

Gene Name	Name of Protein (UniProtID_SPECIES, protein name, origin species, gene name)	Peptide Count (IP)	Peptide Count (IgG)	Fold Enrichment
MATR3	P43243IMATR3_HUMAN Matrin-3 OS=Homo sapiens GN=MATR3 PE=1 SV=2	89	0	Not in IgG control
MCCC1	Q96RQ3IMCCA_HUMAN Methylcrotonoyl-CoA carboxylase subunit alpha, mitochondrial OS=Homo sapiens GN=MCCC1 PE=1 SV=3	28	0	Not in IgG control
MCCC2	Q9HCCOIMCCB_HUMAN Methylcrotonoyl-CoA carboxylase beta chain, mitochondrial OS=Homo sapiens GN=MCCC2 PE=1 SV=1	10	0	Not in IgG control
UBA1	P22314IUBA1_HUMAN Ubiquitin-like modifier-activating enzyme 1 OS=Homo sapiens GN=UBA1 PE=1 SV=3	10	0	Not in IgG control
FN1	P02751FINC_HUMAN Fibronectin OS=Homo sapiens GN=FN1 PE=1 SV=4	8	0	Not in IgG control
ATP2A2	P16615IAT2A2_HUMAN Sarcoplasmic/endoplasmic reticulum calcium ATPase 2 OS=Homo sapiens GN=ATP2A2 PE=1 SV=1	7	0	Not in IgG control
PC	P11498IPYC_HUMAN Pyruvate carboxylase, mitochondrial OS=Homo sapiens GN=PC PE=1 SV=2	7	0	Not in IgG control
ILF3	Q12906ILF3_HUMAN Interleukin enhancer-binding factor 3 OS=Homo sapiens GN=ILF3 PE=1 SV=3	7	0	Not in IgG control
SFPQ	P23246ISFPQ_HUMAN Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPQ PE=1 SV=2	6	0	Not in IgG control
SND1	Q7KZF4ISND1_HUMAN Staphylococcal nuclease domain-containing protein 1 OS=Homo sapiens GN=SND1 PE=1 SV=1	5	0	Not in IgG control
No_GeneID	IA0A180GU03_HUMAN Uncharacterized protein OS=Homo sapiens PE=4 SV=1	5	0	Not in IgG control
ANK2	Q01484IANK2_HUMAN Ankyrin-2 OS=Homo sapiens GN=ANK2 PE=1 SV=4	5	0	Not in IgG control
DSC1	Q08554IDSC1_HUMAN Desmocollin-1 OS=Homo sapiens GN=DSC1 PE=1 SV=2	5	0	Not in IgG control
COPB1	P53618ICOPB_HUMAN Coatamer subunit beta OS=Homo sapiens GN=COPB1 PE=1 SV=3	5	0	Not in IgG control
EIF3B	P55884IEIF3B_HUMAN Eukaryotic translation initiation factor 3 subunit B OS=Homo sapiens GN=EIF3B PE=1 SV=3	5	0	Not in IgG control
EML4	Q9HC35IEMAL4_HUMAN Echinoderm microtubule-associated protein-like 4 OS=Homo sapiens GN=EML4 PE=1 SV=3	5	0	Not in IgG control
CCAR2	Q8N163ICCAR2_HUMAN Cell cycle and apoptosis regulator protein 2 OS=Homo sapiens GN=CCAR2 PE=1 SV=2	4	0	Not in IgG control
EFTUD2	Q15029IUS11_HUMAN 116 kDa U5 small nuclear ribonucleoprotein component OS=Homo sapiens GN=EFTUD2 PE=1 SV=1	4	0	Not in IgG control
DHX15	O43143IDHX15_HUMAN Pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15 OS=Homo sapiens GN=DHX15 PE=1 SV=2	4	0	Not in IgG control
HSPA1A	PODMV8IHS71A_HUMAN Heat shock 70 kDa protein 1A OS=Homo sapiens GN=HSPA1A PE=1 SV=1	4	0	Not in IgG control
ARG1	P05089IARG1_HUMAN Arginase-1 OS=Homo sapiens GN=ARG1 PE=1 SV=2	4	0	Not in IgG control
ANXA2	PO7355IANXA2_HUMAN Annexin A2 OS=Homo sapiens GN=ANXA2 PE=1 SV=2	4	0	Not in IgG control
XRCC5	P13010IXRCC5_HUMAN X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3	4	0	Not in IgG control
SYNE1	Q8NF91ISYNE1_HUMAN Nesprin-1 OS=Homo sapiens GN=SYNE1 PE=1 SV=4	4	0	Not in IgG control
VPS35	Q96QK1IVPS35_HUMAN Vacuolar protein sorting-associated protein 35 OS=Homo sapiens GN=VPS35 PE=1 SV=2	3	0	Not in IgG control
PRPF6	O94906IPRP6_HUMAN Pre-mRNA-processing factor 6 OS=Homo sapiens GN=PRPF6 PE=1 SV=1	3	0	Not in IgG control
KIF1B	O60333KIF1B_HUMAN Kinesin-like protein KIF1B OS=Homo sapiens GN=KIF1B PE=1 SV=5	3	0	Not in IgG control
ESYT2	A0FGR8IESYT2_HUMAN Extended synaptotagmin-2 OS=Homo sapiens GN=ESYT2 PE=1 SV=1	3	0	Not in IgG control
KIF5B	P33176IKINH_HUMAN Kinesin-1 heavy chain OS=Homo sapiens GN=KIF5B PE=1 SV=1	3	0	Not in IgG control
AZGP1	P25311IZA2G_HUMAN Zinc-alpha-2-glycoprotein OS=Homo sapiens GN=AZGP1 PE=1 SV=2	3	0	Not in IgG control
WDR81	Q562E7IHDR81_HUMAN WD repeat-containing protein 81 OS=Homo sapiens GN=WDR81 PE=1 SV=2	3	0	Not in IgG control
RBM14	Q96PK6IRBM14_HUMAN RNA-binding protein 14 OS=Homo sapiens GN=RBM14 PE=1 SV=2	3	0	Not in IgG control
HNRNPM	P52272IHNRPM_HUMAN Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3	3	0	Not in IgG control
GIMAP8	Q8ND71IGIMAP8_HUMAN GTPase IMAP family member 8 OS=Homo sapiens GN=GIMAP8 PE=2 SV=2	2	0	Not in IgG control
IGHA1	PO1876IIGHA1_HUMAN Ig alpha-1 chain C region OS=Homo sapiens GN=IGHA1 PE=1 SV=2	2	0	Not in IgG control
PKM	P14618IKPYM_HUMAN Pyruvate kinase PKM OS=Homo sapiens GN=PKM PE=1 SV=4	2	0	Not in IgG control
HSPA5	P11021IGRP78_HUMAN 78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2	2	0	Not in IgG control
COL4A1	P02462IC04A1_HUMAN Collagen alpha-1(IV) chain OS=Homo sapiens GN=COL4A1 PE=1 SV=3	2	0	Not in IgG control
SPDYE1	Q8NFV5ISPDYE1_HUMAN Speedy protein E1 OS=Homo sapiens GN=SPDYE1 PE=2 SV=3	2	0	Not in IgG control
PDZRN4	Q6ZMN7IPZRN4_HUMAN PDZ domain-containing RING finger protein 4 OS=Homo sapiens GN=PDZRN4 PE=2 SV=3	2	0	Not in IgG control
DHX9	Q08211IDHX9_HUMAN ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4	2	0	Not in IgG control
CAT	PO4040ICATA_HUMAN Catalase OS=Homo sapiens GN=CAT PE=1 SV=3	2	0	Not in IgG control
MICALL2	Q8IY33IMILK2_HUMAN MICAL-like protein 2 OS=Homo sapiens GN=MICALL2 PE=1 SV=1	2	0	Not in IgG control
MYBPC3	Q14896IMYPC3_HUMAN Myosin-binding protein C, cardiac-type OS=Homo sapiens GN=MYBPC3 PE=1 SV=4	2	0	Not in IgG control
SERPINB12	Q96P63ISPB12_HUMAN Serpin B12 OS=Homo sapiens GN=SERPINB12 PE=1 SV=1	2	0	Not in IgG control
TADA2B	Q86TJ2ITAD2B_HUMAN Transcriptional adaptor 2-beta OS=Homo sapiens GN=TADA2B PE=1 SV=2	2	0	Not in IgG control
SEC23A	Q15436IS23A_HUMAN Protein transport protein Sec23A OS=Homo sapiens GN=SEC23A PE=1 SV=2	2	0	Not in IgG control
SMG1	Q96Q15ISMG1_HUMAN Serine/threonine-protein kinase SMG1 OS=Homo sapiens GN=SMG1 PE=1 SV=3	2	0	Not in IgG control
COL28A1	Q2UY09ICOSA1_HUMAN Collagen alpha-1 (XXVIII) chain OS=Homo sapiens GN=COL28A1 PE=2 SV=2	2	0	Not in IgG control
ZNF804B	A4D1E1IZ804B_HUMAN Zinc finger protein 804B OS=Homo sapiens GN=ZNF804B PE=2 SV=2	2	0	Not in IgG control
SHROOM2	Q13796ISHRM2_HUMAN Protein Shroom2 OS=Homo sapiens GN=SHROOM2 PE=1 SV=1	2	0	Not in IgG control
ICE2	Q659A1ICE2_HUMAN Little elongation complex subunit 2 OS=Homo sapiens GN=ICE2 PE=1 SV=2	2	0	Not in IgG control
CYP2J2	P51589ICP2J2_HUMAN Cytochrome P450 2J2 OS=Homo sapiens GN=CYP2J2 PE=1 SV=2	2	0	Not in IgG control
BOD1L1	Q8NFC6IBD1L1_HUMAN Biorientation of chromosomes in cell division protein 1-like 1 OS=Homo sapiens GN=BOD1L1 PE=1 SV=2	2	0	Not in IgG control
CHD3	Q12873ICHD3_HUMAN Chromodomain-helicase-DNA-binding protein 3 OS=Homo sapiens GN=CHD3 PE=1 SV=3	2	0	Not in IgG control
HK2	P52789IHXK2_HUMAN Hexokinase-2 OS=Homo sapiens GN=HK2 PE=1 SV=2	2	0	Not in IgG control
TGM1	P22735ITGM1_HUMAN Protein-glutamine gamma-glutamyltransferase K OS=Homo sapiens GN=TGM1 PE=1 SV=4	2	0	Not in IgG control
TRIM28	Q13263ITIF1B_HUMAN Transcription intermediary factor 1-beta OS=Homo sapiens GN=TRIM28 PE=1 SV=5	2	0	Not in IgG control
PTGFRN	Q9P2B2IFPRP_HUMAN Prostaglandin F2 receptor negative regulator OS=Homo sapiens GN=PTGFRN PE=1 SV=2	1	0	Not in IgG control
STAT1	P42224ISTAT1_HUMAN Signal transducer and activator of transcription 1-alpha/beta OS=Homo sapiens GN=STAT1 PE=1 SV=2	1	0	Not in IgG control
RBM28	Q9NW13IRBM28_HUMAN RNA-binding protein 28 OS=Homo sapiens GN=RBM28 PE=1 SV=3	1	0	Not in IgG control
ATAD3B	Q5T9A4IATD3B_HUMAN ATPase family AAA domain-containing protein 3B OS=Homo sapiens GN=ATAD3B PE=1 SV=1	1	0	Not in IgG control
EWSR1	Q01844IEW5_HUMAN RNA-binding protein EWS OS=Homo sapiens GN=EWSR1 PE=1 SV=1	1	0	Not in IgG control
KATNAL2	Q8IYT4IKATL2_HUMAN Katanin p60 ATPase-containing subunit A-like 2 OS=Homo sapiens GN=KATNAL2 PE=2 SV=3	1	0	Not in IgG control
HBA1	P69905IHBA_HUMAN Hemoglobin subunit alpha OS=Homo sapiens GN=HBA1 PE=1 SV=2	1	0	Not in IgG control
SLC05A1	Q9H2Y9IS05A1_HUMAN Solute carrier organic anion transporter family member 5A1 OS=Homo sapiens GN=SLC05A1 PE=2 SV=2	1	0	Not in IgG control
No_GeneID	MOQZK8_HUMAN Uncharacterized protein OS=Homo sapiens PE=4 SV=1	1	0	Not in IgG control

BLID	QBIZ5YBLID_HUMAN BH3-like motif-containing cell death inducer OS=Homo sapiens GN=BLID PE=1 SV=2	1	0	Not in IgG control
FCGR1B	Q92637IFCGRB_HUMAN High affinity immunoglobulin gamma Fc receptor 1B OS=Homo sapiens GN=FCGR1B PE=2 SV=1	1	0	Not in IgG control
APOD	P05090APOD_HUMAN Apolipoprotein D OS=Homo sapiens GN=APOD PE=1 SV=1	1	0	Not in IgG control
DUSP28	Q460W2IDUS28_HUMAN Dual specificity phosphatase 28 OS=Homo sapiens GN=DUSP28 PE=2 SV=1	1	0	Not in IgG control
VAT1	Q99536VAT1_HUMAN Synaptic vesicle membrane protein VAT-1 homolog OS=Homo sapiens GN=VAT1 PE=1 SV=2	1	0	Not in IgG control
HDDC2	Q724H3HDDC2_HUMAN HD domain-containing protein 2 OS=Homo sapiens GN=HDDC2 PE=1 SV=1	1	0	Not in IgG control
CYP26B1	Q9NR63ICP26B_HUMAN Cytochrome P450 26B1 OS=Homo sapiens GN=CYP26B1 PE=1 SV=1	1	0	Not in IgG control
SLC22A10	Q63ZE4IS22AA_HUMAN Solute carrier family 22 member 10 OS=Homo sapiens GN=SLC22A10 PE=2 SV=2	1	0	Not in IgG control
CLDN12	P56749ICLD12_HUMAN Claudin-12 OS=Homo sapiens GN=CLDN12 PE=1 SV=1	1	0	Not in IgG control
HSPD1	P10809ICH60_HUMAN 60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2	1	0	Not in IgG control
ZNF839	A8KOR7IZN839_HUMAN Zinc finger protein 839 OS=Homo sapiens GN=ZNF839 PE=2 SV=1	1	0	Not in IgG control
QSOX2	Q6ZRP7IQSOX2_HUMAN Sulfhydryl oxidase 2 OS=Homo sapiens GN=QSOX2 PE=1 SV=3	1	0	Not in IgG control
LYZ	P61626ILYSC_HUMAN Lysozyme C OS=Homo sapiens GN=LYZ PE=1 SV=1	1	0	Not in IgG control
PRDX2	P32119IPRDX2_HUMAN Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5	1	0	Not in IgG control
C9orf139	Q6ZV77IC139_HUMAN Uncharacterized protein C9orf139 OS=Homo sapiens GN=C9orf139 PE=2 SV=1	1	0	Not in IgG control
PXMP2	Q9NR77IPXMP2_HUMAN Peroxisomal membrane protein 2 OS=Homo sapiens GN=PXMP2 PE=1 SV=3	1	0	Not in IgG control
PLOD1	Q02809IPLD1_HUMAN Procollagen-lysine,2-oxoglutarate 5-dioxygenase 1 OS=Homo sapiens GN=PLOD1 PE=1 SV=2	1	0	Not in IgG control
TEX10	Q9NXF1ITEX10_HUMAN Testis-expressed sequence 10 protein OS=Homo sapiens GN=TEX10 PE=1 SV=2	1	0	Not in IgG control
DDX5	P17844IDDX5_HUMAN Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 PE=1 SV=1	1	0	Not in IgG control
TNPO1	Q92973ITNPO1_HUMAN Transportin-1 OS=Homo sapiens GN=TNPO1 PE=1 SV=2	1	0	Not in IgG control
UPP2	095045IUPP2_HUMAN Uridine phosphorylase 2 OS=Homo sapiens GN=UPP2 PE=1 SV=1	1	0	Not in IgG control
ASB4	Q9Y574IASB4_HUMAN Ankyrin repeat and SOCS box protein 4 OS=Homo sapiens GN=ASB4 PE=2 SV=1	1	0	Not in IgG control
AXL	P30530IUF0_HUMAN Tyrosine-protein kinase receptor UFO OS=Homo sapiens GN=AXL PE=1 SV=3	1	0	Not in IgG control
MOV10L1	Q9BXT6IM10L1_HUMAN RNA helicase Mov10L1 OS=Homo sapiens GN=MOV10L1 PE=2 SV=1	1	0	Not in IgG control
CREBRF	Q8IUR6ICREBF_HUMAN CREB3 regulatory factor OS=Homo sapiens GN=CREBRF PE=1 SV=2	1	0	Not in IgG control
REXO1L1P	Q8IX06IGOR_HUMAN Putative exonuclease GOR OS=Homo sapiens GN=REXO1L1P PE=5 SV=2	1	0	Not in IgG control
MSH2	P43246IMSH2_HUMAN DNA mismatch repair protein Msh2 OS=Homo sapiens GN=MSH2 PE=1 SV=1	1	0	Not in IgG control
S100A7	P31151IS10A7_HUMAN Protein S100-A7 OS=Homo sapiens GN=S100A7 PE=1 SV=4	1	0	Not in IgG control
NAT10	Q9H0A0INAT10_HUMAN RNA cytidine acetyltransferase OS=Homo sapiens GN=NAT10 PE=1 SV=2	1	0	Not in IgG control
TGM3	Q08188ITGM3_HUMAN Protein-glutamine gamma-glutamyltransferase E OS=Homo sapiens GN=TGM3 PE=1 SV=4	4	1	4.00
CHD3	Q12873ICH3_HUMAN Chromodomain-helicase-DNA-binding protein 3 OS=Homo sapiens GN=CHD3 PE=1 SV=3	4	1	4.00
VCP	P55072ITERA_HUMAN Transitional endoplasmic reticulum ATPase OS=Homo sapiens GN=VCP PE=1 SV=4	9	3	3.00
CASP14	P31944ICASPE_HUMAN Caspase-14 OS=Homo sapiens GN=CASP14 PE=1 SV=2	3	1	3.00
UBB	POCG47IUBB_HUMAN Polyubiquitin-B OS=Homo sapiens GN=UBB PE=1 SV=1	3	1	3.00
DSP	P15924IDESP_HUMAN Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3	17	7	2.43
DCD	P81605IDCD_HUMAN Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2	7	3	2.33
MTHFD1	P11586ICTC_HUMAN C-1-tetrahydrofolate synthase, cytoplasmic OS=Homo sapiens GN=MTHFD1 PE=1 SV=3	7	3	2.33
JUP	P14923IPLAK_HUMAN Junction plakoglobin OS=Homo sapiens GN=JUP PE=1 SV=3	9	4	2.25
GAPDH	P04406IG3P_HUMAN Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	4	2	2.00
CDSN	Q15517ICDSN_HUMAN Corneodesmosin OS=Homo sapiens GN=CDSN PE=1 SV=3	2	1	2.00
CDK16	Q0536ICDK16_HUMAN Cyclin-dependent kinase 16 OS=Homo sapiens GN=CDK16 PE=1 SV=1	2	1	2.00
SERPINA12	Q8IW75ISPA12_HUMAN Serpin A12 OS=Homo sapiens GN=SERPINA12 PE=1 SV=1	2	1	2.00
S100A9	P06702IS10A9_HUMAN Protein S100-A9 OS=Homo sapiens GN=S100A9 PE=1 SV=1	2	1	2.00
TUBA1B	P68363ITBA1B_HUMAN Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1	2	1	2.00
CSTA	P01040ICYTA_HUMAN Cystatin-A OS=Homo sapiens GN=CSTA PE=1 SV=1	2	1	2.00
EEF2	P13639IEF2_HUMAN Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4	15	9	1.67
DSG1	Q02413IDSG1_HUMAN Desmoglein-1 OS=Homo sapiens GN=DSG1 PE=1 SV=2	10	6	1.67
FHAD1	B1AJZ9IFHAD1_HUMAN Forkhead-associated domain-containing protein 1 OS=Homo sapiens GN=FHAD1 PE=2 SV=2	3	2	1.50
ACTB	P60709IACTB_HUMAN Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	4	3	1.33
HSP90AA1	P07900IHS90A_HUMAN Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5	4	3	1.33
ALDH18A1	P54886IPSCS_HUMAN Delta-1-pyrroline-5-carboxylate synthase OS=Homo sapiens GN=ALDH18A1 PE=1 SV=2	5	4	1.25
HNRNPU	Q00839IHNRPU_HUMAN Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU PE=1 SV=6	6	6	1.00
PIP	P12273IPIP_HUMAN Prolactin-inducible protein OS=Homo sapiens GN=PIP PE=1 SV=1	3	3	1.00
S100A8	P05109IS10A8_HUMAN Protein S100-A8 OS=Homo sapiens GN=S100A8 PE=1 SV=1	2	2	1.00
FLG2	Q5D862IFLA2_HUMAN Filaggrin-2 OS=Homo sapiens GN=FLG2 PE=1 SV=1	1	1	1.00
TRAP1	Q12931ITRAP1_HUMAN Heat shock protein 75 kDa, mitochondrial OS=Homo sapiens GN=TRAP1 PE=1 SV=3	1	1	1.00
NFIB	O00712INFIB_HUMAN Nuclear factor 1 B-type OS=Homo sapiens GN=NFIB PE=1 SV=2	1	1	1.00
SHC1	P29353ISHC1_HUMAN SHC-transforming protein 1 OS=Homo sapiens GN=SHC1 PE=1 SV=4	1	1	1.00
GSG2	Q8TF76IHASP_HUMAN Serine/threonine-protein kinase haspin OS=Homo sapiens GN=GSG2 PE=1 SV=3	1	1	1.00
ARID3B	Q8IYV6IARI3B_HUMAN AT-rich interactive domain-containing protein 3B OS=Homo sapiens GN=ARID3B PE=1 SV=2	1	1	1.00
IARS2	Q9NSE4ISYM_HUMAN Isoleucine-tRNA ligase, mitochondrial OS=Homo sapiens GN=IARS2 PE=1 SV=2	1	1	1.00
RASSF10	A6NK89IRASFA_HUMAN Ras association domain-containing protein 10 OS=Homo sapiens GN=RASSF10 PE=2 SV=3	1	1	1.00
HSP90AB1	P08238IHS90B_HUMAN Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4	21	22	0.95
IGHG1	P01857IGHG1_HUMAN Ig gamma-1 chain C region OS=Homo sapiens GN=IGHG1 PE=1 SV=1	3	4	0.75
MARS	P56192ISYMC_HUMAN Methionine-tRNA ligase, cytoplasmic OS=Homo sapiens GN=MARS PE=1 SV=2	2	3	0.67
GANAB	Q14697IGANAB_HUMAN Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB PE=1 SV=3	2	3	0.67
HSP90B1	P14625IENPL_HUMAN Endoplasmic reticulum chaperone protein OS=Homo sapiens GN=HSP90B1 PE=1 SV=1	8	13	0.62
NCL	P19338INCL_HUMAN Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3	4	8	0.50

HRNR	Q86Y23IHORN_HUMAN Hornerin OS=Homo sapiens GN=HRNR PE=1 SV=2	3	6	0.50
ZC3H12C	Q9C0D7IZC12C_HUMAN Probable ribonuclease ZC3H12C OS=Homo sapiens GN=ZC3H12C PE=1 SV=2	1	2	0.50
C2CD2L	O14523IC2C2L_HUMAN C2 domain-containing protein 2-like OS=Homo sapiens GN=C2CD2L PE=1 SV=3	1	2	0.50
ALB	P02768IALBU_HUMAN Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	7	22	0.32
ATP1A1	P05023IAT1A1_HUMAN Sodium/potassium-transporting ATPase subunit alpha-1 OS=Homo sapiens GN=ATP1A1 PE=1 SV=1	2	8	0.25
PAH	P00439IPH4H_HUMAN Phenylalanine-4-hydroxylase OS=Homo sapiens GN=PAH PE=1 SV=1	0	1	0.00
PSD	A5PKW4IPSD1_HUMAN PH and SEC7 domain-containing protein 1 OS=Homo sapiens GN=PSD PE=1 SV=2	0	1	0.00
ABCA12	Q86UK0IABAC_C_HUMAN ATP-binding cassette sub-family A member 12 OS=Homo sapiens GN=ABCA12 PE=1 SV=3	0	1	0.00
KPNB1	Q14974IIMB1_HUMAN Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2	0	1	0.00
PRLH	P81277IPRRP_HUMAN Prolactin-releasing peptide OS=Homo sapiens GN=PRLH PE=2 SV=1	0	1	0.00
MTTP	P55157IMTP_HUMAN Microsomal triglyceride transfer protein large subunit OS=Homo sapiens GN=MTTP PE=1 SV=1	0	3	0.00
HBD	P02042IHBD_HUMAN Hemoglobin subunit delta OS=Homo sapiens GN=HBD PE=1 SV=2	0	2	0.00
NUGGC	Q68CJ6ISLIP_HUMAN Nuclear GTPase SLIP-GC OS=Homo sapiens GN=NUGGC PE=2 SV=3	0	1	0.00
ZCCHC16	Q6ZR62IZCH16_HUMAN Zinc finger CCHC domain-containing protein 16 OS=Homo sapiens GN=ZCCHC16 PE=2 SV=2	0	1	0.00
TFRC	P02786ITFR1_HUMAN Transferrin receptor protein 1 OS=Homo sapiens GN=TFRC PE=1 SV=2	0	2	0.00
TGFB1	Q1582IBGH3_HUMAN Transforming growth factor-beta-induced protein ig-h3 OS=Homo sapiens GN=TGFB1 PE=1 SV=1	0	1	0.00
ADGRL1	O94910IAGRL1_HUMAN Adhesion G protein-coupled receptor L1 OS=Homo sapiens GN=ADGRL1 PE=1 SV=1	0	2	0.00
DST	Q03001IDYST_HUMAN Dystonin OS=Homo sapiens GN=DST PE=1 SV=4	0	3	0.00
DYNC2H1	Q8NCM8IDYHC2_HUMAN Cytoplasmic dynein 2 heavy chain 1 OS=Homo sapiens GN=DYNC2H1 PE=1 SV=4	0	2	0.00
NR4A2	P43354INR4A2_HUMAN Nuclear receptor subfamily 4 group A member 2 OS=Homo sapiens GN=NR4A2 PE=1 SV=1	0	1	0.00
CALML5	Q9NZT1ICALL5_HUMAN Calmodulin-like protein 5 OS=Homo sapiens GN=CALML5 PE=1 SV=2	0	1	0.00
No_GeneID	F8W031_HUMAN Uncharacterized protein (Fragment) OS=Homo sapiens PE=1 SV=1	0	2	0.00
MUC5AC	P98088IMUC5A_HUMAN Mucin-5AC OS=Homo sapiens GN=MUC5AC PE=1 SV=4	0	1	0.00
PRF1	P14222IPERF_HUMAN Perforin-1 OS=Homo sapiens GN=PRF1 PE=1 SV=1	0	1	0.00
PCLO	Q9Y6V0IPCLO_HUMAN Protein piccolo OS=Homo sapiens GN=PCLO PE=1 SV=4	0	3	0.00
MDGA1	Q8NFP4IMDGA1_HUMAN MAM domain-containing glycosylphosphatidylinositol anchor protein 1 OS=Homo sapiens GN=MDGA1 PE=1 SV=1	0	2	0.00
CDC158	Q5M9N0ICD158_HUMAN Coiled-coil domain-containing protein 158 OS=Homo sapiens GN=CDC158 PE=2 SV=2	0	1	0.00
PSMD2	Q13200IPSM2_HUMAN 26S proteasome non-ATPase regulatory subunit 2 OS=Homo sapiens GN=PSMD2 PE=1 SV=3	0	1	0.00
MTMR11	A4FU01IMTMRB_HUMAN Myotubularin-related protein 11 OS=Homo sapiens GN=MTMR11 PE=2 SV=2	0	1	0.00
VIL1	P09327IVIL1_HUMAN Villin-1 OS=Homo sapiens GN=VIL1 PE=1 SV=4	0	2	0.00
SH3RF3	Q8TEJ3ISH3R3_HUMAN SH3 domain-containing RING finger protein 3 OS=Homo sapiens GN=SH3RF3 PE=1 SV=2	0	1	0.00
SLC3A2	P08195I4F2_HUMAN 4F2 cell-surface antigen heavy chain OS=Homo sapiens GN=SLC3A2 PE=1 SV=3	0	2	0.00
TRIM56	Q9BR22ITRI56_HUMAN E3 ubiquitin-protein ligase TRIM56 OS=Homo sapiens GN=TRIM56 PE=1 SV=3	0	1	0.00
COPG1	Q9Y678ICOPG1_HUMAN Coatamer subunit gamma-1 OS=Homo sapiens GN=COPG1 PE=1 SV=1	0	1	0.00
POTED	Q86YR6IPOTED_HUMAN POTE ankyrin domain family member D OS=Homo sapiens GN=POTED PE=2 SV=2	0	1	0.00
FAM186A	AGNE01IF186A_HUMAN Protein FAM186A OS=Homo sapiens GN=FAM186A PE=2 SV=3	0	2	0.00
ACTN4	O43707IACNT4_HUMAN Alpha-actinin-4 OS=Homo sapiens GN=ACTN4 PE=1 SV=2	0	3	0.00
FOXL2	P58012IFOXL2_HUMAN Forkhead box protein L2 OS=Homo sapiens GN=FOXL2 PE=1 SV=1	0	1	0.00
GLS	O94925IGLSK_HUMAN Glutaminase kidney isoform, mitochondrial OS=Homo sapiens GN=GLS PE=1 SV=1	0	1	0.00
UBTF	P17480IUBF1_HUMAN Nucleolar transcription factor 1 OS=Homo sapiens GN=UBTF PE=1 SV=1	0	2	0.00
ZFXH3	Q15911IZFXH3_HUMAN Zinc finger homeobox protein 3 OS=Homo sapiens GN=ZFXH3 PE=1 SV=2	0	1	0.00
IGSF21	Q96ID5IGS21_HUMAN Immunoglobulin superfamily member 21 OS=Homo sapiens GN=IGSF21 PE=2 SV=1	0	1	0.00
UBA3	Q8TBC4IUBA3_HUMAN NEDD8-activating enzyme E1 catalytic subunit OS=Homo sapiens GN=UBA3 PE=1 SV=2	0	1	0.00
LAMB1	P07942ILAMB1_HUMAN Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	0	2	0.00
DECR2	Q9NUJ1IDECR2_HUMAN Peroxisomal 2,4-dienoyl-CoA reductase OS=Homo sapiens GN=DECR2 PE=1 SV=1	0	1	0.00
LILRB4	Q8NHJ6ILIRB4_HUMAN Leukocyte immunoglobulin-like receptor subfamily B member 4 OS=Homo sapiens GN=LILRB4 PE=1 SV=3	0	2	0.00
TNC	P24821ITENA_HUMAN Tenascin OS=Homo sapiens GN=TNC PE=1 SV=3	0	1	0.00
SOD2	P04179ISODM_HUMAN Superoxide dismutase [Mn], mitochondrial OS=Homo sapiens GN=SOD2 PE=1 SV=2	0	1	0.00
SHROOM3	Q8TF72ISHRM3_HUMAN Protein Shroom3 OS=Homo sapiens GN=SHROOM3 PE=1 SV=2	0	1	0.00
GINS3	Q9BRX5IPSF3_HUMAN DNA replication complex GINS protein PSF3 OS=Homo sapiens GN=GINS3 PE=1 SV=1	0	1	0.00
ACTN2	P35609IACNT2_HUMAN Alpha-actinin-2 OS=Homo sapiens GN=ACTN2 PE=1 SV=1	0	2	0.00
PSMG3-AS1	Q96PY0IK1908_HUMAN Putative uncharacterized protein PSMG3-AS1 OS=Homo sapiens GN=PSMG3-AS1 PE=5 SV=2	0	1	0.00
TMEM266	Q2M3C6ITM266_HUMAN Transmembrane protein 266 OS=Homo sapiens GN=TMEM266 PE=2 SV=2	0	1	0.00
ZFYVE28	Q9HCC9ILST2_HUMAN Lateral signaling target protein 2 homolog OS=Homo sapiens GN=ZFYVE28 PE=1 SV=3	0	1	0.00
SECISBP2	Q96T21ISEBP2_HUMAN Selenocysteine insertion sequence-binding protein 2 OS=Homo sapiens GN=SECISBP2 PE=1 SV=2	0	1	0.00
EEF1A1	P68104IEF1A1_HUMAN Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1	0	2	0.00
MALRD1	Q5VYJ5IMALR1_HUMAN MAM and LDL-receptor class A domain-containing protein 1 OS=Homo sapiens GN=MALRD1 PE=1 SV=4	0	2	0.00