

Gene Name	Name of Protein (UniProtID, SPECIES, protein name, origin species, gene name)	Peptide Count (IP)	Peptide Count (IgG)	Fold Enrichment
NIP7	Q9Y221NIP7_HUMAN 60S ribosome subunit biogenesis protein NIP7 homolog OS=Homo sapiens GN=NIP7 PE=1 SV=1	14	0	Not in IgG control
HIST1H2BK	O60814H2BK_HUMAN Histone H2B type 1-K OS=Homo sapiens GN=HIST1H2BK PE=1 SV=3	7	0	Not in IgG control
RAB1B	Q9H0U4RAB1B_HUMAN Ras-related protein Rab-1B OS=Homo sapiens GN=RAB1B PE=1 SV=1	6	0	Not in IgG control
ALB	P02768IALBU_HUMAN Serum albumin OS=Homo sapiens GN=ALB PE=1 SV=2	4	0	Not in IgG control
AHNAK	Q09666AHNAK_HUMAN Neuroblast differentiation-associated protein AHNAK OS=Homo sapiens GN=AHNAK PE=1 SV=2	4	0	Not in IgG control
RPS15A	P62244IRS15A_HUMAN 40S ribosomal protein S15a OS=Homo sapiens GN=RPS15A PE=1 SV=2	4	0	Not in IgG control
DSP	P15924IDESP_HUMAN Desmoplakin OS=Homo sapiens GN=DSP PE=1 SV=3	4	0	Not in IgG control
HIST2H2AC	Q16777H2A2C_HUMAN Histone H2A type 2-C OS=Homo sapiens GN=HIST2H2AC PE=1 SV=4	3	0	Not in IgG control
PKN2	Q16513PKN2_HUMAN Serine/threonine-protein kinase N2 OS=Homo sapiens GN=PKN2 PE=1 SV=1	3	0	Not in IgG control
PLCH1	Q4KWH8PLCH1_HUMAN 1-phosphatidylinositol 4,5-bisphosphate phosphodiesterase eta-1 OS=Homo sapiens GN=PLCH1 PE=1 SV=1	3	0	Not in IgG control
SRSF7	Q16629SRSF7_HUMAN Serine/arginine-rich splicing factor 7 OS=Homo sapiens GN=SRSF7 PE=1 SV=1	3	0	Not in IgG control
PRPF8	Q6P2Q9PRPF8_HUMAN Pre-mRNA-processing-splicing factor 8 OS=Homo sapiens GN=PRPF8 PE=1 SV=2	3	0	Not in IgG control
IGLL5	B9A064IGLL5_HUMAN Immunoglobulin lambda-like polypeptide 5 OS=Homo sapiens GN=IGLL5 PE=2 SV=2	3	0	Not in IgG control
RPL17-C18orf32	A0A0A6YLL6A0A0A6YLL6_HUMAN Protein RPL17-C18orf32 OS=Homo sapiens GN=RPL17-C18orf32 PE=3 SV=1	2	0	Not in IgG control
MA3	Q5JRA6MA3_HUMAN Melanoma inhibitory activity protein 3 OS=Homo sapiens GN=MA3 PE=1 SV=1	2	0	Not in IgG control
SYNE1	Q8NF91SYNE1_HUMAN Nesprin-1 OS=Homo sapiens GN=SYNE1 PE=1 SV=4	2	0	Not in IgG control
RAP1B	P61224RAP1B_HUMAN Ras-related protein Rap-1b OS=Homo sapiens GN=RAP1B PE=1 SV=1	2	0	Not in IgG control
DNHD1	Q96M86DNHD1_HUMAN Dynein heavy chain domain-containing protein 1 OS=Homo sapiens GN=DNHD1 PE=2 SV=2	2	0	Not in IgG control
ZC3H12C	Q3COD7ZC12C_HUMAN Probable ribonuclease ZC3H12C OS=Homo sapiens GN=ZC3H12C PE=1 SV=2	2	0	Not in IgG control
SHROOM3	Q8TF72SHROOM3_HUMAN Protein Shroom3 OS=Homo sapiens GN=SHROOM3 PE=1 SV=2	2	0	Not in IgG control
No_GeneID	P01708ILV205_HUMAN Iq lambda chain V-II region BUR OS=Homo sapiens PE=1 SV=1	2	0	Not in IgG control
INPP5J	Q15735IP5JA_HUMAN Phosphatidylinositol 4,5-bisphosphate 5-phosphatase 4 OS=Homo sapiens GN=INPP5J PE=1 SV=3	2	0	Not in IgG control
APEX1	P27695APEX1_HUMAN DNA-(apurinic or apyrimidinic site) lyase OS=Homo sapiens GN=APEX1 PE=1 SV=2	2	0	Not in IgG control
DNAH9	Q9NYC9IDYH9_HUMAN Dynein heavy chain 9, axonemal OS=Homo sapiens GN=DNAH9 PE=1 SV=3	2	0	Not in IgG control
NFIB	Q00712NFIB_HUMAN Nuclear factor 1 B-type OS=Homo sapiens GN=NFIB PE=1 SV=2	2	0	Not in IgG control
HIST1H2BJ	P06899H2BJ_HUMAN Histone H2B type 1-J OS=Homo sapiens GN=HIST1H2BJ PE=1 SV=3	2	0	Not in IgG control
RPL21	P46778RPL21_HUMAN 60S ribosomal protein L21 OS=Homo sapiens GN=RPL21 PE=1 SV=2	2	0	Not in IgG control
ZC2HC1A	Q96GY0ZC21A_HUMAN Zinc finger C2HC domain-containing protein 1A OS=Homo sapiens GN=ZC2HC1A PE=1 SV=2	1	0	Not in IgG control
IGKV1-8	A0A0C4DH67IA0A0C4DH67_HUMAN Protein IGKV1-8 (Fragment) OS=Homo sapiens GN=IGKV1-8 PE=1 SV=1	1	0	Not in IgG control
TRAJ56	A0A075B622IA0A075B622_HUMAN Protein TRAJ56 (Fragment) OS=Homo sapiens GN=TRAJ56 PE=4 SV=1	1	0	Not in IgG control
IGKV1-6	A0A0C4DH72IA0A0C4DH72_HUMAN Protein IGKV1-6 (Fragment) OS=Homo sapiens GN=IGKV1-6 PE=1 SV=1	1	0	Not in IgG control
JAKMP2	Q96AABJKIP2_HUMAN Janus kinase and microtubule-interacting protein 2 OS=Homo sapiens GN=JAKMP2 PE=1 SV=1	1	0	Not in IgG control
TMF1	P82094TMF1_HUMAN TATA element modulatory factor OS=Homo sapiens GN=TMF1 PE=1 SV=2	1	0	Not in IgG control
No_GeneID	P80748BLV302_HUMAN Iq lambda chain V-III region LOI OS=Homo sapiens PE=1 SV=1	1	0	Not in IgG control
MOGAT2	Q3SYC2IMOGT2_HUMAN 2-acylglycerol O-acyltransferase 2 OS=Homo sapiens GN=MOGAT2 PE=1 SV=2	1	0	Not in IgG control
EEF1E1-BLOC15S	C9J1V9IC9J1V9_HUMAN HCG2043275 OS=Homo sapiens GN=EEF1E1-BLOC15S PE=4 SV=2	1	0	Not in IgG control
FAN1	Q9Y2M0IFAN1_HUMAN Fanconi-associated nuclease 1 OS=Homo sapiens GN=FAN1 PE=1 SV=4	1	0	Not in IgG control
DMRTC2	Q8IXT2DMRTC2_HUMAN Doublesex- and mab-3-related transcription factor C2 OS=Homo sapiens GN=DMRTC2 PE=2 SV=2	1	0	Not in IgG control
AMMECR1	Q9Y4X0JAMMR1_HUMAN AMME syndrome candidate gene 1 protein OS=Homo sapiens GN=AMMECR1 PE=1 SV=1	1	0	Not in IgG control
TRIP10	Q15642ICIP4_HUMAN Cdc42-interacting protein 4 OS=Homo sapiens GN=TRIP10 PE=1 SV=3	1	0	Not in IgG control
TCEA3	O75764ITCEA3_HUMAN Transcription elongation factor A protein 3 OS=Homo sapiens GN=TCEA3 PE=1 SV=2	1	0	Not in IgG control
SRP72	O76094SRP72_HUMAN Signal recognition particle subunit SRP72 OS=Homo sapiens GN=SRP72 PE=1 SV=3	1	0	Not in IgG control
BRK1	Q8WUW1BRK1_HUMAN Protein BRCK1 OS=Homo sapiens GN=BRK1 PE=1 SV=1	1	0	Not in IgG control
IGHG3	P01860IGHG3_HUMAN Iq gamma-3 chain C region OS=Homo sapiens GN=IGHG3 PE=1 SV=2	1	0	Not in IgG control
STEAP3	Q658P3ITEA3_HUMAN Metalloreductase STEAP3 OS=Homo sapiens GN=STEAP3 PE=1 SV=2	1	0	Not in IgG control
RAB6A	P20340RAB6A_HUMAN Ras-related protein Rab-6A OS=Homo sapiens GN=RAB6A PE=1 SV=3	1	0	Not in IgG control
PRDM1	O75626IPRDM1_HUMAN PR domain zinc finger protein 1 OS=Homo sapiens GN=PRDM1 PE=1 SV=2	1	0	Not in IgG control
RAB10	P61026RAB10_HUMAN Ras-related protein Rab-10 OS=Homo sapiens GN=RAB10 PE=1 SV=1	1	0	Not in IgG control
No_GeneID	H7C0C1IH7C0C1_HUMAN Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=2	1	0	Not in IgG control
SH3PX2B	A1X283ISP2B_HUMAN SH3 and PX domain-containing protein 2B OS=Homo sapiens GN=SH3PX2B PE=1 SV=3	1	0	Not in IgG control
LAMTOR4	Q0VGL1LTOR4_HUMAN Regulator complex protein LAMTOR4 OS=Homo sapiens GN=LAMTOR4 PE=1 SV=1	1	0	Not in IgG control
ARPC4-TTL3	A0A0A6Y9YG9A0A0A6Y9YG9_HUMAN Protein ARPC4-TTL3 OS=Homo sapiens GN=ARPC4-TTL3 PE=4 SV=1	1	0	Not in IgG control
LPIN3	Q9BQK6LPIN3_HUMAN Phosphatidate phosphatase LPIN3 OS=Homo sapiens GN=LPIN3 PE=1 SV=3	1	0	Not in IgG control
TUBGCP6	Q96R71GCP6_HUMAN Gamma-tubulin complex component 6 OS=Homo sapiens GN=TUBGCP6 PE=1 SV=3	1	0	Not in IgG control
SURF4	O15260SURF4_HUMAN Surflet locus protein 4 OS=Homo sapiens GN=SURF4 PE=1 SV=3	3	1	3.00
RPL18	Q07020RPL18_HUMAN 60S ribosomal protein L18 OS=Homo sapiens GN=RPL18 PE=1 SV=2	2	1	2.00
COL1A1	P02452COL1A1_HUMAN Collagen alpha-1(I) chain OS=Homo sapiens GN=COL1A1 PE=1 SV=5	2	1	2.00
PTRH2	Q9Y3E5PTH2_HUMAN Peptidyl-tRNA hydrolase 2, mitochondrial OS=Homo sapiens GN=PTRH2 PE=1 SV=1	3	2	1.50
RAB2A	P61019RAB2A_HUMAN Ras-related protein Rab-2A OS=Homo sapiens GN=RAB2A PE=1 SV=1	4	3	1.33
SPCS2	Q15005SPCS2_HUMAN Signal peptidase complex subunit 2 OS=Homo sapiens GN=SPCS2 PE=1 SV=3	4	3	1.33
RPL22	P35268RPL22_HUMAN 60S ribosomal protein L22 OS=Homo sapiens GN=RPL22 PE=1 SV=2	4	3	1.33
TMED10	P49755TMEDA_HUMAN Transmembrane emp24 domain-containing protein 10 OS=Homo sapiens GN=TMED10 PE=1 SV=2	6	5	1.20
RPS16	P62249IRS16_HUMAN 40S ribosomal protein S16 OS=Homo sapiens GN=RPS16 PE=1 SV=2	6	6	1.00
RPL32	P62910RPL32_HUMAN 60S ribosomal protein L32 OS=Homo sapiens GN=RPL32 PE=1 SV=2	4	4	1.00
APRT	P07741APRT_HUMAN Adenine phosphoribosyltransferase OS=Homo sapiens GN=APRT PE=1 SV=2	4	4	1.00
RPS20	P60866IRS20_HUMAN 40S ribosomal protein S20 OS=Homo sapiens GN=RPS20 PE=1 SV=1	3	3	1.00
RPS26	P62854IRS26_HUMAN 40S ribosomal protein S26 OS=Homo sapiens GN=RPS26 PE=1 SV=3	3	3	1.00
CYB5B	O43169CYB5B_HUMAN Cytochrome b5 type B OS=Homo sapiens GN=CYB5B PE=1 SV=2	3	3	1.00
TOMM22	Q9NS69ITOM22_HUMAN Mitochondrial import receptor subunit TOM22 homolog OS=Homo sapiens GN=TOMM22 PE=1 SV=3	3	3	1.00
MGST1	P10620MGST1_HUMAN Microsomal glutathione S-transferase 1 OS=Homo sapiens GN=MGST1 PE=1 SV=1	3	3	1.00
RPS24	P62847IRS24_HUMAN 40S ribosomal protein S24 OS=Homo sapiens GN=RPS24 PE=1 SV=1	2	2	1.00
RPL35	P42766RPL35_HUMAN 60S ribosomal protein L35 OS=Homo sapiens GN=RPL35 PE=1 SV=2	2	2	1.00
RPL30	P62888RPL30_HUMAN 60S ribosomal protein L30 OS=Homo sapiens GN=RPL30 PE=1 SV=2	2	2	1.00
GLUD1	P00367IDHE3_HUMAN Glutamate dehydrogenase 1, mitochondrial OS=Homo sapiens GN=GLUD1 PE=1 SV=2	2	2	1.00
ACTB	P60709IACTB_HUMAN Actin, cytoplasmic 1 OS=Homo sapiens GN=ACTB PE=1 SV=1	2	2	1.00
AP2S1	P53680AP2S1_HUMAN AP-2 complex subunit sigma OS=Homo sapiens GN=AP2S1 PE=1 SV=2	2	2	1.00
MEX3D	Q86XN8MEX3D_HUMAN RNA-binding protein MEX3D OS=Homo sapiens GN=MEX3D PE=1 SV=3	2	2	1.00
RPL34	P49207RPL34_HUMAN 60S ribosomal protein L34 OS=Homo sapiens GN=RPL34 PE=1 SV=3	2	2	1.00
EBP	Q11512IEBP_HUMAN 3-beta-hydroxysteroid-Delta(8),Delta(7)-isomerase OS=Homo sapiens GN=EBP PE=1 SV=3	2	2	1.00
OSTC	Q9NRPOIOSTC_HUMAN Oligosaccharyltransferase complex subunit OSTC OS=Homo sapiens GN=OSTC PE=1 SV=1	1	1	1.00
MT-ATP6	P00846IATP6_HUMAN ATP synthase subunit a OS=Homo sapiens GN=MT-ATP6 PE=1 SV=1	1	1	1.00
KLB	Q86214KLOTB_HUMAN Beta-klotho OS=Homo sapiens GN=KLB PE=1 SV=1	1	1	1.00
No_GeneID	P01621IKV303_HUMAN Iq kappa chain V-III region N69 (Fragment) OS=Homo sapiens PE=2 SV=1	1	1	1.00
NCBP2	P52298INCBP2_HUMAN Nuclear cap-binding protein subunit 2 OS=Homo sapiens GN=NCBP2 PE=1 SV=1	1	1	1.00
IGKV3-7	A0A075B6H7IA0A075B6H7_HUMAN Protein IGKV3-7 (Fragment) OS=Homo sapiens GN=IGKV3-7 PE=4 SV=1	1	1	1.00
TRPM8	Q772W7ITRPM8_HUMAN Transient receptor potential cation channel subfamily M member 8 OS=Homo sapiens GN=TRPM8 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

IGKV4-1	P06312IKV401_HUMAN Immunoglobulin kappa variable 4-1 OS=Homo sapiens GN=IGKV4-1 PE=1 SV=1	1	1	1.00
MRPL18	Q9H0U6IRM18_HUMAN 39S ribosomal protein L18, mitochondrial OS=Homo sapiens GN=MRPL18 PE=1 SV=1	1	1	1.00
MGST2	Q99735MGST2_HUMAN Microsomal glutathione S-transferase 2 OS=Homo sapiens GN=MGST2 PE=1 SV=1	1	1	1.00
MRPL48	Q966C5IRM48_HUMAN 39S ribosomal protein L48, mitochondrial OS=Homo sapiens GN=MRPL48 PE=1 SV=2	1	1	1.00
RAB7A	P51149RAB7A_HUMAN Ras-related protein Rab-7a OS=Homo sapiens GN=RAB7A PE=1 SV=1	7	8	0.88
RPS11	P62280IRS11_HUMAN 40S ribosomal protein S11 OS=Homo sapiens GN=RPS11 PE=1 SV=3	7	8	0.88
PRDX1	Q06830PRDX1_HUMAN Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1	9	11	0.82
HIST1H4A	P62805IH4_HUMAN Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2	4	5	0.80
SSR4	P51571ISSRD_HUMAN Translocon-associated protein subunit delta OS=Homo sapiens GN=SSR4 PE=1 SV=1	4	5	0.80
RPS14	P62263IRS14_HUMAN 40S ribosomal protein S14 OS=Homo sapiens GN=RPS14 PE=1 SV=3	4	5	0.80
PRDX3	P30048IPRDX3_HUMAN Thioredoxin-dependent peroxide reductase, mitochondrial OS=Homo sapiens GN=PRDX3 PE=1 SV=3	3	4	0.75
No_GeneID	J3QQ9UJ3QQ9_HUMAN Uncharacterized protein OS=Homo sapiens PE=4 SV=1	3	4	0.75
FAM213A	Q98RX8IF213A_HUMAN Redox-regulatory protein FAM213A OS=Homo sapiens GN=FAM213A PE=1 SV=3	3	4	0.75
PPIB	P23284IPPIB_HUMAN Peptidyl-prolyl cis-trans isomerase B OS=Homo sapiens GN=PPIB PE=1 SV=2	11	15	0.73
ATP5O	P48047IATPO_HUMAN ATP synthase subunit O, mitochondrial OS=Homo sapiens GN=ATP5O PE=1 SV=1	5	7	0.71
RPL12	P30050IRL12_HUMAN 60S ribosomal protein L12 OS=Homo sapiens GN=RPL12 PE=1 SV=1	5	7	0.71
RPS13	P62277IRS13_HUMAN 40S ribosomal protein S13 OS=Homo sapiens GN=RPS13 PE=1 SV=2	5	7	0.71
PIPA	P62937IPPIA_HUMAN Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PIPA PE=1 SV=2	6	9	0.67
COP21	P61923ICOP21_HUMAN Coatomer subunit zeta-1 OS=Homo sapiens GN=COP21 PE=1 SV=1	2	3	0.67
RPLP2	P05387IRL2_HUMAN 60S acidic ribosomal protein P2 OS=Homo sapiens GN=RPLP2 PE=1 SV=1	2	3	0.67
TMED2	Q15363ITMED2_HUMAN Transmembrane emp24 domain-containing protein 2 OS=Homo sapiens GN=TMED2 PE=1 SV=1	2	3	0.67
MUC5AC	P98088IMUC5A_HUMAN Mucin-5AC OS=Homo sapiens GN=MUC5AC PE=1 SV=4	2	3	0.67
UBB	POC647IUBB_HUMAN Polyubiquitin-B OS=Homo sapiens GN=UBB PE=1 SV=1	2	3	0.67
SEC11A	P67812IS11A_HUMAN Signal peptidase complex catalytic subunit SEC11A OS=Homo sapiens GN=SEC11A PE=1 SV=1	2	3	0.67
MRPS25	P82663IRT25_HUMAN 28S ribosomal protein S25, mitochondrial OS=Homo sapiens GN=MRPS25 PE=1 SV=1	2	3	0.67
ARL1	P40616IARL1_HUMAN ADP-ribosylation factor-like protein 1 OS=Homo sapiens GN=ARL1 PE=1 SV=1	2	3	0.67
RPL18A	Q02543IRL18A_HUMAN 60S ribosomal protein L18a OS=Homo sapiens GN=RPL18A PE=1 SV=2	5	8	0.63
NME1	P15531INDKA_HUMAN Nucleoside diphosphate kinase A OS=Homo sapiens GN=NME1 PE=1 SV=1	5	8	0.63
CFL1	P23528ICOF1_HUMAN Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3	6	10	0.60
No_GeneID	F5H423IF5H423_HUMAN Uncharacterized protein OS=Homo sapiens PE=3 SV=1	3	5	0.60
RPS23	P62266IRS23_HUMAN 40S ribosomal protein S23 OS=Homo sapiens GN=RPS23 PE=1 SV=3	3	5	0.60
RPL27	P61353IRL27_HUMAN 60S ribosomal protein L27 OS=Homo sapiens GN=RPL27 PE=1 SV=2	3	5	0.60
RPL31	P62899IRL31_HUMAN 60S ribosomal protein L31 OS=Homo sapiens GN=RPL31 PE=1 SV=1	3	5	0.60
RPL27A	P46777IRL27A_HUMAN 60S ribosomal protein L27a OS=Homo sapiens GN=RPL27A PE=1 SV=2	3	5	0.60
ACP1	P24666IPAC1_HUMAN Low molecular weight phosphotyrosine protein phosphatase OS=Homo sapiens GN=ACP1 PE=1 SV=3	3	5	0.60
RAC1	P63000IRAC1_HUMAN Ras-related C3 botulinum toxin substrate 1 OS=Homo sapiens GN=RAC1 PE=1 SV=1	4	7	0.57
RPL23	P62829IRL23_HUMAN 60S ribosomal protein L23 OS=Homo sapiens GN=RPL23 PE=1 SV=1	4	7	0.57
ARF4	P18085IARF4_HUMAN ADP-ribosylation factor 4 OS=Homo sapiens GN=ARF4 PE=1 SV=3	6	12	0.50
RPS7	P62081IRS7_HUMAN 40S ribosomal protein S7 OS=Homo sapiens GN=RPS7 PE=1 SV=1	4	8	0.50
MT-CO2	P00403ICOX2_HUMAN Cytochrome c oxidase subunit 2 OS=Homo sapiens GN=MT-CO2 PE=1 SV=1	3	6	0.50
RPL11	P62913IRL11_HUMAN 60S ribosomal protein L11 OS=Homo sapiens GN=RPL11 PE=1 SV=2	3	6	0.50
MRPL22	Q9NWU5IRM22_HUMAN 39S ribosomal protein L22, mitochondrial OS=Homo sapiens GN=MRPL22 PE=1 SV=1	2	4	0.50
NDUFA13	Q9POJ0INDUFA13_HUMAN NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 13 OS=Homo sapiens GN=NDUFA13 PE=1 SV=3	2	4	0.50
UBE2L3	P68036IUBE2L3_HUMAN Ubiquitin-conjugating enzyme E2 L3 OS=Homo sapiens GN=UBE2L3 PE=1 SV=1	2	4	0.50
TMEM205	QBUW68ITM205_HUMAN Transmembrane protein 205 OS=Homo sapiens GN=TMEM205 PE=1 SV=1	2	4	0.50
FAM162A	Q96A26IF162A_HUMAN Protein FAM162A OS=Homo sapiens GN=FAM162A PE=1 SV=2	2	4	0.50
PIP	P12273IPIP_HUMAN Prolactin-inducible protein OS=Homo sapiens GN=PIP PE=1 SV=1	2	4	0.50
SUB1	P53999ITCP4_HUMAN Activated RNA polymerase II transcriptional coactivator p15 OS=Homo sapiens GN=SUB1 PE=1 SV=3	2	4	0.50
MAGOH	P61326IMGN_HUMAN Protein mago nashi homolog OS=Homo sapiens GN=MAGOH PE=1 SV=1	2	4	0.50
DCD	P81605IDCD_HUMAN Dermicidin OS=Homo sapiens GN=DCD PE=1 SV=2	2	4	0.50
TMEM97	Q5BJF2ITMM97_HUMAN Transmembrane protein 97 OS=Homo sapiens GN=TMEM97 PE=1 SV=1	2	4	0.50
TMEM11	P17152ITMM11_HUMAN Transmembrane protein 11, mitochondrial OS=Homo sapiens GN=TMEM11 PE=1 SV=1	1	2	0.50
CDK17	Q00537ICDK17_HUMAN Cyclin-dependent kinase 17 OS=Homo sapiens GN=CDK17 PE=1 SV=2	1	2	0.50
No_GeneID	E9PLN8IE9PLN8_HUMAN Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=1	1	2	0.50
ACOT13	Q9NPJ3IACOT13_HUMAN Acyl-coenzyme A thioesterase 13 OS=Homo sapiens GN=ACOT13 PE=1 SV=1	1	2	0.50
MRPL47	Q9HD33IRML47_HUMAN 39S ribosomal protein L47, mitochondrial OS=Homo sapiens GN=MRPL47 PE=1 SV=2	1	2	0.50
METTL7A	Q9H8H3IMET7A_HUMAN Methyltransferase-like protein 7A OS=Homo sapiens GN=METTL7A PE=1 SV=1	1	2	0.50
RHOA	P61568IRHOA_HUMAN Transforming protein RhoA OS=Homo sapiens GN=RHOA PE=1 SV=1	1	2	0.50
MRPS11	P82912IRT11_HUMAN 28S ribosomal protein S11, mitochondrial OS=Homo sapiens GN=MRPS11 PE=1 SV=2	1	2	0.50
HIST2H3PS2	Q5TEC6IQ5TEC6_HUMAN Histone H3 OS=Homo sapiens GN=HIST2H3PS2 PE=1 SV=1	1	2	0.50
HIST1H3A	P68431IH3A_HUMAN Histone H3.1 OS=Homo sapiens GN=HIST1H3A PE=1 SV=2	4	9	0.44
PRDX5	P30044IPRDX5_HUMAN Peroxiredoxin-5, mitochondrial OS=Homo sapiens GN=PRDX5 PE=1 SV=4	3	7	0.43
RPL23A	P62750IRL23A_HUMAN 60S ribosomal protein L23a OS=Homo sapiens GN=RPL23A PE=1 SV=1	3	7	0.43
RAB1A	P62820IRAB1A_HUMAN Ras-related protein Rab-1A OS=Homo sapiens GN=RAB1A PE=1 SV=3	3	8	0.38
MRPL13	Q9BYD1IRM13_HUMAN 39S ribosomal protein L13, mitochondrial OS=Homo sapiens GN=MRPL13 PE=1 SV=1	2	6	0.33
IMP3	Q9NV3IIMP3_HUMAN U3 small nucleolar ribonucleoprotein protein IMP3 OS=Homo sapiens GN=IMP3 PE=1 SV=1	1	3	0.33
MRPL21	Q722W9IRM21_HUMAN 39S ribosomal protein L21, mitochondrial OS=Homo sapiens GN=MRPL21 PE=1 SV=2	1	3	0.33
MRPL23	Q16540IRM23_HUMAN 39S ribosomal protein L23, mitochondrial OS=Homo sapiens GN=MRPL23 PE=1 SV=1	1	3	0.33
RPL24	P83731IRL24_HUMAN 60S ribosomal protein L24 OS=Homo sapiens GN=RPL24 PE=1 SV=1	1	3	0.33
PSMB6	P28072IPSB6_HUMAN Proteasome subunit beta type-6 OS=Homo sapiens GN=PSMB6 PE=1 SV=4	1	3	0.33
SAR1A	Q9NR31ISAR1A_HUMAN GTP-binding protein SAR1a OS=Homo sapiens GN=SAR1A PE=1 SV=1	1	3	0.33
RPS18	P62269IRS18_HUMAN 40S ribosomal protein S18 OS=Homo sapiens GN=RPS18 PE=1 SV=3	3	12	0.25
RPS19	P39019IRS19_HUMAN 40S ribosomal protein S19 OS=Homo sapiens GN=RPS19 PE=1 SV=2	2	9	0.22
UBE2N	P61088IUBE2N_HUMAN Ubiquitin-conjugating enzyme E2 N OS=Homo sapiens GN=UBE2N PE=1 SV=1	2	9	0.22
PEBP1	P30086IPEBP1_HUMAN Phosphatidylethanolamine-binding protein 1 OS=Homo sapiens GN=PEBP1 PE=1 SV=3	2	9	0.22
RPL28	P46779IRL28_HUMAN 60S ribosomal protein L28 OS=Homo sapiens GN=RPL28 PE=1 SV=3	1	5	0.20
NME2	P22392INDKB_HUMAN Nucleoside diphosphate kinase B OS=Homo sapiens GN=NME2 PE=1 SV=1	1	5	0.20
CDC42	P60953ICDC42_HUMAN Cell division control protein 42 homolog OS=Homo sapiens GN=CDC42 PE=1 SV=2	1	6	0.17
COX4I1	P13073ICOX4I1_HUMAN Cytochrome c oxidase subunit 4 isoform 1, mitochondrial OS=Homo sapiens GN=COX4I1 PE=1 SV=1	1	6	0.17
RPS25	P62851IRS25_HUMAN 40S ribosomal protein S25 OS=Homo sapiens GN=RPS25 PE=1 SV=1	1	6	0.17
RAB18	Q9NP72IRAB18_HUMAN Ras-related protein Rab-18 OS=Homo sapiens GN=RAB18 PE=1 SV=1	1	6	0.17
CFAP20	Q9Y6A4ICFA20_HUMAN Cilia- and flagella-associated protein 20 OS=Homo sapiens GN=CFAP20 PE=1 SV=1	0	2	0.00
PSMB2	P49721IPSB2_HUMAN Proteasome subunit beta type-2 OS=Homo sapiens GN=PSMB2 PE=1 SV=1	0	2	0.00
IL1RN	P18510IIL1RA_HUMAN Interleukin-1 receptor antagonist protein OS=Homo sapiens GN=IL1RN PE=1 SV=1	0	3	0.00
H2AFZ	POCO55IH2AFZ_HUMAN Histone H2A.Z OS=Homo sapiens GN=H2AFZ PE=1 SV=2	0	3	0.00
PTMS	P20962IPTMS_HUMAN Parathyroidin OS=Homo sapiens GN=PTMS PE=1 SV=2	0	2	0.00
ITPA	Q9BY32ITPA_HUMAN Inosine triphosphate pyrophosphatase OS=Homo sapiens GN=ITPA PE=1 SV=2	0	1	0.00
RNASET2	Q00584IRNT2_HUMAN Ribonuclease T2 OS=Homo sapiens GN=RNASET2 PE=1 SV=2	0	1	0.00

SUMO3	P55854ISUMO3_HUMAN Small ubiquitin-related modifier 3 OS=Homo sapiens GN=SUMO3 PE=1 SV=2	0	1	0.00
SOD1	P00441SODC_HUMAN Superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD1 PE=1 SV=2	0	3	0.00
ISG15	P05161ISG15_HUMAN Ubiquitin-like protein ISG15 OS=Homo sapiens GN=ISG15 PE=1 SV=5	0	2	0.00
CEP131	Q9UPN4ICP131_HUMAN Centrosomal protein of 131 kDa OS=Homo sapiens GN=CEP131 PE=1 SV=3	0	2	0.00
TRIM23	P36406ITR23_HUMAN E3 ubiquitin-protein ligase TRIM23 OS=Homo sapiens GN=TRIM23 PE=1 SV=1	0	2	0.00
THEM6	Q8WUY1THEM6_HUMAN Protein THEM6 OS=Homo sapiens GN=THEM6 PE=1 SV=2	0	2	0.00
PTGES3	Q15185ITEBP_HUMAN Prostaglandin E synthase 3 OS=Homo sapiens GN=PTGES3 PE=1 SV=1	0	4	0.00
H3F3A	P84243IH33_HUMAN Histone H3.3 OS=Homo sapiens GN=H3F3A PE=1 SV=2	0	1	0.00
SNX12	Q9UMY4ISNX12_HUMAN Sorting nexin-12 OS=Homo sapiens GN=SNX12 PE=1 SV=3	0	2	0.00
TPT1	P13693ITCTP_HUMAN Translationally-controlled tumor protein OS=Homo sapiens GN=TPT1 PE=1 SV=1	0	2	0.00
CALM1	P62158ICALM_HUMAN Calmodulin OS=Homo sapiens GN=CALM1 PE=1 SV=2	0	3	0.00
TMM22	Q9Y584ITM22_HUMAN Mitochondrial import inner membrane translocase subunit Tim22 OS=Homo sapiens GN=TMM22 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
SRSF3	P84103ISRSF3_HUMAN Serine/arginine-rich splicing factor 3 OS=Homo sapiens GN=SRSF3 PE=1 SV=1	0	4	0.00
MCEE	Q96PE7MCEE_HUMAN Methylmalonyl-CoA epimerase, mitochondrial OS=Homo sapiens GN=MCEE PE=1 SV=1	0	2	0.00
VKORC1L1	Q8N0UBVKORL_HUMAN Vitamin K epoxide reductase complex subunit 1-like protein 1 OS=Homo sapiens GN=VKORC1L1 PE=1 SV=2	0	3	0.00
ZZEF1	Q43149IZZEF1_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
SPEF2	Q9C093SPEF2_HUMAN Sperm flagellar protein 2 OS=Homo sapiens GN=SPEF2 PE=1 SV=2	0	3	0.00
WRN	Q14191WRN_HUMAN Werner syndrome ATP-dependent helicase OS=Homo sapiens GN=WRN PE=1 SV=2	0	1	0.00
CDC58	Q4VC31ICD58_HUMAN Coiled-coil domain-containing protein 58 OS=Homo sapiens GN=CDC58 PE=1 SV=1	0	2	0.00
RHOB	P62745IRHOB_HUMAN Rho-related GTP-binding protein RhoB OS=Homo sapiens GN=RHOB PE=1 SV=1	0	3	0.00
ARL2BP	Q9Y2Y0IAR2BP_HUMAN ADP-ribosylation factor-like protein 2-binding protein OS=Homo sapiens GN=ARL2BP PE=1 SV=1	0	1	0.00
RAB14	P61106IRAB14_HUMAN Ras-related protein Rab-14 OS=Homo sapiens GN=RAB14 PE=1 SV=4	0	1	0.00
SNRPC	P09234IRU1C_HUMAN U1 small nuclear ribonucleoprotein C OS=Homo sapiens GN=SNRPC PE=1 SV=1	0	1	0.00
PTPMT1	Q8WLK0IPTPM1_HUMAN Phosphatidyglycerophosphatase and protein-tyrosine phosphatase 1 OS=Homo sapiens GN=PTPMT1 PE=1 SV=1	0	2	0.00
NDUFS8	Q00217INDUS8_HUMAN NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial OS=Homo sapiens GN=NDUFS8 PE=1 SV=1	0	1	0.00
TXN	P10599ITHIO_HUMAN Thioedoxin OS=Homo sapiens GN=TXN PE=1 SV=3	0	1	0.00
CD59	P13987ICD59_HUMAN CD59 glycoprotein OS=Homo sapiens GN=CD59 PE=1 SV=1	0	1	0.00
CUTA	Q60888ICUTA_HUMAN Protein CutA OS=Homo sapiens GN=CUTA PE=1 SV=2	0	1	0.00
ARPC3	O15145IARPC3_HUMAN Actin-related protein 2/3 complex subunit 3 OS=Homo sapiens GN=ARPC3 PE=1 SV=3	0	1	0.00
GOLGA4	Q13439IGOGA4_HUMAN Golgin subfamily A member 4 OS=Homo sapiens GN=GOLGA4 PE=1 SV=1	0	7	0.00
SAP18	O00422ISAP18_HUMAN Histone deacetylase complex subunit SAP18 OS=Homo sapiens GN=SAP18 PE=1 SV=1	0	3	0.00
SOD2	P04179ISODM_HUMAN Superoxide dismutase [Mn], mitochondrial OS=Homo sapiens GN=SOD2 PE=1 SV=2	0	4	0.00
MPDU1	O75352IMPU1_HUMAN Mannose-6-phosphate utilization defect 1 protein OS=Homo sapiens GN=MPDU1 PE=1 SV=2	0	1	0.00
ZC3H13	Q5T200C3HD_HUMAN Zinc finger CCCH domain-containing protein 13 OS=Homo sapiens GN=ZC3H13 PE=1 SV=1	0	1	0.00
NPC2	P61916INPC2_HUMAN Epididymal secretory protein E1 OS=Homo sapiens GN=NPC2 PE=1 SV=1	0	2	0.00
TMEM126A	Q9H061IT126A_HUMAN Transmembrane protein 126A OS=Homo sapiens GN=TMEM126A PE=1 SV=1	0	3	0.00
CLDN6	P56747ICLD6_HUMAN Claudin-6 OS=Homo sapiens GN=CLDN6 PE=1 SV=2	0	2	0.00
UBE2M	P61081IUBC12_HUMAN NEDD8-conjugating enzyme Ubc12 OS=Homo sapiens GN=UBE2M PE=1 SV=1	0	3	0.00
NPM3	O75607INPM3_HUMAN Nucleoplasmin-3 OS=Homo sapiens GN=NPM3 PE=1 SV=3	0	2	0.00
S100A9	P06702IS10A9_HUMAN Protein S100-A9 OS=Homo sapiens GN=S100A9 PE=1 SV=1	0	2	0.00
S100A8	P05109IS10A8_HUMAN Protein S100-A8 OS=Homo sapiens GN=S100A8 PE=1 SV=1	0	3	0.00
RTN4	Q9NQC3IRTN4_HUMAN Reticulon-4 OS=Homo sapiens GN=RTN4 PE=1 SV=2	0	1	0.00
LYZ	P61626ILYSY_HUMAN Lysozyme C OS=Homo sapiens GN=LYZ PE=1 SV=1	0	1	0.00
TMEM109	Q9BVC6ITM109_HUMAN Transmembrane protein 109 OS=Homo sapiens GN=TMEM109 PE=1 SV=1	0	2	0.00
PTP4A2	Q12974ITP4A2_HUMAN Protein tyrosine phosphatase type IVA 2 OS=Homo sapiens GN=PTP4A2 PE=1 SV=1	0	1	0.00
APOO	Q9BUR5IMIC26_HUMAN MICOS complex subunit MIC26 OS=Homo sapiens GN=APOO PE=1 SV=1	0	1	0.00
ATP5H	O75947IATP5H_HUMAN ATP synthase subunit d, mitochondrial OS=Homo sapiens GN=ATP5H PE=1 SV=3	0	6	0.00
MYDGF	Q969H8IMYDGF_HUMAN Myeloid-derived growth factor OS=Homo sapiens GN=MYDGF PE=1 SV=1	0	5	0.00
CFL2	Q9Y281ICOF2_HUMAN Cofilin-2 OS=Homo sapiens GN=CFL2 PE=1 SV=1	0	2	0.00
HNRNPK	P61978IHNRPK_HUMAN Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK PE=1 SV=1	0	2	0.00
BOK	Q9UMX3IBOK_HUMAN Bcl-2-related ovarian killer protein OS=Homo sapiens GN=BOK PE=1 SV=1	0	1	0.00
KDEL1R	P24390IERD21_HUMAN ER lumen protein-retaining receptor 1 OS=Homo sapiens GN=KDEL1R PE=1 SV=1	0	2	0.00
SKP1	P63208ISKP1_HUMAN S-phase kinase-associated protein 1 OS=Homo sapiens GN=SKP1 PE=1 SV=2	0	4	0.00
COMT	P21964ICOMT_HUMAN Catechol O-methyltransferase OS=Homo sapiens GN=COMT PE=1 SV=2	0	2	0.00
NDUFB10	O96000INDUBA_HUMAN NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=Homo sapiens GN=NDUFB10 PE=1 SV=3	0	3	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
CHST7	Q9NS84ICHST7_HUMAN Carbohydrate sulfotransferase 7 OS=Homo sapiens GN=CHST7 PE=1 SV=2	0	2	0.00
NDUFB5	O43674INDUB5_HUMAN NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 5, mitochondrial OS=Homo sapiens GN=NDUFB5 PE=1 SV=1	0	2	0.00
No_GeneID	L7N2F9L7N2F9_HUMAN Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=1	0	1	0.00
UQC2C	Q9BRT2IUQC2C_HUMAN Ubiquinol-cytochrome-c reductase complex assembly factor 2 OS=Homo sapiens GN=UQC2C PE=1 SV=1	0	1	0.00
PGRM2C	O15173IPGR2C_HUMAN Membrane-associated progesterone receptor component 2 OS=Homo sapiens GN=PGRM2C PE=1 SV=1	0	1	0.00
RABAC1	Q9U14IPRAF1_HUMAN Prenylated Rab acceptor protein 1 OS=Homo sapiens GN=RABAC1 PE=1 SV=1	0	1	0.00
HMG2A	P52926IHMG2A_HUMAN High mobility group protein HMGI-C OS=Homo sapiens GN=HMG2A PE=1 SV=1	0	3	0.00
NDUFB9	Q9Y6M9INDUB9_HUMAN NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 9 OS=Homo sapiens GN=NDUFB9 PE=1 SV=3	0	1	0.00
RPS27	P42677IRPS27_HUMAN 40S ribosomal protein S27 OS=Homo sapiens GN=RPS27 PE=1 SV=3	0	1	0.00
SNRPD3	P62318ISMD3_HUMAN Small nuclear ribonucleoprotein Sm D3 OS=Homo sapiens GN=SNRPD3 PE=1 SV=1	0	2	0.00
MYL6	P60660IMYL6_HUMAN Myosin light polypeptide 6 OS=Homo sapiens GN=MYL6 PE=1 SV=2	0	2	0.00
MYL12B	O14950IML12B_HUMAN Myosin regulatory light chain 12B OS=Homo sapiens GN=MYL12B PE=1 SV=2	0	5	0.00
No_GeneID	P01608IKV116_HUMAN Ig kappa chain V-1 region Roy OS=Homo sapiens PE=1 SV=1	0	1	0.00
PSMB8	P28062IPSB8_HUMAN Proteasome subunit beta type-8 OS=Homo sapiens GN=PSMB8 PE=1 SV=3	0	4	0.00
TRAP1	Q12931ITRAP1_HUMAN Heat shock protein 75 kDa, mitochondrial OS=Homo sapiens GN=TRAP1 PE=1 SV=3	0	4	0.00
RPL36A	P83881IRL36A_HUMAN 60S ribosomal protein L36a OS=Homo sapiens GN=RPL36A PE=1 SV=2	0	4	0.00
NDUFS7	O75251INDUS7_HUMAN NADH dehydrogenase [ubiquinone] iron-sulfur protein 7, mitochondrial OS=Homo sapiens GN=NDUFS7 PE=1 SV=3	0	4	0.00
ARF6	P62330IARF6_HUMAN ADP-ribosylation factor 6 OS=Homo sapiens GN=ARF6 PE=1 SV=2	0	3	0.00
DSG1	Q02413IDSG1_HUMAN Desmoglein-1 OS=Homo sapiens GN=DSG1 PE=1 SV=2	0	6	0.00
GAA	P10253ILYAG_HUMAN Lysosomal alpha-glucosidase OS=Homo sapiens GN=GAA PE=1 SV=4	0	4	0.00
No_GeneID	P01594IKV102_HUMAN Ig kappa chain V-1 region AU OS=Homo sapiens PE=1 SV=1	0	1	0.00
RPL11	Q9Y387IRM11_HUMAN 39S ribosomal protein L11, mitochondrial OS=Homo sapiens GN=RPL11 PE=1 SV=1	0	2	0.00
CDH23	Q9H251ICAD23_HUMAN Cadherin-23 OS=Homo sapiens GN=CDH23 PE=1 SV=2	0	1	0.00
PPH	O43447IPPH_HUMAN Peptidyl-prolyl cis-trans isomerase H OS=Homo sapiens GN=PPH PE=1 SV=1	0	2	0.00
MPV17	P39210IMPV17_HUMAN Protein Mpv17 OS=Homo sapiens GN=MPV17 PE=1 SV=1	0	1	0.00
RAP2C	Q9Y3LSRAP2C_HUMAN Ras-related protein Rap-2c OS=Homo sapiens GN=RAP2C PE=1 SV=1	0	1	0.00
TMM17B	O60830ITM17B_HUMAN Mitochondrial import inner membrane translocase subunit Tim17-B OS=Homo sapiens GN=TMM17B PE=1 SV=1	0	1	0.00
NDUFB11	Q9NX14INDUBB_HUMAN NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 11, mitochondrial OS=Homo sapiens GN=NDUFB11 PE=1 SV=1	0	3	0.00
PHB	P35232IPHB_HUMAN Prohibitin OS=Homo sapiens GN=PHB PE=1 SV=1	0	2	0.00
NDUFAF3	Q9BU61INDUF3_HUMAN NADH dehydrogenase [ubiquinone] 1 alpha subcomplex assembly factor 3 OS=Homo sapiens GN=NDUFAF3 PE=1 SV=1	0	3	0.00
HMG1A	P17096IHMG1A_HUMAN High mobility group protein HMGI-I/HMG-Y OS=Homo sapiens GN=HMG1A PE=1 SV=3	0	4	0.00

DSTN	P60981IDEST_HUMAN Dextrin OS=Homo sapiens GN=DSTN PE=1 SV=3	0	3	0.00
CIRBP	Q14011CIRBP_HUMAN Cold-inducible RNA-binding protein OS=Homo sapiens GN=CIRBP PE=1 SV=1	0	3	0.00
ATP5D	P30049IATPD_HUMAN ATP synthase subunit delta, mitochondrial OS=Homo sapiens GN=ATP5D PE=1 SV=2	0	3	0.00
No_GeneID	H3BN98H3BN98_HUMAN Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=2	0	4	0.00
EMC4	Q5J8M3EMC4_HUMAN ER membrane protein complex subunit 4 OS=Homo sapiens GN=EMC4 PE=1 SV=2	0	2	0.00
PFN1	P07737PROF1_HUMAN Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2	0	3	0.00
GSG2	Q8TF76HASP_HUMAN Serine/threonine-protein kinase haspin OS=Homo sapiens GN=GSG2 PE=1 SV=3	0	1	0.00
SPCS3	P61009ISPCS3_HUMAN Signal peptidase complex subunit 3 OS=Homo sapiens GN=SPCS3 PE=1 SV=1	0	1	0.00
SH3RF3	Q8TEJ3SH3RF3_HUMAN SH3 domain-containing RING finger protein 3 OS=Homo sapiens GN=SH3RF3 PE=1 SV=2	0	1	0.00
TMEM67	Q5HYABMK53_HUMAN Meckelin OS=Homo sapiens GN=TMEM67 PE=1 SV=2	0	1	0.00
CDC42BP6	Q6DT37MRCKG_HUMAN Serine/threonine-protein kinase MRCK gamma OS=Homo sapiens GN=CDC42BP6 PE=1 SV=2	0	2	0.00
COP58	Q99627CSN8_HUMAN COP9 signalosome complex subunit 8 OS=Homo sapiens GN=COP58 PE=1 SV=1	0	4	0.00
COL11A1	P12107ICOB1A1_HUMAN Collagen alpha-1(X) chain OS=Homo sapiens GN=COL11A1 PE=1 SV=4	0	5	0.00
CHCHD2	Q9Y6H1IICH2_HUMAN Coiled-coil-helix-coiled-coil-helix domain-containing protein 2 OS=Homo sapiens GN=CHCHD2 PE=1 SV=1	0	2	0.00
PPIF	P30405PIPF_HUMAN Peptidyl-prolyl cis-trans isomerase F, mitochondrial OS=Homo sapiens GN=PPIF PE=1 SV=1	0	5	0.00
ISOC2	Q96AB3ISOC2_HUMAN Isochorismatase domain-containing protein 2 OS=Homo sapiens GN=ISOC2 PE=1 SV=1	0	2	0.00
No_GeneID	FBW031FBW031_HUMAN Uncharacterized protein (Fragment) OS=Homo sapiens PE=1 SV=1	0	3	0.00
PPLI3	Q9H2H8PPLI3_HUMAN Peptidyl-prolyl cis-trans isomerase-like 3 OS=Homo sapiens GN=PPLI3 PE=1 SV=1	0	1	0.00
UBE2D3	P61077IUB2D3_HUMAN Ubiquitin-conjugating enzyme E2 D3 OS=Homo sapiens GN=UBE2D3 PE=1 SV=1	0	1	0.00
EEF1E1	Q43324MCA3_HUMAN Eukaryotic translation elongation factor 1 epsilon-1 OS=Homo sapiens GN=EEF1E1 PE=1 SV=1	0	5	0.00
DR1	Q01658INC2B_HUMAN Protein Dr1 OS=Homo sapiens GN=DR1 PE=1 SV=1	0	1	0.00
NDUFAF2	Q8N183MMIT_HUMAN Mimitin, mitochondrial OS=Homo sapiens GN=NDUFAF2 PE=1 SV=1	0	2	0.00
CTSA	P10619IPPG_B_HUMAN Lysosomal protective protein OS=Homo sapiens GN=CTSA PE=1 SV=2	0	2	0.00
AP3S1	Q92572IAP3S1_HUMAN AP-3 complex subunit sigma-1 OS=Homo sapiens GN=AP3S1 PE=1 SV=1	0	1	0.00
H2AFX	P16104H2AX_HUMAN Histone H2AX OS=Homo sapiens GN=H2AFX PE=1 SV=2	0	1	0.00
FAM129B	Q96TA1INBL1_HUMAN Niban-like protein 1 OS=Homo sapiens GN=FAM129B PE=1 SV=3	0	1	0.00
IGKV1D-13	AOA0B4J2D9IAO0B4J2D9_HUMAN Protein IGKV1D-13 (Fragment) OS=Homo sapiens GN=IGKV1D-13 PE=1 SV=1	0	1	0.00
EVA1A	Q9H8M9IEVA1A_HUMAN Protein eva-1 homolog A OS=Homo sapiens GN=EVA1A PE=1 SV=1	0	1	0.00
HSP90B1	P14625IENPL_HUMAN Endoplasmic reticulum protein OS=Homo sapiens GN=HSP90B1 PE=1 SV=1	0	4	0.00
RAB39A	Q14964R39A_HUMAN Ras-related protein Rab-39A OS=Homo sapiens GN=RAB39A PE=1 SV=2	0	1	0.00
NDUFB6	O95139INDUB6_HUMAN NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 6 OS=Homo sapiens GN=NDUFB6 PE=1 SV=3	0	2	0.00
BTF3	P20290BTF3_HUMAN Transcription factor BTF3 OS=Homo sapiens GN=BTF3 PE=1 SV=1	0	2	0.00
MGST3	O14880MGST3_HUMAN Microsomal glutathione S-transferase 3 OS=Homo sapiens GN=MGST3 PE=1 SV=1	0	2	0.00
TCEB2	Q15370IELOB_HUMAN Transcription elongation factor B polypeptide 2 OS=Homo sapiens GN=TCEB2 PE=1 SV=1	0	1	0.00
SHROOM2	Q17396ISHRM2_HUMAN Protein Shroom2 OS=Homo sapiens GN=SHROOM2 PE=1 SV=1	0	5	0.00
SSR3	Q9UNL2ISSRG_HUMAN Translocon-associated protein subunit gamma OS=Homo sapiens GN=SSR3 PE=1 SV=1	0	2	0.00
SERPINA12	Q8IW75SPA12_HUMAN Serpin A12 OS=Homo sapiens GN=SERPINA12 PE=1 SV=1	0	1	0.00
HIST1H2BD	P58876IH2BD1_HUMAN Histone H2B type 1-D OS=Homo sapiens GN=HIST1H2BD PE=1 SV=2	0	15	0.00
COX20	Q5R115ICOX20_HUMAN Cytochrome c oxidase protein 20 homolog OS=Homo sapiens GN=COX20 PE=1 SV=2	0	1	0.00
RAB19	A4D155IRAB19_HUMAN Ras-related protein Rab-19 OS=Homo sapiens GN=RAB19 PE=2 SV=2	0	1	0.00
REEP5	Q00765IREEP5_HUMAN Receptor expression-enhancing protein 5 OS=Homo sapiens GN=REEP5 PE=1 SV=3	0	1	0.00
MAN2B1	O00754IMA2B1_HUMAN Lysosomal alpha-mannosidase OS=Homo sapiens GN=MAN2B1 PE=1 SV=3	0	2	0.00
PRDX2	P32119IPRDX2_HUMAN Peroxiredoxin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5	0	10	0.00
CANX	P27824ICALX_HUMAN Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2	0	1	0.00
MADD	Q8WXG6MADD_HUMAN MAP kinase-activating death domain protein OS=Homo sapiens GN=MADD PE=1 SV=2	0	2	0.00
CASP14	P31944CASPE_HUMAN Caspase-14 OS=Homo sapiens GN=CASP14 PE=1 SV=2	0	1	0.00
HIST1H2AG	P0C058IH2A1_HUMAN Histone H2A type 1 OS=Homo sapiens GN=HIST1H2AG PE=1 SV=2	0	7	0.00
PKM	P14618IKPYM_HUMAN Pyruvate kinase PKM OS=Homo sapiens GN=PKM PE=1 SV=4	0	2	0.00
ARL5B	Q96K2IARL5B_HUMAN ADP-ribosylation factor-like protein 5B OS=Homo sapiens GN=ARL5B PE=1 SV=1	0	1	0.00
HJURP	Q8NCD3IHJURP_HUMAN Holliday junction recognition protein OS=Homo sapiens GN=HJURP PE=1 SV=2	0	1	0.00
HIST1H2AA	Q96QV6IH2A1A_HUMAN Histone H2A type 1-A OS=Homo sapiens GN=HIST1H2AA PE=1 SV=3	0	1	0.00
RPLP1	P05386RLA1_HUMAN 60S acidic ribosomal protein P1 OS=Homo sapiens GN=RPLP1 PE=1 SV=1	0	2	0.00
REEP6	Q96H93REEP6_HUMAN Receptor expression-enhancing protein 6 OS=Homo sapiens GN=REEP6 PE=1 SV=1	0	2	0.00
AVL9	Q8NBF6IAVL9_HUMAN Late secretory pathway protein AVL9 homolog OS=Homo sapiens GN=AVL9 PE=1 SV=1	0	1	0.00
CKAP4	Q07065ICKAP4_HUMAN Cytoskeleton-associated protein 4 OS=Homo sapiens GN=CKAP4 PE=1 SV=2	0	4	0.00
RER1	O15258RER1_HUMAN Protein RER1 OS=Homo sapiens GN=RER1 PE=1 SV=1	0	3	0.00
RPS10	P46783RPS10_HUMAN 40S ribosomal protein S10 OS=Homo sapiens GN=RPS10 PE=1 SV=1	0	7	0.00
FLG2	Q5D862IFILA2_HUMAN Filaggrin-2 OS=Homo sapiens GN=FLG2 PE=1 SV=1	0	1	0.00
LAMB1	P07942ILAMB1_HUMAN Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	0	1	0.00
RBM8A	Q9Y59IRBM8A_HUMAN RNA-binding protein 8A OS=Homo sapiens GN=RBM8A PE=1 SV=1	0	4	0.00
HYAL4	Q2M3T9IHYAL4_HUMAN Hyaluronidase-4 OS=Homo sapiens GN=HYAL4 PE=1 SV=2	0	1	0.00
SMIM1	B2RU24SMIM1_HUMAN Small integral membrane protein 1 OS=Homo sapiens GN=SMIM1 PE=1 SV=1	0	1	0.00
NDUFB8	O95169INDUB8_HUMAN NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 8, mitochondrial OS=Homo sapiens GN=NDUFB8 PE=1 SV=1	0	2	0.00
ENO1	P06733IENOA_HUMAN Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	0	2	0.00
ARHGAP42	A6N28IRHG42_HUMAN Rho GTPase-activating protein 42 OS=Homo sapiens GN=ARHGAP42 PE=1 SV=3	0	1	0.00
POLR2G	P62487IRPB7_HUMAN DNA-directed RNA polymerase II subunit RPB7 OS=Homo sapiens GN=POLR2G PE=1 SV=1	0	1	0.00
OLR1	P78380IOLR1_HUMAN Oxidized low-density lipoprotein receptor 1 OS=Homo sapiens GN=OLR1 PE=1 SV=1	0	1	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
GMFB	P60983IGMFB_HUMAN Glia maturation factor beta OS=Homo sapiens GN=GMFB PE=1 SV=2	0	1	0.00
ARL2	P36404IARL2_HUMAN ADP-ribosylation factor-like protein 2 OS=Homo sapiens GN=ARL2 PE=1 SV=4	0	1	0.00
MRPL49	Q13405IRML49_HUMAN 39S ribosomal protein L49, mitochondrial OS=Homo sapiens GN=MRPL49 PE=1 SV=1	0	4	0.00
RPS17	P08708IRS17_HUMAN 40S ribosomal protein S17 OS=Homo sapiens GN=RPS17 PE=1 SV=2	0	4	0.00
TMEM189-UBE2V1	I3LOA0I3LOA0_HUMAN HCG2044781 OS=Homo sapiens GN=TMEM189-UBE2V1 PE=4 SV=1	0	2	0.00
RPS10-NUDT3	S4R435I54R435_HUMAN Protein RPS10-NUDT3 (Fragment) OS=Homo sapiens GN=RPS10-NUDT3 PE=4 SV=1	0	1	0.00
PSMB3	P49720IPSB3_HUMAN Proteasome subunit beta type-3 OS=Homo sapiens GN=PSMB3 PE=1 SV=2	0	3	0.00
MRPS18A	Q9NVS2IR18A_HUMAN 28S ribosomal protein S18a, mitochondrial OS=Homo sapiens GN=MRPS18A PE=1 SV=1	0	1	0.00
STMN1	P16949ISTMN1_HUMAN Stathmin OS=Homo sapiens GN=STMN1 PE=1 SV=3	0	3	0.00
ARPC4	P59998IARPC4_HUMAN Actin-related protein 2/3 complex subunit 4 OS=Homo sapiens GN=ARPC4 PE=1 SV=3	0	5	0.00
ALG13	Q9N973IALG13_HUMAN Putative bifunctional UDP-N-acetylglucosamine transferase and deubiquitinase ALG13 OS=Homo sapiens GN=ALG13 PE=1 SV=2	0	1	0.00
ADCK3	Q8N60IADCK3_HUMAN Atypical kinase ADCK3, mitochondrial OS=Homo sapiens GN=ADCK3 PE=1 SV=1	0	1	0.00
GNPNAT1	Q96EK6IGNA1_HUMAN Glucosamine 6-phosphate N-acetyltransferase OS=Homo sapiens GN=GNPNAT1 PE=1 SV=1	0	3	0.00
RPS6	P62753IRS6_HUMAN 40S ribosomal protein S6 OS=Homo sapiens GN=RPS6 PE=1 SV=1	0	1	0.00
SRP14	P37108ISR14_HUMAN Signal recognition particle 14 kDa protein OS=Homo sapiens GN=SRP14 PE=1 SV=2	0	4	0.00
UFC1	Q9Y3C8IUCF1_HUMAN Ubiquitin-fold modifier-conjugating enzyme 1 OS=Homo sapiens GN=UFC1 PE=1 SV=3	0	2	0.00
NDUFS4	O43181INDUS4_HUMAN NADH dehydrogenase [ubiquinone] iron-sulfur protein 4, mitochondrial OS=Homo sapiens GN=NDUFS4 PE=1 SV=1	0	3	0.00
PSMB1	P20618IPSB1_HUMAN Proteasome subunit beta type-1 OS=Homo sapiens GN=PSMB1 PE=1 SV=2	0	1	0.00
DIABLO	Q9NR28IDBLO_HUMAN Diablo homolog, mitochondrial OS=Homo sapiens GN=DIABLO PE=1 SV=1	0	4	0.00
UBE2I	P63279IUBC9_HUMAN SUMO-conjugating enzyme UBC9 OS=Homo sapiens GN=UBE2I PE=1 SV=1	0	3	0.00

RPS5	P46782IR55_HUMAN 40S ribosomal protein S5 OS=Homo sapiens GN=RPS5 PE=1 SV=4	0	8	0.00
NTRK2	Q16620INTRK2_HUMAN BDNF/NT-3 growth factors receptor OS=Homo sapiens GN=NTRK2 PE=1 SV=1	0	2	0.00
CSTA	P01040CYTA_HUMAN Cystatin-A OS=Homo sapiens GN=CSTA PE=1 SV=1	0	2	0.00
NTPCR	Q9BSD7INTPCR_HUMAN Cancer-related nucleoside-triphosphatase OS=Homo sapiens GN=NTPCR PE=1 SV=1	0	2	0.00
DCTP1	Q9H773DCTP1_HUMAN dCTP pyrophosphatase 1 OS=Homo sapiens GN=DCTP1 PE=1 SV=1	0	6	0.00
USMG5	Q96X5IUSMG5_HUMAN Up-regulated during skeletal muscle growth protein 5 OS=Homo sapiens GN=USMG5 PE=1 SV=1	0	1	0.00
PFDN5	Q99471PFDN5_HUMAN Prefoldin subunit 5 OS=Homo sapiens GN=PFDN5 PE=1 SV=2	0	1	0.00
PCSK9	Q8NB77PCSK9_HUMAN Proprotein convertase subtilisin/kexin type 9 OS=Homo sapiens GN=PCSK9 PE=1 SV=3	0	1	0.00
NDUFA12	Q9U09INDUAC_HUMAN NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 12 OS=Homo sapiens GN=NDUFA12 PE=1 SV=1	0	4	0.00
VPS29	Q9URQ0VPS29_HUMAN Vacuolar protein sorting-associated protein 29 OS=Homo sapiens GN=VPS29 PE=1 SV=1	0	3	0.00
LRIT1	Q9P2V4LRIT1_HUMAN Leucine-rich repeat, immunoglobulin-like domain and transmembrane domain-containing protein 1 OS=Homo sapiens GN=LRIT1 PE=2 SV=1	0	2	0.00
DUSP3	P51452IDUS3_HUMAN Dual specificity protein phosphatase 3 OS=Homo sapiens GN=DUSP3 PE=1 SV=1	0	5	0.00
No_GeneID	E9PLD3IE9PLD3_HUMAN Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0	2	0.00
RPS19BP1	Q86WX3IAROS_HUMAN Active regulator of SIRT1 OS=Homo sapiens GN=RPS19BP1 PE=1 SV=1	0	3	0.00
RHEB	Q15382RHEB_HUMAN GTP-binding protein Rheb OS=Homo sapiens GN=RHEB PE=1 SV=1	0	2	0.00
MRPL40	Q9NQ50IRM40_HUMAN 39S ribosomal protein L40, mitochondrial OS=Homo sapiens GN=MRPL40 PE=1 SV=1	0	3	0.00
TMM23	Q14925TMM23_HUMAN Mitochondrial import inner membrane translocase subunit Tim23 OS=Homo sapiens GN=TMM23 PE=1 SV=1	0	1	0.00
HSPD1	P10809ICH60_HUMAN 60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2	0	10	0.00
CTSC	P53634ICATC_HUMAN Dipeptidyl peptidase 1 OS=Homo sapiens GN=CTSC PE=1 SV=2	0	4	0.00
BAX	Q07812IBAX_HUMAN Apoptosis regulator BAX OS=Homo sapiens GN=BAX PE=1 SV=1	0	1	0.00
RPS27L	Q71UM5IRS27L_HUMAN 40S ribosomal protein S27-like OS=Homo sapiens GN=RPS27L PE=1 SV=3	0	1	0.00
NDUFB8	P51970INDUA8_HUMAN NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 8 OS=Homo sapiens GN=NDUFB8 PE=1 SV=3	0	6	0.00
MRPL43	Q8N983IRM43_HUMAN 39S ribosomal protein L43, mitochondrial OS=Homo sapiens GN=MRPL43 PE=1 SV=1	0	2	0.00
TMED5	Q9Y3A6ITMED5_HUMAN Transmembrane emp24 domain-containing protein 5 OS=Homo sapiens GN=TMED5 PE=1 SV=1	0	1	0.00
MRPS28	Q9Y2Q9IRT28_HUMAN 28S ribosomal protein S28, mitochondrial OS=Homo sapiens GN=MRPS28 PE=1 SV=1	0	1	0.00
CMTM3	Q96MX0ICKLF3_HUMAN CKLF-like MARVEL transmembrane domain-containing protein 3 OS=Homo sapiens GN=CMTM3 PE=1 SV=1	0	1	0.00
NPM1	P06748NPM_HUMAN Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2	0	1	0.00
RLIM	Q9NVV2IRNF12_HUMAN E3 ubiquitin-protein ligase RLIM OS=Homo sapiens GN=RLIM PE=1 SV=3	0	1	0.00
LEMD2	Q8NC56LEMD2_HUMAN LEM domain-containing protein 2 OS=Homo sapiens GN=LEMD2 PE=1 SV=1	0	2	0.00
TMM21	Q9BVV7ITM21_HUMAN Mitochondrial import inner membrane translocase subunit Tim21 OS=Homo sapiens GN=TMM21 PE=1 SV=1	0	2	0.00
BRI3BP	Q8WY22BRI3B_HUMAN BRI3-binding protein OS=Homo sapiens GN=BRI3BP PE=1 SV=1	0	2	0.00
HIST1H2BO	P23527IH2B10_HUMAN Histone H2B type 1-O OS=Homo sapiens GN=HIST1H2BO PE=1 SV=3	0	2	0.00
AP151	P61966IAP151_HUMAN AP-1 complex subunit sigma-1A OS=Homo sapiens GN=AP151 PE=1 SV=1	0	1	0.00
RAP2B	P61225RAP2B_HUMAN Ras-related protein Rap-2b OS=Homo sapiens GN=RAP2B PE=1 SV=1	0	4	0.00
RPL37A	P61513IRL37A_HUMAN 60S ribosomal protein L37a OS=Homo sapiens GN=RPL37A PE=1 SV=2	0	1	0.00
HIST1H1D	P16402IH13_HUMAN Histone H1.3 OS=Homo sapiens GN=HIST1H1D PE=1 SV=2	0	4	0.00
RAP1A	P62834IRAP1A_HUMAN Ras-related protein Rap-1A OS=Homo sapiens GN=RAP1A PE=1 SV=1	0	3	0.00
CBX3	Q13185ICBX3_HUMAN Chromobox protein homolog 3 OS=Homo sapiens GN=CBX3 PE=1 SV=4	0	3	0.00
DOLP1	Q86YN1IDOPP1_HUMAN Dolichylidiphosphatase 1 OS=Homo sapiens GN=DOLP1 PE=2 SV=1	0	1	0.00
NUDCD2	Q8WVJ2INDUC2_HUMAN NudC domain-containing protein 2 OS=Homo sapiens GN=NUDCD2 PE=1 SV=1	0	1	0.00
GPR107	Q5VW38IGP107_HUMAN Protein GPR107 OS=Homo sapiens GN=GPR107 PE=1 SV=1	0	1	0.00
C20orf27	Q9GZN8ICTO27_HUMAN UPF0687 protein C20orf27 OS=Homo sapiens GN=C20orf27 PE=1 SV=3	0	1	0.00
No_GeneID	MQ0Z8BIMOQZK8_HUMAN Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0	1	0.00
DNAJC5	Q9H3Z4IDNJCS_HUMAN DnaJ homolog subfamily C member 5 OS=Homo sapiens GN=DNAJC5 PE=1 SV=1	0	1	0.00
NDUFB7	P17568INDUB7_HUMAN NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 7 OS=Homo sapiens GN=NDUFB7 PE=1 SV=4	0	4	0.00
PCDHA6	Q9UN73IPCD6_HUMAN Protocadherin alpha-6 OS=Homo sapiens GN=PCDHA6 PE=2 SV=1	0	1	0.00
EDF1	Q60869EDF1_HUMAN Endothelial differentiation-related factor 1 OS=Homo sapiens GN=EDF1 PE=1 SV=1	0	1	0.00
AGTRAP	Q6RW13IATRAP_HUMAN Type-1 angiotensin II receptor-associated protein OS=Homo sapiens GN=AGTRAP PE=1 SV=1	0	2	0.00
PFDN2	Q9UH99IPFD2_HUMAN Prefoldin subunit 2 OS=Homo sapiens GN=PFDN2 PE=1 SV=1	0	3	0.00
PSMB5	P28074IPSB5_HUMAN Proteasome subunit beta type-5 OS=Homo sapiens GN=PSMB5 PE=1 SV=3	0	8	0.00
JAGN1	Q8NSM9JAGN1_HUMAN Protein jagunal homolog 1 OS=Homo sapiens GN=JAGN1 PE=1 SV=1	0	2	0.00
MRPS10	P82664IRT10_HUMAN 28S ribosomal protein S10, mitochondrial OS=Homo sapiens GN=MRPS10 PE=1 SV=1	0	3	0.00
UBE2V2	Q15819IUBE2V2_HUMAN Ubiquitin-conjugating enzyme E2 variant 2 OS=Homo sapiens GN=UBE2V2 PE=1 SV=4	0	4	0.00
PCDH11X	Q9BZA7IPC11X_HUMAN Protocadherin-11 X-linked OS=Homo sapiens GN=PCDH11X PE=2 SV=1	0	1	0.00
HIST2H2AB	Q8LUE6IH2A2B_HUMAN Histone H2A type 2-B OS=Homo sapiens GN=HIST2H2AB PE=1 SV=3	0	1	0.00
TRMT2B	Q96GJ1TRM2_HUMAN tRNA (uracil(54)-C(5))-methyltransferase homolog OS=Homo sapiens GN=TRMT2B PE=1 SV=1	0	1	0.00
RNASEH2B	Q5TBB1IRNH2B_HUMAN Ribonuclease H2 subunit B OS=Homo sapiens GN=RNASEH2B PE=1 SV=1	0	1	0.00
HPICAL1	P37235HPICAL1_HUMAN Hippocalcin-like protein 1 OS=Homo sapiens GN=HPICAL1 PE=1 SV=3	0	3	0.00
FUNDC2	Q9BWH2IFUND2_HUMAN FUN14 domain-containing protein 2 OS=Homo sapiens GN=FUNDC2 PE=1 SV=2	0	2	0.00
POLR2H	P52434IRPAB3_HUMAN DNA-directed RNA polymerases I, II, and III subunit RPABC3 OS=Homo sapiens GN=POLR2H PE=1 SV=4	0	3	0.00
ICT1	Q14197ICT1_HUMAN Peptidyl-tRNA hydrolase ICT1, mitochondrial OS=Homo sapiens GN=ICT1 PE=1 SV=1	0	2	0.00
RBP4	P02753IRET4_HUMAN Retinol-binding protein 4 OS=Homo sapiens GN=RBP4 PE=1 SV=3	0	3	0.00
RAB6B	Q9NRW1IRAB6B_HUMAN Ras-related protein Rab-6B OS=Homo sapiens GN=RAB6B PE=1 SV=1	0	1	0.00
EIF1AY	O14602EIF1AY_HUMAN Eukaryotic translation initiation factor 1A, Y-chromosomal OS=Homo sapiens GN=EIF1AY PE=1 SV=4	0	4	0.00
RHOC	P08134IRHOC_HUMAN Rho-related GTP-binding protein Rhc OS=Homo sapiens GN=RHOC PE=1 SV=1	0	6	0.00
PDCD6	O75340PDCD6_HUMAN Programmed cell death protein 6 OS=Homo sapiens GN=PDCD6 PE=1 SV=1	0	2	0.00
CLCF1	Q9UBD9ICLCF1_HUMAN Cardiotrophin-like cytokine factor 1 OS=Homo sapiens GN=CLCF1 PE=1 SV=1	0	1	0.00
SNRPD1	P62314ISMD1_HUMAN Small nuclear ribonucleoprotein Sm D1 OS=Homo sapiens GN=SNRPD1 PE=1 SV=1	0	2	0.00
ARF5	P84085IARF5_HUMAN ADP-ribosylation factor 5 OS=Homo sapiens GN=ARF5 PE=1 SV=2	0	2	0.00
VSNL1	P62760IVSL1_HUMAN Visinin-like protein 1 OS=Homo sapiens GN=VSNL1 PE=1 SV=2	0	7	0.00
KIAA1671	Q9BY89IKI671_HUMAN Uncharacterized protein KIAA1671 OS=Homo sapiens GN=KIAA1671 PE=1 SV=2	0	5	0.00
PNKD	Q8N490IPNKD_HUMAN Probable hydrolase PNKD OS=Homo sapiens GN=PNKD PE=1 SV=2	0	1	0.00
FAM47DP	AGNHR8IFA47D_HUMAN Putative protein FAM47D OS=Homo sapiens GN=FAM47DP PE=5 SV=3	0	2	0.00
SNX3	O60493ISNX3_HUMAN Sorting nexin-3 OS=Homo sapiens GN=SNX3 PE=1 SV=3	0	4	0.00
CYBSA	P00167ICYBSA_HUMAN Cytochrome b5 OS=Homo sapiens GN=CYBSA PE=1 SV=2	0	5	0.00
MCTS1	Q9ULC4IMCTS1_HUMAN Malignant T-cell amplified sequence 1 OS=Homo sapiens GN=MCTS1 PE=1 SV=1	0	1	0.00
TMCO1	Q9UM00TMCO1_HUMAN Calcium load-activated calcium channel OS=Homo sapiens GN=TMCO1 PE=1 SV=1	0	3	0.00
CTS5B	P07858ICATB_HUMAN Cathepsin B OS=Homo sapiens GN=CTS5B PE=1 SV=3	0	2	0.00
PARK7	Q99497IPARK7_HUMAN Protein deglycase DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2	0	4	0.00
EIF5A	P63241IF5A1_HUMAN Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A PE=1 SV=2	0	3	0.00
TRAPPC3	O43617ITPPC3_HUMAN Trafficking protein particle complex subunit 3 OS=Homo sapiens GN=TRAPPC3 PE=1 SV=1	0	2	0.00
NHP2	Q9NX24NHP2_HUMAN H/ACA ribonucleoprotein complex subunit 2 OS=Homo sapiens GN=NHP2 PE=1 SV=1	0	3	0.00
MANF	P55145IMANF_HUMAN Mesencephalic astrocyte-derived neurotrophic factor OS=Homo sapiens GN=MANF PE=1 SV=3	0	5	0.00
HINT2	Q98X68IHINT2_HUMAN Histidine triad nucleotide-binding protein 2, mitochondrial OS=Homo sapiens GN=HINT2 PE=1 SV=1	0	2	0.00
SNRPD2	P62316ISMD2_HUMAN Small nuclear ribonucleoprotein Sm D2 OS=Homo sapiens GN=SNRPD2 PE=1 SV=1	0	1	0.00
RHEB	Q15382RHEB_HUMAN GTP-binding protein Rheb OS=Homo sapiens GN=RHEB PE=1 SV=1	0	1	0.00
RPS15	P62841IRS15_HUMAN 40S ribosomal protein S15 OS=Homo sapiens GN=RPS15 PE=1 SV=2	0	4	0.00
SPRYD4	Q8WWW5ISPRY4_HUMAN SPRY domain-containing protein 4 OS=Homo sapiens GN=SPRYD4 PE=1 SV=2	0	5	0.00

ARPC5	O15511ARPC5_HUMAN Actin-related protein 2/3 complex subunit 5 OS=Homo sapiens GN=ARPC5 PE=1 SV=3	0	2	0.00
GAPDH	P04406G3P_HUMAN Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	0	4	0.00
MRPL12	P5281SIRM12_HUMAN 39S ribosomal protein L12, mitochondrial OS=Homo sapiens GN=MRPL12 PE=1 SV=2	0	4	0.00
HSPAS	P11021GRP78_HUMAN 78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPAS PE=1 SV=2	0	3	0.00
ORMDL2	Q53FV1ORML2_HUMAN ORM1-like protein 2 OS=Homo sapiens GN=ORMDL2 PE=1 SV=2	0	1	0.00
CREBRF	Q8IUR6ICRERF_HUMAN CREB3 regulatory factor OS=Homo sapiens GN=CREBRF PE=1 SV=2	0	2	0.00
STRN3	Q13033STRN3_HUMAN Striatin-3 OS=Homo sapiens GN=STRN3 PE=1 SV=3	0	1	0.00
MRPL17	Q9NRX2IRM17_HUMAN 39S ribosomal protein L17, mitochondrial OS=Homo sapiens GN=MRPL17 PE=1 SV=1	0	3	0.00
RSL24D1	Q9LHA3IRLP24_HUMAN Probable ribosome biogenesis protein RLP24 OS=Homo sapiens GN=RSL24D1 PE=1 SV=1	0	1	0.00
TXNDC12	Q95881ITXD12_HUMAN Thioredoxin domain-containing protein 12 OS=Homo sapiens GN=TXNDC12 PE=1 SV=1	0	7	0.00
MRPL20	Q9BYC9IRM20_HUMAN 39S ribosomal protein L20, mitochondrial OS=Homo sapiens GN=MRPL20 PE=1 SV=1	0	2	0.00
EEF1A1	P68104IEF1A1_HUMAN Elongation factor 1-alpha 1 OS=Homo sapiens GN=EEF1A1 PE=1 SV=1	0	3	0.00
BTF3L4	Q96K17IBT3L4_HUMAN Transcription factor BTF3 homolog 4 OS=Homo sapiens GN=BTF3L4 PE=1 SV=1	0	3	0.00
PFND4	Q9NQ4IIPFD4_HUMAN Prefoldin subunit 4 OS=Homo sapiens GN=PFND4 PE=1 SV=1	0	1	0.00
ARL8B	Q9NVJ2IARL8B_HUMAN ADP-ribosylation factor-like protein 8B OS=Homo sapiens GN=ARL8B PE=1 SV=1	0	2	0.00
QDPR	P09417IDHPR_HUMAN Dihydropyridine reductase OS=Homo sapiens GN=QDPR PE=1 SV=2	0	1	0.00
DNPH1	O43598DNPH1_HUMAN 2'-deoxynucleoside 5'-phosphate N-hydrolase 1 OS=Homo sapiens GN=DNPH1 PE=1 SV=1	0	1	0.00
NRAS	P01111IRASN_HUMAN GTPase Nras OS=Homo sapiens GN=NRAS PE=1 SV=1	0	3	0.00
PLLP	Q9Y342IPLP_HUMAN Plasmalpin OS=Homo sapiens GN=PLLP PE=1 SV=1	0	1	0.00
S100A7	P31151S10A7_HUMAN Protein S100-A7 OS=Homo sapiens GN=S100A7 PE=1 SV=4	0	2	0.00
ARPC5L	Q9BPXSJARPSL_HUMAN Actin-related protein 2/3 complex subunit 5-like protein OS=Homo sapiens GN=ARPC5L PE=1 SV=1	0	2	0.00
ACOX1	Q15067IACOX1_HUMAN Peroxisomal acyl-coenzyme A oxidase 1 OS=Homo sapiens GN=ACOX1 PE=1 SV=3	0	4	0.00
FAM210A	Q96NDOIF210A_HUMAN Protein FAM210A OS=Homo sapiens GN=FAM210A PE=1 SV=2	0	1	0.00
CARHSP1	Q9Y2V2ICHSP1_HUMAN Calcium-regulated heat-stable protein 1 OS=Homo sapiens GN=CARHSP1 PE=1 SV=2	0	1	0.00