

Gene Name	Name of Protein (UniProtID, SPECIES, protein name, origin species, gene name)	Peptide Count in IP	Peptide Count in IgG	Fold Enrichment
ACTN1	P12814IACNT1_HUMAN Alpha-actinin-1 OS=Homo sapiens GN=ACTN1 PE=1 SV=2	15	0	Not in IgG control
ZC3H11A	O75152ZC11A_HUMAN Zinc finger CCH domain-containing protein 11A OS=Homo sapiens GN=ZC3H11A PE=1 SV=3	11	0	Not in IgG control
PLEC	Q15149PLEC_HUMAN Plectin OS=Homo sapiens GN=PLEC PE=1 SV=3	9	0	Not in IgG control
UBA1	P22314UBA1_HUMAN Ubiquitin-like modifier-activating enzyme 1 OS=Homo sapiens GN=UBA1 PE=1 SV=3	9	0	Not in IgG control
KMT2D	O14686KMT2D_HUMAN Histone-lysine N-methyltransferase 2D OS=Homo sapiens GN=KMT2D PE=1 SV=2	7	0	Not in IgG control
GART	P22102IPUR2_HUMAN Trifunctional purine biosynthetic protein adenosine-3 OS=Homo sapiens GN=GART PE=1 SV=1	6	0	Not in IgG control
DSP	P15924IDESP_HUMAN Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3	6	0	Not in IgG control
FLG	P20930FLA_HUMAN Flaggrin OS=Homo sapiens GN=FLG PE=1 SV=3	6	0	Not in IgG control
DHX15	O43143DHX15_HUMAN Pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15 OS=Homo sapiens GN=DHX15 PE=1 SV=2	5	0	Not in IgG control
KDM1A	O60341KDM1A_HUMAN Lysine-specific histone demethylase 1A OS=Homo sapiens GN=KDM1A PE=1 SV=2	5	0	Not in IgG control
EF3B	P55884EF3B_HUMAN Eukaryotic translation initiation factor 3 subunit B OS=Homo sapiens GN=EF3B PE=1 SV=3	5	0	Not in IgG control
GAPDH	P04406G3P_HUMAN Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	5	0	Not in IgG control
ABCA1	O95477ABCA1_HUMAN ATP-binding cassette sub-family A member 1 OS=Homo sapiens GN=ABCA1 PE=1 SV=3	5	0	Not in IgG control
COPB1	P53618COPB_HUMAN Coatomer subunit beta OS=Homo sapiens GN=COPB1 PE=1 SV=3	4	0	Not in IgG control
MCCC1	Q96RQ3MCCA_HUMAN Methylcrotonoyl-CoA carboxylase subunit alpha, mitochondrial OS=Homo sapiens GN=MCCC1 PE=1 SV=3	4	0	Not in IgG control
MCM2	P49736MCM2_HUMAN DNA replication licensing factor MCM2 OS=Homo sapiens GN=MCM2 PE=1 SV=4	4	0	Not in IgG control
MCM3	P25205MCM3_HUMAN DNA replication licensing factor MCM3 OS=Homo sapiens GN=MCM3 PE=1 SV=3	4	0	Not in IgG control
DSG1	Q02413DSG1_HUMAN Desmoglein-1 OS=Homo sapiens GN=DSG1 PE=1 SV=2	4	0	Not in IgG control
ENO1	P06733ENO1_HUMAN Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	4	0	Not in IgG control
COL28A1	QZUY09COSA1_HUMAN Collagen alpha-1(XXVIII) chain OS=Homo sapiens GN=COL28A1 PE=2 SV=2	4	0	Not in IgG control
ACLY	P53964ACLY_HUMAN ATP-citrate synthase OS=Homo sapiens GN=ACLY PE=1 SV=3	4	0	Not in IgG control
DSC1	Q08554DSC1_HUMAN Desmocollin-1 OS=Homo sapiens GN=DSC1 PE=1 SV=2	4	0	Not in IgG control
PPP	P12727PPP_HUMAN Prolactin-inducible protein OS=Homo sapiens GN=PPP PE=1 SV=1	3	0	Not in IgG control
NOLC1	Q14978NOLC1_HUMAN Nucleolar and colloid-body phosphoprotein 1 OS=Homo sapiens GN=NOLC1 PE=1 SV=2	3	0	Not in IgG control
STRADA	Q7RTN6ISTRAA_HUMAN STE20-related kinase adapter protein alpha OS=Homo sapiens GN=STRADA PE=1 SV=1	3	0	Not in IgG control
MTHFD1	P11586MTHFD1_HUMAN C-1-tetrahydrofolate synthase, cytoplasmic OS=Homo sapiens GN=MTHFD1 PE=1 SV=3	3	0	Not in IgG control
UBB	POC647UBB_HUMAN Polyubiquitin-B OS=Homo sapiens GN=UBB PE=1 SV=1	3	0	Not in IgG control
CLTC	Q00610CLH1_HUMAN Clathrin heavy chain 1 OS=Homo sapiens GN=CLTC PE=1 SV=5	3	0	Not in IgG control
EEA1	Q15075EEA1_HUMAN Early endosome antigen 1 OS=Homo sapiens GN=EEA1 PE=1 SV=2	3	0	Not in IgG control
GIMAP8	Q8ND71GIMAP8_HUMAN GTPase IMAP family member 8 OS=Homo sapiens GN=GIMAP8 PE=2 SV=2	3	0	Not in IgG control
TNPO1	Q92973TNPO1_HUMAN Transportin-1 OS=Homo sapiens GN=TNPO1 PE=1 SV=2	3	0	Not in IgG control
CASP14	P31944CASP14_HUMAN Caspase-14 OS=Homo sapiens GN=CASP14 PE=1 SV=2	3	0	Not in IgG control
CAMSAP2	Q08AD1CAMP2_HUMAN Calmodulin-regulated spectrin-associated protein 2 OS=Homo sapiens GN=CAMSAP2 PE=1 SV=3	3	0	Not in IgG control
SEC24C	P53992SEC24C_HUMAN Protein transport protein Sec24C OS=Homo sapiens GN=SEC24C PE=1 SV=3	3	0	Not in IgG control
MCM6	Q14566MCM6_HUMAN DNA replication licensing factor MCM6 OS=Homo sapiens GN=MCM6 PE=1 SV=1	3	0	Not in IgG control
PSMD1	Q99460PSMD1_HUMAN 26S proteasome non-ATPase regulatory subunit 1 OS=Homo sapiens GN=PSMD1 PE=1 SV=2	3	0	Not in IgG control
DCD	P81605DCD_HUMAN Dermcidin OS=Homo sapiens GN=DCD PE=1 SV=2	3	0	Not in IgG control
EFTUD2	Q15029EFTUD2_HUMAN 116 kDa US small nuclear ribonucleoprotein component OS=Homo sapiens GN=EFTUD2 PE=1 SV=1	3	0	Not in IgG control
DNAH1	Q96DT5IDYH11_HUMAN Dynein heavy chain 11, axonemal OS=Homo sapiens GN=DNAH1 PE=1 SV=4	3	0	Not in IgG control
MSH2	P43246MSH2_HUMAN DNA mismatch repair protein Msh2 OS=Homo sapiens GN=MSH2 PE=1 SV=1	3	0	Not in IgG control
SFPQ	P23246SFPQ_HUMAN Splicing factor, proline- and glutamine-rich OS=Homo sapiens GN=SFPQ PE=1 SV=2	3	0	Not in IgG control
XPTOT	O43592XPTOT_HUMAN Exportin-T OS=Homo sapiens GN=XPTOT PE=1 SV=2	3	0	Not in IgG control
COPG1	Q9Y678COPG1_HUMAN Coatomer subunit gamma-1 OS=Homo sapiens GN=COPG1 PE=1 SV=1	3	0	Not in IgG control
CACNA1S	Q13698CAC1S_HUMAN Voltage-dependent L-type calcium channel subunit alpha-1S OS=Homo sapiens GN=CACNA1S PE=1 SV=4	3	0	Not in IgG control
ZSWIM3	Q06MP5ZSWIM3_HUMAN Zinc finger SWIM domain-containing protein 3 OS=Homo sapiens GN=ZSWIM3 PE=2 SV=2	2	0	Not in IgG control
HNRNPUL1	Q9BUJ2HNRNP1_HUMAN Heterogeneous nuclear ribonucleoprotein U-like protein 1 OS=Homo sapiens GN=HNRNPUL1 PE=1 SV=2	2	0	Not in IgG control
GPLD1	P80108PHLD_HUMAN Phosphatidylinositol-glycan-specific phospholipase D OS=Homo sapiens GN=GPLD1 PE=1 SV=3	2	0	Not in IgG control
MCM5	P33992MCM5_HUMAN DNA replication licensing factor MCM5 OS=Homo sapiens GN=MCM5 PE=1 SV=5	2	0	Not in IgG control
S100A8	P05109S100A8_HUMAN Protein S100-A8 OS=Homo sapiens GN=S100A8 PE=1 SV=1	2	0	Not in IgG control
COL28A1	QZUY09COSA1_HUMAN Collagen alpha-1(XXVIII) chain OS=Homo sapiens GN=COL28A1 PE=2 SV=2	2	0	Not in IgG control
ARG1	P05089ARG1_HUMAN Arginase-1 OS=Homo sapiens GN=ARG1 PE=1 SV=2	2	0	Not in IgG control
CHD3	Q12873CHD3_HUMAN Chromodomain-helicase-DNA-binding protein 3 OS=Homo sapiens GN=CHD3 PE=1 SV=3	2	0	Not in IgG control
AMBR1	Q9C0C7AMR1_HUMAN Activating molecule in BECN1-regulated autophagy protein 1 OS=Homo sapiens GN=AMBR1 PE=1 SV=2	2	0	Not in IgG control
MCM4	P33991MCM4_HUMAN DNA replication licensing factor MCM4 OS=Homo sapiens GN=MCM4 PE=1 SV=5	2	0	Not in IgG control
AP1B1	Q10567AP1B1_HUMAN AP-1 complex subunit beta-1 OS=Homo sapiens GN=AP1B1 PE=1 SV=2	2	0	Not in IgG control
WDR62	O43379WDR62_HUMAN WD repeat-containing protein 62 OS=Homo sapiens GN=WDR62 PE=1 SV=4	2	0	Not in IgG control
UBA2	Q9UBT2IAE2_HUMAN SUMO-activating enzyme subunit 2 OS=Homo sapiens GN=UBA2 PE=1 SV=2	2	0	Not in IgG control
HNRNPUL2	Q1KMD3HNRNP2_HUMAN Heterogeneous nuclear ribonucleoprotein U-like protein 2 OS=Homo sapiens GN=HNRNPUL2 PE=1 SV=1	2	0	Not in IgG control
HMGN3	Q15651HMGN3_HUMAN High mobility group nucleosome-binding domain-containing protein 3 OS=Homo sapiens GN=HMGN3 PE=1 SV=2	2	0	Not in IgG control
RANBP17	Q9H277RBP17_HUMAN Ran-binding protein 17 OS=Homo sapiens GN=RANBP17 PE=2 SV=1	1	0	Not in IgG control
GSDMA	Q96QASGSDMA_HUMAN Gasdermin-A OS=Homo sapiens GN=GSDMA PE=1 SV=4	1	0	Not in IgG control
TRIM68	Q6AZ21TRIM68_HUMAN E3 ubiquitin-protein ligase TRIM68 OS=Homo sapiens GN=TRIM68 PE=1 SV=1	1	0	Not in IgG control
EF3CL	B5ME19EF3CL_HUMAN Eukaryotic translation initiation factor 3 subunit C-like protein OS=Homo sapiens GN=EF3CL PE=3 SV=1	1	0	Not in IgG control
TXN	P10599THIO_HUMAN Thioredoxin OS=Homo sapiens GN=TXN PE=1 SV=3	1	0	Not in IgG control
MS4A10	Q96PG2MS4A10_HUMAN Membrane-spanning 4-domains subfamily A member 10 OS=Homo sapiens GN=MS4A10 PE=2 SV=3	1	0	Not in IgG control
TOP1	P11387TOP1_HUMAN DNA topoisomerase 1 OS=Homo sapiens GN=TOP1 PE=1 SV=2	1	0	Not in IgG control
CHUK	O15111IKKA_HUMAN Inhibitor of nuclear factor kappa-B kinase subunit alpha OS=Homo sapiens GN=CHUK PE=1 SV=2	1	0	Not in IgG control
PXMP2	Q9NR77PXMP2_HUMAN Peroxisomal membrane protein 2 OS=Homo sapiens GN=PXMP2 PE=1 SV=3	1	0	Not in IgG control
CDSN	Q15517CDSN_HUMAN Corneodesmosin OS=Homo sapiens GN=CDSN PE=1 SV=3	1	0	Not in IgG control
LONP1	P36776LONP1_HUMAN Lon protease homolog, mitochondrial OS=Homo sapiens GN=LONP1 PE=1 SV=2	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
USP30	Q70CQ3UBP30_HUMAN Ubiquitin carboxyl-terminal hydrolase 30 OS=Homo sapiens GN=USP30 PE=1 SV=1	1	0	Not in IgG control
TMEM26	Q6ZUK4TMEM26_HUMAN Transmembrane protein 26 OS=Homo sapiens GN=TMEM26 PE=1 SV=1	1	0	Not in IgG control
RAB6B	Q9NRW1RAB6B_HUMAN Ras-related protein Rab-6B OS=Homo sapiens GN=RAB6B PE=1 SV=1	1	0	Not in IgG control
MATR3	P43243MATR3_HUMAN Matrin-3 OS=Homo sapiens GN=MATR3 PE=1 SV=3	70	9	7.78
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	5	1	5.00
FLG2	Q50862FLA2_HUMAN Flaggrin-2 OS=Homo sapiens GN=FLG2 PE=1 SV=1	4	1	4.00
SYNE1	Q0NF91SYNE1_HUMAN Nesprin-1 OS=Homo sapiens GN=SYNE1 PE=1 SV=4	3	2	3.50
HSP90AB2P	Q58FF8HSP90B2_HUMAN Putative heat shock protein HSP 90-beta 2 OS=Homo sapiens GN=HSP90AB2P PE=1 SV=2	7	3	3.00
HSP90AB4P	Q58FF6HSP90B4_HUMAN Putative heat shock protein HSP 90-beta 4 OS=Homo sapiens GN=HSP90AB4P PE=5 SV=1	3	1	3.00
TRIM28	Q13263TIF1B_HUMAN Transcription intermediary factor 1-beta OS=Homo sapiens GN=TRIM28 PE=1 SV=5	11	4	2.75
DDX1	Q92499DDX1_HUMAN ATP-dependent RNA helicase DDX1 OS=Homo sapiens GN=DDX1 PE=1 SV=2	5	2	2.50
ATP2A2	P16615AT2A2_HUMAN Sarcoplasmic/endoplasmic reticulum calcium ATPase 2 OS=Homo sapiens GN=ATP2A2 PE=1 SV=1	5	2	2.50
SND1	Q7KZF4SND1_HUMAN Staphylococcal nuclease domain-containing protein 1 OS=Homo sapiens GN=SND1 PE=1 SV=1	11	5	2.20
XPO1	O14980XPO1_HUMAN Exportin-1 OS=Homo sapiens GN=XPO1 PE=1 SV=1	4	2	2.00
HSPA4	P34932HSP74_HUMAN Heat shock 70 kDa protein 4 OS=Homo sapiens GN=HSPA4 PE=1 SV=4	10	5	2.00
KATNAL2	Q8RYT4KATL2_HUMAN Katanin p60 ATPase-containing subunit A-like 2 OS=Homo sapiens GN=KATNAL2 PE=2 SV=3	2	1	2.00
MUC5AC	P98088MUC5A_HUMAN Mucin-5AC OS=Homo sapiens GN=MUC5AC PE=1 SV=4	4	2	2.00
ILF3	Q12906ILF3_HUMAN Interleukin enhancer-binding factor 3 OS=Homo sapiens GN=ILF3 PE=1 SV=3	4	2	2.00
FAAH2	Q6GMR7FAAH2_HUMAN Fatty-acid amide hydrolase 2 OS=Homo sapiens GN=FAAH2 PE=2 SV=1	2	1	2.00

DDX21	Q9NR30DDX21_HUMAN Nucleolar RNA helicase 2 OS=Homo sapiens GN=DDX21 PE=1 SV=5	9	5	1.80
AARS	P49588SIYAC_HUMAN Alanine-tRNA ligase, cytoplasmic OS=Homo sapiens GN=AARS PE=1 SV=2	7	4	1.75
TFR3	P02786ITR1_HUMAN Transferin receptor protein 1 OS=Homo sapiens GN=TFR3 PE=1 SV=2	6	4	1.50
MARS	P56192IYMC_HUMAN Methionine-tRNA ligase, cytoplasmic OS=Homo sapiens GN=MARS PE=1 SV=2	6	4	1.50
GANAB	Q14697GANAB_HUMAN Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB PE=1 SV=3	9	6	1.50
HSP90AA1	P07900H90A_HUMAN Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1 PE=1 SV=5	18	14	1.29
PDCD6IP	Q8WUM4PDC6L_HUMAN Programmed cell death 6-interacting protein OS=Homo sapiens GN=PDCD6IP PE=1 SV=1	5	4	1.25
HNRNPU	Q00839HNRPL_HUMAN Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU PE=1 SV=6	9	8	1.13
HSP90AB1	P08238H90B_HUMAN Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4	29	27	1.07
ZC3H12C	Q9C0D7IZC12C_HUMAN Probable ribonuclease ZC3H12C OS=Homo sapiens GN=ZC3H12C PE=1 SV=2	1	1	1.00
VCP	P55072ITERA_HUMAN Transitional endoplasmic reticulum ATPase OS=Homo sapiens GN=VCP PE=1 SV=4	13	13	1.00
XRCC5	P13010XRCC5_HUMAN X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3	3	3	1.00
PSMD2	Q13200PSMD2_HUMAN 26S proteasome non-ATPase regulatory subunit 2 OS=Homo sapiens GN=PSMD2 PE=1 SV=3	2	2	1.00
SHC1	P29353SHC1_HUMAN SHC-transforming protein 1 OS=Homo sapiens GN=SHC1 PE=1 SV=4	1	1	1.00
CREBRF	Q8UR6ICREBF_HUMAN CREB3 regulatory factor OS=Homo sapiens GN=CREBRF PE=1 SV=2	1	1	1.00
PATL2	C9JAE4PATL2_HUMAN Protein PAT1 homolog 2 OS=Homo sapiens GN=PATL2 PE=1 SV=1	1	1	1.00
SASH1	O94885SASH1_HUMAN SAM and SH3 domain-containing protein 1 OS=Homo sapiens GN=SASH1 PE=1 SV=3	2	2	1.00
ALB	P02768IALBU_HUMAN Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	5	5	1.00
CAP2	P40123ICAP2_HUMAN Adenylyl cyclase-associated protein 2 OS=Homo sapiens GN=CAP2 PE=1 SV=1	1	1	1.00
EEF1A2	Q05639EEF1A2_HUMAN Elongation factor 1-alpha 2 OS=Homo sapiens GN=EEF1A2 PE=1 SV=1	1	1	1.00
TRAP1	Q12931TRAP1_HUMAN Heat shock protein 75 kDa, mitochondrial OS=Homo sapiens GN=TRAP1 PE=1 SV=3	2	2	1.00
HSP90B1	P14625IENPL_HUMAN Endoplasmic reticulum chaperone OS=Homo sapiens GN=HSP90B1 PE=1 SV=1	19	21	0.90
ACTN4	O43707IACNT4_HUMAN Alpha-actinin-4 OS=Homo sapiens GN=ACTN4 PE=1 SV=2	8	9	0.89
KHSRP	Q92945FUBP2_HUMAN Far upstream element-binding protein 2 OS=Homo sapiens GN=KHSRP PE=1 SV=4	3	4	0.75
EEF2	P13639IEF2_HUMAN Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4	17	24	0.71
LAMB1	P07942LAMB1_HUMAN Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	2	3	0.67
SPECC1L-ADORA2A	FBWAN1_HUMAN Protein SPECC1L-ADORA2A OS=Homo sapiens GN=SPECC1L-ADORA2A PE=4 SV=2	2	3	0.67
CSE1L	P55060XPO2_HUMAN Exportin-2 OS=Homo sapiens GN=CSE1L PE=1 SV=3	3	5	0.60
NCL	P19338NCL_HUMAN Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3	6	10	0.60
FAHD2A	Q96GK7FAHD2A_HUMAN Fumarylacetoacetate hydrolase domain-containing protein 2A OS=Homo sapiens GN=FAHD2A PE=1 SV=1	1	2	0.50
CDK16	Q00536CDK16_HUMAN Cyclin-dependent kinase 16 OS=Homo sapiens GN=CDK16 PE=1 SV=1	1	2	0.50
Cys-GrB-Fc-RSP01	IA0A024R160_HUMAN Cys-GrB-Fc-RSP01 OS=Homo sapiens GN=Cys-GrB-Fc-RSP01 PE=3 SV=1	1	3	0.33
KPNB1	Q14974KPNB1_HUMAN Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2	1	4	0.25
VP535	Q96QK1VP535_HUMAN Vacuolar protein sorting-associated protein 35 OS=Homo sapiens GN=VP535 PE=1 SV=2	0	1	0.00
ZZEF1	O43149ZZEF1_HUMAN Zinc finger ZZ-type and EF-hand domain-containing protein 1 OS=Homo sapiens GN=ZZEF1 PE=1 SV=6	0	2	0.00
KTNI	Q86UP2KTNI_HUMAN Kinectin OS=Homo sapiens GN=KTNI PE=1 SV=1	0	5	0.00
RAB3GAP1	Q15042R83GP_HUMAN Rab3 GTPase-activating protein catalytic subunit OS=Homo sapiens GN=RAB3GAP1 PE=1 SV=3	0	2	0.00
NDUFS8	O00217INDU58_HUMAN NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial OS=Homo sapiens GN=NDUFS8 PE=1 SV=1	0	1	0.00
HJURP	Q8NCD3HJURP_HUMAN Holliday junction recognition protein OS=Homo sapiens GN=HJURP PE=1 SV=2	0	2	0.00
ALDH18A1	P54886P5CS_HUMAN Delta-1-pyrroline-5-carboxylate synthase OS=Homo sapiens GN=ALDH18A1 PE=1 SV=2	0	2	0.00
EVIS	O60447EVIS_HUMAN Ecotropic viral integration site 5 protein homolog OS=Homo sapiens GN=EVIS PE=1 SV=3	0	3	0.00
C11orf84	Q8BUA3ICK084_HUMAN Uncharacterized protein C11orf84 OS=Homo sapiens GN=C11orf84 PE=1 SV=3	0	2	0.00
IQCH	Q86V53IQCH_HUMAN IQ domain-containing protein H OS=Homo sapiens GN=IQCH PE=2 SV=2	0	1	0.00
CEP350	Q5VT06ICE350_HUMAN Centrosome-associated protein 350 OS=Homo sapiens GN=CEP350 PE=1 SV=1	0	6	0.00
SBF1	O95248MTMR5_HUMAN Myotubularin-related protein 5 OS=Homo sapiens GN=SBF1 PE=1 SV=3	0	2	0.00
ADH5	P11766IADH5_HUMAN Alcohol dehydrogenase class-3 OS=Homo sapiens GN=ADH5 PE=1 SV=4	0	1	0.00
MECOM	Q03112IEV1_HUMAN MDS1 and EVI1 complex locus protein EVI1 OS=Homo sapiens GN=MECOM PE=1 SV=2	0	3	0.00
ACTB	P60709IACTB_HUMAN Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	0	2	0.00
TTN	Q8WZ42TTIN_HUMAN Titin OS=Homo sapiens GN=TTN PE=1 SV=4	0	27	0.00
PKN2	Q16513PKN2_HUMAN Serine/threonine-protein kinase N2 OS=Homo sapiens GN=PKN2 PE=1 SV=1	0	2	0.00
ICK	Q9UPZ9ICK_HUMAN Serine/threonine-protein kinase ICK OS=Homo sapiens GN=ICK PE=1 SV=1	0	2	0.00
RPS27A	P62979RPS27A_HUMAN Ubiquitin-40S ribosomal protein S27a OS=Homo sapiens GN=RPS27A PE=1 SV=2	0	2	0.00
C17orf53	Q8N3J3CQ053_HUMAN Uncharacterized protein C17orf53 OS=Homo sapiens GN=C17orf53 PE=1 SV=1	0	2	0.00
YDCJ	ABM57YDJC_HUMAN Carbohydrate desacylase OS=Homo sapiens GN=YDCJ PE=1 SV=1	0	2	0.00
VIMP	Q9BQE4ISEL_HUMAN Selenoprotein 5 OS=Homo sapiens GN=VIMP PE=1 SV=3	0	1	0.00
FER	P16591FER_HUMAN Tyrosine-protein kinase Fer OS=Homo sapiens GN=FER PE=1 SV=2	0	1	0.00
GP68	Q15743OGR1_HUMAN Ovarian cancer G-protein coupled receptor 1 OS=Homo sapiens GN=GP68 PE=1 SV=1	0	1	0.00
UBA3	Q8TBC4UBA3_HUMAN NEDD8-activating enzyme E1 catalytic subunit OS=Homo sapiens GN=UBA3 PE=1 SV=2	0	1	0.00
FAM47DP	ASNHR8FAM47D_HUMAN Putative protein FAM47D OS=Homo sapiens GN=FAM47DP PE=5 SV=3	0	1	0.00
IRAK2	O43187IRAK2_HUMAN Interleukin-1 receptor-associated kinase-like 2 OS=Homo sapiens GN=IRAK2 PE=1 SV=2	0	4	0.00
LRMP	Q12912LRMP_HUMAN Lymphoid-restricted membrane protein OS=Homo sapiens GN=LRMP PE=1 SV=3	0	3	0.00
SRP72	O76094SRP72_HUMAN Signal recognition particle subunit SRP72 OS=Homo sapiens GN=SRP72 PE=1 SV=3	0	1	0.00
TUBA1A	Q71U36TBA1A_HUMAN Tubulin alpha-1A chain OS=Homo sapiens GN=TUBA1A PE=1 SV=1	0	2	0.00
MTRF1L	Q9H019MTRF1L_HUMAN Mitochondrial fission regulator 1-like OS=Homo sapiens GN=MTRF1L PE=1 SV=2	0	1	0.00
WDR81	Q562E7WDR81_HUMAN WD repeat-containing protein 81 OS=Homo sapiens GN=WDR81 PE=1 SV=2	0	4	0.00
LILRB4	Q8NHJLILRB4_HUMAN Leukocyte immunoglobulin-like receptor subfamily B member 4 OS=Homo sapiens GN=LILRB4 PE=1 SV=3	0	1	0.00
ATPIA1	P05023IAT1A1_HUMAN Sodium/potassium-transporting ATPase subunit alpha-1 OS=Homo sapiens GN=ATPIA1 PE=1 SV=1	0	7	0.00
TBC1D8	O95759ITBCD8_HUMAN TBC1 domain family member 8 OS=Homo sapiens GN=TBC1D8 PE=1 SV=3	0	4	0.00
DUSP28	Q4G0W2DUS28_HUMAN Dual specificity phosphatase 28 OS=Homo sapiens GN=DUSP28 PE=2 SV=1	0	1	0.00
SNAI1	O95863SNAI1_HUMAN Zinc finger protein SNAI1 OS=Homo sapiens GN=SNAI1 PE=1 SV=2	0	1	0.00
RASSF10	A6NK89IRASFA_HUMAN Ras association domain-containing protein 10 OS=Homo sapiens GN=RASSF10 PE=2 SV=3	0	1	0.00
CD163L1	Q9NR16C163B_HUMAN Scavenger receptor cysteine-rich type 1 protein M160 OS=Homo sapiens GN=CD163L1 PE=1 SV=2	0	2	0.00
IGLL5	B9A064IGLL5_HUMAN Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2	0	1	0.00
LPN3	Q9BQK8LPN3_HUMAN Phosphatidate phosphatase LPN3 OS=Homo sapiens GN=LPN3 PE=1 SV=3	0	2	0.00
CHRNA10	Q9GZ6IACH10_HUMAN Neuronal acetylcholine receptor subunit alpha-10 OS=Homo sapiens GN=CHRNA10 PE=1 SV=1	0	1	0.00
CARS	P49589SIYCC_HUMAN Cysteine-tRNA ligase, cytoplasmic OS=Homo sapiens GN=CARS PE=1 SV=3	0	1	0.00
ZNF423	Q2M19IZNF423_HUMAN Zinc finger protein 423 OS=Homo sapiens GN=ZNF423 PE=1 SV=1	0	1	0.00
CANX	P27824ICALX_HUMAN Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2	0	7	0.00
TRIM21	P19474IROS2_HUMAN E3 ubiquitin-protein ligase TRIM21 OS=Homo sapiens GN=TRIM21 PE=1 SV=1	0	1	0.00
PAICS	P22234IPUR6_HUMAN Multifunctional protein ADE2 OS=Homo sapiens GN=PAICS PE=1 SV=3	0	1	0.00
TRIM25	Q14258ITR25_HUMAN E3 ubiquitin/ISG15 ligase TRIM25 OS=Homo sapiens GN=TRIM25 PE=1 SV=2	0	2	0.00
BRCA1	P38398BRCA1_HUMAN Breast cancer type 1 susceptibility protein OS=Homo sapiens GN=BRCA1 PE=1 SV=2	0	1	0.00
HSPAS1	P11021GRP78_HUMAN 78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPAS1 PE=1 SV=2	0	1	0.00
ATG16L2	Q8NAA4IA16L2_HUMAN Autophagy-related protein 16-2 OS=Homo sapiens GN=ATG16L2 PE=1 SV=2	0	2	0.00
CDC88A	Q3V6T2IGRDN_HUMAN Girdin OS=Homo sapiens GN=CDC88A PE=1 SV=2	0	1	0.00
PFKL	P17858IPFKL_HUMAN ATP-dependent 6-phosphofructokinase, liver type OS=Homo sapiens GN=PFKL PE=1 SV=6	0	3	0.00
AMHR2	Q16671IAMHR2_HUMAN Anti-Muellerian hormone type-2 receptor OS=Homo sapiens GN=AMHR2 PE=1 SV=1	0	1	0.00