

Supporting Information Table 1. List of 1740 proteins identified from the three independent replicates.

Accession No.	Uniprot ID	Protein description	Protein Probability	Log2 Ratio (Treat/Control)
Q709C8-2	VP13C_HUMAN	Isoform 2 of Vacuolar protein sorting-associated protein 13C OS=Homo sapiens GN=VPS13C	1	9.964340868
Q8TEP8	CE192_HUMAN	Centrosomal protein of 192 kDa OS=Homo sapiens GN=CEP192 PE=1 SV=2	0.9957	9.964340868
A8MPV3	YR010_HUMAN	Putative uncharacterized protein ENSP0000344348 OS=Homo sapiens PE=2 SV=2	0.9786	9.964340868
Q5VX85	Q5VX85_HUMAN	Poly (ADP-ribose) polymerase 1 OS=Homo sapiens GN=PARP1 PE=2 SV=1	0.9715	9.964340868
A2A3K4-2	PTPC1_HUMAN	Isoform 2 of Protein tyrosine phosphatase domain-containing protein 1 OS=Homo sapiens GN=PTPDC1	0.9374	9.964340868
Q9Y4R8	TELO2_HUMAN	Telomere length regulation protein TEL2 homolog OS=Homo sapiens GN=TELO2 PE=1 SV=2	0.9322	9.964340868
O75962-2	TRIO_HUMAN	Isoform 2 of Triple functional domain protein OS=Homo sapiens GN=TRIO	0.9775	6.759289016
Q10571	MN1_HUMAN	Probable tumor suppressor protein MN1 OS=Homo sapiens GN=MN1 PE=1 SV=3	0.9953	6.431121473
Q63ZY3-2	KANK2_HUMAN	Isoform 2 of KN motif and ankyrin repeat domain-containing protein 2 OS=Homo sapiens GN=KANK2	0.8881	6.036503334
Q86X12-2	CNDG2_HUMAN	Isoform 2 of Condensin-2 complex subunit G2 OS=Homo sapiens GN=NCAPG2	0.9971	5.462379572
Q99490	AGAP2_HUMAN	Arf-GAP with GTPase	0.9935	5.136683578
O60282	KIF5C_HUMAN	Kinesin heavy chain isoform 5C OS=Homo sapiens GN=KIF5C PE=1 SV=1	0.8201	5.09212285
Q86VM9	ZCH18_HUMAN	Zinc finger CCCH domain-containing protein 18 OS=Homo sapiens GN=ZC3H18 PE=1 SV=2	0.9994	5.001351893
Q9Y6Q5-2	AP1M2_HUMAN	Isoform 2 of AP-1 complex subunit mu-2 OS=Homo sapiens GN=AP1M2	0.9996	4.987775616
F8WDE9	F8WDE9_HUMAN	Inosine-5'-monophosphate dehydrogenase 1 OS=Homo sapiens GN=IMPDH1 PE=4 SV=1	0.9583	4.961623328
Q96HA7-2	TONSL_HUMAN	Isoform 2 of Tonsoku-like protein OS=Homo sapiens GN=TONSL	0.9979	4.886062338
Q52LD8	RFTN2_HUMAN	Raftlin-2 OS=Homo sapiens GN=RFTN2 PE=1 SV=3	0.9998	4.827819025
Q8N2S1-2	LTBP4_HUMAN	Isoform 2 of Latent-transforming growth factor beta-binding protein 4 OS=Homo sapiens GN=LTBP4	0.9967	4.640389956
F5H450	F5H450_HUMAN	Frizzled-10 OS=Homo sapiens GN=FZD10 PE=4 SV=1	0.9794	4.583759754
Q9H4Z2-2	ZN335_HUMAN	Isoform 2 of Zinc finger protein 335 OS=Homo sapiens GN=ZNF335	0.8543	4.337711092
Q9H8E8-2	CSR2B_HUMAN	Isoform 2 of Cysteine-rich protein 2-binding protein OS=Homo sapiens GN=CSR2BP	0.945	4.279471296
Q14571	ITPR2_HUMAN	Inositol 1	1	4.266786541
A8K8P3-10	SF11_HUMAN	Isoform 10 of Protein SF11 homolog OS=Homo sapiens GN=SF11	0.9628	4.240314329
Q86YV0-2	RASL3_HUMAN	Isoform 2 of RAS protein activator like-3 OS=Homo sapiens GN=RASAL3	0.9876	4.213347282
Q9BX26	SYCP2_HUMAN	Synaptonemal complex protein 2 OS=Homo sapiens GN=SYCP2 PE=2 SV=2	0.9997	3.412781525
Q9P2E9-2	RRBP1_HUMAN	Isoform 1 of Ribosome-binding protein 1 OS=Homo sapiens GN=RRBP1	0.9999	3.289834465
Q96HP0	DOCK6_HUMAN	Dedicator of cytokinesis protein 6 OS=Homo sapiens GN=DOCK6 PE=1 SV=3	0.9055	3.173127433
Q9UF83	YM012_HUMAN	Uncharacterized protein DKFZp434B061 OS=Homo sapiens PE=2 SV=2	0.9996	3.163498732
Q6RI45	BRWD3_HUMAN	Bromodomain and WD repeat-containing protein 3 OS=Homo sapiens GN=BRWD3 PE=1 SV=2	0.998	3.148934105
O95714	HERC2_HUMAN	E3 ubiquitin-protein ligase HERC2 OS=Homo sapiens GN=HERC2 PE=1 SV=2	1	3.147306699
Q5T5J6	SWT1_HUMAN	Transcriptional protein SWT1 OS=Homo sapiens GN=SWT1 PE=2 SV=1	0.9789	3.061776198
Q8NCT3-6	K0895_HUMAN	Isoform 6 of Uncharacterized protein KIAA0895 OS=Homo sapiens GN=KIAA0895	0.8001	3.058316496
Q9NY37	ASIC5_HUMAN	Acid-sensing ion channel 5 OS=Homo sapiens GN=ASIC5 PE=1 SV=1	0.9137	3.016139703
Q14432	PDE3A_HUMAN	cGMP-inhibited 3'	0.9993	2.98550043
Q02880-2	TOP2B_HUMAN	Isoform Beta-1 of DNA topoisomerase 2-beta OS=Homo sapiens GN=TOP2B	0.9363	2.853995647
O00418	EEF2K_HUMAN	Eukaryotic elongation factor 2 kinase OS=Homo sapiens GN=EEF2K PE=1 SV=2	0.9562	2.825785627
B7Z1V0	B7Z1V0_HUMAN	Beta-chimaerin OS=Homo sapiens GN=CHN2 PE=2 SV=1	0.9992	2.782408565
Q92620	PRP16_HUMAN	Pre-mRNA-splicing factor ATP-dependent RNA helicase PRP16 OS=Homo sapiens GN=DHX38 PE=1 SV=	0.9996	2.769771739
F5H5Q9	F5H5Q9_HUMAN	Armadillo repeat-containing protein 4 OS=Homo sapiens GN=ARMC4 PE=4 SV=1	0.9512	2.711494907
Q9H254	SPTN4_HUMAN	Spectrin beta chain	1	2.596935142
Q385D2-2	LRRK1_HUMAN	Isoform 2 of Leucine-rich repeat serine/threonine-protein kinase 1 OS=Homo sapiens GN=LRRK1	0.9618	2.570462931
H3BQI5	H3BQI5_HUMAN	Protein FAM65A (Fragment) OS=Homo sapiens GN=FAM65A PE=4 SV=1	0.9363	2.44625623
Q9UH60	DCDC2_HUMAN	Doublecortin domain-containing protein 2 OS=Homo sapiens GN=DCDC2 PE=1 SV=2	0.9256	2.381283373
Q9NZC9	SMAL1_HUMAN	SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily A-like protein 1 OS	0.9342	2.292781749
Q7Z6Z7-2	HUWE1_HUMAN	Isoform 2 of E3 ubiquitin-protein ligase HUWE1 OS=Homo sapiens GN=HUWE1	1	2.253989266
Q86X10-2	RLGPB_HUMAN	Isoform 2 of Ral GTPase-activating protein subunit beta OS=Homo sapiens GN=RALGAPB	0.9503	2.195347598
A7M8M2	DISP2_HUMAN	Protein dispatched homolog 2 OS=Homo sapiens GN=DISP2 PE=2 SV=2	0.9127	2.17951105
P13928	ANXA8_HUMAN	Annexin A8 OS=Homo sapiens GN=ANXA8 PE=1 SV=3	0.949	2.09085343
P84550	SKOR1_HUMAN	SKI family transcriptional corepressor 1 OS=Homo sapiens GN=SKOR1 PE=1 SV=1	0.9375	2.084064265
G3V2J8	G3V2J8_HUMAN	Heat shock protein HSP 90-alpha (Fragment) OS=Homo sapiens GN=HSP90AA1 PE=4 SV=1	0.9886	2.006576663
Q9H967	WDR76_HUMAN	WD repeat-containing protein 76 OS=Homo sapiens GN=WDR76 PE=1 SV=2	0.973	2.070389328
Q5TG53-2	SEAS1_HUMAN	Isoform 2 of Putative uncharacterized protein SERTAD4-AS1 OS=Homo sapiens GN=SERTAD4-AS1	0.9769	2.049630768
Q3KQU3-2	MA7D1_HUMAN	Isoform 2 of MAP7 domain-containing protein 1 OS=Homo sapiens GN=MAP7D1	0.9131	2.028569152
I3L326	I3L326_HUMAN	Hepatocyte growth factor-regulated tyrosine kinase substrate (Fragment) OS=Homo sapiens GN=HGS PE	0.9442	2.010779839
O00764-2	PDXK_HUMAN	Isoform 2 of Pyridoxal kinase OS=Homo sapiens GN=PDXK	0.9939	1.967168608
P18583-3	SON_HUMAN	Isoform B of Protein SON OS=Homo sapiens GN=SON	0.9402	1.959770155
Q9ULT8	HECTD1_HUMAN	E3 ubiquitin-protein ligase HECTD1 OS=Homo sapiens GN=HECTD1 PE=1 SV=3	0.9305	1.761285273
Q8N3X1-2	FNBP4_HUMAN	Isoform 2 of Formin-binding protein 4 OS=Homo sapiens GN=FNBP4	0.9988	1.726831217
Q13315	ATM_HUMAN	Serine-protein kinase ATM OS=Homo sapiens GN=ATM PE=1 SV=3	0.9882	1.68448284
Q8NE28	SGK71_HUMAN	Protein kinase-like protein SqK071 OS=Homo sapiens GN=SGK071 PE=2 SV=4	0.9888	1.627606838
P16402	H13_HUMAN	Histone H1.3 OS=Homo sapiens GN=HIST1H1D PE=1 SV=2	0.9977	1.599317794
A4UGR9-2	XIRP2_HUMAN	Isoform 2 of Xin actin-binding repeat-containing protein 2 OS=Homo sapiens GN=XIRP2	0.9241	1.550900665
O00217	NDU58_HUMAN	NADH dehydrogenase [ubiquinone] iron-sulfur protein 8	1	1.5360529
J3KNE3	J3KNE3_HUMAN	Platelet-activating factor acetylhydrolase IB subunit beta (Fragment) OS=Homo sapiens GN=PFAH1B2 F	0.9828	1.531069493
P17480-2	UBF1_HUMAN	Isoform UBF2 of Nucleolar transcription factor 1 OS=Homo sapiens GN=UBTF	0.9834	1.469885976
Q9P281	BAHC1_HUMAN	BAH and coiled-coil domain-containing protein 1 OS=Homo sapiens GN=BAHCC1 PE=1 SV=3	0.9996	1.448900951
P60900	PSA6_HUMAN	Proteasome subunit alpha type-6 OS=Homo sapiens GN=PSMA6 PE=1 SV=1	1	1.316145742
O43464-2	HTRA2_HUMAN	Isoform 2 of Serine protease HTRA2	0.9996	1.286881148
Q6P6C2-3	ALKB5_HUMAN	Isoform 3 of Probable alpha-ketoglutarate-dependent dioxygenase ABH5 OS=Homo sapiens GN=ALKBH	0.9974	1.22650853
O43497-10	CAC1G_HUMAN	Isoform 10 of Voltage-dependent T-type calcium channel subunit alpha-1G OS=Homo sapiens GN=CACI	0.978	1.22650853
Q9UIA9	XPO7_HUMAN	Exportin-7 OS=Homo sapiens GN=XPO7 PE=1 SV=3	0.9524	1.22650853
Q86T03-3	TM55B_HUMAN	Isoform 3 of Transmembrane protein 55B OS=Homo sapiens GN=TMEM55B	0.9725	1.214124805
P51991-2	ROA3_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein A3 OS=Homo sapiens GN=HNRNPA3	0.9975	1.182692298
B7Z235	B7Z235_HUMAN	Uncharacterized protein OS=Homo sapiens PE=2 SV=1	0.9443	1.176322773
O43598-2	RCL_HUMAN	Isoform 2 of Deoxyribonucleoside 5'-monophosphate N-glycosidase OS=Homo sapiens GN=RCL	0.9986	1.169925001
Q8N6H7	ARFG2_HUMAN	ADP-ribosylation factor GTPase-activating protein 2 OS=Homo sapiens GN=ARFGAP2 PE=1 SV=1	1	1.137503524
H7BY33	H7BY33_HUMAN	Coiled-coil domain-containing protein 88B OS=Homo sapiens GN=CCDC88B PE=4 SV=1	1	1.13093087
Q3MU16-2	PA24E_HUMAN	Isoform 2 of Cytosolic phospholipase A2 epsilon OS=Homo sapiens GN=PLA2G4E	0.8599	1.13093087
Q96RW7-2	HMCN1_HUMAN	Isoform 2 of Hemicentin-1 OS=Homo sapiens GN=HMCN1	0.9947	1.124328135
Q9P035	HACD3_HUMAN	3-hydroxyacyl-CoA dehydratase 3 OS=Homo sapiens GN=PTPLAD1 PE=1 SV=2	0.9985	1.10433666
Q8WZA9	IRGQ_HUMAN	Immunity-related GTPase family Q protein OS=Homo sapiens GN=IRGQ PE=1 SV=1	0.9176	1.077242999
J3KT39	J3KT39_HUMAN	Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0.9993	1.049630768
Q9H1A6	Q9H1A6_HUMAN	DNA-directed RNA polymerase II subunit RPB11-a OS=Homo sapiens GN=POLR2J1 PE=2 SV=1	0.9483	1.042644337
Q09FC8-2	ZN415_HUMAN	Isoform 2 of Zinc finger protein 415 OS=Homo sapiens GN=ZNF415	1	1
Q12874	SF3A3_HUMAN	Splicing factor 3A subunit 3 OS=Homo sapiens GN=SF3A3 PE=1 SV=1	1	0.992768431
Q9NYV4-2	CDK12_HUMAN	Isoform 2 of Cyclin-dependent kinase 12 OS=Homo sapiens GN=CDK12	0.9993	0.992768431
Q9UL54-3	TAOK2_HUMAN	Isoform 3 of Serine/threonine-protein kinase TAO2 OS=Homo sapiens GN=TAOK2	0.9724	0.992768431
P59510	ATS20_HUMAN	A disintegrin and metalloproteinase with thrombospondin motifs 20 OS=Homo sapiens GN=ADAMTS20	0.9573	0.992768431

P10176	COX8A_HUMAN	Cytochrome c oxidase subunit 8A	0.9617	0.963474124
Q8TC05	MDM1_HUMAN	Nuclear protein MDM1 OS=Homo sapiens GN=MDM1 PE=1 SV=2	0.9879	0.948600847
Q9C0E4-2	GRIP2_HUMAN	Isoform 2 of Glutamate receptor-interacting protein 2 OS=Homo sapiens GN=GRIP2	0.9847	0.90303827
Q14164-2	IKKE_HUMAN	Isoform 2 of Inhibitor of nuclear factor kappa-B kinase subunit epsilon OS=Homo sapiens GN=IKBKE	0.9823	0.855989697
Q5F1R6-2	DJC21_HUMAN	Isoform 2 of DnaJ homolog subfamily C member 21 OS=Homo sapiens GN=DNAJC21	0.9888	0.847996907
Q13797	ITA9_HUMAN	Integrin alpha-9 OS=Homo sapiens GN=ITGA9 PE=1 SV=2	0.9991	0.82374936
Q9BT67	NFIP1_HUMAN	NEDD4 family-interacting protein 1 OS=Homo sapiens GN=NFIP1 PE=1 SV=1	0.9673	0.82374936
Q6UW56-3	ARAID_HUMAN	Isoform 3 of All-trans retinoic acid-induced differentiation factor OS=Homo sapiens GN=ATRAID	0.9492	0.782408565
Q9P1Y6-2	PHRF1_HUMAN	Isoform 2 of PHD and RING finger domain-containing protein 1 OS=Homo sapiens GN=PHRF1	0.9969	0.773996325
Q86UP2-2	KTN1_HUMAN	Isoform 2 of Kinectin OS=Homo sapiens GN=KTN1	0.9924	0.765534746
Q6DN12-2	MCTP2_HUMAN	Isoform 2 of Multiple C2 and transmembrane domain-containing protein 2 OS=Homo sapiens GN=MCT1	0.9522	0.757023247
Q09028-2	RBBP4_HUMAN	Isoform 2 of Histone-binding protein RBBP4 OS=Homo sapiens GN=RBBP4	1	0.731183242
Q14683	SMC1A_HUMAN	Structural maintenance of chromosomes protein 1A OS=Homo sapiens GN=SMC1A PE=1 SV=2	1	0.731183242
Q9NZM5	GSCR2_HUMAN	Glioma tumor suppressor candidate region gene 2 protein OS=Homo sapiens GN=GLTSCR2 PE=1 SV=2	0.9986	0.731183242
Q01469	FABP5_HUMAN	Fatty acid-binding protein	1	0.722466024
O14559-11	RHG33_HUMAN	Isoform 3 of Rho GTPase-activating protein 33 OS=Homo sapiens GN=ARHGAP33	0.994	0.722466024
P22105	TENX_HUMAN	Tenascin-X OS=Homo sapiens GN=TNXB PE=1 SV=3	1	0.713695815
Q8TAT6-2	NPL4_HUMAN	Isoform 2 of Nuclear protein localization protein 4 homolog OS=Homo sapiens GN=NPLOC4	0.999	0.713695815
Q58FF6	H90B4_HUMAN	Putative heat shock protein HSP 90-beta 4 OS=Homo sapiens GN=HSP90AB4 PE=5 SV=1	0.9905	0.713695815
HOYKE0	HOYKE0_HUMAN	WD repeat-containing protein 72 OS=Homo sapiens GN=WDR72 PE=4 SV=1	0.9668	0.704871964
Q96Q15	SMG1_HUMAN	Serine/threonine-protein kinase SMG1 OS=Homo sapiens GN=SMG1 PE=1 SV=3	0.9981	0.695993813
P0C055	H2AZ_HUMAN	Histone H2A.Z OS=Homo sapiens GN=H2AFZ PE=1 SV=2	0.9983	0.687060688
P49591	SYSC_HUMAN	Serine--tRNA ligase	1	0.678071905
Q86SP6	GP149_HUMAN	Probable G-protein coupled receptor 149 OS=Homo sapiens GN=GPR149 PE=2 SV=2	0.9634	0.659924558
J3KR12	J3KR12_HUMAN	Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0.9949	0.659924558
Q9Y4A5-2	TRRAP_HUMAN	Isoform 2 of Transformation/transcription domain-associated protein OS=Homo sapiens GN=TRRAP	0.9728	0.632268215
P98160	PGBM_HUMAN	Basement membrane-specific heparan sulfate proteoglycan core protein OS=Homo sapiens GN=HSPG2	0.9939	0.613531653
Q96AE4-2	FUBP1_HUMAN	Isoform 2 of Far upstream element-binding protein 1 OS=Homo sapiens GN=FUBP1	0.998	0.604071324
P69849	NOMO3_HUMAN	Nodal modulator 3 OS=Homo sapiens GN=NOMO3 PE=2 SV=2	0.9997	0.584962501
Q12972	PP1R8_HUMAN	Nuclear inhibitor of protein phosphatase 1 OS=Homo sapiens GN=PPP1R8 PE=1 SV=2	0.9984	0.584962501
Q9BWW7	SCR11_HUMAN	Transcriptional repressor scratch 1 OS=Homo sapiens GN=SCR11 PE=1 SV=1	0.9997	0.575312331
Q969S2-2	NEIL2_HUMAN	Isoform 2 of Endonuclease 8-like 2 OS=Homo sapiens GN=NEIL2	0.9644	0.555816155
Q15758	AAAT_HUMAN	Neutral amino acid transporter B(0) OS=Homo sapiens GN=SLC1A5 PE=1 SV=2	1	0.545968369
O15230	LAMA5_HUMAN	Laminin subunit alpha-5 OS=Homo sapiens GN=LAMA5 PE=1 SV=8	0.9651	0.50589093
P53597	SUCA_HUMAN	Succinyl-CoA ligase [ADP/GDP-forming] subunit alpha	0.9878	0.495695163
O15399	NMDE4_HUMAN	Glutamate [NMDA] receptor subunit epsilon-4 OS=Homo sapiens GN=GRIN2D PE=2 SV=2	0.8864	0.495695163
Q00059	TFAM_HUMAN	Transcription factor A	0.8997	0.475084883
Q8WZ42-2	TITIN_HUMAN	Isoform 2 of Titin OS=Homo sapiens GN=TTN	1	0.454175893
Q9NR99	MXRA5_HUMAN	Matrix-remodeling-associated protein 5 OS=Homo sapiens GN=MXRA5 PE=2 SV=3	0.9996	0.443606651
Q9Y617	SERC_HUMAN	Phosphoserine aminotransferase OS=Homo sapiens GN=PSAT1 PE=1 SV=2	1	0.432959407
O14787-2	TNPO2_HUMAN	Isoform 2 of Transportin-2 OS=Homo sapiens GN=TNPO2	0.982	0.432959407
Q9Y613	FHOD1_HUMAN	FH1/FH2 domain-containing protein 1 OS=Homo sapiens GN=FHOD1 PE=1 SV=3	0.9502	0.432959407
P08243	ASNS_HUMAN	Asparagine synthetase [glutamine-hydrolyzing] OS=Homo sapiens GN=ASNS PE=1 SV=4	0.9977	0.389566812
P02545-2	LMNA_HUMAN	Isoform C of Prelamin-A/C OS=Homo sapiens GN=LMNA	0.8529	0.389566812
P83731	RL24_HUMAN	60S ribosomal protein L24 OS=Homo sapiens GN=RPL24 PE=1 SV=1	1	0.378511623
Q92769	HDAC2_HUMAN	Histone deacetylase 2 OS=Homo sapiens GN=HDAC2 PE=1 SV=2	0.963	0.378511623
Q9P242	NYAP2_HUMAN	Neuronal tyrosine-phosphorylated phosphoinositide-3-kinase adapter 2 OS=Homo sapiens GN=NYAP2 F	0.972	0.367371066
P62750	RL23A_HUMAN	60S ribosomal protein L23a OS=Homo sapiens GN=RPL23A PE=1 SV=1	0.9996	0.35614381
A6ND36-2	FAM83G_HUMAN	Isoform 2 of Protein FAM83G OS=Homo sapiens GN=FAM83G	0.9975	0.35614381
P00491	PNPH_HUMAN	Purine nucleoside phosphorylase OS=Homo sapiens GN=PNP PE=1 SV=2	1	0.344828497
H7BZJ3	H7BZJ3_HUMAN	Thioredoxin (Fragment) OS=Homo sapiens GN=PDIA3 PE=3 SV=1	1	0.344828497
P07355-2	ANXA2_HUMAN	Isoform 2 of Annexin A2 OS=Homo sapiens GN=ANXA2	1	0.344828497
P18754-2	RCC1_HUMAN	Isoform 2 of Regulator of chromosome condensation OS=Homo sapiens GN=RCC1	1	0.344828497
P53999	TCP4_HUMAN	Activated RNA polymerase II transcriptional coactivator p15 OS=Homo sapiens GN=SUB1 PE=1 SV=3	1	0.333423734
Q9HAV7	GRPE1_HUMAN	GrpE protein homolog 1	1	0.333423734
O15144	ARPC2_HUMAN	Actin-related protein 2/3 complex subunit 2 OS=Homo sapiens GN=ARPC2 PE=1 SV=1	0.9945	0.333423734
Q12791-2	KCMA1_HUMAN	Isoform 2 of Calcium-activated potassium channel subunit alpha-1 OS=Homo sapiens GN=KCNMA1	0.9346	0.333423734
Q9BZ55-1	API5_HUMAN	Isoform 1 of Apoptosis inhibitor 5 OS=Homo sapiens GN=API5	1	0.321928095
Q9YZZ0-2	SUGT1_HUMAN	Isoform 2 of Suppressor of G2 allele of SKP1 homolog OS=Homo sapiens GN=SUGT1	1	0.321928095
Q5XX13-2	FBW10_HUMAN	Isoform 2 of F-box/WD repeat-containing protein 10 OS=Homo sapiens GN=FBXW10	0.933	0.321928095
P14866	HNRPL_HUMAN	Heterogeneous nuclear ribonucleoprotein L OS=Homo sapiens GN=HNRPL PE=1 SV=2	1	0.310340121
Q92688-2	AN32B_HUMAN	Isoform 2 of Acidic leucine-rich nuclear phosphoprotein 32 family member B OS=Homo sapiens GN=AN	1	0.310340121
P62191	PRS4_HUMAN	26S protease regulatory subunit 4 OS=Homo sapiens GN=PSMC1 PE=1 SV=1	1	0.286881148
P62424	RL7A_HUMAN	60S ribosomal protein L7a OS=Homo sapiens GN=RPL7A PE=1 SV=2	1	0.286881148
P08195-2	4F2_HUMAN	Isoform 2 of 4F2 cell-surface antigen heavy chain OS=Homo sapiens GN=SLC3A2	1	0.286881148
H0Y858	H0Y858_HUMAN	Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=1	0.9999	0.286881148
P50502	F10A1_HUMAN	Hsc70-interacting protein OS=Homo sapiens GN=ST13 PE=1 SV=2	0.9998	0.286881148
B4DWW1	B4DWW1_HUMAN	ADP-ribosylation factor-like protein 1 OS=Homo sapiens GN=ARL1 PE=2 SV=1	0.9991	0.286881148
Q9C0K0-2	BC11B_HUMAN	Isoform 2 of B-cell lymphoma/leukemia 11B OS=Homo sapiens GN=BCL11B	0.9859	0.286881148
P46926	GNPI1_HUMAN	Glucosamine-6-phosphate isomerase 1 OS=Homo sapiens GN=GNPDA1 PE=1 SV=1	1	0.275007047
P61313	RL15_HUMAN	60S ribosomal protein L15 OS=Homo sapiens GN=RPL15 PE=1 SV=2	1	0.275007047
J3KN67	J3KN67_HUMAN	Tropomyosin alpha-3 chain OS=Homo sapiens GN=TPM3 PE=3 SV=1	1	0.275007047
P61163	ACTZ_HUMAN	Alpha-centractin OS=Homo sapiens GN=ACTR1A PE=1 SV=1	1	0.275007047
Q9Y3D6	FIS1_HUMAN	Mitochondrial fission 1 protein OS=Homo sapiens GN=FIS1 PE=1 SV=2	1	0.263034406
P54920	SNAA_HUMAN	Alpha-soluble NSF attachment protein OS=Homo sapiens GN=NAPA PE=1 SV=3	1	0.263034406
Q5SSJ5-2	HP1B3_HUMAN	Isoform 2 of Heterochromatin protein 1-binding protein 3 OS=Homo sapiens GN=HP1BP3	0.9997	0.263034406
Q92820	GGH_HUMAN	Gamma-glutamyl hydrolase OS=Homo sapiens GN=GGH PE=1 SV=2	0.9976	0.263034406
Q5SNV9-2	CA167_HUMAN	Isoform 2 of Uncharacterized protein C1orf167 OS=Homo sapiens GN=C1orf167	0.9434	0.263034406
P60981-2	DEST_HUMAN	Isoform 2 of Destrin OS=Homo sapiens GN=DESTN	1	0.250961574
Q8N2E6	TOR2X_HUMAN	Prosalusin OS=Homo sapiens GN=TOR2A PE=1 SV=1	0.9888	0.250961574
Q08170	SRSF4_HUMAN	Serine/arginine-rich splicing factor 4 OS=Homo sapiens GN=SRSF4 PE=1 SV=2	0.9517	0.250961574
O15355	PPM1G_HUMAN	Protein phosphatase 1G OS=Homo sapiens GN=PPM1G PE=1 SV=1	1	0.23878686
P13995	MTDC_HUMAN	Bifunctional methylenetetrahydrofolate dehydrogenase/cyclohydrolase	1	0.23878686
P41227	NAA10_HUMAN	N-alpha-acetyltransferase 10 OS=Homo sapiens GN=NAA10 PE=1 SV=1	1	0.23878686
P82979	SARNP_HUMAN	SAP domain-containing ribonucleoprotein OS=Homo sapiens GN=SARNP PE=1 SV=3	1	0.23878686
Q13085-2	ACACA_HUMAN	Isoform 2 of Acetyl-CoA carboxylase 1 OS=Homo sapiens GN=ACACA	1	0.23878686
Q9BQ04	RBM4B_HUMAN	RNA-binding protein 4B OS=Homo sapiens GN=RBM4B PE=1 SV=1	1	0.23878686
Q13158	FADD_HUMAN	Protein FADD OS=Homo sapiens GN=FADD PE=1 SV=1	0.9902	0.23878686
P35268	RL22_HUMAN	60S ribosomal protein L22 OS=Homo sapiens GN=RPL22 PE=1 SV=2	1	0.22650853
P28074	PSB5_HUMAN	Proteasome subunit beta type-5 OS=Homo sapiens GN=PSMB5 PE=1 SV=3	1	0.22650853
Q9C0H9-5	SRCN1_HUMAN	Isoform 4 of SRC kinase signaling inhibitor 1 OS=Homo sapiens GN=SRCIN1	1	0.22650853
Q9H3K6-2	BOLA2_HUMAN	Isoform 2 of BOLA-like protein 2 OS=Homo sapiens GN=BOLA2	0.9997	0.22650853

Q9P210	CPSF2_HUMAN	Cleavage and polyadenylation specificity factor subunit 2 OS=Homo sapiens GN=CPSF2 PE=1 SV=2	0.9997	0.22650853
Q09161	NCBP1_HUMAN	Nuclear cap-binding protein subunit 1 OS=Homo sapiens GN=NCBP1 PE=1 SV=1	0.9993	0.22650853
Q6PI77	BHLH9_HUMAN	Protein BHLHb9 OS=Homo sapiens GN=BHLHb9 PE=2 SV=1	0.9946	0.22650853
P53007	TXTP_HUMAN	Tricarboxylate transport protein	0.9823	0.22650853
Q9COG0-2	ZN407_HUMAN	Isoform 2 of Zinc finger protein 407 OS=Homo sapiens GN=ZNF407	0.942	0.22650853
O75131	CPNE3_HUMAN	Copine-3 OS=Homo sapiens GN=CPNE3 PE=1 SV=1	0.9149	0.22650853
Q9H773	DCTP1_HUMAN	dCTP pyrophosphatase 1 OS=Homo sapiens GN=DCTPP1 PE=1 SV=1	1	0.214124805
O00148	DX39A_HUMAN	ATP-dependent RNA helicase DDX39A OS=Homo sapiens GN=DDX39A PE=1 SV=2	1	0.214124805
Q96AG4	LRC59_HUMAN	Leucine-rich repeat-containing protein 59 OS=Homo sapiens GN=LRRRC59 PE=1 SV=1	1	0.214124805
P35580-2	MYH10_HUMAN	Isoform 2 of Myosin-10 OS=Homo sapiens GN=MYH10	1	0.214124805
Q8N9K3	YA015_HUMAN	Putative uncharacterized protein FUJ37004 OS=Homo sapiens PE=2 SV=1	0.9984	0.214124805
Q15459	SF3A1_HUMAN	Splicing factor 3A subunit 1 OS=Homo sapiens GN=SF3A1 PE=1 SV=1	0.9976	0.214124805
Q96M63	CC114_HUMAN	Coiled-coil domain-containing protein 114 OS=Homo sapiens GN=CCDC114 PE=2 SV=3	0.9157	0.214124805
P05455	LA_HUMAN	Lupus La protein OS=Homo sapiens GN=SSB PE=1 SV=2	1	0.201633861
P08133	ANXA6_HUMAN	Annexin A6 OS=Homo sapiens GN=ANXA6 PE=1 SV=3	1	0.201633861
P20700	LMNB1_HUMAN	Lamin-B1 OS=Homo sapiens GN=LMNB1 PE=1 SV=2	1	0.201633861
P46778	RL21_HUMAN	60S ribosomal protein L21 OS=Homo sapiens GN=RPL21 PE=1 SV=2	1	0.201633861
Q8N5C6	SRBD1_HUMAN	S1 RNA-binding domain-containing protein 1 OS=Homo sapiens GN=SRBD1 PE=1 SV=2	0.9935	0.201633861
P45974-2	UBP5_HUMAN	Isoform Short of Ubiquitin carboxyl-terminal hydrolase 5 OS=Homo sapiens GN=USP5	0.888	0.201633861
P62280	RS11_HUMAN	40S ribosomal protein S11 OS=Homo sapiens GN=RPS11 PE=1 SV=3	1	0.189033824
Q8NBT2	SPC24_HUMAN	Kinetochore protein Spc24 OS=Homo sapiens GN=SPC24 PE=1 SV=2	1	0.189033824
P14868	SYDC_HUMAN	Aspartate--tRNA ligase	1	0.189033824
Q9Y224	CN166_HUMAN	UPF0568 protein C14orf166 OS=Homo sapiens GN=C14orf166 PE=1 SV=1	1	0.189033824
P31946-2	1433B_HUMAN	Isoform Short of 14-3-3 protein beta/alpha OS=Homo sapiens GN=YWHAB	1	0.189033824
P40763-2	STAT3_HUMAN	Isoform Del-701 of Signal transducer and activator of transcription 3 OS=Homo sapiens GN=STAT3	0.9998	0.189033824
Q06323	PSME1_HUMAN	Proteasome activator complex subunit 1 OS=Homo sapiens GN=PSME1 PE=1 SV=1	0.9839	0.189033824
P62316	SMD2_HUMAN	Small nuclear ribonucleoprotein Sm D2 OS=Homo sapiens GN=SNRNP2 PE=1 SV=1	0.8743	0.189033824
Q04760-2	LGUL_HUMAN	Isoform 2 of Lactoylglutathione lyase OS=Homo sapiens GN=GL01	1	0.176322773
Q86V81	THOC4_HUMAN	THO complex subunit 4 OS=Homo sapiens GN=ALYREF PE=1 SV=3	1	0.176322773
Q9BWD1	THIC_HUMAN	Acetyl-CoA acetyltransferase	1	0.176322773
O00299	CLIC1_HUMAN	Chloride intracellular channel protein 1 OS=Homo sapiens GN=CLIC1 PE=1 SV=4	1	0.176322773
P25788-2	PSA3_HUMAN	Isoform 2 of Proteasome subunit alpha type-3 OS=Homo sapiens GN=PSMA3	1	0.176322773
Q02978	M2OM_HUMAN	Mitochondrial 2-oxoglutarate/malate carrier protein OS=Homo sapiens GN=SLC25A11 PE=1 SV=3	1	0.176322773
Q07955	SRSF1_HUMAN	Serine/arginine-rich splicing factor 1 OS=Homo sapiens GN=SRSF1 PE=1 SV=2	1	0.176322773
Q13428-2	TCOF_HUMAN	Isoform 2 of Treacle protein OS=Homo sapiens GN=TCOF1	1	0.176322773
Q8NE71-2	ABCF1_HUMAN	Isoform 2 of ATP-binding cassette sub-family F member 1 OS=Homo sapiens GN=ABCF1	1	0.176322773
Q9BQG0-2	MBB1A_HUMAN	Isoform 2 of Myb-binding protein 1A OS=Homo sapiens GN=MYBBP1A	1	0.176322773
Q04917	1433F_HUMAN	14-3-3 protein eta OS=Homo sapiens GN=YWHAH PE=1 SV=4	1	0.176322773
O75526	RMXL2_HUMAN	RNA-binding motif protein	0.995	0.163498732
P49792	RBP2_HUMAN	E3 SUMO-protein ligase RanBP2 OS=Homo sapiens GN=RANBP2 PE=1 SV=2	0.9466	0.163498732
Q8TEV9-2	SMCR8_HUMAN	Isoform 2 of Smith-Magenis syndrome chromosomal region candidate gene 8 protein OS=Homo sapiens	0.9039	0.163498732
O43175	SERA_HUMAN	D-3-phosphoglycerate dehydrogenase OS=Homo sapiens GN=PHGDH PE=1 SV=4	1	0.150559677
P30044-2	PRDX5_HUMAN	Isoform Cytoplasmic+peroxisomal of Peroxiredoxin-5	1	0.150559677
Q13011	ECH1_HUMAN	Delta(3)	1	0.150559677
Q99729-2	ROAA_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein A/B OS=Homo sapiens GN=HNRNPAB	1	0.150559677
P12270	TPR_HUMAN	Nucleoprotein TPR OS=Homo sapiens GN=TPR PE=1 SV=3	1	0.150559677
P34897-2	GLYM_HUMAN	Isoform 2 of Serine hydroxymethyltransferase	1	0.150559677
P52948-2	NUP98_HUMAN	Isoform 2 of Nuclear pore complex protein Nup98-Nup96 OS=Homo sapiens GN=NUP98	1	0.150559677
Q99613	EIF3C_HUMAN	Eukaryotic translation initiation factor 3 subunit C OS=Homo sapiens GN=EIF3C PE=1 SV=1	1	0.150559677
A5PL33-3	KRBA1_HUMAN	Isoform 3 of Protein KRBA1 OS=Homo sapiens GN=KRBA1	0.97	0.150559677
Q9BZP3-2	CR002_HUMAN	Isoform 2 of Putative uncharacterized protein encoded by LINC00470 OS=Homo sapiens GN=LINC00470	0.9663	0.150559677
Q01484-2	ANK2_HUMAN	Isoform 2 of Ankyrin-2 OS=Homo sapiens GN=ANK2	0.937	0.150559677
P22102	PUR2_HUMAN	Trifunctional purine biosynthetic protein adenosine-3 OS=Homo sapiens GN=GART PE=1 SV=1	1	0.137503524
P09012	SNRPA_HUMAN	U1 small nuclear ribonucleoprotein A OS=Homo sapiens GN=SNRPA PE=1 SV=3	1	0.137503524
Q8NB05	DHB11_HUMAN	Estradiol 17-beta-dehydrogenase 11 OS=Homo sapiens GN=HSD17B11 PE=1 SV=3	0.9832	0.137503524
A8K855-2	EFCB7_HUMAN	Isoform 2 of EF-hand calcium-binding domain-containing protein 7 OS=Homo sapiens GN=EFCAB7	0.9676	0.137503524
Q9H6R3	ACSS3_HUMAN	Acyl-CoA synthetase short-chain family member 3	0.9599	0.137503524
Q9Y266	NUDC_HUMAN	Nuclear migration protein nudC OS=Homo sapiens GN=NUDC PE=1 SV=1	1	0.124328135
P13797	PLST_HUMAN	Plastin-3 OS=Homo sapiens GN=PLS3 PE=1 SV=4	1	0.124328135
Q9Y490	TLN1_HUMAN	Talin-1 OS=Homo sapiens GN=TLN1 PE=1 SV=3	1	0.124328135
P55084	ECHB_HUMAN	Trifunctional enzyme subunit beta	1	0.124328135
P62495	ERF1_HUMAN	Eukaryotic peptide chain release factor subunit 1 OS=Homo sapiens GN=ETF1 PE=1 SV=3	1	0.124328135
P62906	RL10A_HUMAN	60S ribosomal protein L10a OS=Homo sapiens GN=RPL10A PE=1 SV=2	1	0.124328135
Q9P2G9	KLHL8_HUMAN	Kelch-like protein 8 OS=Homo sapiens GN=KLHL8 PE=2 SV=4	0.9994	0.124328135
Q9Y3C8	UFC1_HUMAN	Ubiquitin-fold modifier-conjugating enzyme 1 OS=Homo sapiens GN=UFC1 PE=1 SV=3	0.9928	0.124328135
O14929	HAT1_HUMAN	Histone acetyltransferase type B catalytic subunit OS=Homo sapiens GN=HAT1 PE=1 SV=1	0.9517	0.124328135
Q9UQE7	SMC3_HUMAN	Structural maintenance of chromosomes protein 3 OS=Homo sapiens GN=SMC3 PE=1 SV=2	0.9471	0.124328135
Q9UJU6-2	DBNL_HUMAN	Isoform 2 of Drebrin-like protein OS=Homo sapiens GN=DBNL	0.8911	0.124328135
Q92522	H1X_HUMAN	Histone H1x OS=Homo sapiens GN=H1FX PE=1 SV=1	1	0.111031312
Q9Y383-2	LCT7L2_HUMAN	Isoform 2 of Putative RNA-binding protein Luc7-like 2 OS=Homo sapiens GN=LUC7L2	1	0.111031312
O75390	CISY_HUMAN	Citrate synthase	1	0.111031312
P09936	UCHL1_HUMAN	Ubiquitin carboxyl-terminal hydrolase isozyme L1 OS=Homo sapiens GN=UCHL1 PE=1 SV=2	1	0.111031312
Q14315-2	FLNC_HUMAN	Isoform 2 of Filamin-C OS=Homo sapiens GN=FLNC	1	0.111031312
P41252	SYIC_HUMAN	Isoleucine--tRNA ligase	1	0.111031312
Q99623	PHB2_HUMAN	Prohibitin-2 OS=Homo sapiens GN=PHB2 PE=1 SV=2	1	0.111031312
C9JRL4	C9JRL4_HUMAN	Malate dehydrogenase	1	0.111031312
Q6PRD1	GP179_HUMAN	Probable G-protein coupled receptor 179 OS=Homo sapiens GN=GPR179 PE=1 SV=2	0.9997	0.111031312
P39748	FEN1_HUMAN	Flap endonuclease 1 OS=Homo sapiens GN=FEN1 PE=1 SV=1	0.9976	0.111031312
P16152	CBR1_HUMAN	Carbonyl reductase [NADPH] 1 OS=Homo sapiens GN=CBR1 PE=1 SV=3	0.9946	0.111031312
O14579-2	COPE_HUMAN	Isoform 2 of Coatomer subunit epsilon OS=Homo sapiens GN=COPE	0.9901	0.111031312
Q8N1F7-2	NUP93_HUMAN	Isoform 2 of Nuclear pore complex protein Nup93 OS=Homo sapiens GN=NUP93	0.9882	0.111031312
P08559-2	ODPA_HUMAN	Isoform 2 of Pyruvate dehydrogenase E1 component subunit alpha	0.9854	0.111031312
O43615	TIM44_HUMAN	Mitochondrial import inner membrane translocase subunit TIM44 OS=Homo sapiens GN=TIMM44 PE=1 SV=1	1	0.097610797
P11279	LAMP1_HUMAN	Lysosome-associated membrane glycoprotein 1 OS=Homo sapiens GN=LAMP1 PE=1 SV=3	1	0.097610797
P55265-2	DSRAD_HUMAN	Isoform 2 of Double-stranded RNA-specific adenosine deaminase OS=Homo sapiens GN=ADAR	1	0.097610797
P09874	PARP1_HUMAN	Poly [ADP-ribose] polymerase 1 OS=Homo sapiens GN=PARP1 PE=1 SV=4	1	0.097610797
Q9UQ80	PA2G4_HUMAN	Proliferation-associated protein 2G4 OS=Homo sapiens GN=PA2G4 PE=1 SV=3	1	0.097610797
P36578	RL4_HUMAN	60S ribosomal protein L4 OS=Homo sapiens GN=RPL4 PE=1 SV=5	1	0.097610797
P62258	1433E_HUMAN	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1	1	0.097610797
P63241-2	IF5A1_HUMAN	Isoform 2 of Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A	1	0.097610797
P62857	RS28_HUMAN	40S ribosomal protein S28 OS=Homo sapiens GN=RPS28 PE=1 SV=1	0.9998	0.097610797
P78417	GSTO1_HUMAN	Glutathione S-transferase omega-1 OS=Homo sapiens GN=GSTO1 PE=1 SV=2	1	0.084064265

P61158	ARP3_HUMAN	Actin-related protein 3 OS=Homo sapiens GN=ACTR3 PE=1 SV=3	1	0.084064265
Q06830	PRDX1_HUMAN	Peroxiredoxin-1 OS=Homo sapiens GN=PRDX1 PE=1 SV=1	1	0.084064265
Q96776-8	MMS19_HUMAN	Isoform 5 of MMS19 nucleotide excision repair protein homolog OS=Homo sapiens GN=MMS19	1	0.084064265
Q9UKN7	MYO15_HUMAN	Unconventional myosin-XV OS=Homo sapiens GN=MYO15A PE=1 SV=2	0.9999	0.084064265
O14979-2	HNRDL_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein D-like OS=Homo sapiens GN=HNRPDL	0.9805	0.084064265
Q99661-2	KIF2C_HUMAN	Isoform 2 of Kinesin-like protein KIF2C OS=Homo sapiens GN=KIF2C	0.9642	0.084064265
O00429-2	DNM1L_HUMAN	Isoform 4 of Dynamin-1-like protein OS=Homo sapiens GN=DNM1L	0.9186	0.084064265
P49207	RL34_HUMAN	60S ribosomal protein L34 OS=Homo sapiens GN=RPL34 PE=1 SV=3	1	0.070389328
P55854	SUMO3_HUMAN	Small ubiquitin-related modifier 3 OS=Homo sapiens GN=SUMO3 PE=1 SV=2	1	0.070389328
P62910	RL32_HUMAN	60S ribosomal protein L32 OS=Homo sapiens GN=RPL32 PE=1 SV=2	1	0.070389328
Q15691	MARE1_HUMAN	Microtubule-associated protein RP/EB family member 1 OS=Homo sapiens GN=MAPRE1 PE=1 SV=3	1	0.070389328
O43390-2	HNRPR_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein R OS=Homo sapiens GN=HNRNPR	1	0.070389328
P54687-3	BCAT1_HUMAN	Isoform 3 of Branched-chain-amino-acid aminotransferase	1	0.070389328
P61088	UBE2N_HUMAN	Ubiquitin-conjugating enzyme E2 N OS=Homo sapiens GN=UBE2N PE=1 SV=1	1	0.070389328
P62244	RS15A_HUMAN	40S ribosomal protein S15a OS=Homo sapiens GN=RPS15A PE=1 SV=2	1	0.070389328
Q9Y230	RUVB2_HUMAN	RuvB-like 2 OS=Homo sapiens GN=RUVBL2 PE=1 SV=3	1	0.070389328
A8K0Z3	WASH1_HUMAN	WAS protein family homolog 1 OS=Homo sapiens GN=WASH1 PE=1 SV=2	0.9998	0.070389328
Q9BXJ9-4	NAA15_HUMAN	Isoform 2 of N-alpha-acetyltransferase 15	0.9998	0.070389328
O43818	U3IP2_HUMAN	U3 small nucleolar RNA-interacting protein 2 OS=Homo sapiens GN=RRP9 PE=1 SV=1	0.9964	0.070389328
Q92979	NEP1_HUMAN	Ribosomal RNA small subunit methyltransferase NEP1 OS=Homo sapiens GN=EMG1 PE=1 SV=4	0.994	0.070389328
Q9Y6Y8	S23IP_HUMAN	SEC23-interacting protein OS=Homo sapiens GN=SEC23IP PE=1 SV=1	0.9744	0.070389328
P12035	K2C3_HUMAN	Keratin	0.9666	0.070389328
P33991	MCM4_HUMAN	DNA replication licensing factor MCM4 OS=Homo sapiens GN=MCM4 PE=1 SV=5	1	0.056583528
Q01081-2	U2AF1_HUMAN	Isoform 2 of Splicing factor U2AF 35 kDa subunit OS=Homo sapiens GN=U2AF1	1	0.056583528
P15121	ALDR_HUMAN	Aldehyde reductase OS=Homo sapiens GN=AKR1B1 PE=1 SV=3	1	0.056583528
P32969	RL9_HUMAN	60S ribosomal protein L9 OS=Homo sapiens GN=RPL9 PE=1 SV=1	1	0.056583528
P49588	SYAC_HUMAN	Alanine--tRNA ligase	1	0.056583528
Q13813-2	SPTN1_HUMAN	Isoform 2 of Spectrin alpha chain	1	0.056583528
Q14498-2	RBM39_HUMAN	Isoform 2 of RNA-binding protein 39 OS=Homo sapiens GN=RBM39	1	0.056583528
Q99436	PSB7_HUMAN	Proteasome subunit beta type-7 OS=Homo sapiens GN=PSMB7 PE=1 SV=1	1	0.056583528
P61981	1433G_HUMAN	14-3-3 protein gamma OS=Homo sapiens GN=YWHAG PE=1 SV=2	1	0.056583528
P55196-1	AFAD_HUMAN	Isoform 2 of Afadin OS=Homo sapiens GN=MLLT4	0.9999	0.056583528
O15371	EIF3D_HUMAN	Eukaryotic translation initiation factor 3 subunit D OS=Homo sapiens GN=EIF3D PE=1 SV=1	0.9992	0.056583528
P08123	CO1A2_HUMAN	Collagen alpha-2(I) chain OS=Homo sapiens GN=COL1A2 PE=1 SV=7	0.9972	0.056583528
D6RIA3	D6RIA3_HUMAN	Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0.9818	0.056583528
P51812	KS6A3_HUMAN	Ribosomal protein S6 kinase alpha-3 OS=Homo sapiens GN=RPS6KA3 PE=1 SV=1	0.9544	0.056583528
Q5H9R7-2	PP6R3_HUMAN	Isoform 2 of Serine/threonine-protein phosphatase 6 regulatory subunit 3 OS=Homo sapiens GN=PPP6R	0.9477	0.056583528
P04632	CPNS1_HUMAN	Calpain small subunit 1 OS=Homo sapiens GN=CAPNS1 PE=1 SV=1	1	0.042644337
P48556	PSMD8_HUMAN	26S proteasome non-ATPase regulatory subunit 8 OS=Homo sapiens GN=PSMD8 PE=1 SV=2	1	0.042644337
Q14566	MCM6_HUMAN	DNA replication licensing factor MCM6 OS=Homo sapiens GN=MCM6 PE=1 SV=1	1	0.042644337
Q5T2A2	CROCC_HUMAN	Rootletin OS=Homo sapiens GN=CROCC PE=1 SV=1	1	0.042644337
P13591-1	NCAM1_HUMAN	Isoform 2 of Neural cell adhesion molecule 1 OS=Homo sapiens GN=NCAM1	1	0.042644337
P15170-2	ERF3A_HUMAN	Isoform 2 of Eukaryotic peptide chain release factor GTP-binding subunit ERF3A OS=Homo sapiens GN=	1	0.042644337
P25205	MCM3_HUMAN	DNA replication licensing factor MCM3 OS=Homo sapiens GN=MCM3 PE=1 SV=3	1	0.042644337
P61204	ARF3_HUMAN	ADP-ribosylation factor 3 OS=Homo sapiens GN=ARF3 PE=1 SV=2	1	0.042644337
P84090	ERH_HUMAN	Enhancer of rudimentary homolog OS=Homo sapiens GN=ERH PE=1 SV=1	1	0.042644337
Q15181	IPYR_HUMAN	Inorganic pyrophosphatase OS=Homo sapiens GN=PPA1 PE=1 SV=2	1	0.042644337
Q9HB71	CYBP_HUMAN	Calycylin-binding protein OS=Homo sapiens GN=CACYBP PE=1 SV=2	1	0.042644337
Q9Y3F4	STRAP_HUMAN	Serine-threonine kinase receptor-associated protein OS=Homo sapiens GN=STRAP PE=1 SV=1	1	0.042644337
Q9NV17-1	FANCI_HUMAN	Isoform 1 of Fanconi anemia group I protein OS=Homo sapiens GN=FANCI	0.9999	0.042644337
Q9BT78	CSN4_HUMAN	COP9 signalosome complex subunit 4 OS=Homo sapiens GN=COPS4 PE=1 SV=1	0.9928	0.042644337
Q70CQ4	UBP31_HUMAN	Ubiquitin carboxyl-terminal hydrolase 31 OS=Homo sapiens GN=USP31 PE=1 SV=2	0.9974	0.042644337
P10606	COX5B_HUMAN	Cytochrome c oxidase subunit 5B	0.9813	0.042644337
Q92830-2	KAT2A_HUMAN	Isoform 2 of Histone acetyltransferase KAT2A OS=Homo sapiens GN=KAT2A	0.9675	0.042644337
P17174	AATC_HUMAN	Aspartate aminotransferase	1	0.028569152
P26368-2	U2AF2_HUMAN	Isoform 2 of Splicing factor U2AF 65 kDa subunit OS=Homo sapiens GN=U2AF2	1	0.028569152
P51648-2	AL3A2_HUMAN	Isoform 2 of Fatty aldehyde dehydrogenase OS=Homo sapiens GN=ALDH3A2	1	0.028569152
Q96P70	IPO9_HUMAN	Importin-9 OS=Homo sapiens GN=IPO9 PE=1 SV=3	1	0.028569152
Q9P2J5	SYLC_HUMAN	Leucine--tRNA ligase	1	0.028569152
P15531-2	NDKA_HUMAN	Isoform 2 of Nucleoside diphosphate kinase A OS=Homo sapiens GN=NME1	1	0.028569152
P25786-2	PSA1_HUMAN	Isoform Long of Proteasome subunit alpha type-1 OS=Homo sapiens GN=PSMA1	1	0.028569152
Q01082	SPTB2_HUMAN	Spectrin beta chain	1	0.028569152
Q12931	TRAP1_HUMAN	Heat shock protein 75 kDa	1	0.028569152
P00492	HPRT_HUMAN	Hypoxanthine-guanine phosphoribosyltransferase OS=Homo sapiens GN=HPRT1 PE=1 SV=2	0.9999	0.028569152
P06454-2	PTMA_HUMAN	Isoform 2 of Prothymosin alpha OS=Homo sapiens GN=PTMA	0.9994	0.028569152
Q9BTE3-2	MCMBP_HUMAN	Isoform 2 of Mini-chromosome maintenance complex-binding protein OS=Homo sapiens GN=MCMBP	0.9929	0.028569152
Q9ULI3	HEG1_HUMAN	Protein HEG homolog 1 OS=Homo sapiens GN=HEG1 PE=1 SV=3	0.9465	0.028569152
Q9H0B3-3	K1683_HUMAN	Isoform 3 of Uncharacterized protein KIAA1683 OS=Homo sapiens GN=KIAA1683	0.9327	0.028569152
Q8N2E2	VWDE_HUMAN	von Willebrand factor D and EGF domain-containing protein OS=Homo sapiens GN=VWDE PE=2 SV=4	0.9099	0.028569152
P21266	GSTM3_HUMAN	Glutathione S-transferase Mu 3 OS=Homo sapiens GN=GSTM3 PE=1 SV=3	0.8802	0.028569152
O00232-2	PSD12_HUMAN	Isoform 2 of 26S proteasome non-ATPase regulatory subunit 12 OS=Homo sapiens GN=PSMD12	1	0.014355293
O43143	DHX15_HUMAN	Putative pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15 OS=Homo sapiens GN=DHX15 I	1	0.014355293
Q14444-2	CAPR1_HUMAN	Isoform 2 of Caprin-1 OS=Homo sapiens GN=CAPRIN1	1	0.014355293
Q9BPX3	CND3_HUMAN	Condensin complex subunit 3 OS=Homo sapiens GN=NCAPG PE=1 SV=1	1	0.014355293
Q9NWW8-3	BABA1_HUMAN	Isoform 3 of BRISC and BRCA1-A complex member 1 OS=Homo sapiens GN=BABAM1	1	0.014355293
O14980	XPO1_HUMAN	Exportin-1 OS=Homo sapiens GN=XPO1 PE=1 SV=1	1	0.014355293
P00367	DHE3_HUMAN	Glutamate dehydrogenase 1	1	0.014355293
P25789	PSA4_HUMAN	Proteasome subunit alpha type-4 OS=Homo sapiens GN=PSMA4 PE=1 SV=1	1	0.014355293
P29401	TKT_HUMAN	Transketolase OS=Homo sapiens GN=TKT PE=1 SV=3	1	0.014355293
P31150	GDIA_HUMAN	Rab GDP dissociation inhibitor alpha OS=Homo sapiens GN=GD1I PE=1 SV=2	1	0.014355293
P33992	MCM5_HUMAN	DNA replication licensing factor MCM5 OS=Homo sapiens GN=MCM5 PE=1 SV=5	1	0.014355293
P38117	ETFB_HUMAN	Electron transfer flavoprotein subunit beta OS=Homo sapiens GN=ETFB PE=1 SV=3	1	0.014355293
P49368-2	TCPG_HUMAN	Isoform 2 of T-complex protein 1 subunit gamma OS=Homo sapiens GN=CCT3	1	0.014355293
P61978-2	HNRPK_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens GN=HNRNPK	1	0.014355293
P78347-2	GTF2I_HUMAN	Isoform 2 of General transcription factor II-I OS=Homo sapiens GN=GTF2I	1	0.014355293
Q15185	TEBP_HUMAN	Prostaglandin E synthase 3 OS=Homo sapiens GN=PTGES3 PE=1 SV=1	1	0.014355293
Q92945	FUBP2_HUMAN	Far upstream element-binding protein 2 OS=Homo sapiens GN=KHSRP PE=1 SV=4	1	0.014355293
Q9HAV4	XPO5_HUMAN	Exportin-5 OS=Homo sapiens GN=XPO5 PE=1 SV=1	1	0.014355293
B4DLN1	B4DLN1_HUMAN	Mitochondrial dicarboxylate carrier OS=Homo sapiens GN=SLC25A10 PE=2 SV=1	1	0.014355293
Q08752	PPID_HUMAN	Peptidyl-prolyl cis-trans isomerase D OS=Homo sapiens GN=PPID PE=1 SV=3	0.9999	0.014355293
P42167-2	LAP2B_HUMAN	Isoform Gamma of Lamina-associated polypeptide 2	0.9998	0.014355293
O95104-2	SFR15_HUMAN	Isoform 2 of Splicing factor	0.9997	0.014355293

P49903	SPS1_HUMAN	Selenide	0.9997	0.014355293
P15927-2	RFA2_HUMAN	Isoform 2 of Replication protein A 32 kDa subunit OS=Homo sapiens GN=RPA2	0.9989	0.014355293
O75915	PRAF3_HUMAN	PRA1 family protein 3 OS=Homo sapiens GN=ARL6IP5 PE=1 SV=1	0.9772	0.014355293
O15553	MEFV_HUMAN	Pyrin OS=Homo sapiens GN=MEFV PE=1 SV=1	0.9494	0.014355293
O75396	SC22B_HUMAN	Vesicle-trafficking protein SEC22b OS=Homo sapiens GN=SEC22B PE=1 SV=4	1	0
Q92499	DDX1_HUMAN	ATP-dependent RNA helicase DDX1 OS=Homo sapiens GN=DDX1 PE=1 SV=2	1	0
P06733	ENOA_HUMAN	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2	1	0
P07900-2	HS90A_HUMAN	Isoform 2 of Heat shock protein HSP 90-alpha OS=Homo sapiens GN=HSP90AA1	1	0
P08758	ANXA5_HUMAN	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2	1	0
P17812	PYRG1_HUMAN	CTP synthase 1 OS=Homo sapiens GN=CTPS1 PE=1 SV=2	1	0
P19338	NUCL_HUMAN	Nucleolin OS=Homo sapiens GN=NCL PE=1 SV=3	1	0
P78527-2	PRKDC_HUMAN	Isoform 2 of DNA-dependent protein kinase catalytic subunit OS=Homo sapiens GN=PRKDC	1	0
Q01105-2	SET_HUMAN	Isoform 2 of Protein SET OS=Homo sapiens GN=SET	1	0
Q12906-2	ILF3_HUMAN	Isoform 2 of Interleukin enhancer-binding factor 3 OS=Homo sapiens GN=ILF3	1	0
Q99829	CPNE1_HUMAN	Copine-1 OS=Homo sapiens GN=CPNE1 PE=1 SV=1	1	0
Q9UBT2	SAE2_HUMAN	SUMO-activating enzyme subunit 2 OS=Homo sapiens GN=UBA2 PE=1 SV=2	1	0
Q9UNZ2-4	NSF1C_HUMAN	Isoform 2 of NSF1L cofactor p47 OS=Homo sapiens GN=NSF1C	1	0
E9PNX1	E9PNX1_HUMAN	Serpin H1 (Fragment) OS=Homo sapiens GN=SERPINH1 PE=3 SV=1	1	0
Q98VA1	TBB2B_HUMAN	Tubulin beta-2B chain OS=Homo sapiens GN=TUBB2B PE=1 SV=1	1	0
Q9Y4G6	TLN2_HUMAN	Talin-2 OS=Homo sapiens GN=TLN2 PE=1 SV=4	0.9999	0
P05091	ALDH2_HUMAN	Aldehyde dehydrogenase	0.9996	0
Q8WXF1-2	PSPC1_HUMAN	Isoform 2 of Paraspeckle component 1 OS=Homo sapiens GN=PSPC1	0.9995	0
O75570	RF1M_HUMAN	Peptide chain release factor 1	0.9987	0
Q9Y276	BCS1_HUMAN	Mitochondrial chaperone BCS1 OS=Homo sapiens GN=BCS1L PE=1 SV=1	0.996	0
O60256	KPRB_HUMAN	Phosphoribosyl pyrophosphate synthase-associated protein 2 OS=Homo sapiens GN=PRPSAP2 PE=1 SV:1	0.9911	0
Q13724	MOGS_HUMAN	Mannosyl-oligosaccharide glucosidase OS=Homo sapiens GN=MOGS PE=1 SV=5	0.986	0
Q9Y2S7	PDIP2_HUMAN	Polymerase delta-interacting protein 2 OS=Homo sapiens GN=POLDIP2 PE=1 SV=1	0.9607	0
Q6ZRY4	RBP52_HUMAN	RNA-binding protein with multiple splicing 2 OS=Homo sapiens GN=RBPMS2 PE=2 SV=1	0.9318	0
O75400-2	PR40A_HUMAN	Isoform 2 of Pre-mRNA-processing factor 40 homolog A OS=Homo sapiens GN=PRPF40A	1	-0.01449957
O95232-2	LC7L3_HUMAN	Isoform 2 of Luc7-like protein 3 OS=Homo sapiens GN=LUC7L3	1	-0.01449957
P07814	SYEP_HUMAN	Bifunctional glutamate/proline-tRNA ligase OS=Homo sapiens GN=EPRS PE=1 SV=5	1	-0.01449957
P54577	SYIC_HUMAN	Tyrosine-tRNA ligase	1	-0.01449957
Q03252	LMNB2_HUMAN	Lamin-B2 OS=Homo sapiens GN=LMNB2 PE=1 SV=3	1	-0.01449957
Q96AB3-2	ISOC2_HUMAN	Isoform 2 of Isochorismatase domain-containing protein 2	1	-0.01449957
P13010	XRCC5_HUMAN	X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3	1	-0.01449957
P46821	MAP1B_HUMAN	Microtubule-associated protein 1B OS=Homo sapiens GN=MAP1B PE=1 SV=2	1	-0.01449957
P55786	PSA_HUMAN	Puromycin-sensitive aminopeptidase OS=Homo sapiens GN=NPEPPS PE=1 SV=2	1	-0.01449957
P55884-2	EIF3B_HUMAN	Isoform 2 of Eukaryotic translation initiation factor 3 subunit B OS=Homo sapiens GN=EIF3B	1	-0.01449957
P63104	1433Z_HUMAN	14-3-3 protein zeta/delta OS=Homo sapiens GN=YWHAZ PE=1 SV=1	1	-0.01449957
Q92973-2	TNPO1_HUMAN	Isoform 2 of Transportin-1 OS=Homo sapiens GN=TNPO1	1	-0.01449957
Q13263	TIF1B_HUMAN	Transcription intermediary factor 1-beta OS=Homo sapiens GN=TRIM28 PE=1 SV=5	1	-0.01449957
Q92621	NU205_HUMAN	Nuclear pore complex protein Nup205 OS=Homo sapiens GN=NUP205 PE=1 SV=3	1	-0.01449957
Q93008-1	USP9X_HUMAN	Isoform 2 of Probable ubiquitin carboxyl-terminal hydrolase FAF-X OS=Homo sapiens GN=USP9X	1	-0.01449957
P27348	1433T_HUMAN	14-3-3 protein theta OS=Homo sapiens GN=YWHAQ PE=1 SV=1	1	-0.01449957
P51665	PSD7_HUMAN	26S proteasome non-ATPase regulatory subunit 7 OS=Homo sapiens GN=PSMD7 PE=1 SV=2	0.9999	-0.01449957
P12236	ADT3_HUMAN	ADP/ATP translocase 3 OS=Homo sapiens GN=SLC25A6 PE=1 SV=4	0.9996	-0.01449957
O15173	PGRMC2_HUMAN	Membrane-associated progesterone receptor component 2 OS=Homo sapiens GN=PGRMC2 PE=1 SV=1	0.9967	-0.01449957
Q7Z2Y8	GVIN1_HUMAN	Interferon-induced very large GTPase 1 OS=Homo sapiens GN=GVINP1 PE=2 SV=2	0.9928	-0.01449957
F8WD12	F8WD12_HUMAN	SET domain-containing protein 5 OS=Homo sapiens GN=SETD5 PE=4 SV=1	0.982	-0.01449957
Q6ZUM4-3	RHG27_HUMAN	Isoform 3 of Rho GTPase-activating protein 27 OS=Homo sapiens GN=ARHGAP27	0.9791	-0.01449957
P29322	EPHA8_HUMAN	Ephrin type-A receptor 8 OS=Homo sapiens GN=EPHA8 PE=1 SV=2	0.9725	-0.01449957
Q86TC9	MYPN_HUMAN	Myopalladin OS=Homo sapiens GN=MYPN PE=1 SV=2	0.9485	-0.01449957
P13637	AT1A3_HUMAN	Sodium/potassium-transporting ATPase subunit alpha-3 OS=Homo sapiens GN=ATP1A3 PE=1 SV=3	0.8444	-0.01449957
O00629	IMA4_HUMAN	Isoform 2 of IMA4 OS=Homo sapiens GN=KPNAA4 PE=1 SV=1	1	-0.029146346
P06744-2	G6PI_HUMAN	Isoform 2 of Glucose-6-phosphate isomerase OS=Homo sapiens GN=GPI	1	-0.029146346
P22061-2	PIMT_HUMAN	Isoform 2 of Protein-L-isoaspartate(D-aspartate) O-methyltransferase OS=Homo sapiens GN=PCMT1	1	-0.029146346
P61081	UBC12_HUMAN	NEDD8-conjugating enzyme Ubc12 OS=Homo sapiens GN=UBE2M PE=1 SV=1	1	-0.029146346
P61353	RL27_HUMAN	60S ribosomal protein L27 OS=Homo sapiens GN=RPL27 PE=1 SV=2	1	-0.029146346
Q14232	E12BA_HUMAN	Translation initiation factor eIF-2B subunit alpha OS=Homo sapiens GN=EIF2B1 PE=1 SV=1	1	-0.029146346
Q15436	SEC23A_HUMAN	Protein transport protein Sec23A OS=Homo sapiens GN=SEC23A PE=1 SV=2	1	-0.029146346
P12277	KCRB_HUMAN	Creatine kinase B-type OS=Homo sapiens GN=CKB PE=1 SV=1	1	-0.029146346
P34932	HSP74_HUMAN	Heat shock 70 kDa protein 4 OS=Homo sapiens GN=HSPA4 PE=1 SV=4	1	-0.029146346
P61086-2	UBE2K_HUMAN	Isoform 2 of Ubiquitin-conjugating enzyme E2 K OS=Homo sapiens GN=UBE2K	1	-0.029146346
Q14103-2	HNRPD_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein D0 OS=Homo sapiens GN=HNRNPD	1	-0.029146346
Q14157-1	UBP2L_HUMAN	Isoform 2 of Ubiquitin-associated protein 2-like OS=Homo sapiens GN=UBAP2L	1	-0.029146346
Q14974	IMB1_HUMAN	Importin subunit beta-1 OS=Homo sapiens GN=KPNB1 PE=1 SV=2	1	-0.029146346
Q9Y6E2	BZW2_HUMAN	Basic leucine zipper and W2 domain-containing protein 2 OS=Homo sapiens GN=BZW2 PE=1 SV=1	1	-0.029146346
Q401N2-2	ZACN_HUMAN	Isoform 2 of Zinc-activated ligand-gated ion channel OS=Homo sapiens GN=ZACN	0.9862	-0.029146346
Q14966-2	ZN638_HUMAN	Isoform 2 of Zinc finger protein 638 OS=Homo sapiens GN=ZNF638	0.9843	-0.029146346
Q9GZ21	NAA50_HUMAN	N-alpha-acetyltransferase 50 OS=Homo sapiens GN=NAA50 PE=1 SV=1	0.9749	-0.029146346
O75347	TBCA_HUMAN	Tubulin-specific chaperone A OS=Homo sapiens GN=TBCA PE=1 SV=3	0.9626	-0.029146346
H0YJP3	H0YJP3_HUMAN	Tryptophan-tRNA ligase	0.9573	-0.029146346
B8ZZB3	B8ZZB3_HUMAN	Coiled-coil domain-containing protein 141 OS=Homo sapiens GN=CCDC141 PE=4 SV=1	0.9479	-0.029146346
P35250-2	RFC2_HUMAN	Isoform 2 of Replication factor C subunit 2 OS=Homo sapiens GN=RFC2	1	-0.043943348
P37802	TAGL2_HUMAN	Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3	1	-0.043943348
P49458	SRP09_HUMAN	Signal recognition particle 9 kDa protein OS=Homo sapiens GN=SRP9 PE=1 SV=2	1	-0.043943348
P49736	MCM2_HUMAN	DNA replication licensing factor MCM2 OS=Homo sapiens GN=MCM2 PE=1 SV=4	1	-0.043943348
Q02543	RL18A_HUMAN	60S ribosomal protein L18a OS=Homo sapiens GN=RPL18A PE=1 SV=2	1	-0.043943348
Q99798	ACON_HUMAN	Aconitate hydratase	1	-0.043943348
Q9HC38-2	GLOD4_HUMAN	Isoform 2 of Glyoxalase domain-containing protein 4 OS=Homo sapiens GN=GLOD4	1	-0.043943348
P04075	ALDOA_HUMAN	Fructose-bisphosphate aldolase A OS=Homo sapiens GN=ALDOA PE=1 SV=2	1	-0.043943348
P40227	TCMZ_HUMAN	T-complex protein 1 subunit zeta OS=Homo sapiens GN=CCT6A PE=1 SV=3	1	-0.043943348
P51532-2	SMCA4_HUMAN	Isoform 2 of Transcription activator BRG1 OS=Homo sapiens GN=SMARCA4	1	-0.043943348
P60891	PRPS1_HUMAN	Ribose-phosphate pyrophosphokinase 1 OS=Homo sapiens GN=PRPS1 PE=1 SV=2	1	-0.043943348
Q16658	FSCN1_HUMAN	Fascin OS=Homo sapiens GN=FSCN1 PE=1 SV=3	1	-0.043943348
Q5T2N8	ATD3C_HUMAN	ATPase family AAA domain-containing protein 3C OS=Homo sapiens GN=ATAD3C PE=1 SV=2	1	-0.043943348
Q99832	TCPH_HUMAN	T-complex protein 1 subunit eta OS=Homo sapiens GN=CCT7 PE=1 SV=2	1	-0.043943348
Q9UN52	CSN3_HUMAN	COP9 signalosome complex subunit 3 OS=Homo sapiens GN=COPS3 PE=1 SV=3	1	-0.043943348
B4DMU0	B4DMU0_HUMAN	Pyrroline-5-carboxylate reductase OS=Homo sapiens GN=PYCR1 PE=2 SV=1	1	-0.043943348
Q9H845	ACAD9_HUMAN	Acyl-CoA dehydrogenase family member 9	0.9999	-0.043943348
Q8IZL8-2	PELP1_HUMAN	Isoform 2 of Proline-	0.9999	-0.043943348
O15498	YKT6_HUMAN	Synaptobrevin homolog YKT6 OS=Homo sapiens GN=YKT6 PE=1 SV=1	0.999	-0.043943348

Q9BUZ4	TRAF4_HUMAN	TNF receptor-associated factor 4 OS=Homo sapiens GN=TRAF4 PE=1 SV=1	0.9963	-0.043943348
P24043	LAMA2_HUMAN	Laminin subunit alpha-2 OS=Homo sapiens GN=LAMA2 PE=1 SV=4	0.9774	-0.043943348
Q96CN9	GCC1_HUMAN	GRIP and coiled-coil domain-containing protein 1 OS=Homo sapiens GN=GCC1 PE=1 SV=1	0.972	-0.043943348
Q9Y4F9-2	FA65B_HUMAN	Isoform 2 of Protein FAM65B OS=Homo sapiens GN=FA65B	0.9577	-0.043943348
P07305-2	H10_HUMAN	Isoform 2 of Histone H1.0 OS=Homo sapiens GN=H1F0	1	-0.058893689
P54105	ICLN_HUMAN	Methylosome subunit pICln OS=Homo sapiens GN=CLNS1A PE=1 SV=1	1	-0.058893689
P62266	RS23_HUMAN	40S ribosomal protein S23 OS=Homo sapiens GN=RP523 PE=1 SV=3	1	-0.058893689
Q96CN7	ISOC1_HUMAN	Isochorismatase domain-containing protein 1 OS=Homo sapiens GN=ISOC1 PE=1 SV=3	1	-0.058893689
Q9UKN8	TF3C4_HUMAN	General transcription factor 3C polypeptide 4 OS=Homo sapiens GN=TF3C4 PE=1 SV=2	1	-0.058893689
Q95373	IPO7_HUMAN	Importin-7 OS=Homo sapiens GN=IPO7 PE=1 SV=1	1	-0.058893689
P08107	HSP71_HUMAN	Heat shock 70 kDa protein 1A/1B OS=Homo sapiens GN=HSPA1A PE=1 SV=5	1	-0.058893689
P15311	EZRI_HUMAN	Ezrin OS=Homo sapiens GN=EZR PE=1 SV=4	1	-0.058893689
P41250	SYG_HUMAN	Glycine--tRNA ligase OS=Homo sapiens GN=GARS PE=1 SV=3	1	-0.058893689
Q16555	DPYL2_HUMAN	Dihydropyrimidinase-related protein 2 OS=Homo sapiens GN=DPYSL2 PE=1 SV=1	1	-0.058893689
Q8IZT6	ASPM_HUMAN	Abnormal spindle-like microcephaly-associated protein OS=Homo sapiens GN=ASPM PE=1 SV=2	0.9999	-0.058893689
Q15631	TSN_HUMAN	Translin OS=Homo sapiens GN=TSN PE=1 SV=1	0.9996	-0.058893689
Q5D1E8	ZC12A_HUMAN	Ribonuclease ZC3H12A OS=Homo sapiens GN=ZC3H12A PE=1 SV=1	0.97	-0.058893689
Q7LBC6	KDM3B_HUMAN	Lysine-specific demethylase 3B OS=Homo sapiens GN=KDM3B PE=1 SV=2	0.9633	-0.058893689
Q2KJY2	KIF26B_HUMAN	Kinesin-like protein KIF26B OS=Homo sapiens GN=KIF26B PE=1 SV=1	0.963	-0.058893689
Q94842	TOX4_HUMAN	TOX high mobility group box family member 4 OS=Homo sapiens GN=TOX4 PE=1 SV=1	0.9476	-0.058893689
Q95271	TNKS1_HUMAN	Tankyrase-1 OS=Homo sapiens GN=TNKS PE=1 SV=2	0.9104	-0.058893689
P06737	PYGL_HUMAN	Glycogen phosphorylase	0.837	-0.058893689
P27797	CALR_HUMAN	Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1	1	-0.074000581
P62318	SMD3_HUMAN	Small nuclear ribonucleoprotein Sm D3 OS=Homo sapiens GN=SNRPD3 PE=1 SV=1	1	-0.074000581
Q15021	CND1_HUMAN	Condensin complex subunit 1 OS=Homo sapiens GN=NCAPD2 PE=1 SV=3	1	-0.074000581
Q9Y262	EIF3L_HUMAN	Eukaryotic translation initiation factor 3 subunit L OS=Homo sapiens GN=EIF3L PE=1 SV=1	1	-0.074000581
Q95831-3	AIFM1_HUMAN	Isoform 3 of Apoptosis-inducing factor 1	1	-0.074000581
P05141	ADT2_HUMAN	ADP/ATP translocase 2 OS=Homo sapiens GN=SLC25A5 PE=1 SV=7	1	-0.074000581
P14678-2	RSMB_HUMAN	Isoform SM-B of Small nuclear ribonucleoprotein-associated proteins B and B' OS=Homo sapiens GN=SI	1	-0.074000581
P16615-2	AT2A2_HUMAN	Isoform 2 of Sarcoplasmic/endoplasmic reticulum calcium ATPase 2 OS=Homo sapiens GN=ATP2A2	1	-0.074000581
P19367-2	HXK1_HUMAN	Isoform 2 of Hexokinase-1 OS=Homo sapiens GN=HK1	1	-0.074000581
P27635	RL10_HUMAN	60S ribosomal protein L10 OS=Homo sapiens GN=RPL10 PE=1 SV=4	1	-0.074000581
P35613-2	BASI_HUMAN	Isoform 2 of Basigin OS=Homo sapiens GN=BSG	1	-0.074000581
P39687	AN32A_HUMAN	Acidic leucine-rich nuclear phosphoprotein 32 family member A OS=Homo sapiens GN=ANP32A PE=1 S	1	-0.074000581
P50990	TCPQ_HUMAN	T-complex protein 1 subunit theta OS=Homo sapiens GN=CCT8 PE=1 SV=4	1	-0.074000581
P52701-2	MSH6_HUMAN	Isoform GTBP-alt of DNA mismatch repair protein Msh6 OS=Homo sapiens GN=MSH6	1	-0.074000581
P53396-2	ACLY_HUMAN	Isoform 2 of ATP-citrate synthase OS=Homo sapiens GN=ACLY	1	-0.074000581
P23284	PIIB_HUMAN	Peptidyl-prolyl cis-trans isomerase B OS=Homo sapiens GN=PIIB PE=1 SV=2	1	-0.074000581
P68371	TBB4B_HUMAN	Tubulin beta-4B chain OS=Homo sapiens GN=TUBB4B PE=1 SV=1	1	-0.074000581
J3KR44	J3KR44_HUMAN	Ubiquitin thioesterase OTUB1 OS=Homo sapiens GN=OTUB1 PE=4 SV=1	1	-0.074000581
P38919	IF4A3_HUMAN	Eukaryotic initiation factor 4A-III OS=Homo sapiens GN=EIF4A3 PE=1 SV=4	1	-0.074000581
Q00688	FKBP3_HUMAN	Peptidyl-prolyl cis-trans isomerase FKBP3 OS=Homo sapiens GN=FKBP3 PE=1 SV=1	0.9997	-0.074000581
Q9UHD1-2	CHRD1_HUMAN	Isoform 2 of Cysteine and histidine-rich domain-containing protein 1 OS=Homo sapiens GN=CHORDC1	0.9997	-0.074000581
Q5T5U3	RHG21_HUMAN	Rho GTPase-activating protein 21 OS=Homo sapiens GN=ARHGAP21 PE=1 SV=1	0.9996	-0.074000581
P12532-2	KCRU_HUMAN	Isoform 2 of Creatine kinase U-type	0.9993	-0.074000581
P20929	NEBU_HUMAN	Nebulin OS=Homo sapiens GN=NEB PE=1 SV=4	0.9991	-0.074000581
P05496	AT5G1_HUMAN	ATP synthase lipid-binding protein	0.9989	-0.074000581
Q68E01-2	INT3_HUMAN	Isoform 2 of Integrator complex subunit 3 OS=Homo sapiens GN=INTS3	0.9938	-0.074000581
H3BM57	H3BM57_HUMAN	Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=1	0.9905	-0.074000581
O75607	NPM3_HUMAN	Nucleoplasmin-3 OS=Homo sapiens GN=NPM3 PE=1 SV=3	1	-0.089267338
P07737	PROF1_HUMAN	Profilin-1 OS=Homo sapiens GN=PFN1 PE=1 SV=2	1	-0.089267338
P20618	PSB1_HUMAN	Proteasome subunit beta type-1 OS=Homo sapiens GN=PSMB1 PE=1 SV=2	1	-0.089267338
P62241	RS8_HUMAN	40S ribosomal protein S8 OS=Homo sapiens GN=RPS8 PE=1 SV=2	1	-0.089267338
P62917	RL8_HUMAN	60S ribosomal protein L8 OS=Homo sapiens GN=RPL8 PE=1 SV=2	1	-0.089267338
Q15005	SPCS2_HUMAN	Signal peptidase complex subunit 2 OS=Homo sapiens GN=SPCS2 PE=1 SV=3	1	-0.089267338
Q8NC51-2	PAIRB_HUMAN	Isoform 2 of Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERBP1	1	-0.089267338
Q9BVC6	TM109_HUMAN	Transmembrane protein 109 OS=Homo sapiens GN=TMEM109 PE=1 SV=1	1	-0.089267338
Q9UMS4	PRP19_HUMAN	Pre-mRNA-processing factor 19 OS=Homo sapiens GN=PRPF19 PE=1 SV=1	1	-0.089267338
P04181	OAT_HUMAN	Ornithine aminotransferase	1	-0.089267338
P18669	PGAM1_HUMAN	Phosphoglycerate mutase 1 OS=Homo sapiens GN=PGAM1 PE=1 SV=2	1	-0.089267338
P23921	RIR1_HUMAN	Ribonucleoside-diphosphate reductase large subunit OS=Homo sapiens GN=RRM1 PE=1 SV=1	1	-0.089267338
P27695	APEX1_HUMAN	DNA-(apurinic or pyrimidinic site) lyase OS=Homo sapiens GN=APEX1 PE=1 SV=2	1	-0.089267338
P29692-2	EEF1D_HUMAN	Elongation factor 1-delta OS=Homo sapiens GN=EEF1D	1	-0.089267338
P45880-1	VDAC2_HUMAN	Isoform 1 of Voltage-dependent anion-selective channel protein 2 OS=Homo sapiens GN=VDAC2	1	-0.089267338
P47897	SYQ_HUMAN	Glutamine--tRNA ligase OS=Homo sapiens GN=QARS PE=1 SV=1	1	-0.089267338
P60174-1	TPIS_HUMAN	Isoform 2 of Triosephosphate isomerase OS=Homo sapiens GN=TP1	1	-0.089267338
P61026	RAB10_HUMAN	Ras-related protein Rab-10 OS=Homo sapiens GN=RAB10 PE=1 SV=1	1	-0.089267338
Q13162	PRDX4_HUMAN	Peroxiredoxin-4 OS=Homo sapiens GN=PRDX4 PE=1 SV=1	1	-0.089267338
O60287	NPA1P_HUMAN	Nucleolar pre-ribosomal-associated protein 1 OS=Homo sapiens GN=URB1 PE=1 SV=4	1	-0.089267338
G3V438	G3V438_HUMAN	Activator of 90 kDa heat shock protein ATPase homolog 1 (Fragment) OS=Homo sapiens GN=AHSA1 PE	1	-0.089267338
Q5HYJ1	TECLR_HUMAN	Trans-2	0.9982	-0.089267338
Q98TV4	TMM43_HUMAN	Transmembrane protein 43 OS=Homo sapiens GN=TMEM43 PE=1 SV=1	0.9897	-0.089267338
Q6AWC2-2	WWC2_HUMAN	Isoform 2 of Protein WWC2 OS=Homo sapiens GN=WWC2	0.9763	-0.089267338
P09211	GSTP1_HUMAN	Glutathione S-transferase P OS=Homo sapiens GN=GSTP1 PE=1 SV=2	1	-0.104697379
P37837	TALDO_HUMAN	Transaldolase OS=Homo sapiens GN=TALDO1 PE=1 SV=2	1	-0.104697379
Q99497	PARK7_HUMAN	Protein DJ-1 OS=Homo sapiens GN=PARK7 PE=1 SV=2	1	-0.104697379
Q9H0D6-2	XRN2_HUMAN	Isoform 2 of 5'-3' exoribonuclease 2 OS=Homo sapiens GN=XRN2	1	-0.104697379
P11387	TOP1_HUMAN	DNA topoisomerase 1 OS=Homo sapiens GN=TOP1 PE=1 SV=2	1	-0.104697379
P26599-2	PTBP1_HUMAN	Isoform 2 of Polypyrimidine tract-binding protein 1 OS=Homo sapiens GN=PTBP1	1	-0.104697379
P26639	SYTC_HUMAN	Threonine--tRNA ligase	1	-0.104697379
P37108	SRP14_HUMAN	Signal recognition particle 14 kDa protein OS=Homo sapiens GN=SRP14 PE=1 SV=2	1	-0.104697379
P40429	RL13A_HUMAN	60S ribosomal protein L13a OS=Homo sapiens GN=RPL13A PE=1 SV=2	1	-0.104697379
P46777	RL5_HUMAN	60S ribosomal protein L5 OS=Homo sapiens GN=RPL5 PE=1 SV=3	1	-0.104697379
P60660-2	MYL6_HUMAN	Isoform Smooth muscle of Myosin light polypeptide 6 OS=Homo sapiens GN=MYL6	1	-0.104697379
P68402	PA1B2_HUMAN	Platelet-activating factor acetylhydrolase IB subunit beta OS=Homo sapiens GN=PFAH1B2 PE=1 SV=1	1	-0.104697379
Q02790	FKBP4_HUMAN	Peptidyl-prolyl cis-trans isomerase FKBP4 OS=Homo sapiens GN=FKBP4 PE=1 SV=3	1	-0.104697379
Q13509	TBB3_HUMAN	Tubulin beta-3 chain OS=Homo sapiens GN=TUBB3 PE=1 SV=2	1	-0.104697379
Q6P2Q9	PRP8_HUMAN	Pre-mRNA-processing-splicing factor 8 OS=Homo sapiens GN=PRPF8 PE=1 SV=2	1	-0.104697379
Q8N163-2	K1967_HUMAN	Isoform 2 of DBIRD complex subunit KIAA1967 OS=Homo sapiens GN=KIAA1967	1	-0.104697379
Q9H892-2	TTC12_HUMAN	Isoform 2 of Tetratricopeptide repeat protein 12 OS=Homo sapiens GN=TTC12	1	-0.104697379
P53992	SEC24C_HUMAN	Protein transport protein Sec24C OS=Homo sapiens GN=SEC24C PE=1 SV=3	0.9996	-0.104697379
Q9NR09	BIRC6_HUMAN	Baculoviral IAP repeat-containing protein 6 OS=Homo sapiens GN=BIRC6 PE=1 SV=2	0.9989	-0.104697379

P67812	SC11A_HUMAN	Signal peptidase complex catalytic subunit SEC11A OS=Homo sapiens GN=SEC11A PE=1 SV=1	0.9937	-0.104697379
Q86Y91-3	K118B_HUMAN	Isoform 3 of Kinesin-like protein KIF18B OS=Homo sapiens GN=KIF18B	0.9905	-0.104697379
O14497-2	ARI1A_HUMAN	Isoform 2 of AT-rich interactive domain-containing protein 1A OS=Homo sapiens GN=ARI1A	0.9808	-0.104697379
F222W0	F222W0_HUMAN	TRIO and F-actin-binding protein OS=Homo sapiens GN=TRIOBP PE=4 SV=1	0.9793	-0.104697379
O15372	EIF3H_HUMAN	Eukaryotic translation initiation factor 3 subunit H OS=Homo sapiens GN=EIF3H PE=1 SV=1	1	-0.120294234
P30048	PRDX3_HUMAN	Thioredoxin-dependent peroxide reductase	1	-0.120294234
P42766	RL35_HUMAN	60S ribosomal protein L35 OS=Homo sapiens GN=RPL35 PE=1 SV=2	1	-0.120294234
Q8NEJ9-2	NGDN_HUMAN	Isoform 2 of Neuroguidin OS=Homo sapiens GN=NGDN	1	-0.120294234
P16949-2	STMN1_HUMAN	Isoform 2 of Stathmin OS=Homo sapiens GN=STMN1	1	-0.120294234
P21333-2	FLNA_HUMAN	Isoform 2 of Filamin-A OS=Homo sapiens GN=FLNA	1	-0.120294234
P22234	PUR6_HUMAN	Multifunctional protein ADE2 OS=Homo sapiens GN=PAICS PE=1 SV=3	1	-0.120294234
P46781	RS9_HUMAN	40S ribosomal protein S9 OS=Homo sapiens GN=RPS9 PE=1 SV=3	1	-0.120294234
P51148	RAB5C_HUMAN	Ras-related protein Rab-5C OS=Homo sapiens GN=RAB5C PE=1 SV=2	1	-0.120294234
P61254	RL26_HUMAN	60S ribosomal protein L26 OS=Homo sapiens GN=RPL26 PE=1 SV=1	1	-0.120294234
Q15019-2	SEPT2_HUMAN	Isoform 2 of Septin-2 OS=Homo sapiens GN=SEPT2	1	-0.120294234
P09429	HMGB1_HUMAN	High mobility group protein B1 OS=Homo sapiens GN=HMGB1 PE=1 SV=3	0.9999	-0.120294234
Q9NR31	SAR1A_HUMAN	GTP-binding protein SAR1a OS=Homo sapiens GN=SAR1A PE=1 SV=1	0.9999	-0.120294234
Q16630-2	CPSF6_HUMAN	Isoform 2 of Cleavage and polyadenylation specificity factor subunit 6 OS=Homo sapiens GN=CPSF6	0.9994	-0.120294234
Q9NXR7-1	BRE_HUMAN	Isoform 1 of BRCA1-A complex subunit BRE OS=Homo sapiens GN=BRE	0.9994	-0.120294234
P22087	FBRL_HUMAN	rRNA 2'-O-methyltransferase fibrillar OS=Homo sapiens GN=FBRL PE=1 SV=2	0.9989	-0.120294234
Q9C0B2-2	K1751_HUMAN	Isoform 2 of Uncharacterized protein KIAA1751 OS=Homo sapiens GN=KIAA1751	0.9981	-0.120294234
Q969H8	CS010_HUMAN	UPF0556 protein C19orf10 OS=Homo sapiens GN=C19orf10 PE=1 SV=1	0.9957	-0.120294234
Q15700-2	DLG2_HUMAN	Isoform 2 of Disks large homolog 2 OS=Homo sapiens GN=DLG2	0.9587	-0.120294234
P42336	PK3CA_HUMAN	Phosphatidylinositol 4	0.9522	-0.120294234
O00487	PSDE_HUMAN	26S proteasome non-ATPase regulatory subunit 14 OS=Homo sapiens GN=PSMD14 PE=1 SV=1	1	-0.13606155
P30084	ECHM_HUMAN	Enoyl-CoA hydratase	1	-0.13606155
P46782	RS5_HUMAN	40S ribosomal protein S5 OS=Homo sapiens GN=RPS5 PE=1 SV=4	1	-0.13606155
P61604	CH10_HUMAN	10 kDa heat shock protein	1	-0.13606155
P62854	RS26_HUMAN	40S ribosomal protein S26 OS=Homo sapiens GN=RPS26 PE=1 SV=3	1	-0.13606155
Q9NSD9	SYFB_HUMAN	Phenylalanine--tRNA ligase beta subunit OS=Homo sapiens GN=FARSB PE=1 SV=3	1	-0.13606155
O15067	PUR4_HUMAN	Phosphoribosylformylglycinamide synthase OS=Homo sapiens GN=PFAS PE=1 SV=4	1	-0.13606155
O94905	ERLN2_HUMAN	Erlin-2 OS=Homo sapiens GN=ERLN2 PE=1 SV=1	1	-0.13606155
P04843	RPN1_HUMAN	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1 OS=Homo sapiens GN=RPN1	1	-0.13606155
P17844	DDX5_HUMAN	Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 PE=1 SV=1	1	-0.13606155
P17987	TCPA_HUMAN	T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1	1	-0.13606155
P18124	RL7_HUMAN	60S ribosomal protein L7 OS=Homo sapiens GN=RPL7 PE=1 SV=1	1	-0.13606155
P23528	COF1_HUMAN	Cofilin-1 OS=Homo sapiens GN=CFL1 PE=1 SV=3	1	-0.13606155
P33176	KINH_HUMAN	Kinesin-1 heavy chain OS=Homo sapiens GN=KIF5B PE=1 SV=1	1	-0.13606155
P35659	DEK_HUMAN	Protein DEK OS=Homo sapiens GN=DEK PE=1 SV=1	1	-0.13606155
P46779	RL28_HUMAN	60S ribosomal protein L28 OS=Homo sapiens GN=RPL28 PE=1 SV=3	1	-0.13606155
P50991	TCPD_HUMAN	T-complex protein 1 subunit delta OS=Homo sapiens GN=CCT4 PE=1 SV=4	1	-0.13606155
P53618	COPB_HUMAN	Coatomer subunit beta OS=Homo sapiens GN=COPB1 PE=1 SV=3	1	-0.13606155
P62701	RS4X_HUMAN	40S ribosomal protein S4	1	-0.13606155
P62826	RAN_HUMAN	GTP-binding nuclear protein Ran OS=Homo sapiens GN=RAN PE=1 SV=3	1	-0.13606155
P63261	ACTG_HUMAN	Actin	1	-0.13606155
Q43432-3	IF4G3_HUMAN	Isoform 3 of Eukaryotic translation initiation factor 4 gamma 3 OS=Homo sapiens GN=EIF4G3	1	-0.13606155
Q07021	C1QBP_HUMAN	Complement component 1 Q subcomponent-binding protein	1	-0.13606155
P14550	AK1A1_HUMAN	Alcohol dehydrogenase [NADP(+)] OS=Homo sapiens GN=AKR1A1 PE=1 SV=3	0.9998	-0.13606155
P09104	ENOG_HUMAN	Gamma-enolase OS=Homo sapiens GN=ENO2 PE=1 SV=3	0.9997	-0.13606155
Q96CS3	FAF2_HUMAN	FAS-associated factor 2 OS=Homo sapiens GN=FAF2 PE=1 SV=2	0.9996	-0.13606155
Q9NXF1-2	TEX10_HUMAN	Isoform 2 of Testis-expressed sequence 10 protein OS=Homo sapiens GN=TEX10	0.9967	-0.13606155
P11137-3	MAP2_HUMAN	Isoform 3 of Microtubule-associated protein 2 OS=Homo sapiens GN=MAP2	0.9943	-0.13606155
Q8NF91-4	SYNE1_HUMAN	Isoform 4 of Nesprin-1 OS=Homo sapiens GN=SYNE1	0.9936	-0.13606155
P57721-4	PCBP3_HUMAN	Isoform 4 of Poly(rC)-binding protein 3 OS=Homo sapiens GN=PCBP3	0.9763	-0.13606155
Q9NYF8-2	BCLF1_HUMAN	Isoform 2 of Bcl-2-associated transcription factor 1 OS=Homo sapiens GN=BCLAF1	0.9673	-0.13606155
Q43242	PSMD3_HUMAN	26S proteasome non-ATPase regulatory subunit 3 OS=Homo sapiens GN=PSMD3 PE=1 SV=2	1	-0.152003093
P13667	PDIA4_HUMAN	Protein disulfide-isomerase A4 OS=Homo sapiens GN=PDIA4 PE=1 SV=2	1	-0.152003093
P30101	PDIA3_HUMAN	Protein disulfide-isomerase A3 OS=Homo sapiens GN=PDIA3 PE=1 SV=4	1	-0.152003093
P39019	RS19_HUMAN	40S ribosomal protein S19 OS=Homo sapiens GN=RPS19 PE=1 SV=2	1	-0.152003093
P62841	RS15_HUMAN	40S ribosomal protein S15 OS=Homo sapiens GN=RPS15 PE=1 SV=2	1	-0.152003093
Q12905	ILF2_HUMAN	Interleukin enhancer-binding factor 2 OS=Homo sapiens GN=ILF2 PE=1 SV=2	1	-0.152003093
Q9NR30-2	DDX21_HUMAN	Isoform 2 of Nucleolar RNA helicase 2 OS=Homo sapiens GN=DDX21	1	-0.152003093
P04406	G3P_HUMAN	Glyceraldehyde-3-phosphate dehydrogenase OS=Homo sapiens GN=GAPDH PE=1 SV=3	1	-0.152003093
P12268	IMDH2_HUMAN	Inosine-5'-monophosphate dehydrogenase 2 OS=Homo sapiens GN=IMPDH2 PE=1 SV=2	1	-0.152003093
P13693	TCTP_HUMAN	Translationally-controlled tumor protein OS=Homo sapiens GN=TPT1 PE=1 SV=1	1	-0.152003093
P23526	SAHH_HUMAN	Adenosylhomocysteinase OS=Homo sapiens GN=AHCY PE=1 SV=4	1	-0.152003093
P25705	ATPA_HUMAN	ATP synthase subunit alpha	1	-0.152003093
P35579	MYH9_HUMAN	Myosin-9 OS=Homo sapiens GN=MYH9 PE=1 SV=4	1	-0.152003093
P39023	RL3_HUMAN	60S ribosomal protein L3 OS=Homo sapiens GN=RPL3 PE=1 SV=2	1	-0.152003093
P46783	RS10_HUMAN	40S ribosomal protein S10 OS=Homo sapiens GN=RPS10 PE=1 SV=1	1	-0.152003093
P62753	RS6_HUMAN	40S ribosomal protein S6 OS=Homo sapiens GN=RPS6 PE=1 SV=1	1	-0.152003093
P63220	RS21_HUMAN	40S ribosomal protein S21 OS=Homo sapiens GN=RPS21 PE=1 SV=1	1	-0.152003093
Q9GZQ3	COMD5_HUMAN	COMM domain-containing protein 5 OS=Homo sapiens GN=COMM5 PE=1 SV=1	1	-0.152003093
P8VSB5	F8VSB5_HUMAN	Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 PE=4 SV=1	1	-0.152003093
P30405	PIIF_HUMAN	Peptidyl-prolyl cis-trans isomerase F	1	-0.152003093
P46776	RL27A_HUMAN	60S ribosomal protein L27a OS=Homo sapiens GN=RPL27A PE=1 SV=2	0.9999	-0.152003093
Q8TD57	DYH3_HUMAN	Dynein heavy chain 3	0.9455	-0.152003093
Q8N1X5	YF001_HUMAN	Uncharacterized protein FLJ37310 OS=Homo sapiens PE=1 SV=1	0.8816	-0.152003093
P62847-2	RS24_HUMAN	Isoform 2 of 40S ribosomal protein S24 OS=Homo sapiens GN=RPS24	0.8447	-0.152003093
Q9NX40	OCAD1_HUMAN	OCIA domain-containing protein 1 OS=Homo sapiens GN=OCIAD1 PE=1 SV=1	0.8339	-0.152003093
P62304	RUXE_HUMAN	Small nuclear ribonucleoprotein E OS=Homo sapiens GN=SNRPE PE=1 SV=1	1	-0.168122759
P05388	RLA0_HUMAN	60S acidic ribosomal protein P0 OS=Homo sapiens GN=RPLP0 PE=1 SV=1	1	-0.168122759
P21796	VDAC1_HUMAN	Voltage-dependent anion-selective channel protein 1 OS=Homo sapiens GN=VDAC1 PE=1 SV=2	1	-0.168122759
P25787	PSA2_HUMAN	Proteasome subunit alpha type-2 OS=Homo sapiens GN=PSMA2 PE=1 SV=2	1	-0.168122759
P26583	HMGB2_HUMAN	High mobility group protein B2 OS=Homo sapiens GN=HMGB2 PE=1 SV=2	1	-0.168122759
P35232	PHB_HUMAN	Prohibitin OS=Homo sapiens GN=PHB PE=1 SV=1	1	-0.168122759
P48643	TCPE_HUMAN	T-complex protein 1 subunit epsilon OS=Homo sapiens GN=CCT5 PE=1 SV=1	1	-0.168122759
Q05639	EFT1A2_HUMAN	Elongation factor 1-alpha 2 OS=Homo sapiens GN=EEF1A2 PE=1 SV=1	1	-0.168122759
P78371	TCBP_HUMAN	T-complex protein 1 subunit beta OS=Homo sapiens GN=CCT2 PE=1 SV=4	1	-0.168122759
Q07020	RL18_HUMAN	60S ribosomal protein L18 OS=Homo sapiens GN=RPL18 PE=1 SV=2	1	-0.168122759
Q99460-2	PSMD1_HUMAN	Isoform 2 of 26S proteasome non-ATPase regulatory subunit 1 OS=Homo sapiens GN=PSMD1	1	-0.168122759
Q96TC7-2	RMD3_HUMAN	Isoform 2 of Regulator of microtubule dynamics protein 3 OS=Homo sapiens GN=FAM82A2	0.9997	-0.168122759

P54136-2	SYRC_HUMAN	Isoform Monomeric of Arginine--tRNA ligase	0.9987	-0.168122759
P68400	CSK21_HUMAN	Casein kinase II subunit alpha OS=Homo sapiens GN=CSNK2A1 PE=1 SV=1	0.9961	-0.168122759
P17980	PRS6A_HUMAN	26S protease regulatory subunit 6A OS=Homo sapiens GN=PSMC3 PE=1 SV=3	0.9923	-0.168122759
Q99719-2	SEPT5_HUMAN	Isoform 2 of Septin-5 OS=Homo sapiens GN=SEPT5	0.9744	-0.168122759
P22626-2	ROA2_HUMAN	Isoform A2 of Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1	1	-0.184424571
P28072	PSB6_HUMAN	Proteasome subunit beta type-6 OS=Homo sapiens GN=PSMB6 PE=1 SV=4	1	-0.184424571
P31942-2	HNRH3_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein H3 OS=Homo sapiens GN=HNRNPH3	1	-0.184424571
Q99598	TSNAX_HUMAN	Translin-associated protein X OS=Homo sapiens GN=TSNAX PE=1 SV=1	1	-0.184424571
P00505	AATM_HUMAN	Aspartate aminotransferase	1	-0.184424571
P08238	HS90B_HUMAN	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90A1 PE=1 SV=4	1	-0.184424571
P08708	RS17_HUMAN	40S ribosomal protein S17 OS=Homo sapiens GN=RPS17 PE=1 SV=2	1	-0.184424571
P23396	RS3_HUMAN	40S ribosomal protein S3 OS=Homo sapiens GN=RPS3 PE=1 SV=2	1	-0.184424571
P26641	EF1G_HUMAN	Elongation factor 1-gamma OS=Homo sapiens GN=EEF1G PE=1 SV=3	1	-0.184424571
P30050	RL12_HUMAN	60S ribosomal protein L12 OS=Homo sapiens GN=RPL12 PE=1 SV=1	1	-0.184424571
P48735	IDHP_HUMAN	Isocitrate dehydrogenase [NADP]	1	-0.184424571
Q00610-2	CLH1_HUMAN	Isoform 2 of Clathrin heavy chain 1 OS=Homo sapiens GN=CLTC	1	-0.184424571
Q15003	CND2_HUMAN	Condensin complex subunit 2 OS=Homo sapiens GN=NCAHP PE=1 SV=3	1	-0.184424571
Q9UNF1	MAGD2_HUMAN	Melanoma-associated antigen D2 OS=Homo sapiens GN=MAGED2 PE=1 SV=2	1	-0.184424571
Q9UNM6-2	PSD13_HUMAN	Isoform 2 of 26S proteasome non-ATPase regulatory subunit 13 OS=Homo sapiens GN=PSMD13	1	-0.184424571
P14625	ENPL_HUMAN	Endoplasmic reticulum protein OS=Homo sapiens GN=HSP90B1 PE=1 SV=1	1	-0.184424571
Q13151	ROA0_HUMAN	Heterogeneous nuclear ribonucleoprotein A0 OS=Homo sapiens GN=HNRNPA0 PE=1 SV=1	0.9988	-0.184424571
Q96HR8	NAF1_HUMAN	H/ACA ribonucleoprotein complex non-core subunit NAF1 OS=Homo sapiens GN=NAF1 PE=1 SV=2	0.997	-0.184424571
B1AJR9	B1AJR9_HUMAN	Protein tyrosine phosphatase	0.9842	-0.184424571
A6NP24	A6NP24_HUMAN	Quinone oxidoreductase (Fragment) OS=Homo sapiens GN=CRYZ PE=4 SV=1	0.9841	-0.184424571
Q9Y2W1	TR150_HUMAN	Thyroid hormone receptor-associated protein 3 OS=Homo sapiens GN=THRAP3 PE=1 SV=2	0.9827	-0.184424571
Q01780-2	EXOSX_HUMAN	Isoform 2 of Exosome component 10 OS=Homo sapiens GN=EXOSC10	0.9674	-0.184424571
P30086	PEBP1_HUMAN	Phosphatidylethanolamine-binding protein 1 OS=Homo sapiens GN=PEBP1 PE=1 SV=3	1	-0.200912694
P50914	RL14_HUMAN	60S ribosomal protein L14 OS=Homo sapiens GN=RPL14 PE=1 SV=4	1	-0.200912694
P51570	GALK1_HUMAN	Galactokinase OS=Homo sapiens GN=GALK1 PE=1 SV=1	1	-0.200912694
P61221	ABCE1_HUMAN	ATP-binding cassette sub-family E member 1 OS=Homo sapiens GN=ABCE1 PE=1 SV=1	1	-0.200912694
P62913-2	RL11_HUMAN	Isoform 2 of 60S ribosomal protein L11 OS=Homo sapiens GN=RPL11	1	-0.200912694
Q04837	SSBP_HUMAN	Single-stranded DNA-binding protein	1	-0.200912694
O00303	EIF3F_HUMAN	Eukaryotic translation initiation factor 3 subunit F OS=Homo sapiens GN=EIF3F PE=1 SV=1	1	-0.200912694
P07195	LDHB_HUMAN	L-lactate dehydrogenase B chain OS=Homo sapiens GN=LDHB PE=1 SV=2	1	-0.200912694
P33316	DUT_HUMAN	Deoxyuridine 5'-triphosphate nucleotidohydrolase	1	-0.200912694
P61224	RAP1B_HUMAN	Ras-related protein Rap-1b OS=Homo sapiens GN=RAP1B PE=1 SV=1	1	-0.200912694
P68363	TBA1B_HUMAN	Tubulin alpha-1B chain OS=Homo sapiens GN=TUBA1B PE=1 SV=1	1	-0.200912694
Q15366-2	PCBP2_HUMAN	Isoform 2 of Poly(rC)-binding protein 2 OS=Homo sapiens GN=PCBP2	1	-0.200912694
Q7KZF4	SND1_HUMAN	Staphylococcal nuclease domain-containing protein 1 OS=Homo sapiens GN=SND1 PE=1 SV=1	1	-0.200912694
Q5T9B7	Q5T9B7_HUMAN	Adenylate kinase F OS=Homo sapiens GN=AK1 PE=2 SV=1	1	-0.200912694
P43243	MATR3_HUMAN	Matrin-3 OS=Homo sapiens GN=MATR3 PE=1 SV=2	1	-0.200912694
O60684	IMAT7_HUMAN	Importin subunit alpha-7 OS=Homo sapiens GN=KPNA6 PE=1 SV=1	0.9999	-0.200912694
Q9NX31	CT111_HUMAN	Uncharacterized protein C20orf111 OS=Homo sapiens GN=C20orf111 PE=2 SV=2	0.8675	-0.200912694
P31040	DHSA_HUMAN	Succinate dehydrogenase [ubiquinone] flavoprotein subunit	1	-0.217591435
P99999	CYC_HUMAN	Cytochrome c OS=Homo sapiens GN=CYCS PE=1 SV=2	1	-0.217591435
O14818	PSA7_HUMAN	Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1	1	-0.217591435
P00558	PGK1_HUMAN	Phosphoglycerate kinase 1 OS=Homo sapiens GN=PGK1 PE=1 SV=3	1	-0.217591435
P22314	UBA1_HUMAN	Ubiquitin-like modifier-activating enzyme 1 OS=Homo sapiens GN=UBA1 PE=1 SV=3	1	-0.217591435
P40939	ECHA_HUMAN	Trifunctional enzyme subunit alpha	1	-0.217591435
P49755	TMEDA_HUMAN	Transmembrane emp24 domain-containing protein 10 OS=Homo sapiens GN=TMED10 PE=1 SV=2	1	-0.217591435
Q15233	NONO_HUMAN	Non-POU domain-containing octamer-binding protein OS=Homo sapiens GN=NONO PE=1 SV=4	1	-0.217591435
Q99873-2	ANM1_HUMAN	Isoform 2 of Protein arginine N-methyltransferase 1 OS=Homo sapiens GN=PRMT1	1	-0.217591435
Q9H0A0	NAT10_HUMAN	N-acetyltransferase 10 OS=Homo sapiens GN=NAT10 PE=1 SV=2	1	-0.217591435
Q5VT25-2	MRCKA_HUMAN	Isoform 2 of Serine/threonine-protein kinase MRCK alpha OS=Homo sapiens GN=CDC42BPA	0.9991	-0.217591435
O60812	HNRCL_HUMAN	Heterogeneous nuclear ribonucleoprotein C-like 1 OS=Homo sapiens GN=HNRNPL1 PE=1 SV=1	0.9915	-0.217591435
A8MVJ3	A8MVJ3_HUMAN	Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0.9796	-0.217591435
F5H6J7	F5H6J7_HUMAN	G-protein-coupled receptor 161 OS=Homo sapiens GN=GPR161 PE=4 SV=1	0.9765	-0.217591435
P31948	STIP1_HUMAN	Stress-induced-phosphoprotein 1 OS=Homo sapiens GN=STIP1 PE=1 SV=1	1	-0.234465254
P42285	SK2L2_HUMAN	Superkiller viralicidal activity 2-like 2 OS=Homo sapiens GN=SKIV2L2 PE=1 SV=3	1	-0.234465254
Q04323-2	UBXN1_HUMAN	Isoform 2 of UBX domain-containing protein 1 OS=Homo sapiens GN=UBXN1	1	-0.234465254
P05023-3	AT1A1_HUMAN	Isoform 3 of Sodium/potassium-transporting ATPase subunit alpha-1 OS=Homo sapiens GN=ATP1A1	1	-0.234465254
P11142	HSP7C_HUMAN	Heat shock cognate 71 kDa protein OS=Homo sapiens GN=HSPA8 PE=1 SV=1	1	-0.234465254
P18077	RL35A_HUMAN	60S ribosomal protein L35a OS=Homo sapiens GN=RPL35A PE=1 SV=2	1	-0.234465254
P42704	LPPRC_HUMAN	Leucine-rich PPR motif-containing protein	1	-0.234465254
P62249	RS16_HUMAN	40S ribosomal protein S16 OS=Homo sapiens GN=RPS16 PE=1 SV=2	1	-0.234465254
P62269	RS18_HUMAN	40S ribosomal protein S18 OS=Homo sapiens GN=RPS18 PE=1 SV=3	1	-0.234465254
Q13200	PSMD2_HUMAN	26S proteasome non-ATPase regulatory subunit 2 OS=Homo sapiens GN=PSMD2 PE=1 SV=3	1	-0.234465254
P18621	RL17_HUMAN	60S ribosomal protein L17 OS=Homo sapiens GN=RPL17 PE=1 SV=3	1	-0.234465254
P29353-2	SHC1_HUMAN	Isoform p52Shc of SHC-transforming protein 1 OS=Homo sapiens GN=SHC1	0.9998	-0.234465254
Q9UL46	PSME2_HUMAN	Proteasome activator complex subunit 2 OS=Homo sapiens GN=PSME2 PE=1 SV=4	0.9993	-0.234465254
P04818	TYSY_HUMAN	Thymidylate synthase OS=Homo sapiens GN=TYMS PE=1 SV=3	0.9988	-0.234465254
P25398	RS12_HUMAN	40S ribosomal protein S12 OS=Homo sapiens GN=RPS12 PE=1 SV=3	1	-0.251538767
P31930	QCR1_HUMAN	Cytochrome b-c1 complex subunit 1	1	-0.251538767
P52272-2	HNRPM_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM	1	-0.251538767
O75643	U520_HUMAN	U5 small nuclear ribonucleoprotein 200 kDa helicase OS=Homo sapiens GN=SNRNP200 PE=1 SV=2	1	-0.251538767
P23246-2	SFPQ_HUMAN	Isoform Short of Splicing factor	1	-0.251538767
P30153	2AAA_HUMAN	Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha isoform OS=Homo sapiens	1	-0.251538767
P32119	PRDX2_HUMAN	Peroxisome oxidoreductin-2 OS=Homo sapiens GN=PRDX2 PE=1 SV=5	1	-0.251538767
P55060-3	XPO2_HUMAN	Isoform 3 of Exportin-2 OS=Homo sapiens GN=CSE1L	1	-0.251538767
P55072	TERA_HUMAN	Transitional endoplasmic reticulum ATPase OS=Homo sapiens GN=VCP PE=1 SV=4	1	-0.251538767
P60842	IF4A1_HUMAN	Eukaryotic initiation factor 4A-I OS=Homo sapiens GN=EIF4A1 PE=1 SV=1	1	-0.251538767
P61586	RHOA_HUMAN	Transforming protein RhoA OS=Homo sapiens GN=RHOA PE=1 SV=1	1	-0.251538767
P84098	RL19_HUMAN	60S ribosomal protein L19 OS=Homo sapiens GN=RPL19 PE=1 SV=1	1	-0.251538767
Q07666-2	KHDR1_HUMAN	Isoform 2 of KH domain-containing	1	-0.251538767
Q13347	EIF3I_HUMAN	Eukaryotic translation initiation factor 3 subunit I OS=Homo sapiens GN=EIF3I PE=1 SV=1	1	-0.251538767
Q14697-2	GANAB_HUMAN	Isoform 2 of Neutral alpha-glucosidase AB OS=Homo sapiens GN=GANAB	1	-0.251538767
P55957-2	BID_HUMAN	Isoform 2 of BH3-interacting domain death agonist OS=Homo sapiens GN=BID	0.9999	-0.251538767
Q536Q0	DHB12_HUMAN	Estradiol 17-beta-dehydrogenase 12 OS=Homo sapiens GN=HSD17B12 PE=1 SV=2	0.9999	-0.251538767
P14314-2	GLU2B_HUMAN	Isoform 2 of Glucosidase 2 subunit beta OS=Homo sapiens GN=PRKCSH	0.9998	-0.251538767
P49770	EI2B8_HUMAN	Translation initiation factor eIF-2B subunit beta OS=Homo sapiens GN=EIF2B2 PE=1 SV=3	0.9998	-0.251538767
Q13459-2	MYO9B_HUMAN	Isoform Short of Unconventional myosin-IXb OS=Homo sapiens GN=MYO9B	0.9974	-0.251538767
P30085	KCY_HUMAN	UMP-CMP kinase OS=Homo sapiens GN=CMK1 PE=1 SV=3	0.9873	-0.251538767

Q96A33-2	CCD47_HUMAN	Isoform 2 of Coiled-coil domain-containing protein 47 OS=Homo sapiens GN=CCDC47	0.9846	-0.251538767
Q9NR77	PXMP2_HUMAN	Peroxisomal membrane protein 2 OS=Homo sapiens GN=PXMP2 PE=1 SV=3	0.946	-0.251538767
Q86Y13	DZIP3_HUMAN	E3 ubiquitin-protein ligase DZIP3 OS=Homo sapiens GN=DZIP3 PE=1 SV=2	0.944	-0.251538767
O60518	RNBP6_HUMAN	Ran-binding protein 6 OS=Homo sapiens GN=RANBP6 PE=1 SV=2	0.9359	-0.251538767
Q8IXS8	F126B_HUMAN	Protein FAM126B OS=Homo sapiens GN=FAM126B PE=1 SV=1	0.9344	-0.251538767
P24666-2	PPAC_HUMAN	Isoform 2 of Low molecular weight phosphotyrosine protein phosphatase OS=Homo sapiens GN=ACP1	1	-0.268816758
P35270	SPRE_HUMAN	Septiapterin reductase OS=Homo sapiens GN=SPR PE=1 SV=1	1	-0.268816758
P49720	PSB3_HUMAN	Proteasome subunit beta type-3 OS=Homo sapiens GN=PSMB3 PE=1 SV=2	1	-0.268816758
Q9Y3B4	PM14_HUMAN	Pre-mRNA branch site protein p14 OS=Homo sapiens GN=SF3B14 PE=1 SV=1	1	-0.268816758
P02786	TFR1_HUMAN	Transferrin receptor protein 1 OS=Homo sapiens GN=TFR1 PE=1 SV=2	1	-0.268816758
P09455	RET1_HUMAN	Retinol-binding protein 1 OS=Homo sapiens GN=RBP1 PE=1 SV=2	1	-0.268816758
P11177-2	ODPB_HUMAN	Isoform 2 of Pyruvate dehydrogenase E1 component subunit beta	1	-0.268816758
P26373	RL13_HUMAN	Isoform 2 of 60S ribosomal protein L13 OS=Homo sapiens GN=RPL13 PE=1 SV=4	1	-0.268816758
P33993	MCM7_HUMAN	DNA replication licensing factor MCM7 OS=Homo sapiens GN=MCM7 PE=1 SV=4	1	-0.268816758
P37268	FDFT_HUMAN	Squalene synthase OS=Homo sapiens GN=FDFT1 PE=1 SV=1	1	-0.268816758
P62899-2	RL31_HUMAN	Isoform 2 of 60S ribosomal protein L31 OS=Homo sapiens GN=RPL31	1	-0.268816758
P62979	RS27A_HUMAN	Ubiquitin-40S ribosomal protein S27a OS=Homo sapiens GN=RPS27A PE=1 SV=2	1	-0.268816758
Q15029-2	US51_HUMAN	Isoform 2 of 116 kDa U5 small nuclear ribonucleoprotein component OS=Homo sapiens GN=EFTUD2	1	-0.268816758
Q9UID3-2	VPS51_HUMAN	Isoform 2 of Vacuolar protein sorting-associated protein 51 homolog OS=Homo sapiens GN=VPS51	1	-0.268816758
HOY2Z6	HOY2Z6_HUMAN	Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0.9999	-0.268816758
Q9UHV9	PFD2_HUMAN	Prefoldin subunit 2 OS=Homo sapiens GN=PFDN2 PE=1 SV=1	0.9991	-0.268816758
P37198	NUP62_HUMAN	Nuclear pore glycoprotein p62 OS=Homo sapiens GN=NUP62 PE=1 SV=3	0.9987	-0.268816758
Q15652	JHD2C_HUMAN	Probable JmjC domain-containing histone demethylation protein 2C OS=Homo sapiens GN=JMJD1C PE=	0.9975	-0.268816758
Q15070-2	OXA1L_HUMAN	Isoform 2 of Mitochondrial inner membrane protein OXA1L OS=Homo sapiens GN=OXA1L	0.9855	-0.268816758
Q00341	VIGLN_HUMAN	Vigilin OS=Homo sapiens GN=HDLBP PE=1 SV=2	0.9823	-0.268816758
Q8NVCV1-2	ADAD2_HUMAN	Isoform 2 of Adenosine deaminase domain-containing protein 2 OS=Homo sapiens GN=ADAD2	0.9754	-0.268816758
P30154-2	2AAB_HUMAN	Isoform 2 of Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A beta isoform OS=Ho	0.9508	-0.268816758
O43684-2	BUB3_HUMAN	Isoform 2 of Mitotic checkpoint protein BUB3 OS=Homo sapiens GN=BUB3	1	-0.286304185
O75608-2	LYPA1_HUMAN	Isoform 2 of Acyl-protein thioesterase 1 OS=Homo sapiens GN=LYPA1	1	-0.286304185
O75694-2	NU155_HUMAN	Isoform 2 of Nuclear pore complex protein Nup155 OS=Homo sapiens GN=NUP155	1	-0.286304185
P08397-2	HEM3_HUMAN	Isoform 2 of Porphobilinogen deaminase OS=Homo sapiens GN=HMBS	1	-0.286304185
P14174	MIF_HUMAN	Macrophage migration inhibitory factor OS=Homo sapiens GN=MIF PE=1 SV=4	1	-0.286304185
P36542-2	ATPG_HUMAN	Isoform Heart of ATP synthase subunit gamma	1	-0.286304185
P49721	PSB2_HUMAN	Proteasome subunit beta type-2 OS=Homo sapiens GN=PSMB2 PE=1 SV=1	1	-0.286304185
Q9NTK5-3	OLA1_HUMAN	Isoform 3 of Obg-like ATPase 1 OS=Homo sapiens GN=OLA1	1	-0.286304185
O75533	SF3B1_HUMAN	Splicing factor 3B subunit 1 OS=Homo sapiens GN=SF3B1 PE=1 SV=3	1	-0.286304185
P04844-2	RPN2_HUMAN	Isoform 2 of Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 2 OS=Homo sapien	1	-0.286304185
P18206-2	VINC_HUMAN	Isoform 1 of Vinculin OS=Homo sapiens GN=VCL	1	-0.286304185
Q92598-2	HS105_HUMAN	Isoform Beta of Heat shock protein 105 kDa OS=Homo sapiens GN=HSPH1	1	-0.286304185
P40926	MDHM_HUMAN	Malate dehydrogenase	1	-0.286304185
P52209	6PGD_HUMAN	6-phosphogluconate dehydrogenase	1	-0.286304185
P60228	EIF3E_HUMAN	Eukaryotic translation initiation factor 3 subunit E OS=Homo sapiens GN=EIF3E PE=1 SV=1	1	-0.286304185
P61289-2	PSME3_HUMAN	Isoform 2 of Proteasome activator complex subunit 3 OS=Homo sapiens GN=PSME3	1	-0.286304185
P62633-2	CNBP_HUMAN	Isoform 2 of Cellular nucleic acid-binding protein OS=Homo sapiens GN=CNBP	1	-0.286304185
P62714	PPP2AB_HUMAN	Serine/threonine-protein phosphatase 2A catalytic subunit beta isoform OS=Homo sapiens GN=PPP2CB	1	-0.286304185
Q04637-3	IF4G1_HUMAN	Isoform B of Eukaryotic translation initiation factor 4 gamma 1 OS=Homo sapiens GN=EIF4G1	1	-0.286304185
Q12789-3	TF3C1_HUMAN	Isoform 2 of General transcription factor 3C polypeptide 1 OS=Homo sapiens GN=GTF3C1	1	-0.286304185
Q14839-2	CHD4_HUMAN	Isoform 2 of Chromodomain-helicase-DNA-binding protein 4 OS=Homo sapiens GN=CHD4	1	-0.286304185
Q16891-2	IMMT_HUMAN	Isoform 2 of Mitochondrial inner membrane protein OS=Homo sapiens GN=IMMT	1	-0.286304185
Q9Y265	RUVB1_HUMAN	RuvB-like 1 OS=Homo sapiens GN=RUVB1 PE=1 SV=1	1	-0.286304185
P40394-2	ADH7_HUMAN	Isoform 2 of Alcohol dehydrogenase class 4 mu/sigma chain OS=Homo sapiens GN=ADH7	0.9954	-0.286304185
O00625	PIR_HUMAN	Pirin OS=Homo sapiens GN=PIR PE=1 SV=1	1	-0.304006187
O43765	SGTA_HUMAN	Small glutamine-rich tetratricopeptide repeat-containing protein alpha OS=Homo sapiens GN=SGTA PE=	1	-0.304006187
P12004	PCNA_HUMAN	Proliferating cell nuclear antigen OS=Homo sapiens GN=PCNA PE=1 SV=1	1	-0.304006187
P30041	PRDX6_HUMAN	Peroxiredoxin-6 OS=Homo sapiens GN=PRDX6 PE=1 SV=3	1	-0.304006187
Q9UPA5	BSN_HUMAN	Protein bassoon OS=Homo sapiens GN=BSN PE=1 SV=4	1	-0.304006187
P06576	ATPB_HUMAN	ATP synthase subunit beta	1	-0.304006187
P06748	NPM_HUMAN	Nucleophosmin OS=Homo sapiens GN=NPM1 PE=1 SV=2	1	-0.304006187
P62879	GBB2_HUMAN	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-2 OS=Homo sapiens GN=GNB2 PE=1 SV	1	-0.304006187
C9JD32	C9JD32_HUMAN	60S ribosomal protein L23 (Fragment) OS=Homo sapiens GN=RPL23 PE=3 SV=1	1	-0.304006187
P61006	RAB8A_HUMAN	Ras-related protein Rab-8A OS=Homo sapiens GN=RAB8A PE=1 SV=1	0.9999	-0.304006187
P54819-2	KAD2_HUMAN	Isoform 2 of Adenylate kinase 2	0.9999	-0.304006187
Q8IWX7-2	UNC45B_HUMAN	Isoform 2 of Protein unc-45 homolog B OS=Homo sapiens GN=UNC45B	0.9996	-0.304006187
Q9P258	RCC2_HUMAN	Protein RCC2 OS=Homo sapiens GN=RCC2 PE=1 SV=2	0.9976	-0.304006187
Q09472	EP300_HUMAN	Histone acetyltransferase p300 OS=Homo sapiens GN=EP300 PE=1 SV=2	0.9936	-0.304006187
Q8N7X0	ADGB_HUMAN	Androglobin OS=Homo sapiens GN=ADGB PE=2 SV=3	0.8681	-0.304006187
O76003	GLRX3_HUMAN	Glutaredoxin-3 OS=Homo sapiens GN=GLRX3 PE=1 SV=2	1	-0.321928095
P54886-2	P5CS_HUMAN	Isoform Short of Delta-1-pyrroline-5-carboxylate synthase OS=Homo sapiens GN=ALDH18A1	1	-0.321928095
P08865	RSSA_HUMAN	40S ribosomal protein SA OS=Homo sapiens GN=RPSA PE=1 SV=4	1	-0.321928095
P11586	C1TC_HUMAN	C-1-tetrahydrofolate synthase	1	-0.321928095
P47755	CAZA2_HUMAN	F-actin-capping protein subunit alpha-2 OS=Homo sapiens GN=CAPZA2 PE=1 SV=3	1	-0.321928095
P84103	SRSF3_HUMAN	Serine/arginine-rich splicing factor 3 OS=Homo sapiens GN=SRSF3 PE=1 SV=1	1	-0.321928095
Q7L2H7	EIF3M_HUMAN	Eukaryotic translation initiation factor 3 subunit M OS=Homo sapiens GN=EIF3M PE=1 SV=1	1	-0.321928095
Q86VP6	CAND1_HUMAN	Cullin-associated NEDD8-dissociated protein 1 OS=Homo sapiens GN=CAND1 PE=1 SV=2	1	-0.321928095
Q8TEX9-2	IPO4_HUMAN	Isoform 2 of Importin-4 OS=Homo sapiens GN=IPO4	1	-0.321928095
O14880	MGST3_HUMAN	Microsomal glutathione S-transferase 3 OS=Homo sapiens GN=MGST3 PE=1 SV=1	0.9999	-0.321928095
P11216	PYGB_HUMAN	Glycogen phosphorylase	0.993	-0.321928095
P62851	RS25_HUMAN	40S ribosomal protein S25 OS=Homo sapiens GN=RPS25 PE=1 SV=1	0.9886	-0.321928095
P82933	RT09_HUMAN	28S ribosomal protein S9	0.9807	-0.321928095
Q9GZM5	YIPF3_HUMAN	Protein YIPF3 OS=Homo sapiens GN=YIPF3 PE=1 SV=1	0.8784	-0.321928095
A6NHG4	DDTL_HUMAN	D-dopachrome decarboxylase-like protein OS=Homo sapiens GN=DDTL PE=2 SV=1	1	-0.340075442
Q96QK1	VPS35_HUMAN	Vacuolar protein sorting-associated protein 35 OS=Homo sapiens GN=VPS35 PE=1 SV=2	1	-0.340075442
Q08211	DHX9_HUMAN	ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4	1	-0.340075442
P27708	PYR1_HUMAN	CAD protein OS=Homo sapiens GN=CAD PE=1 SV=3	1	-0.340075442
P31943	HNRH1_HUMAN	Heterogeneous nuclear ribonucleoprotein H OS=Homo sapiens GN=HNRNPH1 PE=1 SV=4	1	-0.340075442
P62937	PPIA_HUMAN	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2	1	-0.340075442
P68366	TBA4A_HUMAN	Tubulin alpha-4A chain OS=Homo sapiens GN=TUBA4A PE=1 SV=1	1	-0.340075442
Q00839-2	HNRPU_HUMAN	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU	1	-0.340075442
Q02878	RL6_HUMAN	60S ribosomal protein L6 OS=Homo sapiens GN=RPL6 PE=1 SV=3	1	-0.340075442
Q10567-2	AP1B1_HUMAN	Isoform B of AP-1 complex subunit beta-1 OS=Homo sapiens GN=AP1B1	1	-0.340075442
Q9NZ18	IF2B1_HUMAN	Insulin-like growth factor 2 mRNA-binding protein 1 OS=Homo sapiens GN=IGF2BP1 PE=1 SV=2	1	-0.340075442
P52788-2	SPSY_HUMAN	Isoform 2 of Spermine synthase OS=Homo sapiens GN=SMS	0.9999	-0.340075442

C9JQ55	C9JQ55_HUMAN	Protein RPS4XP21 OS=Homo sapiens GN=RPS4XP21 PE=4 SV=1	0.9993	-0.340075442
P49756-2	RBM25_HUMAN	Isoform 2 of RNA-binding protein 25 OS=Homo sapiens GN=RBM25	0.9986	-0.340075442
Q9B2M5	N2DL2_HUMAN	NKG2D ligand 2 OS=Homo sapiens GN=ULBP2 PE=1 SV=1	0.9924	-0.340075442
P61923	COPZ1_HUMAN	Coatomer subunit zeta-1 OS=Homo sapiens GN=COPZ1 PE=1 SV=1	1	-0.358453971
Q15084-2	PDIA6_HUMAN	Isoform 2 of Protein disulfide-isomerase A6 OS=Homo sapiens GN=PDIA6	1	-0.358453971
O00571-2	DDX3X_HUMAN	Isoform 2 of ATP-dependent RNA helicase DDX3X OS=Homo sapiens GN=DDX3X	1	-0.358453971
P10809	CH60_HUMAN	60 kDa heat shock protein	1	-0.358453971
P12956	XRCC6_HUMAN	X-ray repair cross-complementing protein 6 OS=Homo sapiens GN=XRCC6 PE=1 SV=2	1	-0.358453971
P13639	EEF2_HUMAN	Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4	1	-0.358453971
P62491	RB11A_HUMAN	Ras-related protein Rab-11A OS=Homo sapiens GN=RB11A PE=1 SV=3	1	-0.358453971
Q16531	DDB1_HUMAN	DNA damage-binding protein 1 OS=Homo sapiens GN=DDB1 PE=1 SV=1	1	-0.358453971
Q9NX63	CHCH3_HUMAN	Coiled-coil-helix-coiled-coil-helix domain-containing protein 3	1	-0.358453971
Q9UBE0	SAE1_HUMAN	SUMO-activating enzyme subunit 1 OS=Homo sapiens GN=SAE1 PE=1 SV=1	1	-0.358453971
Q9Y6C9	MTCH2_HUMAN	Mitochondrial carrier homolog 2 OS=Homo sapiens GN=MTCH2 PE=1 SV=1	1	-0.358453971
O95336	6PGL_HUMAN	6-phosphogluconolactonase OS=Homo sapiens GN=6PGL PE=1 SV=2	0.9987	-0.358453971
J3QQT0	J3QQT0_HUMAN	Protein-methionine sulfoxide oxidase MICAL3 (Fragment) OS=Homo sapiens GN=MICAL3 PE=4 SV=1	0.9975	-0.358453971
O75417-2	DPOLQ_HUMAN	Isoform 2 of DNA polymerase theta OS=Homo sapiens GN=POLQ	0.9915	-0.358453971
O60341-2	KDM1A_HUMAN	Isoform 2 of Lysine-specific histone demethylase 1A OS=Homo sapiens GN=KDM1A	0.9891	-0.358453971
Q00154-2	BACH_HUMAN	Isoform 2 of Cytosolic acyl coenzyme A thioester hydrolase OS=Homo sapiens GN=ACOT7	0.9601	-0.358453971
Q9B2H6	WDR11_HUMAN	WD repeat-containing protein 11 OS=Homo sapiens GN=WDR11 PE=1 SV=1	0.958	-0.358453971
P23368-2	MAOM_HUMAN	Isoform 2 of NAD-dependent malic enzyme	1	-0.377069649
Q9UBQ0-2	VPS29_HUMAN	Isoform 2 of Vacuolar protein sorting-associated protein 29 OS=Homo sapiens GN=VPS29	1	-0.377069649
P24534	EEF1B_HUMAN	Elongation factor 1-beta OS=Homo sapiens GN=EEF1B2 PE=1 SV=3	1	-0.377069649
P36776	LONM_HUMAN	Lon protease homolog	1	-0.377069649
P38646	GRP75_HUMAN	Stress-70 protein	1	-0.377069649
P49411	EFTU_HUMAN	Elongation factor Tu	1	-0.377069649
P62081	RS7_HUMAN	40S ribosomal protein S7 OS=Homo sapiens GN=RPS7 PE=1 SV=1	1	-0.377069649
Q15417	CNN3_HUMAN	Calponin-3 OS=Homo sapiens GN=CNN3 PE=1 SV=1	1	-0.377069649
Q8NCW5	NNRE_HUMAN	NAD(P)H-hydrate epimerase OS=Homo sapiens GN=APOA1B1P PE=1 SV=2	1	-0.377069649
P57678	GEM14_HUMAN	Gem-associated protein 4 OS=Homo sapiens GN=GEMIN4 PE=1 SV=2	0.9999	-0.377069649
Q9Y606-2	TRUA_HUMAN	Isoform 2 of tRNA pseudouridine synthase A	0.9961	-0.377069649
Q8TDW7-2	FAT3_HUMAN	Isoform 2 of Protocadherin Fat 3 OS=Homo sapiens GN=FAT3	0.9804	-0.377069649
Q13576-2	IQGA2_HUMAN	Isoform 2 of Ras GTPase-activating-like protein IQGAP2 OS=Homo sapiens GN=IQGAP2	0.9742	-0.377069649
Q9H0M0-2	WWP1_HUMAN	Isoform 2 of NEDD4-like E3 ubiquitin-protein ligase WWP1 OS=Homo sapiens GN=WWP1	0.9328	-0.377069649
Q6UB35	C1TM_HUMAN	Monofunctional C1-tetrahydrofolate synthase	0.891	-0.377069649
P07339	CATD_HUMAN	Cathepsin D OS=Homo sapiens GN=CTSD PE=1 SV=1	1	-0.395928676
P09884	DPOLA_HUMAN	DNA polymerase alpha catalytic subunit OS=Homo sapiens GN=POLA1 PE=1 SV=2	1	-0.395928676
Q16836-2	HCDH_HUMAN	Isoform 2 of Hydroxyacyl-coenzyme A dehydrogenase	1	-0.395928676
P05387	RLA2_HUMAN	60S acidic ribosomal protein P2 OS=Homo sapiens GN=RPLP2 PE=1 SV=1	1	-0.395928676
P31939	PUR9_HUMAN	Bifunctional purine biosynthesis protein PURH OS=Homo sapiens GN=ATIC PE=1 SV=3	1	-0.395928676
P51571	SSRD_HUMAN	Translocan-associated protein subunit delta OS=Homo sapiens GN=SSR4 PE=1 SV=1	1	-0.395928676
P68104	EF1A1_HUMAN	Elongation factor 1-alpha 1 OS=Homo sapiens GN=EF1A1 PE=1 SV=1	1	-0.395928676
Q01130	SRSF2_HUMAN	Serine/arginine-rich splicing factor 2 OS=Homo sapiens GN=SRSF2 PE=1 SV=4	1	-0.395928676
O43795-2	MYO1B_HUMAN	Isoform 2 of Unconventional myosin-Ib OS=Homo sapiens GN=MYO1B	0.9999	-0.395928676
Q8TB22-2	SPT20_HUMAN	Isoform 2 of Spermatogenesis-associated protein 20 OS=Homo sapiens GN=SPATA20	0.9979	-0.395928676
O60841	IF2P_HUMAN	Eukaryotic translation initiation factor 5B OS=Homo sapiens GN=EIF5B PE=1 SV=4	0.9953	-0.395928676
O43823	AKAP8_HUMAN	A-kinase anchor protein 8 OS=Homo sapiens GN=AKAP8 PE=1 SV=1	0.9919	-0.395928676
O14782	KIF3C_HUMAN	Kinesin-like protein KIF3C OS=Homo sapiens GN=KIF3C PE=1 SV=3	0.98	-0.395928676
Q00534	CDK6_HUMAN	Cyclin-dependent kinase 6 OS=Homo sapiens GN=CDK6 PE=1 SV=1	0.9795	-0.395928676
B0UXE9	B0UXE9_HUMAN	Collagen alpha-2(XI) chain OS=Homo sapiens GN=COL11A2 PE=4 SV=1	0.9574	-0.395928676
P56134-2	ATPK_HUMAN	Isoform 2 of ATP synthase subunit f	1	-0.415037499
P07741	APT_HUMAN	Adenine phosphoribosyltransferase OS=Homo sapiens GN=APRT PE=1 SV=2	1	-0.415037499
Q9BUJ2-2	HNRL1_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein U-like protein 1 OS=Homo sapiens GN=HNRNPU	1	-0.415037499
Q9UBQ5	EIF3K_HUMAN	Eukaryotic translation initiation factor 3 subunit K OS=Homo sapiens GN=EIF3K PE=1 SV=1	1	-0.415037499
Q9Y4L1	HYOU1_HUMAN	Hypoxia up-regulated protein 1 OS=Homo sapiens GN=HYOU1 PE=1 SV=1	1	-0.415037499
E9PAV3	E9PAV3_HUMAN	Nascent polypeptide-associated complex subunit alpha OS=Homo sapiens GN=NACA PE=4 SV=1	1	-0.415037499
P11021	GRP78_HUMAN	78 kDa glucose-regulated protein OS=Homo sapiens GN=HSPA5 PE=1 SV=2	1	-0.415037499
P62888	RL30_HUMAN	60S ribosomal protein L30 OS=Homo sapiens GN=RPL30 PE=1 SV=2	1	-0.415037499
G3V2X9	G3V2X9_HUMAN	Alpha-actinin-1 (Fragment) OS=Homo sapiens GN=ACTN1 PE=4 SV=1	0.9997	-0.415037499
P31689	DNJA1_HUMAN	DnaJ homolog subfamily A member 1 OS=Homo sapiens GN=DNJA1 PE=1 SV=2	0.9925	-0.415037499
Q9B2K3	NACP1_HUMAN	Putative nascent polypeptide-associated complex subunit alpha-like protein OS=Homo sapiens GN=NACP	0.9916	-0.415037499
O14686-3	MLL2_HUMAN	Isoform 3 of Histone-lysine N-methyltransferase MLL2 OS=Homo sapiens GN=MLL2	0.9891	-0.415037499
Q13283	G3BP1_HUMAN	Ras GTPase-activating protein-binding protein 1 OS=Homo sapiens GN=G3BP1 PE=1 SV=1	0.9186	-0.415037499
Q92667-2	AKAP1_HUMAN	Isoform 2 of A-kinase anchor protein 1	0.916	-0.415037499
O43399-2	TPD54_HUMAN	Isoform 2 of Tumor protein D54 OS=Homo sapiens GN=TPD52L2	1	-0.434402824
P23258	TBG1_HUMAN	Tubulin gamma-1 chain OS=Homo sapiens GN=TUBG1 PE=1 SV=2	1	-0.434402824
P55157	MTP_HUMAN	Microsomal triglyceride transfer protein large subunit OS=Homo sapiens GN=MTPP PE=1 SV=1	1	-0.434402824
Q9BQ67	GRWD1_HUMAN	Glutamate-rich WD repeat-containing protein 1 OS=Homo sapiens GN=GRWD1 PE=1 SV=1	1	-0.434402824
Q9Y295	DRG1_HUMAN	Developmentally-regulated GTP-binding protein 1 OS=Homo sapiens GN=DRG1 PE=1 SV=1	1	-0.434402824
P46060	RAGP1_HUMAN	Ran GTPase-activating protein 1 OS=Homo sapiens GN=RANGAP1 PE=1 SV=1	1	-0.434402824
P49773	HINT1_HUMAN	Histidine triad nucleotide-binding protein 1 OS=Homo sapiens GN=HINT1 PE=1 SV=2	1	-0.434402824
P51149	RAB7A_HUMAN	Ras-related protein Rab-7a OS=Homo sapiens GN=RAB7A PE=1 SV=1	1	-0.434402824
P53621	COPA_HUMAN	Coatomer subunit alpha OS=Homo sapiens GN=COPA PE=1 SV=2	1	-0.434402824
P55209	NP1L1_HUMAN	Nucleosome assembly protein 1-like 1 OS=Homo sapiens GN=NP1L1 PE=1 SV=1	1	-0.434402824
P56192	SYMC_HUMAN	Methionine--tRNA ligase	1	-0.434402824
Q92616	GCN1L_HUMAN	Translational activator GCN1 OS=Homo sapiens GN=GCN1L1 PE=1 SV=6	1	-0.434402824
Q9Y678	COPG1_HUMAN	Coatomer subunit gamma-1 OS=Homo sapiens GN=COPG1 PE=1 SV=1	1	-0.434402824
Q53G59	SNUT2_HUMAN	U4/U6.U5 tri-snRNP-associated protein 2 OS=Homo sapiens GN=USP39 PE=1 SV=2	0.9959	-0.434402824
Q9H2M9	RBGPR_HUMAN	Rab3 GTPase-activating protein non-catalytic subunit OS=Homo sapiens GN=RAB3GAP2 PE=1 SV=1	0.9868	-0.434402824
P11532-2	DMD_HUMAN	Isoform 1 of Dystrophin OS=Homo sapiens GN=DMD	0.954	-0.434402824
Q96737-2	RBM15_HUMAN	Isoform 2 of Putative RNA-binding protein 15 OS=Homo sapiens GN=RBM15	0.9082	-0.434402824
Q9B5J8-2	ESYT1_HUMAN	Isoform 2 of Extended synaptotagmin-1 OS=Homo sapiens GN=ESYT1	1	-0.454031631
Q9N569	TOM22_HUMAN	Mitochondrial import receptor subunit TOM22 homolog OS=Homo sapiens GN=TOMM22 PE=1 SV=3	1	-0.454031631
P39656	OST48_HUMAN	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit OS=Homo sapiens GN=L	1	-0.454031631
P48047	ATPO_HUMAN	ATP synthase subunit O	1	-0.454031631
P61970	NTF2_HUMAN	Nuclear transport factor 2 OS=Homo sapiens GN=NUTF2 PE=1 SV=1	1	-0.454031631
P62277	RS13_HUMAN	40S ribosomal protein S13 OS=Homo sapiens GN=RPS13 PE=1 SV=2	1	-0.454031631
Q9H984	SFXN1_HUMAN	Sideroflexin-1 OS=Homo sapiens GN=SFXN1 PE=1 SV=4	1	-0.454031631
P52292	IMA2_HUMAN	Importin subunit alpha-2 OS=Homo sapiens GN=KPNA2 PE=1 SV=1	1	-0.473931188
Q14978-2	NOLC1_HUMAN	Isoform Beta of Nucleolar and coiled-body phosphoprotein 1 OS=Homo sapiens GN=NOLC1	1	-0.473931188
Q14152	EIF3A_HUMAN	Eukaryotic translation initiation factor 3 subunit A OS=Homo sapiens GN=EIF3A PE=1 SV=1	1	-0.473931188
Q9C005	DPY30_HUMAN	Protein dpy-30 homolog OS=Homo sapiens GN=DPY30 PE=1 SV=1	1	-0.473931188

Q6P0N0	M18BP_HUMAN	Mis18-binding protein 1 OS=Homo sapiens GN=MIS18BP1 PE=1 SV=1	0.999	-0.473931188
Q969G6	RIFK_HUMAN	Riboflavin kinase OS=Homo sapiens GN=RFK PE=1 SV=2	0.9979	-0.473931188
Q14651	PLS1_HUMAN	Plastin-1 OS=Homo sapiens GN=PLS1 PE=1 SV=2	0.9928	-0.473931188
Q5JU85-2	IQEC2_HUMAN	Isoform 2 of IQ motif and SEC7 domain-containing protein 2 OS=Homo sapiens GN=IQSEC2	0.9927	-0.473931188
P06899	H2B1J_HUMAN	Histone H2B type 1-J OS=Homo sapiens GN=HIST1H2BJ PE=1 SV=3	0.9831	-0.473931188
Q13825-2	AUHM_HUMAN	Isoform 2 of Methylglutaconyl-CoA hydratase	0.9794	-0.473931188
Q9H0S4	DDX47_HUMAN	Probable ATP-dependent RNA helicase DDX47 OS=Homo sapiens GN=DDX47 PE=1 SV=1	0.8967	-0.473931188
P09622	DLDH_HUMAN	Dihydrolipoyl dehydrogenase	1	-0.49410907
P52732	KIF11_HUMAN	Kinesin-like protein KIF11 OS=Homo sapiens GN=KIF11 PE=1 SV=2	1	-0.49410907
O96019	ACTL6A_HUMAN	Actin-like protein 6A OS=Homo sapiens GN=ACTL6A PE=1 SV=1	1	-0.49410907
P22695	QCR2_HUMAN	Cytochrome b-c1 complex subunit 2	1	-0.49410907
P27824	CALX_HUMAN	Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2	1	-0.49410907
P52907	CAZA1_HUMAN	F-actin-capping protein subunit alpha-1 OS=Homo sapiens GN=CAPZA1 PE=1 SV=3	1	-0.49410907
Q6S8J3	POTEE_HUMAN	POTE ankyrin domain family member E OS=Homo sapiens GN=POTEE PE=1 SV=3	0.9988	-0.49410907
O00231-2	PSD11_HUMAN	Isoform 2 of 26S proteasome non-ATPase regulatory subunit 11 OS=Homo sapiens GN=PSMD11	0.9927	-0.49410907
Q8N857-2	ENAH_HUMAN	Isoform 2 of Protein enabled homolog OS=Homo sapiens GN=ENAH	0.9873	-0.49410907
P30040	ERP29_HUMAN	Endoplasmic reticulum resident protein 29 OS=Homo sapiens GN=ERP29 PE=1 SV=4	1	-0.514573173
P49327	FAS_HUMAN	Fatty acid synthase OS=Homo sapiens GN=FASN PE=1 SV=3	1	-0.514573173
Q13148-2	TADBP_HUMAN	Isoform 2 of TAR DNA-binding protein 43 OS=Homo sapiens GN=TARDBP	1	-0.514573173
P68036	UB2L3_HUMAN	Ubiquitin-conjugating enzyme E2 L3 OS=Homo sapiens GN=UBE2L3 PE=1 SV=1	0.9915	-0.514573173
Q14247-2	SRC8_HUMAN	Isoform 2 of Src substrate cortactin OS=Homo sapiens GN=CTTN	0.9891	-0.514573173
Q8NCM8-2	DYHC2_HUMAN	Isoform 2 of Cytoplasmic dynein 2 heavy chain 1 OS=Homo sapiens GN=DYNC2H1	0.9538	-0.514573173
P16401	H15_HUMAN	Histone H1.5 OS=Homo sapiens GN=HIST1H1B PE=1 SV=3	1	-0.535331733
O95202	LETM1_HUMAN	LETM1 and EF-hand domain-containing protein 1	1	-0.535331733
P15880	RS2_HUMAN	40S ribosomal protein S2 OS=Homo sapiens GN=RPS2 PE=1 SV=2	1	-0.535331733
P24752	THIL_HUMAN	Acetyl-CoA acetyltransferase	1	-0.535331733
P61758	VPD3_HUMAN	Prefoldin subunit 3 OS=Homo sapiens GN=VBP1 PE=1 SV=3	1	-0.535331733
Q99714	HCD2_HUMAN	3-hydroxyacyl-CoA dehydrogenase type-2 OS=Homo sapiens GN=HSD17B10 PE=1 SV=3	1	-0.535331733
P10599	THIO_HUMAN	Thioredoxin OS=Homo sapiens GN=TXN PE=1 SV=3	0.9998	-0.535331733
Q92922	SMRC1_HUMAN	SWI/SNF complex subunit SMARCC1 OS=Homo sapiens GN=SMARCC1 PE=1 SV=3	0.9986	-0.535331733
Q96FT7-2	ASIC4_HUMAN	Isoform 2 of Acid-sensing ion channel 4 OS=Homo sapiens GN=ASIC4	0.9961	-0.535331733
Q9BRX8-2	F213A_HUMAN	Isoform 2 of Redox-regulatory protein FAM213A OS=Homo sapiens GN=FAM213A	1	-0.556393349
P00387-3	NB5R3_HUMAN	Isoform 3 of NADH-cytochrome b5 reductase 3 OS=Homo sapiens GN=CYB5R3	1	-0.556393349
P13804-2	ETFA_HUMAN	Isoform 2 of Electron transfer flavoprotein subunit alpha	1	-0.556393349
P62140	PP1B_HUMAN	Serine/threonine-protein phosphatase PP1-beta catalytic subunit OS=Homo sapiens GN=PPP1CB PE=1 SV=1	1	-0.556393349
Q15365	PCBP1_HUMAN	Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2	1	-0.556393349
Q15125	EBP_HUMAN	3-beta-hydroxysteroid-Delta(8)	0.9781	-0.556393349
Q5T946	Q5T946_HUMAN	Glyoxylate reductase/hydroxypyruvate reductase OS=Homo sapiens GN=GRHPR PE=2 SV=1	0.9636	-0.556393349
Q13435	SF3B2_HUMAN	Splicing factor 3B subunit 2 OS=Homo sapiens GN=SF3B2 PE=1 SV=2	0.942	-0.556393349
O95347-2	SMC2_HUMAN	Isoform 2 of Structural maintenance of chromosomes protein 2 OS=Homo sapiens GN=SMC2	1	-0.577766999
P08670	VIME_HUMAN	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4	1	-0.577766999
P78344-2	IF4G2_HUMAN	Isoform 2 of Eukaryotic translation initiation factor 4 gamma 2 OS=Homo sapiens GN=EIF4G2	1	-0.577766999
Q92974-2	ARHG2_HUMAN	Isoform 2 of Rho guanine nucleotide exchange factor 2 OS=Homo sapiens GN=ARHG2	0.9896	-0.577766999
P00441	SODC_HUMAN	Superoxide dismutase [Cu-Zn] OS=Homo sapiens GN=SOD1 PE=1 SV=2	1	-0.59946207
Q9UJZ1	STML2_HUMAN	Stomatin-like protein 2 OS=Homo sapiens GN=STOML2 PE=1 SV=1	1	-0.59946207
O15020-2	SPTN2_HUMAN	Isoform 2 of Spectrin beta chain	0.9995	-0.59946207
P30566-2	PUR8_HUMAN	Isoform 2 of Adenylosuccinate lyase OS=Homo sapiens GN=ADSL	0.9991	-0.59946207
Q9C0C9	UBE2O_HUMAN	Ubiquitin-conjugating enzyme E2 O OS=Homo sapiens GN=UBE2O PE=1 SV=3	0.9912	-0.59946207
Q96LP4	YS017_HUMAN	Putative uncharacterized protein FU25328 OS=Homo sapiens PE=2 SV=1	0.9808	-0.59946207
Q7Z4W1	DCXR_HUMAN	L-xylulose reductase OS=Homo sapiens GN=DCXR PE=1 SV=2	1	-0.621488377
P08621-2	RU17_HUMAN	Isoform 2 of U1 small nuclear ribonucleoprotein 70 kDa OS=Homo sapiens GN=SNRNP70	1	-0.621488377
Q9BXW7-2	CECR5_HUMAN	Isoform 1 of Cat eye syndrome critical region protein 5 OS=Homo sapiens GN=CECR5	1	-0.621488377
C9JDM3	C9JDM3_HUMAN	Ran-specific GTPase-activating protein (Fragment) OS=Homo sapiens GN=RANBP1 PE=4 SV=1	0.9872	-0.621488377
P21439-2	MDR3_HUMAN	Isoform 2 of Multidrug resistance protein 3 OS=Homo sapiens GN=ABCB4	0.9856	-0.621488377
P53814-2	SMTN_HUMAN	Isoform A of Smoothelin OS=Homo sapiens GN=SMTN	0.9842	-0.621488377
Q9ULC4-2	MCTS1_HUMAN	Isoform 2 of Malignant T-cell-amplified sequence 1 OS=Homo sapiens GN=MCTS1	0.9778	-0.621488377
H0YAE5	H0YAE5_HUMAN	Matrix metalloproteinase-23	0.975	-0.621488377
P05386	RLA1_HUMAN	60S acidic ribosomal protein P1 OS=Homo sapiens GN=RPLP1 PE=1 SV=1	1	-0.64385619
Q01518	CAP1_HUMAN	Adenylyl cyclase-associated protein 1 OS=Homo sapiens GN=CAP1 PE=1 SV=5	1	-0.64385619
Q86US8	EST1A_HUMAN	Telomerase-binding protein EST1A OS=Homo sapiens GN=SMG6 PE=1 SV=2	0.9994	-0.64385619
P10412	H14_HUMAN	Histone H1.4 OS=Homo sapiens GN=HIST1H1E PE=1 SV=2	1	-0.666572626
Q5SW79-3	CE170_HUMAN	Isoform 3 of Centrosomal protein of 170 kDa OS=Homo sapiens GN=CEP170	0.9993	-0.666572626
P27487	DPP4_HUMAN	Dipeptidyl peptidase 4 OS=Homo sapiens GN=DPP4 PE=1 SV=2	0.979	-0.666572626
Q9BYV6-3	TRI55_HUMAN	Isoform 3 of Tripartite motif-containing protein 55 OS=Homo sapiens GN=TRIM55	0.915	-0.666572626
Q7Z7H8	RM10_HUMAN	39S ribosomal protein L10	1	-0.689659879
O60814	H2B1K_HUMAN	Histone H2B type 1-K OS=Homo sapiens GN=HIST1H2BK PE=1 SV=3	1	-0.689659879
P20290-2	BTF3_HUMAN	Isoform 2 of Transcription factor BTF3 OS=Homo sapiens GN=BTF3	1	-0.689659879
Q9UHD8-5	SEPT9_HUMAN	Isoform 5 of Septin-9 OS=Homo sapiens GN=SEPT9	0.9995	-0.689659879
Q8WYA6	CTBL1_HUMAN	Beta-catenin-like protein 1 OS=Homo sapiens GN=CTNBL1 PE=1 SV=1	0.9951	-0.689659879
P47756-2	CAPZB_HUMAN	Isoform 2 of F-actin-capping protein subunit beta OS=Homo sapiens GN=CAPZB	1	-0.713118852
Q86UE4	LYRIC_HUMAN	Protein LYRIC OS=Homo sapiens GN=MTDH PE=1 SV=2	1	-0.713118852
P24539	AT5F1_HUMAN	ATP synthase subunit b	1	-0.713118852
Q7Z6E9-2	RBBP6_HUMAN	Isoform 2 of E3 ubiquitin-protein ligase RBBP6 OS=Homo sapiens GN=RBBP6	1	-0.713118852
Q86XP3-2	DDX42_HUMAN	Isoform 2 of ATP-dependent RNA helicase DDX42 OS=Homo sapiens GN=DDX42	0.9993	-0.713118852
O60869-2	EDF1_HUMAN	Isoform 2 of Endothelial differentiation-related factor 1 OS=Homo sapiens GN=EDF1	0.9981	-0.713118852
P20674	COX5A_HUMAN	Cytochrome c oxidase subunit 5A	0.9957	-0.713118852
Q96BW9	TAM41_HUMAN	Mitochondrial translocator assembly and maintenance protein 41 homolog OS=Homo sapiens GN=TAMI	0.988	-0.713118852
Q14204	DYHC1_HUMAN	Cytoplasmic dynein 1 heavy chain 1 OS=Homo sapiens GN=DYNC1H1 PE=1 SV=5	1	-0.736965594
P53004	BIEA_HUMAN	Biliverdin reductase A OS=Homo sapiens GN=BLVRA PE=1 SV=2	0.9998	-0.736965594
Q9Y5K5-2	UCHL5_HUMAN	Isoform 2 of Ubiquitin carboxyl-terminal hydrolase isozyme L5 OS=Homo sapiens GN=UCHL5	0.9996	-0.736965594
F5H0L1	F5H0L1_HUMAN	Cullin-7 OS=Homo sapiens GN=CUL7 PE=3 SV=1	0.9797	-0.736965594
P54709	AT1B3_HUMAN	Sodium/potassium-transporting ATPase subunit beta-3 OS=Homo sapiens GN=ATP1B3 PE=1 SV=1	1	-0.76121314
O60885-2	BRD4_HUMAN	Isoform 2 of Bromodomain-containing protein 4 OS=Homo sapiens GN=BRD4	0.9885	-0.76121314
Q9GZR7	DDX24_HUMAN	ATP-dependent RNA helicase DDX24 OS=Homo sapiens GN=DDX24 PE=1 SV=1	0.9671	-0.76121314
O96008	TOM40_HUMAN	Mitochondrial import receptor subunit TOM40 homolog OS=Homo sapiens GN=TOMM40 PE=1 SV=1	1	-0.785875195
P16403	H12_HUMAN	Histone H1.2 OS=Homo sapiens GN=HIST1H1C PE=1 SV=2	1	-0.785875195
P62314	SMD1_HUMAN	Small nuclear ribonucleoprotein Sm D1 OS=Homo sapiens GN=SNRPD1 PE=1 SV=1	1	-0.785875195
Q8N3T6	T132C_HUMAN	Transmembrane protein 132C OS=Homo sapiens GN=TMEM132C PE=2 SV=3	0.9994	-0.785875195
I3L1E1	I3L1E1_HUMAN	Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0.9196	-0.785875195
Q9BYN8	RT26_HUMAN	28S ribosomal protein S26	0.8861	-0.785875195
Q7Z2W9	RM21_HUMAN	39S ribosomal protein L21	0.808	-0.785875195
Q8NBX0	SCPDL_HUMAN	Saccharopine dehydrogenase-like oxidoreductase OS=Homo sapiens GN=SCCPDH PE=1 SV=1	0.9999	-0.810966176

P84243	H33_HUMAN	Histone H3.3 OS=Homo sapiens GN=H3F3A PE=1 SV=2	0.9997	-0.810966176
Q9UBU8-2	MO4L1_HUMAN	Isoform 2 of Mortality factor 4-like protein 1 OS=Homo sapiens GN=MORF4L1	0.9974	-0.836501268
P38935	SMBP2_HUMAN	DNA-binding protein SMBP2 OS=Homo sapiens GN=IGHMBP2 PE=1 SV=3	0.9492	-0.836501268
P62805	H4_HUMAN	Histone H4 OS=Homo sapiens GN=HIST1H4A PE=1 SV=2	1	-0.862496476
O43809	CPSF5_HUMAN	Cleavage and polyadenylation specificity factor subunit 5 OS=Homo sapiens GN=NUDT21 PE=1 SV=1	1	-0.862496476
O75489	NDU53_HUMAN	NADH dehydrogenase [ubiquinone] iron-sulfur protein 3	1	-0.862496476
Q16795	NDUA9_HUMAN	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 9	1	-0.862496476
Q9BPW8	NIPS1_HUMAN	Protein NipSnap homolog 1 OS=Homo sapiens GN=NIPSNAP1 PE=1 SV=1	1	-0.862496476
Q9P000-2	COMD9_HUMAN	Isoform 2 of COMM domain-containing protein 9 OS=Homo sapiens GN=COMMD9	0.9505	-0.862496476
P13796	PLSL_HUMAN	Plastin-2 OS=Homo sapiens GN=LCP1 PE=1 SV=6	1	-0.888968688
Q9Y5A9-2	YTHD2_HUMAN	Isoform 2 of YTH domain family protein 2 OS=Homo sapiens GN=YTHDF2	0.9994	-0.888968688
P00403	COX2_HUMAN	Cytochrome c oxidase subunit 2 OS=Homo sapiens GN=MT-CO2 PE=1 SV=1	0.9981	-0.888968688
Q965B4-3	SRPK1_HUMAN	Isoform 1 of SRSF protein kinase 1 OS=Homo sapiens GN=SRPK1	0.9276	-0.915935735
Q6ZP65	BICR1_HUMAN	Bicaudal D-related protein 1 OS=Homo sapiens GN=CCDC64 PE=2 SV=2	0.9888	-0.943416472
P42771-2	CD2A1_HUMAN	Isoform 2 of Cyclin-dependent kinase inhibitor 2A	0.9852	-0.943416472
P07225	PROS_HUMAN	Vitamin K-dependent protein S OS=Homo sapiens GN=PROS1 PE=1 SV=1	0.9415	-0.943416472
Q96T58	MINT_HUMAN	Msx2-interacting protein OS=Homo sapiens GN=SPEN PE=1 SV=1	0.9394	-0.943416472
Q9UPY3-2	DICER_HUMAN	Isoform 2 of Endoribonuclease Dicer OS=Homo sapiens GN=DICER1	0.8812	-0.943416472
Q14008-2	CKAP5_HUMAN	Isoform 2 of Cytoskeleton-associated protein 5 OS=Homo sapiens GN=CKAP5	1	-0.971430848
Q16763	UBE2S_HUMAN	Ubiquitin-conjugating enzyme E2 S OS=Homo sapiens GN=UBE2S PE=1 SV=2	0.9999	-0.971430848
P28161	GSTM2_HUMAN	Glutathione S-transferase Mu 2 OS=Homo sapiens GN=GSTM2 PE=1 SV=2	0.9994	-0.971430848
P32004-2	L1CAM_HUMAN	Isoform 2 of Neural cell adhesion molecule L1 OS=Homo sapiens GN=L1CAM	0.9989	-0.971430848
Q6IQ23-2	PKHA7_HUMAN	Isoform 2 of Pleckstrin homology domain-containing family A member 7 OS=Homo sapiens GN=PLEKH7	0.9938	-0.971430848
Q9BXP5-2	SRRT_HUMAN	Isoform 2 of Serrate RNA effector molecule homolog OS=Homo sapiens GN=SRRT	0.9669	-0.971430848
Q9BTC0	DIDO1_HUMAN	Death-inducer obliterator 1 OS=Homo sapiens GN=DIDO1 PE=1 SV=5	0.9845	-1
Q9H9J2	RM44_HUMAN	39S ribosomal protein L44	0.8585	-1
O14874	BCKD_HUMAN	[3-methyl-2-oxobutanoate dehydrogenase [lipamide]] kinase	0.8107	-1.029146346
P0C7U0	PPR28_HUMAN	Protein phosphatase 1 regulatory subunit 28 OS=Homo sapiens GN=ELFN1 PE=1 SV=1	1	-1.089267338
Q9Y277-2	VDAC3_HUMAN	Isoform 2 of Voltage-dependent anion-selective channel protein 3 OS=Homo sapiens GN=VDAC3	1	-1.089267338
Q13009	TIAM1_HUMAN	T-lymphoma invasion and metastasis-inducing protein 1 OS=Homo sapiens GN=TIAM1 PE=1 SV=2	0.9999	-1.089267338
Q00796	DHSO_HUMAN	Sorbitol dehydrogenase OS=Homo sapiens GN=SORD PE=1 SV=4	0.9997	-1.089267338
Q8NI27	THOC2_HUMAN	THO complex subunit 2 OS=Homo sapiens GN=THOC2 PE=1 SV=2	0.9985	-1.089267338
P25325	THTM_HUMAN	3-mercaptopyruvate sulfurtransferase OS=Homo sapiens GN=MPST PE=1 SV=3	1	-1.152003093
Q14517	FAT1_HUMAN	Protocadherin Fat 1 OS=Homo sapiens GN=FAT1 PE=1 SV=2	0.9998	-1.152003093
Q16363-2	LAMA4_HUMAN	Isoform 2 of Laminin subunit alpha-4 OS=Homo sapiens GN=LAMA4	0.9984	-1.152003093
Q12849-5	GRSF1_HUMAN	Isoform 2 of G-rich sequence factor 1 OS=Homo sapiens GN=GRSF1	0.994	-1.152003093
F8WAI1	F8WAI1_HUMAN	Immunoglobulin-like and fibronectin type III domain-containing protein 1 OS=Homo sapiens GN=IGFN1	1	-1.184424571
P35908	K22E_HUMAN	Keratin	0.9985	-1.184424571
Q8IVL6-2	P3H3_HUMAN	Isoform 2 of Prolyl 3-hydroxylase 3 OS=Homo sapiens GN=LEPREL2	0.8109	-1.184424571
Q8N1G2	MTR1_HUMAN	Cap-specific mRNA (nucleoside-2'-O-)-methyltransferase 1 OS=Homo sapiens GN=FTSD2 PE=1 SV=1	0.9964	-1.251538767
Q5JTZ9	SYAM_HUMAN	Alanine--tRNA ligase	0.9843	-1.251538767
Q8TEK3-2	DOT1L_HUMAN	Isoform 1 of Histone-Lysine N-methyltransferase	0.9762	-1.286304185
Q14203	DCTN1_HUMAN	Dynactin subunit 1 OS=Homo sapiens GN=DCTN1 PE=1 SV=3	0.9703	-1.321928095
Q5VUUA	ZN318_HUMAN	Zinc finger protein 318 OS=Homo sapiens GN=ZNF318 PE=1 SV=2	0.9989	-1.358453971
Q4VXP2	Q4VXP2_HUMAN	Potassium voltage-gated channel subfamily KQT member 2 OS=Homo sapiens GN=KCNQ2 PE=2 SV=1	0.9993	-1.395928676
O43805	SSNA1_HUMAN	Sjogren syndrome nuclear autoantigen 1 OS=Homo sapiens GN=SSNA1 PE=1 SV=2	0.9974	-1.395928676
P04908	H2A1B_HUMAN	Histone H2A type 1-B/E OS=Homo sapiens GN=HIST1H2AB PE=1 SV=2	1	-1.434402824
Q9Y2G4-1	ANKR6_HUMAN	Isoform 2 of Ankyrin repeat domain-containing protein 6 OS=Homo sapiens GN=ANKRD6	0.9171	-1.434402824
P20648	ATP4A_HUMAN	Potassium-transporting ATPase alpha chain 1 OS=Homo sapiens GN=ATP4A PE=2 SV=5	1	-1.473931188
Q15056-2	IF4H_HUMAN	Isoform Short of Eukaryotic translation initiation factor 4H OS=Homo sapiens GN=EIF4H	0.9813	-1.473931188
B2RTY4-2	MYO9A_HUMAN	Isoform 2 of Unconventional myosin-IXa OS=Homo sapiens GN=MYO9A	0.9577	-1.473931188
Q86V15	CASZ1_HUMAN	Zinc finger protein castor homolog 1 OS=Homo sapiens GN=CASZ1 PE=2 SV=4	0.9996	-1.514573173
Q9NWW4	CD027_HUMAN	UPF0609 protein C4orf27 OS=Homo sapiens GN=C4orf27 PE=1 SV=2	0.9993	-1.514573173
Q8NEM7	FA48A_HUMAN	Protein FAM48A OS=Homo sapiens GN=FAM48A PE=1 SV=2	0.9886	-1.514573173
Q96NP5	Q96NP5_HUMAN	HCG2008157 OS=Homo sapiens GN=hCG_2008157 PE=2 SV=1	0.9659	-1.514573173
P09001	RM03_HUMAN	39S ribosomal protein L3	0.9586	-1.514573173
Q6N043-2	Z280D_HUMAN	Isoform 2 of Zinc finger protein 280D OS=Homo sapiens GN=ZNF280D	0.9951	-1.556393349
Q9NP61	ARFG3_HUMAN	ADP-ribosylation factor GTPase-activating protein 3 OS=Homo sapiens GN=ARFGAP3 PE=1 SV=1	0.9819	-1.556393349
O95983-2	MBD3_HUMAN	Isoform 2 of Methyl-CpG-binding domain protein 3 OS=Homo sapiens GN=MBD3	0.9995	-1.689659879
O15481	MAGB4_HUMAN	Melanoma-associated antigen B4 OS=Homo sapiens GN=MAGEB4 PE=1 SV=1	0.9769	-1.689659879
Q9UPQ7-2	PZRN3_HUMAN	Isoform 2 of E3 ubiquitin-protein ligase PDZRN3 OS=Homo sapiens GN=PDZRN3	0.9761	-1.689659879
P04264	K2C1_HUMAN	Keratin	0.9955	-1.785875195
H7C3E0	H7C3E0_HUMAN	Carboxy-terminal domain RNA polymerase II polypeptide A small phosphatase 1 (Fragment) OS=Homo sapiens	0.9722	-1.785875195
O00602	FCN1_HUMAN	Ficolin-1 OS=Homo sapiens GN=FCN1 PE=1 SV=2	0.9507	-1.836501268
D6RGV7	D6RGV7_HUMAN	Immunoglobulin superfamily member 22 OS=Homo sapiens GN=IGSF22 PE=4 SV=1	0.9679	-1.943416472
Q9BR39-2	JPH2_HUMAN	Isoform 2 of Junctophilin-2 OS=Homo sapiens GN=JPH2	0.9492	-1.943416472
B7ZM82	B7ZM82_HUMAN	Protein Wiz OS=Homo sapiens GN=WIZ PE=2 SV=1	0.9583	-2
Q9Y618-3	NCOR2_HUMAN	Isoform 3 of Nuclear receptor corepressor 2 OS=Homo sapiens GN=NCOR2	0.9037	-2
H3BPPE1	H3BPPE1_HUMAN	Microtubule-actin cross-linking factor 1	0.9999	-2.120294234
Q3MIP1	IPIL2_HUMAN	Inositol 1	0.9962	-2.120294234
Q9H6D7-3	HAUS4_HUMAN	Isoform 3 of HAUS augmin-like complex subunit 4 OS=Homo sapiens GN=HAUS4	0.9887	-2.120294234
Q8IYB8	SUV3_HUMAN	ATP-dependent RNA helicase SUPV3L1	0.9485	-2.120294234
Q9H1H9-2	KIF13A_HUMAN	Isoform 2 of Kinesin-like protein KIF13A OS=Homo sapiens GN=KIF13A	1	-2.184424571
Q9UULM3	YEYS2_HUMAN	YEATS domain-containing protein 2 OS=Homo sapiens GN=YEATS2 PE=1 SV=2	0.9997	-2.184424571
Q9NV56	MRGBP_HUMAN	MRG-binding protein OS=Homo sapiens GN=MRGBP PE=1 SV=1	0.9706	-2.184424571
O15021-1	MAST4_HUMAN	Isoform 5 of Microtubule-associated serine/threonine-protein kinase 4 OS=Homo sapiens GN=MAST4	0.9993	-2.251538767
Q8N302	AGGF1_HUMAN	Angiogenic factor with G patch and FHA domains 1 OS=Homo sapiens GN=AGGF1 PE=1 SV=2	0.9887	-2.251538767
E9PBE3	E9PBE3_HUMAN	Bardet-Biedl syndrome 5 protein OS=Homo sapiens GN=BB55 PE=4 SV=1	0.9818	-2.251538767
Q5JSZ5	PRRC2B_HUMAN	Protein PRRC2B OS=Homo sapiens GN=PRRC2B PE=1 SV=2	0.9693	-2.251538767
HOYMT1	HOYMT1_HUMAN	Talin-2 (Fragment) OS=Homo sapiens GN=TLN2 PE=4 SV=1	0.9419	-2.251538767
P02458-1	CO2A1_HUMAN	Isoform 1 of Collagen alpha-1(II) chain OS=Homo sapiens GN=COL2A1	1	-2.473931188
P20930	FILA_HUMAN	Filaggrin OS=Homo sapiens GN=FLG PE=1 SV=3	0.9999	-2.473931188
Q9P227	RHG23_HUMAN	Rho GTPase-activating protein 23 OS=Homo sapiens GN=ARHGAP23 PE=1 SV=2	0.9999	-2.473931188
O14827	RGRF2_HUMAN	Ras-specific guanine nucleotide-releasing factor 2 OS=Homo sapiens GN=RASGRF2 PE=1 SV=2	0.9914	-2.473931188
Q9BYB0-2	SHANK3_HUMAN	Isoform 2 of SH3 and multiple ankyrin repeat domains protein 3 OS=Homo sapiens GN=SHANK3	0.999	-2.56393349
Q9Y2V7-2	COG6_HUMAN	Isoform 2 of Conserved oligomeric Golgi complex subunit 6 OS=Homo sapiens GN=COG6	0.9807	-2.64385619
Q9Y5B9	SP16H_HUMAN	FACT complex subunit SPT16 OS=Homo sapiens GN=SPT16H PE=1 SV=1	0.9782	-2.64385619
F6QF33	F6QF33_HUMAN	Killer cell immunoglobulin-like receptor 3DL1 OS=Homo sapiens GN=KIR3DL1 PE=4 SV=1	0.9658	-2.64385619
Q9UDT6-2	CLIP2_HUMAN	Isoform 2 of CAP-Gly domain-containing linker protein 2 OS=Homo sapiens GN=CLIP2	0.9959	-2.736965594
P51587	BRCA2_HUMAN	Breast cancer type 2 susceptibility protein OS=Homo sapiens GN=BRCA2 PE=1 SV=2	0.988	-2.736965594
Q5VTQ0-3	TT39B_HUMAN	Isoform 3 of Tetratricopeptide repeat protein 39B OS=Homo sapiens GN=TTC39B	0.9286	-2.736965594
Q14055	CO9A2_HUMAN	Collagen alpha-2(IX) chain OS=Homo sapiens GN=COL9A2 PE=1 SV=2	0.9997	-2.943416472

Q63HN8-4	RN213_HUMAN	Isoform 2 of E3 ubiquitin-protein ligase RNF213 OS=Homo sapiens GN=RNF213	0.928	-2.943416472
Q99062-2	CSF3R_HUMAN	Isoform 2 of Granulocyte colony-stimulating factor receptor OS=Homo sapiens GN=CSF3R	1	-3.058893689
Q08J23-2	NSUN2_HUMAN	Isoform 2 of tRNA (cytosine(34)-C(5))-methyltransferase OS=Homo sapiens GN=NSUN2	0.9997	-3.058893689
Q16254	Q16254	Transcription factor E2F4 OS=Homo sapiens GN=E2F4 PE=1 SV=2	0.9752	-3.058893689
O15381-2	NVL_HUMAN	Isoform 2 of Nuclear valosin-containing protein-like OS=Homo sapiens GN=NVL	0.921	-3.184424571
O75592-2	MYCB2_HUMAN	Isoform 2 of Probable E3 ubiquitin-protein ligase MYCBP2 OS=Homo sapiens GN=MYCBP2	0.9976	-3.321928095
Q5VZE5	NAA35_HUMAN	N-alpha-acetyltransferase 35	0.9969	-3.321928095
Q9Y5T5-2	UBP16_HUMAN	Isoform 2 of Ubiquitin carboxyl-terminal hydrolase 16 OS=Homo sapiens GN=USP16	0.9862	-3.321928095
Q15046-2	SYK_HUMAN	Isoform Mitochondrial of Lysine-tRNA ligase OS=Homo sapiens GN=KARS	0.9648	-3.321928095
Q96L91-2	EP400_HUMAN	Isoform 2 of E1A-binding protein p400 OS=Homo sapiens GN=EP400	0.9489	-3.321928095
Q7Z2K8	GRIN1_HUMAN	G protein-regulated inducer of neurite outgrowth 1 OS=Homo sapiens GN=GPRIN1 PE=1 SV=2	1	-3.473931188
Q99996-2	AKAP9_HUMAN	Isoform 2 of A-kinase anchor protein 9 OS=Homo sapiens GN=AKAP9	1	-3.473931188
O15417-2	TNC18_HUMAN	Isoform 2 of Trinucleotide repeat-containing gene 18 protein OS=Homo sapiens GN=TNRC18	0.9987	-3.473931188
Q13562	NDF1_HUMAN	Neurogenic differentiation factor 1 OS=Homo sapiens GN=NEUROD1 PE=1 SV=3	0.9969	-3.473931188
Q5V706	CE350_HUMAN	Centrosome-associated protein 350 OS=Homo sapiens GN=CEP350 PE=1 SV=1	1	-3.64385619
F8VXG0	F8VXG0_HUMAN	Homeobox protein Hox-B3 OS=Homo sapiens GN=HOXB3 PE=3 SV=1	0.8305	-3.64385619
Q5VWJ9	SNX30_HUMAN	Sorting nexin-30 OS=Homo sapiens GN=SNX30 PE=1 SV=1	0.9637	-3.836501268
Q8IVL1-10	NAV2_HUMAN	Isoform 10 of Neuron navigator 2 OS=Homo sapiens GN=NAV2	0.9889	-4.058893689
P07942	LAMB1_HUMAN	Laminin subunit beta-1 OS=Homo sapiens GN=LAMB1 PE=1 SV=2	0.9974	-4.321928095
C9JLR9	CK095_HUMAN	Uncharacterized protein C11orf95 OS=Homo sapiens GN=C11orf95 PE=2 SV=1	0.9794	-4.321928095
Q92618	ZN516_HUMAN	Zinc finger protein 516 OS=Homo sapiens GN=ZNF516 PE=1 SV=1	0.9626	-4.321928095
Q8N535	CB052_HUMAN	Putative uncharacterized protein encoded by LINC00471 OS=Homo sapiens GN=LINC00471 PE=5 SV=1	0.9956	-4.64385619
Q9BRK4	LZTS2_HUMAN	Leucine zipper putative tumor suppressor 2 OS=Homo sapiens GN=LZTS2 PE=1 SV=2	0.9612	-4.64385619
O00423-3	EMAL1_HUMAN	Isoform 3 of Echinoderm microtubule-associated protein-like 1 OS=Homo sapiens GN=EML1	0.9386	-4.64385619
Q7KZ85	SPT6_HUMAN	Transcription elongation factor SPT6 OS=Homo sapiens GN=SPT6H PE=1 SV=2	0.9985	-5.058893689
Q02539	H11_HUMAN	Histone H1.1 OS=Homo sapiens GN=HIST1H1A PE=1 SV=3	0.9974	-5.058893689
A6NMZ7	CO6A6_HUMAN	Collagen alpha-6(VI) chain OS=Homo sapiens GN=COL6A6 PE=1 SV=2	0.9985	-5.64385619
Q9BX12	CCG6_HUMAN	Voltage-dependent calcium channel gamma-6 subunit OS=Homo sapiens GN=CACNG6 PE=2 SV=1	0.9451	-5.64385619
O75475	PSIP1_HUMAN	PC4 and SFRS1-interacting protein OS=Homo sapiens GN=PSIP1 PE=1 SV=1	0.9985	-6.64385619
Q6ZN55-2	ZN574_HUMAN	Isoform 2 of Zinc finger protein 574 OS=Homo sapiens GN=ZNF574	0.9951	-6.64385619
P36957	ODO2_HUMAN	Dihydrodipolyllysine-residue succinyltransferase component of 2-oxoglutarate dehydrogenase complex	0.9728	-6.64385619
Q92736-2	RYR2_HUMAN	Isoform 2 of Ryanodine receptor 2 OS=Homo sapiens GN=RYR2	0.8284	-6.64385619
Q659C4-2	LAR1B_HUMAN	Isoform 2 of La-related protein 1B OS=Homo sapiens GN=LARP1B	0.9909	9.964340868
Q14562	DHX8_HUMAN	ATP-dependent RNA helicase DHX8 OS=Homo sapiens GN=DHX8 PE=1 SV=1	0.9465	6.453517579
I3L404	I3L404_HUMAN	40S ribosomal protein S2 (Fragment) OS=Homo sapiens GN=RPS2 PE=3 SV=1	0.9596	6.036942845
A0AVI2	FR1L5_HUMAN	Fer-1-like protein 5 OS=Homo sapiens GN=FR1L5 PE=2 SV=2	0.9961	5.283921772
Q9UL58	ZN215_HUMAN	Zinc finger protein 215 OS=Homo sapiens GN=ZNF215 PE=2 SV=2	0.9299	5.098032083
Q86TB3	ALPK2_HUMAN	Coiled-coil protein kinase 2 OS=Homo sapiens GN=ALPK2 PE=1 SV=3	0.9663	4.307428525
Q8NDH2	CC168_HUMAN	Coiled-coil domain-containing protein 168 OS=Homo sapiens GN=CCDC168 PE=2 SV=2	0.9685	4.041768665
Q8IUD2-2	RB612_HUMAN	Isoform 2 of ELKS/Rab6-interacting/CAST family member 1 OS=Homo sapiens GN=ERC1	0.8806	3.959770155
P25391	LAMA1_HUMAN	Laminin subunit alpha-1 OS=Homo sapiens GN=LAMA1 PE=1 SV=2	0.9994	3.014355293
F5GWR6	F5GWR6_HUMAN	Homeobox protein cut-like 2 (Fragment) OS=Homo sapiens GN=CUX2 PE=4 SV=2	0.7888	2.969195503
Q86XW9-2	TXND6_HUMAN	Isoform 2 of Thioredoxin domain-containing protein 6 OS=Homo sapiens GN=NME9	0.9593	2.965322548
Q6E0U4-2	DMKN_HUMAN	Isoform 2 of Dermokine OS=Homo sapiens GN=DMKN	0.9899	2.944858446
Q69VZ2	T200B_HUMAN	Transmembrane protein 200B OS=Homo sapiens GN=TMEM200B PE=2 SV=1	0.9671	2.769771739
J3KT41	J3KT41_HUMAN	SRC kinase-signaling inhibitor 1 (Fragment) OS=Homo sapiens GN=SRCIN1 PE=4 SV=1	0.863	2.50589093
Q9NY28	GALT8_HUMAN	Probable polypeptide N-acetylglucosaminyltransferase 8 OS=Homo sapiens GN=GALNT8 PE=2 SV=1	0.9535	2.480265122
Q96PE2	ARHGH_HUMAN	Rho guanine nucleotide exchange factor 17 OS=Homo sapiens GN=ARHGEF17 PE=1 SV=1	0.9943	2.451540833
Q9Y6X6	MYO16_HUMAN	Unconventional myosin-XVI OS=Homo sapiens GN=MYO16 PE=1 SV=3	1	2.13093087
Q5T3J0-3	GPTC4_HUMAN	Isoform 3 of G patch domain-containing protein 4 OS=Homo sapiens GN=GPATCH4	0.9423	2.010779839
P54259	ATN1_HUMAN	Atrophin-1 OS=Homo sapiens GN=ATN1 PE=1 SV=3	0.923	1.956056652
Q9BQ69	MACD1_HUMAN	O-acetyl-ADP-ribose deacetylase MACROD1 OS=Homo sapiens GN=MACROD1 PE=1 SV=2	0.9861	1.90303827
Q9NYU1	UGGT2_HUMAN	UDP-glucose:glycoprotein glucosyltransferase 2 OS=Homo sapiens GN=UGGT2 PE=1 SV=4	0.9621	1.89917563
Q7LS91	DOK3_HUMAN	Docking protein 3 OS=Homo sapiens GN=DOK3 PE=1 SV=2	0.9677	1.879705766
Q2MKA7-2	RSPO1_HUMAN	Isoform 2 of R-spondin-1 OS=Homo sapiens GN=RSPO1	0.9139	1.761285273
P40937-2	RFC5_HUMAN	Isoform 2 of Replication factor C subunit 5 OS=Homo sapiens GN=RFC5	0.991	1.459431619
Q5VST9-2	OBSCN_HUMAN	Isoform 2 of Obscurin OS=Homo sapiens GN=OBSCN	1	1.350497247
E9PJL5	E9PJL5_HUMAN	Uncharacterized protein C12orf55 OS=Homo sapiens GN=C12orf55 PE=4 SV=1	0.9996	1.316145742
Q9NTK5-2	OLA1_HUMAN	Isoform 2 of Obg-like ATPase 1 OS=Homo sapiens GN=OLA1	0.8459	1.269033146
Q8IVF2-3	AHNAK2_HUMAN	Isoform 3 of Protein AHNAK2 OS=Homo sapiens GN=AHNAK2	0.9805	1.244887059
Q8TDG2	ACTT1_HUMAN	Actin-related protein T1 OS=Homo sapiens GN=ACTRT1 PE=2 SV=2	0.977	1.220329955
Q05C16	LRC63_HUMAN	Leucine-rich repeat-containing protein 63 OS=Homo sapiens GN=LRR63 PE=2 SV=2	0.9329	1.214124805
P02452	CO1A1_HUMAN	Collagen alpha-1(I) chain OS=Homo sapiens GN=COL1A1 PE=1 SV=5	0.9619	1.176322773
Q8TDX9-2	PK1L1_HUMAN	Isoform 2 of Polycystic kidney disease protein 1-like 1 OS=Homo sapiens GN=PKD1L1	0.9671	1.14404637
Q9P1Z9-2	C1174_HUMAN	Isoform 2 of Uncharacterized protein C9orf174 OS=Homo sapiens GN=C9orf174	0.8997	1.137503524
Q9Y3T9	NOC2L_HUMAN	Nucleolar complex protein 2 homolog OS=Homo sapiens GN=NOC2L PE=1 SV=4	0.9631	1.137503524
Q9NS37	ZHANG_HUMAN	CREB/ATF bZIP transcription factor OS=Homo sapiens GN=CREBZF PE=1 SV=2	0.9922	1.042644337
Q96L96	ALPK3_HUMAN	Alpha-protein kinase 3 OS=Homo sapiens GN=ALPK3 PE=2 SV=2	0.985	1.021479727
H7C4U7	H7C4U7_HUMAN	Membrane-associated guanylate kinase	0.9587	0.970853654
P05997	CO5A2_HUMAN	Collagen alpha-2(V) chain OS=Homo sapiens GN=COL5A2 PE=1 SV=3	0.9781	0.963474124
Q70Z35	PREX2_HUMAN	Phosphatidylinositol 3	0.9581	0.782408565
P54802	ANAG_HUMAN	Alpha-N-acetylglucosaminidase OS=Homo sapiens GN=NAGLU PE=1 SV=2	0.9377	0.782408565
Q9NYH9	UTP6_HUMAN	U3 small nucleolar RNA-associated protein 6 homolog OS=Homo sapiens GN=UTP6 PE=1 SV=2	0.9292	0.773996325
P52789	HXK2_HUMAN	Hexokinase-2 OS=Homo sapiens GN=HK2 PE=1 SV=2	0.9927	0.739848103
H7C244	H7C244_HUMAN	DDB1- and CUL4-associated factor 17 (Fragment) OS=Homo sapiens GN=DCAF17 PE=4 SV=1	0.8418	0.739848103
F5HQH8	F5HQH8_HUMAN	WD repeat-containing protein 74 OS=Homo sapiens GN=WDR74 PE=4 SV=1	0.8979	0.669026766
P12955-2	PEPD_HUMAN	Isoform 2 of Xaa-Pro dipeptidase OS=Homo sapiens GN=PEPD	1	0.632268215
Q9NWZ5-4	UCKL1_HUMAN	Isoform 4 of Uridine-cytidine kinase-like 1 OS=Homo sapiens GN=UCKL1	0.9511	0.604071324
Q5TF21	SOGA3_HUMAN	Protein SOGA3 OS=Homo sapiens GN=SOGA3 PE=2 SV=1	0.962	0.555816155
P22680	CPTA1_HUMAN	Cholesterol 7-alpha-monooxygenase OS=Homo sapiens GN=CYP7A1 PE=1 SV=2	0.9016	0.545968369
J3KQD6	J3KQD6_HUMAN	PWWP domain-containing protein MUM1 OS=Homo sapiens GN=MUM1 PE=4 SV=1	0.9988	0.526068812
P19623	SPEE_HUMAN	Spermidine synthase OS=Homo sapiens GN=SRM PE=1 SV=1	0.9999	0.495695163
Q9Y5Y6	ST14_HUMAN	Suppressor of tumorigenicity 14 protein OS=Homo sapiens GN=ST14 PE=1 SV=2	0.9593	0.495695163
Q9UBG3	CRNN_HUMAN	Cornulin OS=Homo sapiens GN=CRNN PE=1 SV=1	0.9976	0.485426827
P09960-2	LKHA4_HUMAN	Isoform 2 of Leukotriene A-4 hydrolase OS=Homo sapiens GN=LTA4H	0.978	0.485426827
Q9NTJ3-2	SMC4_HUMAN	Isoform 2 of Structural maintenance of chromosomes protein 4 OS=Homo sapiens GN=SMC4	1	0.475084883
Q9BUL8	PDC10_HUMAN	Programmed cell death protein 10 OS=Homo sapiens GN=PDC10 PE=1 SV=1	0.9998	0.475084883
P43004-2	EAA2_HUMAN	Isoform 2 of Excitatory amino acid transporter 2 OS=Homo sapiens GN=SLC1A2	0.9981	0.454175893
O94903	PROSC_HUMAN	Proline synthase co-transcribed bacterial homolog protein OS=Homo sapiens GN=PROSC PE=1 SV=1	0.9999	0.443606651
P55010	IF5_HUMAN	Eukaryotic translation initiation factor 5 OS=Homo sapiens GN=EIF5 PE=1 SV=2	1	0.40053793
P46459	NSF_HUMAN	Vesicle-fusing ATPase OS=Homo sapiens GN=NSF PE=1 SV=3	0.9997	0.40053793
Q9Y5S9-2	RBM8A_HUMAN	Isoform 2 of RNA-binding protein 8A OS=Homo sapiens GN=RBM8A	0.9997	0.40053793

A8MYV0	DCDC2C_HUMAN	Doublecortin domain-containing protein 2C OS=Homo sapiens GN=DCDC2C PE=4 SV=3	0.9838	0.40053793
A9YTQ3-2	AHRR_HUMAN	Isoform 2 of Aryl hydrocarbon receptor repressor OS=Homo sapiens GN=AHRR	0.9464	0.40053793
P28290-2	SSFA2_HUMAN	Isoform 2 of Sperm-specific antigen 2 OS=Homo sapiens GN=SSFA2	0.9981	0.389566812
P50225	ST1A1_HUMAN	Sulfotransferase 1A1 OS=Homo sapiens GN=SULT1A1 PE=1 SV=3	0.8414	0.367371066
P40925-2	MDHC_HUMAN	Isoform 2 of Malate dehydrogenase	1	0.344828497
Q14684-2	RRP1B_HUMAN	Isoform 2 of Ribosomal RNA processing protein 1 homolog B OS=Homo sapiens GN=RRP1B	0.9999	0.344828497
Q5VIR6-2	VP53_HUMAN	Isoform 2 of Vacuolar protein sorting-associated protein 53 homolog OS=Homo sapiens GN=VP53	0.9999	0.344828497
Q16881-2	TRXR1_HUMAN	Isoform 2 of Thioredoxin reductase 1	0.9996	0.333423734
P28066	PSA5_HUMAN	Proteasome subunit alpha type-5 OS=Homo sapiens GN=PSMA5 PE=1 SV=3	1	0.298658316
Q5VZU9	Q5VZU9_HUMAN	Tripeptidyl peptidase II OS=Homo sapiens GN=TPP2 PE=2 SV=1	0.9999	0.298658316
P31153	METK2_HUMAN	S-adenosylmethionine synthase isoform type-2 OS=Homo sapiens GN=MAT2A PE=1 SV=1	0.9993	0.298658316
B4E283	B4E283_HUMAN	Apoptosis inhibitor 5 OS=Homo sapiens GN=API5 PE=2 SV=1	0.9979	0.298658316
F8VWV3	F8VWV3_HUMAN	Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0.989	0.298658316
Q9Y238-2	DLEC1_HUMAN	Isoform 2 of Deleted in lung and esophageal cancer protein 1 OS=Homo sapiens GN=DLEC1	0.8871	0.286881148
Q16352	AINX_HUMAN	Alpha-internexin OS=Homo sapiens GN=INA PE=1 SV=2	1	0.275007047
Q6P2E9	EDC4_HUMAN	Enhancer of mRNA-decapping protein 4 OS=Homo sapiens GN=EDC4 PE=1 SV=1	1	0.275007047
Q9BXJ9	NAA15_HUMAN	N-alpha-acetyltransferase 15	1	0.263034406
P38432	COIL_HUMAN	Coilin OS=Homo sapiens GN=COIL PE=1 SV=1	0.9475	0.263034406
A2AB15	A2AB15_HUMAN	DEAD/H (Asp-Glu-Ala-Asp/His) box polypeptide 16 (Fragment) OS=Homo sapiens GN=DHX16 PE=4 SV=	0.9305	0.263034406
Q14669-2	TRIP_C_HUMAN	Isoform 2 of E3 ubiquitin-protein ligase TRIP12 OS=Homo sapiens GN=TRIP12	0.95	0.250961574
P57088	TMM33_HUMAN	Transmembrane protein 33 OS=Homo sapiens GN=TMEM33 PE=1 SV=2	0.8531	0.250961574
P42126-2	ECI1_HUMAN	Isoform 2 of Enoyl-CoA delta isomerase 1	1	0.23878686
P01111	RASN_HUMAN	GTPase NRas OS=Homo sapiens GN=NRAS PE=1 SV=1	0.9998	0.23878686
Q9NVP1	DDX18_HUMAN	ATP-dependent RNA helicase DDX18 OS=Homo sapiens GN=DDX18 PE=1 SV=2	0.9997	0.23878686
Q99583	MNT_HUMAN	Max-binding protein MNT OS=Homo sapiens GN=MNT PE=1 SV=1	0.9553	0.22650853
Q71H61	ILDR2_HUMAN	Immunoglobulin-like domain-containing receptor 2 OS=Homo sapiens GN=ILDR2 PE=2 SV=1	0.9539	0.22650853
Q9BQ39	DDX50_HUMAN	ATP-dependent RNA helicase DDX50 OS=Homo sapiens GN=DDX50 PE=1 SV=1	0.8465	0.22650853
P35637-2	FUS_HUMAN	Isoform Short of RNA-binding protein FUS OS=Homo sapiens GN=FUS	1	0.214124805
Q8N684-2	CPSF7_HUMAN	Isoform 2 of Cleavage and polyadenylation specificity factor subunit 7 OS=Homo sapiens GN=CPSF7	1	0.214124805
Q96NU0	CNT3B_HUMAN	Contactin-associated protein-like 3B OS=Homo sapiens GN=CNTNAP3B PE=2 SV=2	0.958	0.201633861
Q9NXG0-2	CNTLN_HUMAN	Isoform 2 of Centelin OS=Homo sapiens GN=CNTLN	0.9529	0.201633861
O75822	EIF3J_HUMAN	Eukaryotic translation initiation factor 3 subunit J OS=Homo sapiens GN=EIF3J PE=1 SV=2	0.9073	0.201633861
Q99426	TBCB_HUMAN	Tubulin-folding cofactor B OS=Homo sapiens GN=TBCB PE=1 SV=2	1	0.189033824
P46940	IQGA1_HUMAN	Ras GTPase-activating-like protein IQGAP1 OS=Homo sapiens GN=IQGAP1 PE=1 SV=1	1	0.189033824
Q12852	M3K12_HUMAN	Mitogen-activated protein kinase kinase kinase 12 OS=Homo sapiens GN=MAP3K12 PE=1 SV=2	0.9607	0.189033824
Q01082-2	SPTB2_HUMAN	Isoform Short of Spectrin beta chain	1	0.176322773
Q3ZCQ8-2	TIM50_HUMAN	Isoform 2 of Mitochondrial import inner membrane translocase subunit TIM50 OS=Homo sapiens GN=TI	0.9995	0.176322773
Q14151	SAFB2_HUMAN	Scaffold attachment factor B2 OS=Homo sapiens GN=SAFB2 PE=1 SV=1	0.9906	0.176322773
Q86519-2	CEL_HUMAN	Isoform 2 of Protein CEI OS=Homo sapiens GN=C5orf38	0.9639	0.176322773
P08579	RU2B_HUMAN	U2 small nuclear ribonucleoprotein B'' OS=Homo sapiens GN=SNRPB2 PE=1 SV=1	1	0.163498732
H0YKF0	H0YKF0_HUMAN	Electron transfer flavoprotein subunit alpha	0.949	0.150559677
Q13428-3	TCOF_HUMAN	Isoform 3 of Treacle protein OS=Homo sapiens GN=TCOF1	1	0.150559677
Q9Y310	RTCB_HUMAN	tRNA-splicing ligase RtcB homolog OS=Homo sapiens GN=C22orf28 PE=1 SV=1	0.8764	0.150559677
P38606	VATA_HUMAN	V-type proton ATPase catalytic subunit A OS=Homo sapiens GN=ATP6V1A PE=1 SV=2	1	0.137503524
Q96A49	SYAP1_HUMAN	Synapse-associated protein 1 OS=Homo sapiens GN=SYAP1 PE=1 SV=1	0.9972	0.137503524
P36871	PGM1_HUMAN	Phosphoglucomutase-1 OS=Homo sapiens GN=PGM1 PE=1 SV=3	0.9949	0.137503524
O15042-2	SR140_HUMAN	Isoform 2 of U2 snRNP-associated SURP motif-containing protein OS=Homo sapiens GN=U2SURP	0.9891	0.137503524
Q96N67-2	DOCK7_HUMAN	Isoform 2 of Dedicator of cytokinesis protein 7 OS=Homo sapiens GN=DOCK7	1	0.124328135
P0CG47	UBB_HUMAN	Polyubiquitin-B OS=Homo sapiens GN=UBB PE=1 SV=1	1	0.124328135
P49756	RBM25_HUMAN	RNA-binding protein 25 OS=Homo sapiens GN=RBM25 PE=1 SV=3	1	0.124328135
Q9UI26-2	IPO11_HUMAN	Isoform 2 of Importin-11 OS=Homo sapiens GN=IPO11	1	0.124328135
P62873	GBB1_HUMAN	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV	0.9995	0.111031312
Q9H583	HEAT1_HUMAN	HEAT repeat-containing protein 1 OS=Homo sapiens GN=HEATR1 PE=1 SV=3	1	0.111031312
Q6ZU80	CE128_HUMAN	Centrosomal protein of 128 kDa OS=Homo sapiens GN=CEP128 PE=1 SV=2	0.9996	0.111031312
Q96176-5	MMS19_HUMAN	Isoform 4 of MMS19 nucleotide excision repair protein homolog OS=Homo sapiens GN=MMS19	0.9997	0.111031312
P48029-2	SC6A8_HUMAN	Isoform 2 of Sodium- and chloride-dependent creatine transporter 1 OS=Homo sapiens GN=SLC6A8	0.8973	0.111031312
O00541-2	PESC_HUMAN	Isoform 2 of Pescadillo homolog OS=Homo sapiens GN=PES1	1	0.097610797
P17858-2	K6PL_HUMAN	Isoform 2 of 6-phosphofructokinase	1	0.097610797
Q8IY67-2	RAVR1_HUMAN	Isoform 2 of Ribonucleoprotein PTB-binding 1 OS=Homo sapiens GN=RAVER1	0.9829	0.097610797
Q9Y2X3	NOP58_HUMAN	Nucleolar protein 58 OS=Homo sapiens GN=NOP58 PE=1 SV=1	1	0.084064265
Q9Y5L0-1	TNPO3_HUMAN	Isoform 1 of Transportin-3 OS=Homo sapiens GN=TNPO3	1	0.084064265
Q8IVF4	DYH10_HUMAN	Dynein heavy chain 10	0.9144	0.084064265
P78527	PRKDC_HUMAN	DNA-dependent protein kinase catalytic subunit OS=Homo sapiens GN=PRKDC PE=1 SV=3	1	0.084064265
Q9UHB9	SRP68_HUMAN	Signal recognition particle 68 kDa protein OS=Homo sapiens GN=SRP68 PE=1 SV=2	0.9998	0.084064265
O14617-2	AP3D1_HUMAN	Isoform 2 of AP-3 complex subunit delta-1 OS=Homo sapiens GN=AP3D1	0.9834	0.084064265
Q6KB66-2	K2C80_HUMAN	Isoform 2 of Keratin	0.9632	0.084064265
P48052	CBPA2_HUMAN	Carboxypeptidase A2 OS=Homo sapiens GN=CPA2 PE=1 SV=3	0.951	0.084064265
Q5SXM2	SNPC4_HUMAN	snRNA-activating protein complex subunit 4 OS=Homo sapiens GN=SNAPC4 PE=1 SV=1	0.9862	0.070389328
O15260-2	SURF4_HUMAN	Isoform 2 of Surfeit locus protein 4 OS=Homo sapiens GN=SURF4	1	0.056583528
Q92841-1	DDX17_HUMAN	Isoform 2 of Probable ATP-dependent RNA helicase DDX17 OS=Homo sapiens GN=DDX17	0.9992	0.056583528
P15153	RAC2_HUMAN	Ras-related C3 botulinum toxin substrate 2 OS=Homo sapiens GN=RAC2 PE=1 SV=1	0.9991	0.056583528
P35221	CTNA1_HUMAN	Catenin alpha-1 OS=Homo sapiens GN=CTNNA1 PE=1 SV=1	0.9973	0.056583528
O43598	RCL_HUMAN	Deoxyribonucleoside 5'-monophosphate N-glycosidase OS=Homo sapiens GN=RCL PE=1 SV=1	0.9886	0.056583528
P07954-2	FUMH_HUMAN	Isoform Cytoplasmic of Fumarate hydratase	0.9793	0.056583528
Q867C9-3	MYPN_HUMAN	Isoform 3 of Myopalladin OS=Homo sapiens GN=MYPN	0.9362	0.056583528
O00178	GTPB1_HUMAN	GTP-binding protein 1 OS=Homo sapiens GN=GTPBP1 PE=1 SV=3	0.9999	0.042644337
P56537	IF6_HUMAN	Eukaryotic translation initiation factor 6 OS=Homo sapiens GN=EIF6 PE=1 SV=1	0.9988	0.028569152
P06730-2	IF4E_HUMAN	Isoform 2 of Eukaryotic translation initiation factor 4E OS=Homo sapiens GN=EIF4E	0.9977	0.028569152
Q5HY18	RABL3_HUMAN	Rab-like protein 3 OS=Homo sapiens GN=RABL3 PE=1 SV=1	0.9966	0.028569152
Q9BW60	ELOV1_HUMAN	Elongation of very long chain fatty acids protein 1 OS=Homo sapiens GN=ELOVL1 PE=1 SV=1	0.8719	0.028569152
P43487	RANG_HUMAN	Ran-specific GTPase-activating protein OS=Homo sapiens GN=RANBP1 PE=1 SV=1	1	0.014355293
O43707	ACTN4_HUMAN	Alpha-actinin-4 OS=Homo sapiens GN=ACTN4 PE=1 SV=2	1	0.014355293
E9PPF2	E9PPF2_HUMAN	Non-receptor tyrosine-protein kinase TYK2 OS=Homo sapiens GN=TYK2 PE=4 SV=1	0.9896	0.014355293
Q9NYU2-2	UGGG1_HUMAN	Isoform 2 of UDP-glucose:glycoprotein glucosyltransferase 1 OS=Homo sapiens GN=UGGT1	1	0
P46977	STT3A_HUMAN	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit STT3A OS=Homo sapiens GN=ST	1	0
P49368	TCPG_HUMAN	T-complex protein 1 subunit gamma OS=Homo sapiens GN=CCT3 PE=1 SV=4	1	0
Q6PKG0-3	LARP1_HUMAN	Isoform 2 of La-related protein 1 OS=Homo sapiens GN=LARP1	1	0
Q9H3U1-2	UN45A_HUMAN	Isoform 2 of Protein unc-45 homolog A OS=Homo sapiens GN=UNC45A	1	0
H3BPJ5	H3BPJ5_HUMAN	Synaptogyrin-3 (Fragment) OS=Homo sapiens GN=SYNGR3 PE=4 SV=1	0.9989	0
P28288-2	ABCD3_HUMAN	Isoform 2 of ATP-binding cassette sub-family D member 3 OS=Homo sapiens GN=ABCD3	0.9943	0
Q6UYE1-2	LEU7_HUMAN	Isoform 2 of Leukemia-associated protein 7 OS=Homo sapiens GN=DLEU7	0.9359	0
O43264	ZW10_HUMAN	Centromere/kinetochore protein zw10 homolog OS=Homo sapiens GN=ZW10 PE=1 SV=3	1	-0.01449957

P35914	HMGCL_HUMAN	Hydroxymethylglutaryl-CoA lyase	1	-0.01449957
P52701	MSH6_HUMAN	DNA mismatch repair protein Msh6 OS=Homo sapiens GN=MSH6 PE=1 SV=2	1	-0.01449957
E7ENF9	E7ENF9_HUMAN	Isochorismatase domain-containing protein 1 (Fragment) OS=Homo sapiens GN=ISOC1 PE=4 SV=1	1	-0.01449957
H0Y2V6	H0Y2V6_HUMAN	Centrosomal protein of 170 kDa (Fragment) OS=Homo sapiens GN=CEP170 PE=4 SV=1	0.9997	-0.01449957
Q9UHX1-2	PUF60_HUMAN	Isoform 2 of Poly(U)-binding-splicing factor PUF60 OS=Homo sapiens GN=PUF60	0.9982	-0.01449957
O43719	HTSF1_HUMAN	HIV Tat-specific factor 1 OS=Homo sapiens GN=HTATSF1 PE=1 SV=1	0.9939	-0.01449957
O94973-2	AP2A2_HUMAN	Isoform 2 of AP-2 complex subunit alpha-2 OS=Homo sapiens GN=AP2A2	0.976	-0.01449957
Q5T0Z8	CF132_HUMAN	Uncharacterized protein C6orf132 OS=Homo sapiens GN=C6orf132 PE=1 SV=4	0.9609	-0.01449957
P49454	CENPF_HUMAN	Centromere protein F OS=Homo sapiens GN=CENPF PE=1 SV=2	0.957	-0.01449957
O95433	AHSA1_HUMAN	Activator of 90 kDa heat shock protein ATPase homolog 1 OS=Homo sapiens GN=AHSA1 PE=1 SV=1	1	-0.029146346
Q9NPD3	EXOS4_HUMAN	Exosome complex component RRP41 OS=Homo sapiens GN=EXOSC4 PE=1 SV=3	1	-0.029146346
P23919	KTHY_HUMAN	Thymidylate kinase OS=Homo sapiens GN=DTYMK PE=1 SV=4	0.9976	-0.029146346
O96000	NDUBA_HUMAN	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=Homo sapiens GN=NDUBF10 PE=	0.9968	-0.029146346
Q8IYW5	RN168_HUMAN	E3 ubiquitin-protein ligase RNF168 OS=Homo sapiens GN=RNF168 PE=1 SV=1	0.9941	-0.029146346
Q8WV44-2	TR141_HUMAN	Isoform 2 of E3 ubiquitin-protein ligase TRIM41 OS=Homo sapiens GN=TRIM41	0.9724	-0.029146346
Q5JRA6-2	MIA3_HUMAN	Isoform 2 of Melanoma inhibitory activity protein 3 OS=Homo sapiens GN=MIA3	1	-0.043943348
Q9NPR9	GP108_HUMAN	Protein GPR108 OS=Homo sapiens GN=GPR108 PE=2 SV=3	0.9949	-0.043943348
P50454	SERPH_HUMAN	Serpin H1 OS=Homo sapiens GN=SERPINH1 PE=1 SV=2	0.9899	-0.043943348
O95934	SPF27_HUMAN	Pre-mRNA-splicing factor SPF27 OS=Homo sapiens GN=BCAS2 PE=1 SV=1	0.9698	-0.043943348
O94826	TOM70_HUMAN	Mitochondrial import receptor subunit TOM70 OS=Homo sapiens GN=TOMM70A PE=1 SV=1	1	-0.058893689
P04179-2	SODM_HUMAN	Isoform 2 of Superoxide dismutase [Mn]	0.9999	-0.058893689
P19388	RPAB1_HUMAN	DNA-directed RNA polymerases I	0.9999	-0.058893689
Q60832	DKC1_HUMAN	H/ACA ribonucleoprotein complex subunit 4 OS=Homo sapiens GN=DKC1 PE=1 SV=3	0.9999	-0.058893689
Q06203	PUR1_HUMAN	Amidophosphoribosyltransferase OS=Homo sapiens GN=PPAT PE=1 SV=1	0.9992	-0.058893689
P09661	RU2A_HUMAN	U2 small nuclear ribonucleoprotein A' OS=Homo sapiens GN=SNRPA1 PE=1 SV=2	1	-0.074005581
O95292	VAPB_HUMAN	Vesicle-associated membrane protein-associated protein B/C OS=Homo sapiens GN=VAPB PE=1 SV=3	1	-0.074005581
G3V3W4	G3V3W4_HUMAN	Proteasome subunit alpha type-3 OS=Homo sapiens GN=PSMA3 PE=4 SV=1	0.809	-0.074005581
Q7L014	DDX46_HUMAN	Probable ATP-dependent RNA helicase DDX46 OS=Homo sapiens GN=DDX46 PE=1 SV=2	1	-0.074005581
Q9NR28-2	DBLOH_HUMAN	Isoform 2 of Diablo homolog	1	-0.074005581
P58107	EPIPL_HUMAN	Epiplakin OS=Homo sapiens GN=EPPK1 PE=1 SV=2	0.9956	-0.074005581
O60610-2	DIAP1_HUMAN	Isoform 2 of Protein diaphanous homolog 1 OS=Homo sapiens GN=DIAPH1	0.9878	-0.074005581
Q15653-2	IKBB_HUMAN	Isoform 2 of NF-kappa-B inhibitor beta OS=Homo sapiens GN=NFKBIB	0.9648	-0.074005581
P05198	IF2A_HUMAN	Eukaryotic translation initiation factor 2 subunit 1 OS=Homo sapiens GN=EIF2S1 PE=1 SV=3	0.846	-0.074005581
Q13642-1	FHL1_HUMAN	Isoform 1 of Four and a half LIM domains protein 1 OS=Homo sapiens GN=FHL1	1	-0.089267338
Q9NZ01-2	TECR_HUMAN	Isoform 2 of Trans-2	1	-0.089267338
P46109	CRKL_HUMAN	Crk-like protein OS=Homo sapiens GN=CRKL PE=1 SV=1	0.9987	-0.089267338
Q8WW12-2	PCNP_HUMAN	Isoform 2 of PEST proteolytic signal-containing nuclear protein OS=Homo sapiens GN=PCNP	0.9866	-0.089267338
Q8IZ81	ELMD2_HUMAN	ELMO domain-containing protein 2 OS=Homo sapiens GN=ELMOD2 PE=1 SV=1	0.8824	-0.089267338
Q9NQW6-2	ANLN_HUMAN	Isoform 2 of Actin-binding protein anillin OS=Homo sapiens GN=ANLN	0.8363	-0.104697379
Q15020-2	SART3_HUMAN	Isoform 2 of Squamous cell carcinoma antigen recognized by T-cells 3 OS=Homo sapiens GN=SART3	1	-0.120294234
O60762	DPM1_HUMAN	Dolichol-phosphate mannosyltransferase OS=Homo sapiens GN=DPM1 PE=1 SV=1	1	-0.120294234
P00568	KAD1_HUMAN	Adenylyl kinase isoenzyme 1 OS=Homo sapiens GN=AK1 PE=1 SV=3	1	-0.120294234
P08237	K6PF_HUMAN	6-phosphofructokinase	1	-0.120294234
P52272	HNRPM_HUMAN	Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens GN=HNRNPM PE=1 SV=3	1	-0.120294234
A8MXQ7	YH010_HUMAN	Putative IQ motif and ankyrin repeat domain-containing protein LOC642574 OS=Homo sapiens PE=5 SV	0.9961	-0.120294234
E9PBQ3	E9PBQ3_HUMAN	Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0.9912	-0.120294234
Q9P2D3-2	HTRS5_HUMAN	Isoform 2 of HEAT repeat-containing protein 5B OS=Homo sapiens GN=HEATR5B	0.989	-0.120294234
Q9HOH3	ENC2_HUMAN	Ectoderm-neural cortex protein 2 OS=Homo sapiens GN=KLHL25 PE=2 SV=1	0.8885	-0.120294234
P51659	DHB4_HUMAN	Peroxisomal multifunctional enzyme type 2 OS=Homo sapiens GN=HSD17B4 PE=1 SV=3	0.8845	-0.120294234
P28070	PSB4_HUMAN	Proteasome subunit beta type-4 OS=Homo sapiens GN=PSMB4 PE=1 SV=4	1	-0.13606155
P48147	PPCE_HUMAN	Prolyl endopeptidase OS=Homo sapiens GN=PREP PE=1 SV=2	1	-0.13606155
Q13126-2	MTAP_HUMAN	Isoform 2 of S-methyl-5'-thioadenosine phosphorylase OS=Homo sapiens GN=MTAP	1	-0.13606155
Q9Y285	SYFA_HUMAN	Phenylalanine--tRNA ligase alpha subunit OS=Homo sapiens GN=FARSA PE=1 SV=3	0.9999	-0.13606155
Q96AC1-2	FERM2_HUMAN	Isoform 2 of Fermitin family homolog 2 OS=Homo sapiens GN=FERMT2	0.9996	-0.13606155
Q53EL6	PDCC4_HUMAN	Programmed cell death protein 4 OS=Homo sapiens GN=PDCC4 PE=1 SV=2	0.9973	-0.13606155
O95071	UBR5_HUMAN	E3 ubiquitin-protein ligase UBR5 OS=Homo sapiens GN=UBR5 PE=1 SV=2	0.9906	-0.13606155
Q9NQJ5	PAK6_HUMAN	Serine/threonine-protein kinase PAK 6 OS=Homo sapiens GN=PAK6 PE=1 SV=1	0.985	-0.13606155
P61599	NAA20_HUMAN	N-alpha-acetyltransferase 20 OS=Homo sapiens GN=NAA20 PE=1 SV=1	0.981	-0.13606155
P62829	RL23_HUMAN	60S ribosomal protein L23 OS=Homo sapiens GN=RPL23 PE=1 SV=1	1	-0.152003093
Q9NTK5	OLA1_HUMAN	Obg-like ATPase 1 OS=Homo sapiens GN=OLA1 PE=1 SV=2	1	-0.152003093
Q92575	UBXN4_HUMAN	UBX domain-containing protein 4 OS=Homo sapiens GN=UBXN4 PE=1 SV=2	0.9969	-0.152003093
P33316-2	DUT_HUMAN	Isoform 2 of Deoxyuridine 5'-triphosphate nucleotidohydrolase	1	-0.168122759
P67936-2	TPM4_HUMAN	Isoform 2 of Tropomyosin alpha-4 chain OS=Homo sapiens GN=TPM4	0.9998	-0.168122759
Q14353	GAMT_HUMAN	Guanidinoacetate N-methyltransferase OS=Homo sapiens GN=GAMT PE=1 SV=1	0.9882	-0.168122759
Q9NTN9-2	SEM4G_HUMAN	Isoform 2 of Semaphorin-4G OS=Homo sapiens GN=SEMA4G	0.9193	-0.168122759
P26038	MOES_HUMAN	Moesin OS=Homo sapiens GN=MSN PE=1 SV=3	1	-0.184424571
P46379-2	BAG6_HUMAN	Isoform 2 of Large proline-rich protein BAG6 OS=Homo sapiens GN=BAG6	1	-0.184424571
Q15393	SF3B3_HUMAN	Splicing factor 3B subunit 3 OS=Homo sapiens GN=SF3B3 PE=1 SV=4	1	-0.184424571
Q8N1F7	NUP93_HUMAN	Nuclear pore complex protein Nup93 OS=Homo sapiens GN=NUP93 PE=1 SV=2	1	-0.184424571
O14925	TIM23_HUMAN	Mitochondrial import inner membrane translocase subunit Tim23 OS=Homo sapiens GN=TIMM23 PE=1	0.999	-0.184424571
Q8N7B9	EFCB3_HUMAN	EF-hand calcium-binding domain-containing protein 3 OS=Homo sapiens GN=EFCAB3 PE=1 SV=1	0.974	-0.184424571
Q9NWX5-2	ECHD1_HUMAN	Isoform 2 of Ethylmalonyl-CoA decarboxylase OS=Homo sapiens GN=ECHDC1	0.8136	-0.184424571
P30838	AL3A1_HUMAN	Aldehyde dehydrogenase	0.8008	-0.184424571
Q6Y1H2	HACD2_HUMAN	3-hydroxyacyl-CoA dehydratase 2 OS=Homo sapiens GN=PTPLB PE=1 SV=1	1	-0.200912694
O00461	GOL14_HUMAN	Golgi integral membrane protein 4 OS=Homo sapiens GN=GOLIM4 PE=1 SV=1	0.9984	-0.200912694
O14813	PHX2A_HUMAN	Paired mesoderm homeobox protein 2A OS=Homo sapiens GN=PHOX2A PE=1 SV=2	0.9779	-0.200912694
F5GZT1	F5GZT1_HUMAN	Exocyst complex component 4 OS=Homo sapiens GN=EXOC4 PE=4 SV=1	0.9644	-0.200912694
Q9UID3	VPS51_HUMAN	Vacuolar protein sorting-associated protein 51 homolog OS=Homo sapiens GN=VPS51 PE=1 SV=2	1	-0.217591435
O75116	ROCK2_HUMAN	Rho-associated protein kinase 2 OS=Homo sapiens GN=ROCK2 PE=1 SV=4	0.9352	-0.217591435
Q14676-2	MDC1_HUMAN	Isoform 2 of Mediator of DNA damage checkpoint protein 1 OS=Homo sapiens GN=MDC1	0.9998	-0.217591435
Q8N1G0	ZN687_HUMAN	Zinc finger protein 687 OS=Homo sapiens GN=ZNF687 PE=1 SV=1	0.9997	-0.217591435
Q9H330	TM245_HUMAN	Transmembrane protein 245 OS=Homo sapiens GN=TMEM245 PE=1 SV=2	0.9476	-0.217591435
O15523	DDX3Y_HUMAN	ATP-dependent RNA helicase DDX3Y OS=Homo sapiens GN=DDX3Y PE=1 SV=2	0.9307	-0.234465254
P00387-2	NB5R3_HUMAN	Isoform 2 of NADH-cytochrome b5 reductase 3 OS=Homo sapiens GN=CYB5R3	1	-0.234465254
Q8TAQ2-2	SMRCC2_HUMAN	Isoform 2 of SWI/SNF complex subunit SMARCC2 OS=Homo sapiens GN=SMARCC2	0.9998	-0.234465254
Q9H7Z7	PGES2_HUMAN	Prostaglandin E synthase 2 OS=Homo sapiens GN=PTGES2 PE=1 SV=1	0.9996	-0.234465254
O15085-2	ARHGB_HUMAN	Isoform 2 of Rho guanine nucleotide exchange factor 11 OS=Homo sapiens GN=ARHGEF11	0.9993	-0.234465254
Q5UIP0-2	RIF1_HUMAN	Isoform 2 of Telomere-associated protein RIF1 OS=Homo sapiens GN=RIF1	0.9982	-0.234465254
Q6YFQ2	CX6B2_HUMAN	Cytochrome c oxidase subunit 6B2 OS=Homo sapiens GN=COX6B2 PE=2 SV=1	0.9863	-0.234465254
Q9Y314	NOSIP_HUMAN	Nitric oxide synthase-interacting protein OS=Homo sapiens GN=NOSIP PE=1 SV=1	0.9718	-0.234465254
O00425	IF2B3_HUMAN	Insulin-like growth factor 2 mRNA-binding protein 3 OS=Homo sapiens GN=IGF2BP3 PE=1 SV=2	1	-0.251538767
O14773-2	TPP1_HUMAN	Isoform 2 of Tripeptidyl-peptidase 1 OS=Homo sapiens GN=TPP1	0.999	-0.251538767

P48681	NEST_HUMAN	Nestin OS=Homo sapiens GN=NES PE=1 SV=2	0.9943	-0.251538767
P42858	HD_HUMAN	Huntingtin OS=Homo sapiens GN=HTT PE=1 SV=2	0.9929	-0.251538767
P06748-2	NPM_HUMAN	Isoform 2 of Nucleophosmin OS=Homo sapiens GN=NPM1	1	-0.268816758
Q8NCW5-2	NNRE_HUMAN	Isoform 2 of NAD(P)H-hydrate epimerase OS=Homo sapiens GN=APOA1BP	1	-0.268816758
Q9Y5Y2	NUBP2_HUMAN	Cytosolic Fe-S cluster assembly factor NUBP2 OS=Homo sapiens GN=NUBP2 PE=1 SV=1	0.9996	-0.268816758
O95359-3	TACC2_HUMAN	Isoform 3 of Transforming acidic coiled-coil-containing protein 2 OS=Homo sapiens GN=TACC2	0.9976	-0.268816758
P00846	ATP6_HUMAN	ATP synthase subunit a OS=Homo sapiens GN=MT-ATP6 PE=1 SV=1	0.9935	-0.268816758
P23588	IF4B_HUMAN	Eukaryotic translation initiation factor 4B OS=Homo sapiens GN=EIF4B PE=1 SV=2	0.9883	-0.268816758
Q68CR1-2	SEL13_HUMAN	Isoform 2 of Protein sel-1 homolog 3 OS=Homo sapiens GN=SEL13	0.9678	-0.268816758
P48741	HSP77_HUMAN	Putative heat shock 70 kDa protein 7 OS=Homo sapiens GN=HSPA7 PE=5 SV=2	0.9975	-0.286304185
Q93009	UBP7_HUMAN	Ubiquitin carboxyl-terminal hydrolase 7 OS=Homo sapiens GN=USP7 PE=1 SV=2	1	-0.286304185
G3XAB4	G3XAB4_HUMAN	Acetyl-CoA acetyltransferase	1	-0.286304185
Q9HQ00	FA49A_HUMAN	Protein FAM49A OS=Homo sapiens GN=FAM49A PE=1 SV=1	0.9996	-0.286304185
P59998	ARPC4_HUMAN	Actin-related protein 2/3 complex subunit 4 OS=Homo sapiens GN=ARPC4 PE=1 SV=3	0.9993	-0.286304185
Q9NVU7-2	SDA1_HUMAN	Isoform 2 of Protein SDA1 homolog OS=Homo sapiens GN=SDAD1	0.9437	-0.286304185
Q9NRG7-2	D39U1_HUMAN	Isoform 2 of Epimerase family protein SDR39U1 OS=Homo sapiens GN=SDR39U1	0.8994	-0.286304185
Q3L8U1-2	CHD9_HUMAN	Isoform 2 of Chromodomain-helicase-DNA-binding protein 9 OS=Homo sapiens GN=CHD9	0.9916	-0.304006187
Q5VYK3	ECM29_HUMAN	Proteasome-associated protein ECM29 homolog OS=Homo sapiens GN=ECM29 PE=1 SV=2	1	-0.304006187
Q8WYA6-2	CTBL1_HUMAN	Isoform 2 of Beta-catenin-like protein 1 OS=Homo sapiens GN=CTNBL1	0.9912	-0.304006187
F2Z2N4	F2Z2N4_HUMAN	Putative uncharacterized protein C1orf213 OS=Homo sapiens GN=C1orf213 PE=4 SV=1	0.9522	-0.304006187
Q9NZA1-2	CLIC5_HUMAN	Isoform 1 of Chloride intracellular channel protein 5 OS=Homo sapiens GN=CLIC5	0.9543	-0.321928095
Q99733	NP1L4_HUMAN	Nucleosome assembly protein 1-like 4 OS=Homo sapiens GN=NP1L4 PE=1 SV=1	1	-0.321928095
Q8N4F7	RN175_HUMAN	RING finger protein 175 OS=Homo sapiens GN=RN175 PE=2 SV=2	0.9888	-0.321928095
P04216	THY1_HUMAN	Thy-1 membrane glycoprotein OS=Homo sapiens GN=THY1 PE=1 SV=2	0.8085	-0.321928095
P26640	SYVC_HUMAN	Valine-tRNA ligase OS=Homo sapiens GN=VAR5 PE=1 SV=4	1	-0.340075442
P36543-2	VATE1_HUMAN	Isoform 2 of V-type proton ATPase subunit E 1 OS=Homo sapiens GN=ATP6V1E1	0.9992	-0.340075442
P61011-2	SRP54_HUMAN	Isoform 2 of Signal recognition particle 54 kDa protein OS=Homo sapiens GN=SRP54	0.9988	-0.340075442
Q96FW1	OTUB1_HUMAN	Ubiquitin thioesterase OTUB1 OS=Homo sapiens GN=OTUB1 PE=1 SV=2	1	-0.358453971
Q96552-2	PIGS_HUMAN	Isoform 2 of GPI transamidase component PIG-S OS=Homo sapiens GN=PIGS	0.9997	-0.358453971
Q7Z333-3	SETX_HUMAN	Isoform 3 of Probable helicase senataxin OS=Homo sapiens GN=SETX	0.9992	-0.358453971
P21964-2	COMT_HUMAN	Isoform Soluble of Catechol O-methyltransferase OS=Homo sapiens GN=COMT	0.9816	-0.358453971
Q9Y3E0	GOT1B_HUMAN	Vesicle transport protein GOT1B OS=Homo sapiens GN=GOT1B PE=1 SV=1	0.9218	-0.358453971
Q16543	CDC37_HUMAN	Hsp90 co-chaperone Cdc37 OS=Homo sapiens GN=CDC37 PE=1 SV=1	0.8947	-0.358453971
Q5VTL8-2	PR38B_HUMAN	Isoform 2 of Pre-mRNA-splicing factor 38B OS=Homo sapiens GN=PRPF38B	0.835	-0.377069649
P13804	ETFA_HUMAN	Electron transfer flavoprotein subunit alpha	1	-0.395928676
Q01484	ANK2_HUMAN	Ankyrin-2 OS=Homo sapiens GN=ANK2 PE=1 SV=4	0.9992	-0.395928676
O14524-2	T194A_HUMAN	Isoform 2 of Transmembrane protein 194A OS=Homo sapiens GN=TMEM194A	0.9903	-0.395928676
P10768	ESTD_HUMAN	S-formylglutathione hydrolase OS=Homo sapiens GN=ESD PE=1 SV=2	1	-0.415037499
P24941-2	CDK2_HUMAN	Isoform 2 of Cyclin-dependent kinase 2 OS=Homo sapiens GN=CDK2	1	-0.415037499
Q8WXH0-2	SYNE2_HUMAN	Isoform 2 of Nesprin-2 OS=Homo sapiens GN=SYNE2	0.9997	-0.415037499
Q14774	HLX_HUMAN	H2.0-like homeobox protein OS=Homo sapiens GN=HLX PE=1 SV=3	0.9677	-0.415037499
Q9P265	DIP2B_HUMAN	Disco-interacting protein 2 homolog B OS=Homo sapiens GN=DIP2B PE=1 SV=3	1	-0.434402824
P49915	GUA4_HUMAN	GMP synthase [glutamine-hydrolyzing] OS=Homo sapiens GN=GMPS PE=1 SV=1	1	-0.434402824
Q6UY18	LIGO4_HUMAN	Leucine-rich repeat and immunoglobulin-like domain-containing nogo receptor-interacting protein 4 OS	0.9466	-0.434402824
O75691	UTP20_HUMAN	Small subunit processome component 20 homolog OS=Homo sapiens GN=UTP20 PE=1 SV=3	1	-0.454031631
P11233	RALA_HUMAN	Ras-related protein Ral-A OS=Homo sapiens GN=RALA PE=1 SV=1	0.9996	-0.454031631
P53814-5	SMTN_HUMAN	Isoform B2 of Smoothelin OS=Homo sapiens GN=SMTN	0.9987	-0.454031631
P10515	ODP2_HUMAN	Dihydropyridyllysine-residue acetyltransferase component of pyruvate dehydrogenase complex	1	-0.473931188
Q8IY57-2	YAF2_HUMAN	Isoform 1 of YY1-associated factor 2 OS=Homo sapiens GN=YAF2	0.9924	-0.473931188
Q6SPF0	SAMD1_HUMAN	Atherin OS=Homo sapiens GN=SAMD1 PE=1 SV=1	0.8996	-0.473931188
P13942-2	COBA2_HUMAN	Isoform 2 of Collagen alpha-2(XI) chain OS=Homo sapiens GN=COL11A2	0.9643	-0.49410907
Q86XL3-2	ANKL2_HUMAN	Isoform 2 of Ankyrin repeat and LEM domain-containing protein 2 OS=Homo sapiens GN=ANKL2	0.9312	-0.49410907
Q9BZX2-2	UCK2_HUMAN	Isoform 2 of Uridine-cytidine kinase 2 OS=Homo sapiens GN=UCK2	0.9998	-0.535331733
Q8TEQ8	PIGO_HUMAN	GPI ethanolamine phosphate transferase 3 OS=Homo sapiens GN=PIGO PE=2 SV=3	0.9992	-0.556393349
Q96ND0	F210A_HUMAN	Protein FAM210A OS=Homo sapiens GN=FAM210A PE=1 SV=2	0.995	-0.556393349
P35222-2	CTNBN1_HUMAN	Isoform 2 of Catenin beta-1 OS=Homo sapiens GN=CTNBN1	0.992	-0.577766999
Q8WWY3-2	PRP31_HUMAN	Isoform 2 of U4/U6 small nuclear ribonucleoprotein Prp31 OS=Homo sapiens GN=PRPF31	0.8046	-0.577766999
Q96RP9-2	EFGM_HUMAN	Isoform 2 of Elongation factor G	0.9959	-0.59946207
Q68D10-2	SPT2_HUMAN	Isoform 2 of Protein SPT2 homolog OS=Homo sapiens GN=SPTY2D1	0.9965	-0.59946207
F8VRG5	F8VRG5_HUMAN	Protein C12orf52 OS=Homo sapiens GN=C12orf52 PE=4 SV=1	0.979	-0.59946207
Q9NSB4	KRT82_HUMAN	Keratin	0.8531	-0.621488377
Q8TEM1	PO210_HUMAN	Nuclear pore membrane glycoprotein 210 OS=Homo sapiens GN=NUP210 PE=1 SV=3	1	-0.64385619
O75431	MTX2_HUMAN	Metaxin-2 OS=Homo sapiens GN=MTX2 PE=1 SV=1	0.9997	-0.64385619
P04792	HSPB1_HUMAN	Heat shock protein beta-1 OS=Homo sapiens GN=HSPB1 PE=1 SV=2	0.9988	-0.64385619
Q8IUG5	MYO18B_HUMAN	Unconventional myosin-XVIIIb OS=Homo sapiens GN=MYO18B PE=1 SV=1	0.9952	-0.64385619
P36915-2	GNL1_HUMAN	Isoform 2 of Guanine nucleotide-binding protein-like 1 OS=Homo sapiens GN=GNL1	0.9808	-0.64385619
Q5CZC0	FSIP2_HUMAN	Fibrous sheath-interacting protein 2 OS=Homo sapiens GN=FSIP2 PE=1 SV=4	0.9579	-0.64385619
O00567	NOP56_HUMAN	Nucleolar protein 56 OS=Homo sapiens GN=NOP56 PE=1 SV=4	0.9999	-0.666576266
Q7L1Q6-2	BZW1_HUMAN	Isoform 2 of Basic leucine zipper and W2 domain-containing protein 1 OS=Homo sapiens GN=BZW1	0.9993	-0.666576266
Q14393-2	GAS6_HUMAN	Isoform 2 of Growth arrest-specific protein 6 OS=Homo sapiens GN=GAS6	0.9987	-0.666576266
Q8WZ42-11	TITIN_HUMAN	Isoform 11 of Titin OS=Homo sapiens GN=TTN	1	-0.689659879
Q13492-2	PICALM_HUMAN	Isoform 2 of Phosphatidylinositol-binding clathrin assembly protein OS=Homo sapiens GN=PICALM	0.9983	-0.689659879
Q99961	SH3G1_HUMAN	Endophilin-A2 OS=Homo sapiens GN=SH3GL1 PE=1 SV=1	0.9942	-0.689659879
Q8TE73	DYH5_HUMAN	Dynein heavy chain 5	0.8932	-0.713118852
Q9UPT8	ZC3H4_HUMAN	Zinc finger CCCH domain-containing protein 4 OS=Homo sapiens GN=ZC3H4 PE=1 SV=3	0.9983	-0.713118852
P21912	DHSB_HUMAN	Succinate dehydrogenase [ubiquinone] iron-sulfur subunit	1	-0.736965594
O75947-2	ATP5H_HUMAN	Isoform 2 of ATP synthase subunit d	1	-0.736965594
P41091	IF2G_HUMAN	Eukaryotic translation initiation factor 2 subunit 3 OS=Homo sapiens GN=EIF2S3 PE=1 SV=3	0.9167	-0.76121314
P50993	AT1A2_HUMAN	Sodium/potassium-transporting ATPase subunit alpha-2 OS=Homo sapiens GN=ATP1A2 PE=1 SV=1	0.8199	-0.785875195
O60333-3	KIF1B_HUMAN	Isoform 3 of Kinesin-like protein KIF1B OS=Homo sapiens GN=KIF1B	0.9904	-0.785875195
P25100	ADA1D_HUMAN	Alpha-1D adrenergic receptor OS=Homo sapiens GN=ADRA1D PE=2 SV=2	0.9381	-0.810966176
Q9BW71	HIRP3_HUMAN	HIRA-interacting protein 3 OS=Homo sapiens GN=HIRP3 PE=1 SV=3	0.9935	-0.862496476
Q9Y696	CLIC4_HUMAN	Chloride intracellular channel protein 4 OS=Homo sapiens GN=CLIC4 PE=1 SV=4	1	-0.888968688
Q15120-2	PKD3_HUMAN	Isoform 2 of [Pyruvate dehydrogenase (lipoamide)] kinase isozyme 3	0.985	-0.888968688
P05177-2	CP1A2_HUMAN	Isoform 2 of Cytochrome P450 1A2 OS=Homo sapiens GN=CYP1A2	0.9085	-0.888968688
P06493-2	CDK1_HUMAN	Isoform 2 of Cyclin-dependent kinase 1 OS=Homo sapiens GN=CDK1	0.9979	-0.943416472
Q9Y4G2	PKHM1_HUMAN	Pleckstrin homology domain-containing family M member 1 OS=Homo sapiens GN=PLEKHM1 PE=1 SV=1	0.9994	-0.971430848
O94988	FA13A_HUMAN	Protein FAM13A OS=Homo sapiens GN=FAM13A PE=1 SV=2	0.9093	-0.971430848
Q9Y4W6	AFG32_HUMAN	AFG32-like protein 2 OS=Homo sapiens GN=AFG32L PE=1 SV=2	0.9746	-0.971430848
Q32MK0-4	MYLK3_HUMAN	Isoform 2 of Myosin light chain kinase 3 OS=Homo sapiens GN=MYLK3	0.9564	-1.029146346
Q15113	PCOC1_HUMAN	Procollagen C-endopeptidase enhancer 1 OS=Homo sapiens GN=PCOLCE PE=1 SV=2	0.919	-1.058893689
P48058	GRIA4_HUMAN	Glutamate receptor 4 OS=Homo sapiens GN=GRIA4 PE=2 SV=2	0.9421	-1.089267338

Q9NRE2	TSH2_HUMAN	Teashirt homolog 2 OS=Homo sapiens GN=TSHZ2 PE=1 SV=3	0.9998	-1.120294234
P26358-2	DNMT1_HUMAN	Isoform 2 of DNA (cytosine-5)-methyltransferase 1 OS=Homo sapiens GN=DNMT1	0.9988	-1.120294234
Q14117	DPYS_HUMAN	Dihydropyrimidinase OS=Homo sapiens GN=DPYS PE=1 SV=1	0.9519	-1.120294234
P31146	COR1A_HUMAN	Coronin-1A OS=Homo sapiens GN=CORO1A PE=1 SV=4	0.9318	-1.120294234
P60953	CDC42_HUMAN	Cell division control protein 42 homolog OS=Homo sapiens GN=CDC42 PE=1 SV=2	1	-1.152003093
P54727	RD23B_HUMAN	UV excision repair protein RAD23 homolog B OS=Homo sapiens GN=RD23B PE=1 SV=1	1	-1.152003093
H3BQ73	H3BQ73_HUMAN	Vacuolar fusion protein MON1 homolog B OS=Homo sapiens GN=MON1B PE=4 SV=1	0.9718	-1.152003093
Q9HCK1	ZDBF2_HUMAN	DBF4-type zinc finger-containing protein 2 OS=Homo sapiens GN=ZDBF2 PE=1 SV=3	0.9656	-1.152003093
Q969Y2-2	GTPB3_HUMAN	Isoform 2 of tRNA modification GTPase GTPBP3	0.9693	-1.184424571
Q01954	BNC1_HUMAN	Zinc finger protein basonuclin-1 OS=Homo sapiens GN=BNC1 PE=1 SV=2	0.9996	-1.217591435
A4D212	A4D212_HUMAN	DKFZP586J0619 protein OS=Homo sapiens GN=DKFZP586J0619 PE=4 SV=1	0.9994	-1.217591435
Q9BZF9	UACA_HUMAN	Uveal autoantigen with coiled-coil domains and ankyrin repeats OS=Homo sapiens GN=UACA PE=1 SV=	0.9879	-1.251538767
P14618	KPYM_HUMAN	Pyruvate kinase isozymes M1/M2 OS=Homo sapiens GN=PKM PE=1 SV=4	1	-1.395928676
Q9UPN3-2	MACF1_HUMAN	Isoform 2 of Microtubule-actin cross-linking factor 1	0.9997	-1.434402824
O60741	HCN1_HUMAN	Potassium/sodium hyperpolarization-activated cyclic nucleotide-gated channel 1 OS=Homo sapiens GN=	0.9911	-1.556393349
P46013-2	KI67_HUMAN	Isoform Short of Antigen KI-67 OS=Homo sapiens GN=MKI67	1	-1.736965594
E7EX42	E7EX42_HUMAN	Pancreatic lipase-related protein 1 OS=Homo sapiens GN=PNLIPRP1 PE=4 SV=1	0.9121	-1.736965594
Q14934-14	NFAC4_HUMAN	Isoform 14 of Nuclear factor of activated T-cells	0.95	-1.785875195
Q9Y281	COF2_HUMAN	Cofilin-2 OS=Homo sapiens GN=CFL2 PE=1 SV=1	0.9898	-1.888968688
Q8NDA2	HMCN2_HUMAN	Hemicentin-2 OS=Homo sapiens GN=HMCN2 PE=2 SV=2	0.9901	-1.888968688
Q9Y261-2	FOXA2_HUMAN	Isoform 2 of Hepatocyte nuclear factor 3-beta OS=Homo sapiens GN=FOXA2	0.9993	-2
Q8NHY3	GAS2L2_HUMAN	GAS2-like protein 2 OS=Homo sapiens GN=GAS2L2 PE=2 SV=1	0.9672	-2.058893689
Q8TF72	SHRM3_HUMAN	Protein Shroom3 OS=Homo sapiens GN=SHROOM3 PE=1 SV=2	0.9976	-2.120294234
Q8TDW7-3	FAT3_HUMAN	Isoform 3 of Protocadherin Fat 3 OS=Homo sapiens GN=FAT3	0.9999	-2.184424571
O15083	ERC2_HUMAN	ERC protein 2 OS=Homo sapiens GN=ERC2 PE=1 SV=3	0.9835	-2.251538767
Q15323	K1H1_HUMAN	Keratin	1	-2.251538767
Q9NQW8-2	CNGB3_HUMAN	Isoform 2 of Cyclic nucleotide-gated cation channel beta-3 OS=Homo sapiens GN=CNGB3	0.9982	-2.251538767
D6R922	D6R922_HUMAN	WD repeat-containing protein 36 OS=Homo sapiens GN=WDR36 PE=4 SV=1	0.9921	-2.251538767
Q7Z2K8-2	GRIN1_HUMAN	Isoform 2 of G protein-regulated inducer of neurite outgrowth 1 OS=Homo sapiens GN=GPRIN1	1	-2.64385619
Q81YR2	SMYD4_HUMAN	SET and MYND domain-containing protein 4 OS=Homo sapiens GN=SMYD4 PE=2 SV=3	0.9999	-2.836501268
A7E2U8	CD047_HUMAN	UPF0602 protein C4orf47 OS=Homo sapiens GN=C4orf47 PE=2 SV=1	0.9362	-2.836501268
P02768-2	ALBU_HUMAN	Isoform 2 of Serum albumin OS=Homo sapiens GN=ALB	0.9989	-3.058893689
Q9UMN6	MLL4_HUMAN	Histone-lysine N-methyltransferase MLL4 OS=Homo sapiens GN=WBP7 PE=1 SV=1	0.9938	-3.058893689
Q92623	TTC9A_HUMAN	Tetratricopeptide repeat protein 9A OS=Homo sapiens GN=TTC9 PE=2 SV=3	0.9678	-3.184424571
E5RJ70	E5RJ70_HUMAN	Eukaryotic translation initiation factor 3 subunit H (Fragment) OS=Homo sapiens GN=EIF3H PE=4 SV=1	1	-3.64385619
Q10471	GALT2_HUMAN	Polypeptide N-acetylgalactosaminyltransferase 2 OS=Homo sapiens GN=GALNT2 PE=1 SV=1	0.9965	-3.836501268
Q9Y6Z5	MLAS1_HUMAN	Putative uncharacterized protein MLLT4-AS1 OS=Homo sapiens GN=MLLT4-AS1 PE=5 SV=1	0.8992	-3.836501268
O15078	CE290_HUMAN	Centrosomal protein of 290 kDa OS=Homo sapiens GN=CEP290 PE=1 SV=2	0.9618	-4.321928095
Q865Q6-1	GP123_HUMAN	Isoform 1 of Probable G-protein coupled receptor 123 OS=Homo sapiens GN=GPR123	0.9373	-4.64385619
P78385	KRT83_HUMAN	Keratin	0.9999	-5.64385619
O75044	SRGP2_HUMAN	SLIT-ROBO Rho GTPase-activating protein 2 OS=Homo sapiens GN=SRGAP2 PE=1 SV=2	0.9924	-6.64385619
Q0VG06-3	FP100_HUMAN	Isoform 3 of Fanconi anemia-associated protein of 100 kDa OS=Homo sapiens GN=FAAP100	0.9987	9.964340868
Q03188	CENPC_HUMAN	Centromere protein C 1 OS=Homo sapiens GN=CENPC1 PE=1 SV=2	0.9897	5.015247774
J3QRW8	J3QRW8_HUMAN	Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=1	0.9438	4.793895883
Q7Z3U7-2	MON2_HUMAN	Isoform 2 of Protein MON2 homolog OS=Homo sapiens GN=MON2	0.9853	3.74343989
P42771-3	CD2A1_HUMAN	Isoform 3 of Cyclin-dependent kinase inhibitor 2A	0.9924	2.769771739
Q12888-2	TP53B_HUMAN	Isoform 2 of Tumor suppressor p53-binding protein 1 OS=Homo sapiens GN=TP53BP1	0.9981	2.488000771
Q9C0G6	DYH6_HUMAN	Dynein heavy chain 6	0.9985	2.339137385
Q8WWW8	ARAP3_HUMAN	Arf-GAP with Rho-GAP domain	0.8591	2.207892852
Q5T5C0-2	STXB5_HUMAN	Isoform 2 of Syntaxin-binding protein 5 OS=Homo sapiens GN=STXB5	0.9901	2.189033824
Q5SR65-2	NU188_HUMAN	Isoform 2 of Nucleoporin NUP188 homolog OS=Homo sapiens GN=NUP188	0.9875	1.827819025
A5D8V7	CC151_HUMAN	Coiled-coil domain-containing protein 151 OS=Homo sapiens GN=CCDC151 PE=2 SV=1	0.9972	1.744161096
P04899-3	GNAI2_HUMAN	Isoform 3 of Guanine nucleotide-binding protein G(i) subunit alpha-2 OS=Homo sapiens GN=GNAI2	0.9938	1.432959407
P13073	COX41_HUMAN	Cytochrome c oxidase subunit 4 isoform 1	0.9446	1.344828497
H0Y138	H0Y138_HUMAN	Uncharacterized protein OS=Homo sapiens PE=4 SV=1	0.9787	1.15704371
F6WGS5	F6WGS5_HUMAN	Steroid receptor RNA activator 1 OS=Homo sapiens GN=SRA1 PE=4 SV=1	0.9841	1.021479727
Q8N201	INT1_HUMAN	Integrator complex subunit 1 OS=Homo sapiens GN=INTS1 PE=1 SV=2	0.9999	1.014355293
F8WCL5	F8WCL5_HUMAN	Nebulin OS=Homo sapiens GN=NEB PE=4 SV=1	0.9999	1.014355293
P55263-2	ADK_HUMAN	Isoform Short of Adenosine kinase OS=Homo sapiens GN=ADK	0.9085	0.963474124
F8W7C6	F8W7C6_HUMAN	60S ribosomal protein L10 OS=Homo sapiens GN=RPL10 PE=4 SV=1	1	0.748461233
Q12904-2	AIMP1_HUMAN	Isoform 2 of Aminoacyl tRNA synthase complex-interacting multifunctional protein 1 OS=Homo sapiens	1	0.731183242
Q99615-2	DNJC7_HUMAN	Isoform 2 of DnaJ homolog subfamily C member 7 OS=Homo sapiens GN=DNAJC7	0.9364	0.731183242
Q8TAA5	GRPE2_HUMAN	GrpE protein homolog 2	0.9999	0.695993813
Q01813	K6PP_HUMAN	6-phosphofructokinase type C OS=Homo sapiens GN=PFKP PE=1 SV=2	0.8455	0.622930351
Q86W42-3	THOC6_HUMAN	Isoform 3 of THO complex subunit 6 homolog OS=Homo sapiens GN=THOC6	1	0.59454855
Q16740	CLPP_HUMAN	Putative ATP-dependent Clp protease proteolytic subunit	0.8778	0.545968369
Q8TF39	ZN483_HUMAN	Zinc finger protein 483 OS=Homo sapiens GN=ZNF483 PE=1 SV=3	0.9575	0.454175893
E7EMK4	E7EMK4_HUMAN	ATP-dependent RNA helicase DDX19B OS=Homo sapiens GN=DDX19B PE=4 SV=1	0.9968	0.443606651
Q8IZT6-2	ASPM_HUMAN	Isoform 2 of Abnormal spindle-like microcephaly-associated protein OS=Homo sapiens GN=ASPM	0.9915	0.443606651
Q07837	SLC31_HUMAN	Neutral and basic amino acid transport protein rBAT OS=Homo sapiens GN=SLC3A1 PE=1 SV=2	0.9975	0.400557993
Q9NPJ3	ACO13_HUMAN	Acyl-coenzyme A thioesterase 13 OS=Homo sapiens GN=ACOT13 PE=1 SV=1	0.9997	0.367371066
Q9NTI5-2	PDS5B_HUMAN	Isoform 2 of Sister chromatid cohesion protein PDS5 homolog B OS=Homo sapiens GN=PDS5B	1	0.35614381
P14866-2	HNRNPL_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein L OS=Homo sapiens GN=HNRNPL	1	0.333423734
P63010-2	AP2B1_HUMAN	Isoform 2 of AP-2 complex subunit beta OS=Homo sapiens GN=AP2B1	1	0.333423734
Q96P48-1	ARAP1_HUMAN	Isoform 1 of Arf-GAP with Rho-GAP domain	0.9937	0.333423734
P20340-2	RAB6A_HUMAN	Isoform 2 of Ras-related protein Rab-6A OS=Homo sapiens GN=RAB6A	0.9581	0.321928095
P35244	RFA3_HUMAN	Replication protein A 14 kDa subunit OS=Homo sapiens GN=RPA3 PE=1 SV=1	0.9999	0.310340121
P61106	RAB14_HUMAN	Ras-related protein Rab-14 OS=Homo sapiens GN=RAB14 PE=1 SV=4	1	0.23878686
Q9UBU9	NXF1_HUMAN	Nuclear RNA export factor 1 OS=Homo sapiens GN=NXF1 PE=1 SV=1	0.9999	0.23878686
Q8IY11	FAM71A_HUMAN	Protein FAM71A OS=Homo sapiens GN=FAM71A PE=2 SV=2	0.9707	0.23878686
Q99720-2	SGMR1_HUMAN	Isoform 2 of Sigma non-opioid intracellular receptor 1 OS=Homo sapiens GN=SIGMAR1	0.9952	0.22650853
G3V248	G3V248_HUMAN	Interferon alpha-inducible protein 27-like protein 1 OS=Homo sapiens GN=IFI27L1 PE=4 SV=1	0.9325	0.22650853
O95197-2	RTN3_HUMAN	Isoform 2 of Reticulon-3 OS=Homo sapiens GN=RTN3	1	0.214124805
F8W031	F8W031_HUMAN	Uncharacterized protein (Fragment) OS=Homo sapiens PE=4 SV=1	0.9863	0.214124805
Q13885	TUBB2A_HUMAN	Tubulin beta-2A chain OS=Homo sapiens GN=TUBB2A PE=1 SV=1	0.9992	0.214124805
Q8TEA8	DTD1_HUMAN	D-tyrosyl-tRNA(Tyr) deacylase 1 OS=Homo sapiens GN=DTD1 PE=1 SV=2	1	0.214124805
C9J7Q0	ANKRD63_HUMAN	Ankyrin repeat domain-containing protein 63 OS=Homo sapiens GN=ANKRD63 PE=4 SV=1	0.9988	0.201633861
Q49A88-5	CCD14_HUMAN	Isoform 5 of Coiled-coil domain-containing protein 14 OS=Homo sapiens GN=CCDC14	0.998	0.201633861
Q02809	PLOD1_HUMAN	Procollagen-lysine	0.9962	0.176322773
Q8TE49-2	OTU7A_HUMAN	Isoform 2 of OTU domain-containing protein 7A OS=Homo sapiens GN=OTUD7A	0.9227	0.176322773
P63173	RL38_HUMAN	60S ribosomal protein L38 OS=Homo sapiens GN=RPL38 PE=1 SV=2	0.995	0.163498732
O95777	NAA38_HUMAN	N-alpha-acetyltransferase 38	0.9997	0.163498732

Q13416	ORC2_HUMAN	Origin recognition complex subunit 2 OS=Homo sapiens GN=ORC2 PE=1 SV=2	0.9882	0.163498732
Q08623-3	HDHD1_HUMAN	Isoform 3 of Pseudouridine-5'-monophosphatase OS=Homo sapiens GN=HDHD1	1	0.150559677
Q5HYC2	K2026_HUMAN	Uncharacterized protein KIAA2026 OS=Homo sapiens GN=KIAA2026 PE=2 SV=2	0.9869	0.150559677
P28838-2	AMPL_HUMAN	Isoform 2 of Cytosol aminopeptidase OS=Homo sapiens GN=LAP3	0.9996	0.150559677
P35080	PROF2_HUMAN	Profilin-2 OS=Homo sapiens GN=PFN2 PE=1 SV=3	1	0.137503524
Q8TDM6-2	DLG5_HUMAN	Isoform 2 of Disks large homolog 5 OS=Homo sapiens GN=DLG5	0.9926	0.124328135
O75832	PSD10_HUMAN	26S proteasome non-ATPase regulatory subunit 10 OS=Homo sapiens GN=PSMD10 PE=1 SV=1	1	0.111031312
P12235	ADT1_HUMAN	ADP/ATP translocase 1 OS=Homo sapiens GN=SLC25A4 PE=1 SV=4	0.989	0.111031312
Q5V266	JKIP3_HUMAN	Janus kinase and microtubule-interacting protein 3 OS=Homo sapiens GN=JAKMIP3 PE=2 SV=2	0.9861	0.111031312
Q7L0J3-2	SV2A_HUMAN	Isoform 2 of Synaptic vesicle glycoprotein 2A OS=Homo sapiens GN=SV2A	0.962	0.111031312
Q9N9R3	NUA4L_HUMAN	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 4-like 2 OS=Homo sapiens GN=NDUFA	0.9606	0.111031312
Q9BUF5	TBB6_HUMAN	Tubulin beta-6 chain OS=Homo sapiens GN=TUBB6 PE=1 SV=1	1	0.097610797
Q9HC16-2	K1586_HUMAN	Isoform 2 of Uncharacterized protein KIAA1586 OS=Homo sapiens GN=KIAA1586	0.9821	0.097610797
Q9NSC7	SIA7A_HUMAN	Alpha-N-acetylgalactosaminide alpha-2	0.9727	0.097610797
P52333-2	JAK3_HUMAN	Isoform 1 of Tyrosine-protein kinase JAK3 OS=Homo sapiens GN=JAK3	0.8629	0.097610797
P35249	RFC4_HUMAN	Replication factor C subunit 4 OS=Homo sapiens GN=RFC4 PE=1 SV=2	1	0.084064265
Q6IBS0	TWF2_HUMAN	Twinfilin-2 OS=Homo sapiens GN=TWF2 PE=1 SV=2	1	0.084064265
Q8N3U4-2	STAG2_HUMAN	Isoform 2 of Cohesin subunit SA-2 OS=Homo sapiens GN=STAG2	0.9995	0.084064265
P07237	PDIA1_HUMAN	Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3	1	0.070389328
Q8TEM1-2	PO210_HUMAN	Isoform 2 of Nuclear pore membrane glycoprotein 210 OS=Homo sapiens GN=NUP210	0.9991	0.070389328
P63172	DYLT1_HUMAN	Dynein light chain Tctex-type 1 OS=Homo sapiens GN=DYNLT1 PE=1 SV=1	1	0.056583528
Q15363	TMED2_HUMAN	Transmembrane emp24 domain-containing protein 2 OS=Homo sapiens GN=TMED2 PE=1 SV=1	1	0.056583528
Q8NBJ5	GT251_HUMAN	Procollagen galactosyltransferase 1 OS=Homo sapiens GN=GLT25D1 PE=1 SV=1	1	0.056583528
P63241	IF5A1_HUMAN	Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A PE=1 SV=2	1	0.056583528
Q9NVH0-2	EXD2_HUMAN	Isoform 2 of Exonuclease 3'-5' domain-containing protein 2 OS=Homo sapiens GN=EXD2	0.9983	0.056583528
O43592	XPOT_HUMAN	Exportin-T OS=Homo sapiens GN=XPOT PE=1 SV=2	0.9999	0.028569152
Q5JTH9-2	RRP12_HUMAN	Isoform 2 of RRP12-like protein OS=Homo sapiens GN=RRP12	0.9999	0.028569152
P14324	FPPS_HUMAN	Farnesyl pyrophosphate synthase OS=Homo sapiens GN=FDPS PE=1 SV=4	0.9987	0.014355293
P17405-2	ASM_HUMAN	Isoform 2 of Sphingomyelin phosphodiesterase OS=Homo sapiens GN=SMPD1	0.9584	0.014355293
P28074-2	PSB5_HUMAN	Isoform 2 of Proteasome subunit beta type-5 OS=Homo sapiens GN=PSMB5	1	0
Q96HS1-2	PGAM5_HUMAN	Isoform 2 of Serine/threonine-protein phosphatase PGAM5	1	0
Q6N069	NAA16_HUMAN	N-alpha-acetyltransferase 16	0.9998	0
Q9NQ78	K113B_HUMAN	Kinesin-like protein KIF13B OS=Homo sapiens GN=KIF13B PE=1 SV=1	0.9547	0
O00116	ADAS_HUMAN	Aldyldihydroxyacetonephosphate synthase	0.9994	-0.01449957
Q6L93-2	K116B_HUMAN	Isoform 2 of Kinesin-like protein KIF16B OS=Homo sapiens GN=KIF16B	0.9992	-0.01449957
O95372	LYP2_HUMAN	Acyl-protein thioesterase 2 OS=Homo sapiens GN=LYPLA2 PE=1 SV=1	0.9982	-0.01449957
Q6ZU69	F205A_HUMAN	Protein FAM205A OS=Homo sapiens GN=FAM205A PE=2 SV=4	0.9911	-0.01449957
Q15435-2	PP1R7_HUMAN	Isoform 2 of Protein phosphatase 1 regulatory subunit 7 OS=Homo sapiens GN=PPP1R7	1	-0.029146346
P32322	P5CR1_HUMAN	Pyrrrole-5-carboxylate reductase 1	1	-0.029146346
Q9Y4P8-2	WIPI2_HUMAN	Isoform 2 of WD repeat domain phosphoinositide-interacting protein 2 OS=Homo sapiens GN=WIPI2	0.9897	-0.029146346
P36551	HEM6_HUMAN	Coproporphyrinogen-III oxidase	0.8411	-0.029146346
Q9UL25	RAB21_HUMAN	Ras-related protein Rab-21 OS=Homo sapiens GN=RAB21 PE=1 SV=3	1	-0.043943348
P30042-2	ES1_HUMAN	Isoform Short of ES1 protein homolog	1	-0.043943348
P52597	HNRPF_HUMAN	Heterogeneous nuclear ribonucleoprotein F OS=Homo sapiens GN=HNRNPF PE=1 SV=3	0.9407	-0.043943348
Q9H9L3	I20L2_HUMAN	Interferon-stimulated 20 kDa exonuclease-like 2 OS=Homo sapiens GN=ISG20L2 PE=1 SV=1	0.9916	-0.043943348
Q75VX8-2	FA59B_HUMAN	Isoform 2 of Protein FAM59B OS=Homo sapiens GN=FAM59B	0.9386	-0.058893689
P18085	ARF4_HUMAN	ADP-ribosylation factor 4 OS=Homo sapiens GN=ARF4 PE=1 SV=3	1	-0.058893689
Q86X4-2	FRAS1_HUMAN	Isoform 2 of Extracellular matrix protein FRAS1 OS=Homo sapiens GN=FRAS1	0.9906	-0.058893689
Q15418-2	KS6A1_HUMAN	Isoform 2 of Ribosomal protein S6 kinase alpha-1 OS=Homo sapiens GN=RPS6KA1	0.9999	-0.074000581
P22307-2	NLTP_HUMAN	Isoform SCP2 of Non-specific lipid-transfer protein OS=Homo sapiens GN=SCP2	0.998	-0.074000581
O60271-2	JIP4_HUMAN	Isoform 2 of C-Jun-amino-terminal kinase-interacting protein 4 OS=Homo sapiens GN=SPAG9	0.9534	-0.074000581
O14950	ML12B_HUMAN	Myosin regulatory light chain 12B OS=Homo sapiens GN=MYL12B PE=1 SV=2	1	-0.089267338
P09651-2	ROA1_HUMAN	Isoform A1-A of Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1	1	-0.089267338
Q99832-3	TCPH_HUMAN	Isoform 3 of T-complex protein 1 subunit eta OS=Homo sapiens GN=CCT7	1	-0.089267338
P55145	MANF_HUMAN	Mesencephalic astrocyte-derived neurotrophic factor OS=Homo sapiens GN=MANF PE=1 SV=3	0.9999	-0.089267338
Q27J81-2	INF2_HUMAN	Isoform 2 of Inverted formin-2 OS=Homo sapiens GN=INF2	0.9946	-0.104697379
F8VR82	F8VR82_HUMAN	Serine/threonine-protein phosphatase OS=Homo sapiens GN=PPP1CC PE=3 SV=1	0.9916	-0.120294234
P62195	PRS8_HUMAN	26S protease regulatory subunit 8 OS=Homo sapiens GN=PSMCS5 PE=1 SV=1	1	-0.13606155
P60709	ACTB_HUMAN	Actin	1	-0.13606155
O43847-2	NRDC_HUMAN	Isoform 2 of Nardilysin OS=Homo sapiens GN=NRD1	0.9975	-0.13606155
Q13404-1	UB2V1_HUMAN	Isoform 1 of Ubiquitin-conjugating enzyme E2 variant 1 OS=Homo sapiens GN=UBE2V1	0.8011	-0.13606155
P53621-2	COPA_HUMAN	Isoform 2 of Coatomer subunit alpha OS=Homo sapiens GN=COPA	1	-0.152003093
Q92797	SYMPK_HUMAN	Symplekin OS=Homo sapiens GN=SYMPK PE=1 SV=2	1	-0.152003093
A8MWP4	YU008_HUMAN	Putative uncharacterized protein ENSP0000401716 OS=Homo sapiens GN=PE=5 SV=1	0.9944	-0.152003093
Q96CX2	KCD12_HUMAN	BTB/POZ domain-containing protein KCTD12 OS=Homo sapiens GN=KCTD12 PE=1 SV=1	0.9937	-0.152003093
P26378-2	ELAV4_HUMAN	Isoform 2 of ELAV-like protein 4 OS=Homo sapiens GN=ELAVL4	0.9912	-0.152003093
Q9NZP6	NPAP1_HUMAN	Nuclear pore-associated protein 1 OS=Homo sapiens GN=NPAP1 PE=1 SV=2	0.9713	-0.152003093
P07910-2	HNRPC_HUMAN	Isoform C1 of Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC	1	-0.168122759
O60506-2	HNRPQ_HUMAN	Isoform 2 of Heterogeneous nuclear ribonucleoprotein Q OS=Homo sapiens GN=SYNCRIP	0.9999	-0.168122759
P09972	ALDOC_HUMAN	Fructose-bisphosphate aldolase C OS=Homo sapiens GN=ALDOC PE=1 SV=2	1	-0.168122759
Q9HCM9	TR139_HUMAN	E3 ubiquitin-protein ligase TRIM39 OS=Homo sapiens GN=TRIM39 PE=1 SV=2	0.9973	-0.168122759
Q9P2R3-2	ANKFY1_HUMAN	Isoform 2 of Ankyrin repeat and FYVE domain-containing protein 1 OS=Homo sapiens GN=ANKFY1	0.9878	-0.168122759
Q09666	AHNAK_HUMAN	Neuroblast differentiation-associated protein AHNAK OS=Homo sapiens GN=AHNAK PE=1 SV=2	0.9553	-0.168122759
O43617	TPPC3_HUMAN	Trafficking protein particle complex subunit 3 OS=Homo sapiens GN=TRAPPC3 PE=1 SV=1	1	-0.184424571
P61086	UBE2K_HUMAN	Ubiquitin-conjugating enzyme E2 K OS=Homo sapiens GN=UBE2K PE=1 SV=3	1	-0.184424571
Q96EY4	TMA16_HUMAN	Translation machinery-associated protein 16 OS=Homo sapiens GN=TMA16 PE=1 SV=2	0.9992	-0.184424571
Q8NC54	ZMYNB_HUMAN	Uncharacterized protein ZMYNB OS=Homo sapiens GN=ZMYNB PE=2 SV=1	0.9758	-0.184424571
P98175-2	RBM10_HUMAN	Isoform 2 of RNA-binding protein 10 OS=Homo sapiens GN=RBM10	0.9012	-0.184424571
P61160	ARP2_HUMAN	Actin-related protein 2 OS=Homo sapiens GN=ACTR2 PE=1 SV=1	0.998	-0.200912694
Q9BTT0	AN32E_HUMAN	Acidic leucine-rich nuclear phosphoprotein 32 family member E OS=Homo sapiens GN=ANP32E PE=1 SV=1	1	-0.200912694
Q9H0U4	RAB1B_HUMAN	Ras-related protein Rab-1B OS=Homo sapiens GN=RAB1B PE=1 SV=1	1	-0.200912694
O14744-2	ANM5_HUMAN	Isoform 2 of Protein arginine N-methyltransferase 5 OS=Homo sapiens GN=PRMT5	0.9996	-0.200912694
Q96AT9-2	RPE_HUMAN	Isoform 2 of Ribulose-phosphate 3-epimerase OS=Homo sapiens GN=RPE	1	-0.217591435
P0CG30	GSTT2_HUMAN	Glutathione S-transferase theta-2B OS=Homo sapiens GN=GSTT2B PE=1 SV=1	1	-0.217591435
Q7L9L4	MOB1B_HUMAN	MOB kinase activator 1B OS=Homo sapiens GN=MOB1B PE=1 SV=3	1	-0.217591435
Q9Y265-2	RUVB1_HUMAN	Isoform 2 of RuvB-like 1 OS=Homo sapiens GN=RUVB1	1	-0.217591435
O75694	NU155_HUMAN	Nuclear pore complex protein Nup155 OS=Homo sapiens GN=NUP155 PE=1 SV=1	1	-0.234465254
O75400-3	PR40A_HUMAN	Isoform 3 of Pre-mRNA-processing factor 40 homolog A OS=Homo sapiens GN=PRPF40A	0.9998	-0.234465254
Q9UHY7	ENOPH_HUMAN	Enolase-phosphatase E1 OS=Homo sapiens GN=ENOPH1 PE=1 SV=1	0.9867	-0.234465254
A8MTJ3	GNAT3_HUMAN	Guanine nucleotide-binding protein G(t) subunit alpha-3 OS=Homo sapiens GN=GNAT3 PE=2 SV=2	1	-0.251538767
P04179	SODM_HUMAN	Superoxide dismutase [Mn]	1	-0.251538767
P50395	GDIB_HUMAN	Rab GDP dissociation inhibitor beta OS=Homo sapiens GN=GD12 PE=1 SV=2	1	-0.251538767

H0Y9N1	H0Y9N1_HUMAN	Programmed cell death protein 6 (Fragment) OS=Homo sapiens GN=PDCD6 PE=4 SV=1	0.9986	-0.251538767
Q99653	CHP1_HUMAN	Calcineurin B homologous protein 1 OS=Homo sapiens GN=CHP1 PE=1 SV=3	0.8949	-0.251538767
P35658-2	NU214_HUMAN	Isoform 2 of Nuclear pore complex protein Nup214 OS=Homo sapiens GN=NUP214	0.8924	-0.251538767
Q02952-2	AKA12_HUMAN	Isoform 2 of A-kinase anchor protein 12 OS=Homo sapiens GN=AKAP12	1	-0.268816758
P49321-2	NASP_HUMAN	Isoform 2 of Nuclear autoantigenic sperm protein OS=Homo sapiens GN=NASP	1	-0.268816758
Q15907	RB11B_HUMAN	Ras-related protein Rab-11B OS=Homo sapiens GN=RB11B PE=1 SV=4	1	-0.268816758
Q13618-2	CUL3_HUMAN	Isoform 2 of Cullin-3 OS=Homo sapiens GN=CUL3	0.9997	-0.268816758
Q9Y512	SAM50_HUMAN	Sorting and assembly machinery component 50 homolog OS=Homo sapiens GN=SAMM50 PE=1 SV=3	0.9995	-0.268816758
Q9Y3D8	KAD6_HUMAN	Adenylate kinase isoenzyme 6 OS=Homo sapiens GN=TAF9 PE=1 SV=1	0.9337	-0.268816758
Q961C7	RMD3_HUMAN	Regulator of microtubule dynamics protein 3 OS=Homo sapiens GN=FAM82A2 PE=1 SV=2	1	-0.286304185
Q9BYD2	RM09_HUMAN	39S ribosomal protein L9	0.9984	-0.286304185
P49753	ACOT2_HUMAN	Acyl-coenzyme A thioesterase 2	0.8754	-0.286304185
Q9Y263	PLAP_HUMAN	Phospholipase A-2-activating protein OS=Homo sapiens GN=PLAA PE=1 SV=2	0.9834	-0.304006187
Q9H3P2	NELFA_HUMAN	Negative elongation factor A OS=Homo sapiens GN=WHSC2 PE=1 SV=3	0.9304	-0.304006187
O76094	SRP72_HUMAN	Signal recognition particle 72 kDa protein OS=Homo sapiens GN=SRP72 PE=1 SV=3	1	-0.321928095
P12109	CO6A1_HUMAN	Collagen alpha-1(VI) chain OS=Homo sapiens GN=COL6A1 PE=1 SV=3	0.9979	-0.321928095
Q13405	RM49_HUMAN	39S ribosomal protein L49	0.9446	-0.321928095
P50213-2	IDH3A_HUMAN	Isoform 2 of Isocitrate dehydrogenase [NAD] subunit alpha	1	-0.340075442
Q76L83	ASXL2_HUMAN	Putative Polycomb group protein ASXL2 OS=Homo sapiens GN=ASXL2 PE=1 SV=1	0.9991	-0.340075442
O60721-2	NCKX1_HUMAN	Isoform 2 of Sodium/potassium/calcium exchanger 1 OS=Homo sapiens GN=SLC24A1	0.9609	-0.340075442
O00264	PGR1_HUMAN	Membrane-associated progesterone receptor component 1 OS=Homo sapiens GN=PGRMC1 PE=1 SV=3	1	-0.358453971
P55060	XPO2_HUMAN	Exportin-2 OS=Homo sapiens GN=CSE1L PE=1 SV=3	1	-0.358453971
Q14324	MYPC2_HUMAN	Myosin-binding protein C	0.9222	-0.358453971
Q9BV86-2	NTM1A_HUMAN	Isoform 2 of N-terminal Xaa-Pro-Lys N-methyltransferase 1 OS=Homo sapiens GN=NTMT1	1	-0.377069649
P42677	RS27_HUMAN	40S ribosomal protein S27 OS=Homo sapiens GN=RP527 PE=1 SV=3	1	-0.377069649
Q8WUW0	NU133_HUMAN	Nuclear pore complex protein Nup133 OS=Homo sapiens GN=NUP133 PE=1 SV=2	0.9998	-0.377069649
Q9NR45	SIAS_HUMAN	Sialic acid synthase OS=Homo sapiens GN=NANS PE=1 SV=2	0.9446	-0.377069649
O15160-2	RPAC1_HUMAN	Isoform 2 of DNA-directed RNA polymerases I and III subunit RPAC1 OS=Homo sapiens GN=POLR1C	0.8568	-0.377069649
Q15102	PA1B3_HUMAN	Platelet-activating factor acetylhydrolase IB subunit gamma OS=Homo sapiens GN=PFAFH1B3 PE=1 SV=	1	-0.395928676
Q9NZA1	CLIC5_HUMAN	Chloride intracellular channel protein 5 OS=Homo sapiens GN=CLIC5 PE=1 SV=3	0.9983	-0.395928676
P30049	ATPD_HUMAN	ATP synthase subunit delta	0.9999	-0.395928676
Q13765	NACA_HUMAN	Nascent polypeptide-associated complex subunit alpha OS=Homo sapiens GN=NACA PE=1 SV=1	1	-0.415037499
J3QLC8	J3QLC8_HUMAN	60S ribosomal protein L17 OS=Homo sapiens GN=RPL17 PE=3 SV=1	1	-0.415037499
P60059	SC61G_HUMAN	Protein transport protein Sec61 subunit gamma OS=Homo sapiens GN=SEC61G PE=2 SV=1	0.9446	-0.415037499
P08237-2	K6PF_HUMAN	Isoform 2 of 6-phosphofructokinase	1	-0.434402824
Q8TDD1-2	DDX54_HUMAN	Isoform 2 of ATP-dependent RNA helicase DDX54 OS=Homo sapiens GN=DDX54	0.9998	-0.434402824
Q16643-2	DREB_HUMAN	Isoform 2 of Drebrin OS=Homo sapiens GN=DNB1	0.9999	-0.454031631
P43304-2	GPDM_HUMAN	Isoform 2 of Glycerol-3-phosphate dehydrogenase	0.9984	-0.473931188
O43708-2	MAAL1_HUMAN	Isoform 2 of Maleylacetoacetate isomerase OS=Homo sapiens GN=GSTZ1	0.8731	-0.473931188
Q6DN03	H2B2C_HUMAN	Putative histone H2B type 2-C OS=Homo sapiens GN=HIST2H2BC PE=5 SV=3	1	-0.49410907
P52565	GDIR1_HUMAN	Rho GDP-dissociation inhibitor 1 OS=Homo sapiens GN=ARHGDI1A PE=1 SV=3	1	-0.49410907
Q8WTV0-2	SCR1B_HUMAN	Isoform 1 of Scavenger receptor class B member 1 OS=Homo sapiens GN=SCR1B	0.9836	-0.49410907
O43660-2	PLRG1_HUMAN	Isoform 2 of Pleiotropic regulator 1 OS=Homo sapiens GN=PLRG1	0.9405	-0.49410907
P05023-2	AT1A1_HUMAN	Isoform 2 of Sodium/potassium-transporting ATPase subunit alpha-1 OS=Homo sapiens GN=ATP1A1	0.992	-0.514573173
P11717	MPRI1_HUMAN	Cation-independent mannose-6-phosphate receptor OS=Homo sapiens GN=IGF2R PE=1 SV=3	0.9904	-0.514573173
P21281	VATB2_HUMAN	V-type proton ATPase subunit B	0.8971	-0.535331733
P67870	CSK2B_HUMAN	Casein kinase II subunit beta OS=Homo sapiens GN=CSNK2B PE=1 SV=1	1	-0.556393349
P62333	PRS10_HUMAN	26S protease regulatory subunit 10B OS=Homo sapiens GN=PSMC6 PE=1 SV=1	0.9999	-0.556393349
Q9NPL8	TIDC1_HUMAN	Translocase of inner mitochondrial membrane domain-containing protein 1 OS=Homo sapiens GN=TIMM	1	-0.577766999
Q13868	EXOS2_HUMAN	Exosome complex component RRP4 OS=Homo sapiens GN=EXOSC2 PE=1 SV=2	0.9419	-0.577766999
Q96A35	RM24_HUMAN	39S ribosomal protein L24	0.9152	-0.577766999
Q15370	ELOB_HUMAN	Transcription elongation factor B polypeptide 2 OS=Homo sapiens GN=TCEB2 PE=1 SV=1	0.9974	-0.59946207
Q96QC0	PP1RA_HUMAN	Serine/threonine-protein phosphatase 1 regulatory subunit 10 OS=Homo sapiens GN=PPP1R10 PE=1 SV	0.9849	-0.621488377
Q5T9A4-3	ATD3B_HUMAN	Isoform 3 of ATPase family AAA domain-containing protein 3B OS=Homo sapiens GN=ATAD3B	0.9933	-0.76121314
Q15061	WDR43_HUMAN	WD repeat-containing protein 43 OS=Homo sapiens GN=WDR43 PE=1 SV=3	1	-0.785875195
O75376-2	NCOR1_HUMAN	Isoform 2 of Nuclear receptor corepressor 1 OS=Homo sapiens GN=NCOR1	0.9813	-0.810966176
Q92665	RT31_HUMAN	28S ribosomal protein S31	1	-0.862496476
O75964	ATP5L_HUMAN	ATP synthase subunit g	1	-0.915935735
Q00325-2	MPCP_HUMAN	Isoform B of Phosphate carrier protein	0.9999	-0.915935735
Q86UP3-2	ZFH4_HUMAN	Isoform 2 of Zinc finger homeobox protein 4 OS=Homo sapiens GN=ZFH4	0.9999	-0.915935735
Q8IV32	CCD71_HUMAN	Coiled-coil domain-containing protein 71 OS=Homo sapiens GN=CCDC71 PE=2 SV=3	0.9958	-0.943416472
P82914	RT15_HUMAN	28S ribosomal protein S15	0.8345	-0.943416472
Q9H8H3	MET7A_HUMAN	Methyltransferase-like protein 7A OS=Homo sapiens GN=METTL7A PE=1 SV=1	0.9502	-1
P51398-2	RT29_HUMAN	Isoform 2 of 28S ribosomal protein S29	0.9998	-1.029146346
Q9Y3D9	RT23_HUMAN	28S ribosomal protein S23	0.9997	-1.029146346
Q9UPX0	TUTLB_HUMAN	Protein turtle homolog B OS=Homo sapiens GN=IGSF9B PE=2 SV=2	0.9601	-1.029146346
P60866-2	RS20_HUMAN	Isoform 2 of 40S ribosomal protein S20 OS=Homo sapiens GN=RPS20	0.9997	-1.058893689
P19404	NDUV2_HUMAN	NADH dehydrogenase [ubiquinone] flavoprotein 2	0.9985	-1.120294234
Q9P0J0	NDUAD_HUMAN	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 13 OS=Homo sapiens GN=NDUFA13 PE=1 SV=1	0.8924	-1.217591435
P21675-2	TAF1_HUMAN	Isoform 2 of Transcription initiation factor TFIID subunit 1 OS=Homo sapiens GN=TAF1	0.957	-1.358453971
Q9NQ50	RM40_HUMAN	39S ribosomal protein L40	0.8433	-1.434402824
Q9NX20	RM16_HUMAN	39S ribosomal protein L16	0.91	-1.473931188
Q9Y570-2	PPME1_HUMAN	Isoform 2 of Protein phosphatase methylesterase 1 OS=Homo sapiens GN=PPME1	1	-1.59946207
E5RH44	E5RH44_HUMAN	RB1-inducible coiled-coil protein 1 (Fragment) OS=Homo sapiens GN=RB1CC1 PE=4 SV=1	0.8737	-1.689659879
Q9P0L0-2	VAPA_HUMAN	Isoform 2 of Vesicle-associated membrane protein-associated protein A OS=Homo sapiens GN=VAPA	0.9921	-1.736965594
Q9HD33-2	RM47_HUMAN	Isoform 2 of 39S ribosomal protein L47	0.8345	-1.736965594
Q96JN8-2	NEUL4_HUMAN	Isoform 2 of Neuralized-like protein 4 OS=Homo sapiens GN=NEURL4	0.9995	-2.058893689
P0C671	CF222_HUMAN	Uncharacterized protein C6orf222 OS=Homo sapiens GN=C6orf222 PE=2 SV=1	0.871	-2.058893689
Q9Y237-2	PIN4_HUMAN	Isoform 2 of Peptidyl-prolyl cis-trans isomerase NIMA-interacting 4 OS=Homo sapiens GN=PIN4	1	-2.184424571
Q9BYD1	RM13_HUMAN	39S ribosomal protein L13	0.9894	-2.251538767
P56715	RP1_HUMAN	Oxygen-regulated protein 1 OS=Homo sapiens GN=RP1 PE=1 SV=1	0.9721	-2.251538767
Q92626	PXDN_HUMAN	Peroxidasin homolog OS=Homo sapiens GN=PXDN PE=1 SV=2	0.9303	-2.321928095
I3L4Z2	I3L4Z2_HUMAN	Uncharacterized protein OS=Homo sapiens GN=PE=4 SV=1	0.9495	-2.556393349
E9PEI6	E9PEI6_HUMAN	Diffuse panbronchiolitis critical region protein 1 OS=Homo sapiens GN=DPCR1 PE=4 SV=1	0.9309	-2.556393349
P02647	APOA1_HUMAN	Apolipoprotein A-I OS=Homo sapiens GN=APOA1 PE=1 SV=1	0.9364	-2.64385619
Q7Z589	EMSY_HUMAN	Protein EMSY OS=Homo sapiens GN=EMSY PE=1 SV=2	0.9635	-4.321928095
Q5T5P2-7	SKT_HUMAN	Isoform 5 of Sickle tail protein homolog OS=Homo sapiens GN=KIAA1217	0.9856	-5.058893689
Q5CZC0-2	FSIP2_HUMAN	Isoform 2 of Fibrous sheath-interacting protein 2 OS=Homo sapiens GN=FSIP2	0.9908	-6.64385619