo sapiens GN=TNIP3 PE=1 SV=2	0	1	0.00
POLR3D	P05423R3PC4_HUMAN DNA-directed RNA polymerase II subunit R3C4 OS=Homo sapiens GN=POLR3D PE=1 SV=2	0	1	0.00

NELFA	Q9H3P2NELFA_HUMAN Negative elongation factor A OS=Homo sapiens GN=NELFA PE=1 SV=3	0	3	0.00
FAM184A	Q8N25F184A_HUMAN Protein FAM184A OS=Homo sapiens GN=FAM184A PE=2 SV=3	0	2	0.00
NACC1	Q96RE7NACC1_HUMAN Nucleus accumbens-associated protein 1 OS=Homo sapiens GN=NACC1 PE=1 SV=1	0	2	0.00
SNX17	Q15036SNX17_HUMAN Sorting nexin-17 OS=Homo sapiens GN=SNX17 PE=1 SV=1	0	2	0.00
DLAT	P1051SIOOP2_HUMAN Dihydropyridine-residue acetyltransferase component of pyruvate dehydrogenase complex, mitochondrial OS=Homo sapiens GN=DLAT PE=1 SV=3	0	2	0.00
DEK	P35659DEK_HUMAN Protein DEK OS=Homo sapiens GN=DEK PE=1 SV=1	0	5	0.00
NAMPT	P43490NAMPT_HUMAN Nicotinamide phosphoribosyltransferase OS=Homo sapiens GN=NAMPT PE=1 SV=1	0	8	0.00
FAF2	Q96CS3FAF2_HUMAN FAS-associated factor 2 OS=Homo sapiens GN=FAF2 PE=1 SV=2	0	3	0.00
NEK3	P51956INEK3_HUMAN Serine/threonine-protein kinase Nek3 OS=Homo sapiens GN=NEK3 PE=1 SV=2	0	2	0.00
SEPT7	Q16181SEPT7_HUMAN Septin-7 OS=Homo sapiens GN=SEPT7 PE=1 SV=2	0	1	0.00
IMPACT	Q9P2X3IMPCT_HUMAN Protein IMPACT OS=Homo sapiens GN=IMPACT PE=1 SV=2	0	1	0.00
SNX4	Q95219ISNX4_HUMAN Sorting nexin-4 OS=Homo sapiens GN=SNX4 PE=1 SV=1	0	2	0.00
TERF2IP	Q9NYB0ITE2IP_HUMAN Telomeric repeat-binding factor 2-interacting protein 1 OS=Homo sapiens GN=TERF2IP PE=1 SV=1	0	2	0.00
SDHA	P31040SDHA_HUMAN Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Homo sapiens GN=SDHA PE=1 SV=2	0	8	0.00
ACSM2B	Q68CK6ACSM2B_HUMAN Acyl-coenzyme A synthetase ACSM2B, mitochondrial OS=Homo sapiens GN=ACSM2B PE=1 SV=2	0	5	0.00
GABPA	Q06546GABPA_HUMAN GA-binding protein alpha chain OS=Homo sapiens GN=GABPA PE=1 SV=1	0	2	0.00
SLC38A3	Q99624IS38A3_HUMAN Sodium-coupled neutral amino acid transporter 3 OS=Homo sapiens GN=SLC38A3 PE=1 SV=1	0	2	0.00
ABCB1	P08183MDR1_HUMAN Multidrug resistance protein 1 OS=Homo sapiens GN=ABCB1 PE=1 SV=3	0	2	0.00
GOPC	Q9HD26IGOPC_HUMAN Golgi-associated PDZ and coiled-coil motif-containing protein OS=Homo sapiens GN=GOPC PE=1 SV=1	0	4	0.00
PPM1G	O15355PPM1G_HUMAN Protein phosphatase 1G OS=Homo sapiens GN=PPM1G PE=1 SV=1	0	3	0.00
TUBB8	Q32CM7TBB8_HUMAN Tubulin beta-8 chain OS=Homo sapiens GN=TUBB8 PE=1 SV=2	0	2	0.00
CRAT	P43155CACP_HUMAN Carnitine O-acetyltransferase OS=Homo sapiens GN=CRAT PE=1 SV=5	0	7	0.00
NOTUM	Q6P988NOTUM_HUMAN Palmitoleoyl-protein carboxylesterase NOTUM OS=Homo sapiens GN=NOTUM PE=1 SV=2	0	1	0.00
TDRKH	Q9Y2W6ITDRKH_HUMAN Tudor and KH domain-containing protein OS=Homo sapiens GN=TDRKH PE=1 SV=2	0	2	0.00
CLPX	O76031CLPX_HUMAN ATP-dependent Clp protease ATP-binding subunit clpX-like, mitochondrial OS=Homo sapiens GN=CLPX PE=1 SV=2	0	9	0.00
EIF2A	Q9B744EIF2A_HUMAN Eukaryotic translation initiation factor 2A OS=Homo sapiens GN=EIF2A PE=1 SV=3	0	8	0.00
ATP1A3	P13637IAT1A3_HUMAN Sodium/potassium-transporting ATPase subunit alpha-3 OS=Homo sapiens GN=ATP1A3 PE=1 SV=3	0	4	0.00
LTV1	Q96GA3LTV1_HUMAN Protein LTV1 homolog OS=Homo sapiens GN=LTV1 PE=1 SV=1	0	1	0.00
CPSF7	Q8N684CPSF7_HUMAN Cleavage and polyadenylation specificity factor subunit 7 OS=Homo sapiens GN=CPSF7 PE=1 SV=1	0	5	0.00
SLC1A5	Q15758IAAAT_HUMAN Neutral amino acid transporter B(O) OS=Homo sapiens GN=SLC1A5 PE=1 SV=2	0	1	0.00
NOC4L	Q9BV4HNOC4L_HUMAN Nucleolar complex protein 4 homolog OS=Homo sapiens GN=NOC4L PE=1 SV=1	0	9	0.00
NCL	P19338NCL_HUMAN Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3	0	7	0.00
PRCC	Q92733PRCC_HUMAN Proline-rich protein PRCC OS=Homo sapiens GN=PRCC PE=1 SV=1	0	2	0.00
PRM2	P49643PRM2_HUMAN DNA primase large subunit OS=Homo sapiens GN=PRM2 PE=1 SV=2	0	1	0.00
STK26	Q9P289STK26_HUMAN Serine/threonine-protein kinase 26 OS=Homo sapiens GN=STK26 PE=1 SV=2	0	3	0.00
PDE12	Q6L8Q7PDE12_HUMAN 2',5'-phosphodiesterase 12 OS=Homo sapiens GN=PDE12 PE=1 SV=2	0	3	0.00
PACSN3	Q9UKS6IPACN3_HUMAN Protein kinase C and casein kinase substrate in neurons protein 3 OS=Homo sapiens GN=PACSN3 PE=1 SV=2	0	4	0.00
CTNNB1	P32222CTNB1_HUMAN Catenin beta-1 OS=Homo sapiens GN=CTNNB1 PE=1 SV=1	0	4	0.00
GSK3A	P49840GSK3A_HUMAN Glycogen synthase kinase-3 alpha OS=Homo sapiens GN=GSK3A PE=1 SV=2	0	2	0.00
SPTAN1	Q13813SPTN1_HUMAN Spectrin alpha chain, non-erythrocytic 1 OS=Homo sapiens GN=SPTAN1 PE=1 SV=3	0	7	0.00
UTS2	O95399UTS2_HUMAN Urotesin-2 OS=Homo sapiens GN=UTS2 PE=1 SV=1	0	1	0.00
SLC27A4	Q6P1M0IS27A4_HUMAN Long-chain fatty acid transport protein 4 OS=Homo sapiens GN=SLC27A4 PE=1 SV=1	0	2	0.00
No_GeneID	ADA0A1RRH7IA0A01RRH7_HUMAN Histone H2A OS=Homo sapiens GN=H2A PE=1 SV=3	0	1	0.00
NASP	P49321NASP_HUMAN Nuclear autoantigenic sperm protein OS=Homo sapiens GN=NASP PE=1 SV=2	0	9	0.00
UBQLN1	Q9UMX0UBQLN1_HUMAN Ubiquilin-1 OS=Homo sapiens GN=UBQLN1 PE=1 SV=2	0	2	0.00
DBNL	Q3UJUGDBNL_HUMAN Drebrin-like protein OS=Homo sapiens GN=DBNL PE=1 SV=1	0	3	0.00
SLC39A4	Q6P5W5IS39A4_HUMAN Zinc transporter ZIP4 OS=Homo sapiens GN=SLC39A4 PE=1 SV=3	0	1	0.00
RNF2	Q99496RNF2_HUMAN E3 ubiquitin-protein ligase RING2 OS=Homo sapiens GN=RNF2 PE=1 SV=1	0	2	0.00
DNAJC7	Q99615IDNAJC7_HUMAN DnaJ homolog subfamily C member 7 OS=Homo sapiens GN=DNAJC7 PE=1 SV=2	0	6	0.00
MPP6	Q9NZW5MPP6_HUMAN MAGUK p55 subfamily member 6 OS=Homo sapiens GN=MPP6 PE=1 SV=2	0	2	0.00
DNAJC1	Q96KCBIDNAJC1_HUMAN DnaJ homolog subfamily C member 1 OS=Homo sapiens GN=DNAJC1 PE=1 SV=1	0	2	0.00
SNTB2	Q13425SNTB2_HUMAN Beta-2-syntrophin OS=Homo sapiens GN=SNTB2 PE=1 SV=1	0	1	0.00
APIS	Q9RZ25APIS_HUMAN Apoptosis inhibitor 5 OS=Homo sapiens GN=APIS PE=1 SV=3	0	6	0.00
HNRNPLL	Q8WV99HNRLL_HUMAN Heterogeneous nuclear ribonucleoprotein L-like OS=Homo sapiens GN=HNRNPLL PE=1 SV=1	0	3	0.00
FASTKD2	Q9NY8FADK2_HUMAN FAST kinase domain-containing protein 2, mitochondrial OS=Homo sapiens GN=FASTKD2 PE=1 SV=1	0	5	0.00
NMD3	Q96D46NMD3_HUMAN 60S ribosomal export protein NMD3 OS=Homo sapiens GN=NMD3 PE=1 SV=1	0	2	0.00
UGT2B7	P16662UGT2B7_HUMAN UDP-glucuronosyltransferase 2B7 OS=Homo sapiens GN=UGT2B7 PE=1 SV=1	0	3	0.00
FAM170A	A1A519F170A_HUMAN Protein FAM170A OS=Homo sapiens GN=FAM170A PE=2 SV=1	0	3	0.00
ATXN10	Q9UBB4ATXN10_HUMAN Ataxin-10 OS=Homo sapiens GN=ATXN10 PE=1 SV=1	0	1	0.00
LMF2	Q9BU23LMF2_HUMAN Lipase maturation factor 2 OS=Homo sapiens GN=LMF2 PE=1 SV=2	0	1	0.00
RSRY1	Q96DX4IRSPY_HUMAN RING finger and SPRY domain-containing protein 1 OS=Homo sapiens GN=RSRY1 PE=1 SV=1	0	1	0.00
NMT1	P30419NMT1_HUMAN Glycylpeptide N-tetradecanoyltransferase 1 OS=Homo sapiens GN=NMT1 PE=1 SV=2	0	4	0.00
TXNDC5	Q9NB59TXNDC5_HUMAN Thioredoxin domain-containing protein 5 OS=Homo sapiens GN=TXNDC5 PE=1 SV=2	0	2	0.00
SDPR	O95810SDPR_HUMAN Serum deprivation-response protein OS=Homo sapiens GN=SDPR PE=1 SV=3	0	2	0.00
MAVS	Q72434MAVS_HUMAN Mitochondrial antiviral-signaling protein OS=Homo sapiens GN=MAVS PE=1 SV=2	0	8	0.00
CLU	P10909CLUS_HUMAN Clusterin OS=Homo sapiens GN=CLU PE=1 SV=1	0	3	0.00
COASY	Q13057COASY_HUMAN Bifunctional coenzyme A synthase OS=Homo sapiens GN=COASY PE=1 SV=4	0	4	0.00
PABPC1	P11940PABP1_HUMAN Polyadenylate-binding protein 1 OS=Homo sapiens GN=PABPC1 PE=1 SV=2	0	4	0.00
MYEF2	Q9P2K3MYEF2_HUMAN Myelin expression factor 2 OS=Homo sapiens GN=MYEF2 PE=1 SV=3	0	3	0.00
ARFGAP3	Q9NP61ARFG3_HUMAN ADP-ribosylation factor GTPase-activating protein 3 OS=Homo sapiens GN=ARFGAP3 PE=1 SV=1	0	3	0.00
SLC2A2	P11168GTR2_HUMAN Solute carrier family 2, facilitated glucose transporter member 2 OS=Homo sapiens GN=SLC2A2 PE=1 SV=1	0	1	0.00
RIC8A	Q9NPQ8RIC8A_HUMAN Symplekxin-A OS=Homo sapiens GN=RIC8A PE=1 SV=3	0	1	0.00
SARS2	Q9NP81ISYSM_HUMAN Serine-tRNA ligase, mitochondrial OS=Homo sapiens GN=SARS2 PE=1 SV=1	0	9	0.00
UBXN4	Q92575UBXN4_HUMAN UBX domain-containing protein 4 OS=Homo sapiens GN=UBXN4 PE=1 SV=2	0	4	0.00
RBBP5	Q15291RBBP5_HUMAN Retinoblastoma-binding protein 5 OS=Homo sapiens GN=RBBP5 PE=1 SV=2	0	3	0.00
LGALS3BP	Q08380LG3BP_HUMAN Galectin-3-binding protein OS=Homo sapiens GN=LGALS3BP PE=1 SV=1	0	2	0.00
PAIP1	Q9H074IPAI1_HUMAN Polyadenylate-binding protein-interacting protein 1 OS=Homo sapiens GN=PAIP1 PE=1 SV=1	0	1	0.00
DYNC1L2	O43237DNC1L2_HUMAN Cytoplasmic dynein 1 light intermediate chain 2 OS=Homo sapiens GN=DYNC1L2 PE=1 SV=1	0	4	0.00
POTEF	A5A3E0POTEF_HUMAN POTE ankyrin domain family member F OS=Homo sapiens GN=POTEF PE=1 SV=2	0	3	0.00
CFAP57	Q96MR6ICAF57_HUMAN Cilia- and flagella-associated protein 57 OS=Homo sapiens GN=CFAP57 PE=2 SV=3	0	2	0.00
TXNRD2	Q9NNW7ITXR2_HUMAN Thioredoxin reductase 2, mitochondrial OS=Homo sapiens GN=TXNRD2 PE=1 SV=3	0	4	0.00
FMOS	P49326FMOS_HUMAN Dimethylamine monoxygenase [N-oxide-forming] 5 OS=Homo sapiens GN=FMOS PE=1 SV=2	0	3	0.00
FLNB	O75369FLNB_HUMAN Filamin-B OS=Homo sapiens GN=FLNB PE=1 SV=2	0	7	0.00
SLC38A2	Q96QD8IS38A2_HUMAN Sodium-coupled neutral amino acid transporter 2 OS=Homo sapiens GN=SLC38A2 PE=1 SV=2	0	3	0.00
FAM179A	Q6ZUX3F179A_HUMAN Protein FAM179A OS=Homo sapiens GN=FAM179A PE=2 SV=2	0	4	0.00
ARSA	P15289IARSA_HUMAN Arylsulfatase A OS=Homo sapiens GN=ARSA PE=1 SV=3	0	1	0.00
ASNS	P08243IASNS_HUMAN Asparagine synthetase [glutamine-hydrolyzing] OS=Homo sapiens GN=ASNS PE=1 SV=4	0	6	0.00
PRCP	P42785IPCP_HUMAN Lysosomal Pro-X carboxypeptidase OS=Homo sapiens GN=PRCP PE=1 SV=1	0	3	0.00

RMDN3	Q96TC7/RMD3_HUMAN Regulator of microtubule dynamics protein 3 OS=Homo sapiens GN=RMDN3 PE=1 SV=2	0	10	0.00
HSPA8	P11142/HSPA7C_HUMAN Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1	0	1	0.00
ALDH9A1	P49189/AL9A1_HUMAN 4-trimethylaminobutyraldehyde dehydrogenase OS=Homo sapiens GN=ALDH9A1 PE=1 SV=3	0	4	0.00
CANX	P27824/CALX_HUMAN Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2	0	4	0.00
ENDOD1	O94919/ENDOD1_HUMAN Endonuclease domain-containing 1 protein OS=Homo sapiens GN=ENDOD1 PE=1 SV=2	0	2	0.00
UGT2A3	QJUMW9/UGT2A3_HUMAN UDP-glucuronosyltransferase 2A3 OS=Homo sapiens GN=UGT2A3 PE=2 SV=2	0	2	0.00
POLR3C	Q9BU4H/RPC3_HUMAN DNA-directed RNA polymerase III subunit RPC3 OS=Homo sapiens GN=POLR3C PE=1 SV=1	0	2	0.00
C14orf159	Q723D6/C159_HUMAN UPF0317 protein C14orf159, mitochondrial OS=Homo sapiens GN=C14orf159 PE=1 SV=2	0	2	0.00
SLC6A8	P48029/SLC6A8_HUMAN Sodium- and chloride-dependent creatine transporter 1 OS=Homo sapiens GN=SLC6A8 PE=1 SV=1	0	1	0.00
G3BP2	Q9UN86/G3BP2_HUMAN Ras GTPase-activating protein-binding protein 2 OS=Homo sapiens GN=G3BP2 PE=1 SV=2	0	1	0.00
LSR	Q86X29/LSR_HUMAN Lipolysis-stimulated lipoprotein receptor OS=Homo sapiens GN=LSR PE=1 SV=4	0	5	0.00
SOWAHC	Q53LP3/SWAHC_HUMAN Ankyrin repeat domain-containing protein SOWAHC OS=Homo sapiens GN=SOWAHC PE=1 SV=1	0	2	0.00
UTP4	Q969X6/UTP4_HUMAN U3 small nucleolar RNA-associated protein 4 homolog OS=Homo sapiens GN=UTP4 PE=1 SV=1	0	2	0.00
ALDOA	P04075/ALDOA_HUMAN Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2	0	1	0.00
SEPT9	Q9UHD8/SEPT9_HUMAN Septin-9 OS=Homo sapiens GN=SEPT9 PE=1 SV=2	0	2	0.00
SFPQ	P23246/SFPQ_HUMAN Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPQ PE=1 SV=2	0	6	0.00
FUS	P35637/FUS_HUMAN RNA-binding protein FUS OS=Homo sapiens GN=FUS PE=1 SV=1	0	7	0.00
ANO10	Q9NWT1/ANO10_HUMAN Anoctamin-10 OS=Homo sapiens GN=ANO10 PE=1 SV=2	0	2	0.00
CXADR	P78310/CXAR_HUMAN Coxsackievirus and adenovirus receptor OS=Homo sapiens GN=CXADR PE=1 SV=1	0	5	0.00
MDH2	P40926/MDHM_HUMAN Malate dehydrogenase, mitochondrial OS=Homo sapiens GN=MDH2 PE=1 SV=3	0	1	0.00
PSMC5	PE2195/PSR8_HUMAN 26S protease regulatory subunit 8 OS=Homo sapiens GN=PSMC5 PE=1 SV=1	0	2	0.00
FAM98A	Q8NCAS/FAM98A_HUMAN Protein FAM98A OS=Homo sapiens GN=FAM98A PE=1 SV=1	0	2	0.00
RLM	Q9NVW2/RNF12_HUMAN E3 ubiquitin-protein ligase RLM OS=Homo sapiens GN=RLM PE=1 SV=3	0	1	0.00
NECTN2	Q92692/NECT2_HUMAN Nectin-2 OS=Homo sapiens GN=NECTN2 PE=1 SV=1	0	1	0.00
LMNB2	Q03252/LMNB2_HUMAN Lamin-B2 OS=Homo sapiens GN=LMNB2 PE=1 SV=4	0	2	0.00
AAAS	Q9NRG9/AAAS_HUMAN Aladin OS=Homo sapiens GN=AAAS PE=1 SV=1	0	4	0.00
ZDHHC8	Q9ULC8/ZDHHC8_HUMAN Probable palmitoyltransferase ZDHHC8 OS=Homo sapiens GN=ZDHHC8 PE=1 SV=3	0	2	0.00
GHDC	Q8N2G8/GHDC_HUMAN GH3 domain-containing protein OS=Homo sapiens GN=GHDC PE=1 SV=2	0	2	0.00
DFNA5	O60443/DFNA5_HUMAN Non-syndromic hearing impairment protein 5 OS=Homo sapiens GN=DFNA5 PE=1 SV=2	0	2	0.00
EML4	Q9HC35/EMAL4_HUMAN Echinoderm microtubule-associated protein-like 4 OS=Homo sapiens GN=EML4 PE=1 SV=3	0	6	0.00
NELL1	Q92832/NELL1_HUMAN Protein kinase C-binding protein NELL1 OS=Homo sapiens GN=NELL1 PE=1 SV=4	0	3	0.00
RRP1	P56182/RRP1_HUMAN Ribosomal RNA processing protein 1 homolog A OS=Homo sapiens GN=RRP1 PE=1 SV=1	0	3	0.00
PPM1F	P49593/PPM1F_HUMAN Protein phosphatase 1F OS=Homo sapiens GN=PPM1F PE=1 SV=3	0	2	0.00
CPSF3L	Q5TA45/INT11_HUMAN Integrator complex subunit 11 OS=Homo sapiens GN=CPSF3L PE=1 SV=2	0	2	0.00
NUP54	Q723B4/NUP54_HUMAN Nucleoporin p54 OS=Homo sapiens GN=NUP54 PE=1 SV=2	0	6	0.00
ELP3	Q9H9T3/ELP3_HUMAN Elongator complex protein 3 OS=Homo sapiens GN=ELP3 PE=1 SV=2	0	3	0.00
APMAP	Q9HDC9/APMAP_HUMAN Adipocyte plasma membrane-associated protein OS=Homo sapiens GN=APMAP PE=1 SV=2	0	1	0.00
IMMT	Q16891/IMC60_HUMAN MICOS complex subunit MIC60 OS=Homo sapiens GN=IMMT PE=1 SV=1	0	1	0.00
SCD	Q00767/ACOD_HUMAN Acyl-CoA desaturase OS=Homo sapiens GN=SCD PE=1 SV=2	0	1	0.00
MAP4	P27816/MAP4_HUMAN Microtubule-associated protein 4 OS=Homo sapiens GN=MAP4 PE=1 SV=3	0	3	0.00
OXCT1	P55809/ISCO11_HUMAN Succinyl-CoA:3-ketoacid coenzyme A transferase 1, mitochondrial OS=Homo sapiens GN=OXCT1 PE=1 SV=1	0	1	0.00
XYLB	O75191/XYLB_HUMAN Xylulose kinase OS=Homo sapiens GN=XYLB PE=1 SV=3	0	6	0.00
TBL1XR1	Q9BZK7/TBL1R_HUMAN F-box-like/WD repeat-containing protein TBL1XR1 OS=Homo sapiens GN=TBL1XR1 PE=1 SV=1	0	3	0.00
EIF5	P55010/EIF5_HUMAN Eukaryotic translation initiation factor 5 OS=Homo sapiens GN=EIF5 PE=1 SV=2	0	6	0.00
DMAP1	Q9NPF5/DMAP1_HUMAN DNA methyltransferase 1-associated protein 1 OS=Homo sapiens GN=DMAP1 PE=1 SV=1	0	5	0.00
CSTF2	P33240/CSTF2_HUMAN Cleavage stimulation factor subunit 2 OS=Homo sapiens GN=CSTF2 PE=1 SV=1	0	4	0.00
ALDOC	P09972/ALDOC_HUMAN Fructose-bisphosphate aldolase C OS=Homo sapiens GN=ALDOC PE=1 SV=2	0	1	0.00
KIAA2013	Q8IYS2/K2013_HUMAN Uncharacterized protein KIAA2013 OS=Homo sapiens GN=KIAA2013 PE=1 SV=1	0	2	0.00
RARS	P54136/SYRC_HUMAN Arginine-tRNA ligase, cytoplasmic OS=Homo sapiens GN=RARS PE=1 SV=2	0	1	0.00
LRPPRC	P42704/LRPPRC_HUMAN Leucine-rich PPR motif-containing protein, mitochondrial OS=Homo sapiens GN=LRPPRC PE=1 SV=3	0	4	0.00
SERBP1	Q8NC51/PAIRB_HUMAN Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERBP1 PE=1 SV=2	0	6	0.00
TOMM70	O94826/TOM70_HUMAN Mitochondrial import receptor subunit TOM70 OS=Homo sapiens GN=TOMM70 PE=1 SV=1	0	1	0.00
RNF149	Q8NC42/RNF149_HUMAN E3 ubiquitin-protein ligase RNF149 OS=Homo sapiens GN=RNF149 PE=2 SV=2	0	2	0.00
DNAJC11	Q9NVH1/DJC11_HUMAN DnaJ homolog subfamily C member 11 OS=Homo sapiens GN=DNAJC11 PE=1 SV=2	0	4	0.00
CBSL	PODN79/CBSL_HUMAN Cystathionine beta-synthase-like protein OS=Homo sapiens GN=CBSL PE=1 SV=1	0	5	0.00
FOXRED1	Q96CU9/IFXR1_HUMAN FAD-dependent oxidoreductase domain-containing protein 1 OS=Homo sapiens GN=FOXRED1 PE=1 SV=2	0	1	0.00
NF1	P21359/NF1_HUMAN Neurofibromin OS=Homo sapiens GN=NF1 PE=1 SV=2	0	4	0.00
CLCC1	Q96566/CLCC1_HUMAN Chloride channel CLIC-like protein 1 OS=Homo sapiens GN=CLCC1 PE=1 SV=1	0	2	0.00
NUP62	P37198/NUP62_HUMAN Nuclear pore glycoprotein p62 OS=Homo sapiens GN=NUP62 PE=1 SV=3	0	3	0.00
GGT3P	A6NGU5/GGT3_HUMAN Putative gamma-glutamyltranspeptidase 3 OS=Homo sapiens GN=GGT3P PE=5 SV=2	0	2	0.00
FXR1	P51114/FXR1_HUMAN Fragile X mental retardation syndrome-related protein 1 OS=Homo sapiens GN=FXR1 PE=1 SV=3	0	2	0.00
EEF1G	P26641/EEF1G_HUMAN Elongation factor 1-gamma OS=Homo sapiens GN=EEF1G PE=1 SV=3	0	2	0.00
TRPC5	Q9UL62/TRPC5_HUMAN Short transient receptor potential channel 5 OS=Homo sapiens GN=TRPC5 PE=1 SV=1	0	3	0.00
SLC25A3	Q00325/SIMPCP_HUMAN Phosphate carrier protein, mitochondrial OS=Homo sapiens GN=SLC25A3 PE=1 SV=2	0	1	0.00
CTTN	Q14247/SRCS_HUMAN Src substrate cortactin OS=Homo sapiens GN=CTTN PE=1 SV=2	0	4	0.00
WARS	P23381/SIYWC_HUMAN Tryptophan-tRNA ligase, cytoplasmic OS=Homo sapiens GN=WARS PE=1 SV=2	0	10	0.00
IGSF1	Q8N6C5/IGSF1_HUMAN Immunoglobulin superfamily member 1 OS=Homo sapiens GN=IGSF1 PE=1 SV=3	0	2	0.00
RCOR1	Q9UKL0/RCOR1_HUMAN REST corepressor 1 OS=Homo sapiens GN=RCOR1 PE=1 SV=1	0	3	0.00
CARS2	Q9HA77/SIYCM_HUMAN Probable cysteine-tRNA ligase, mitochondrial OS=Homo sapiens GN=CARS2 PE=1 SV=1	0	3	0.00
ACAD10	Q6JQN1/ACAD10_HUMAN Acyl-CoA dehydrogenase family member 10 OS=Homo sapiens GN=ACAD10 PE=1 SV=1	0	3	0.00
AHNAK	Q09666/AHNAK_HUMAN Neuroblast differentiation-associated protein AHNAK OS=Homo sapiens GN=AHNAK PE=1 SV=2	0	8	0.00
SRSF5	Q13243/SRSF5_HUMAN Serine/arginine-rich splicing factor 5 OS=Homo sapiens GN=SRSF5 PE=1 SV=1	0	3	0.00
ATL2	Q8NH99/ATLA2_HUMAN Atlastin-2 OS=Homo sapiens GN=ATL2 PE=1 SV=2	0	3	0.00
ZNF815P	ABK554/ZNF815_HUMAN Putative protein ZNF815 OS=Homo sapiens GN=ZNF815P PE=5 SV=1	0	1	0.00
MGAT2	Q10469/MGAT2_HUMAN Alpha-1,6-mannosyl-glycoprotein 2-beta-N-acetylglucosaminyltransferase OS=Homo sapiens GN=MGAT2 PE=1 SV=1	0	2	0.00