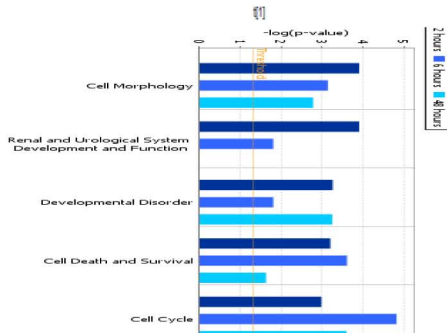
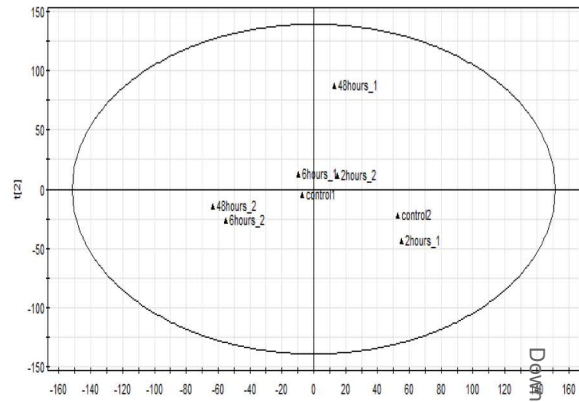
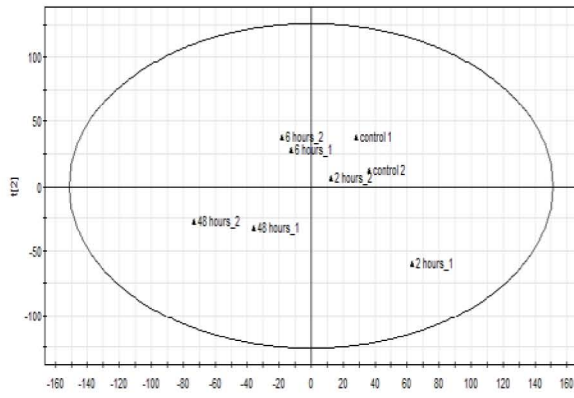
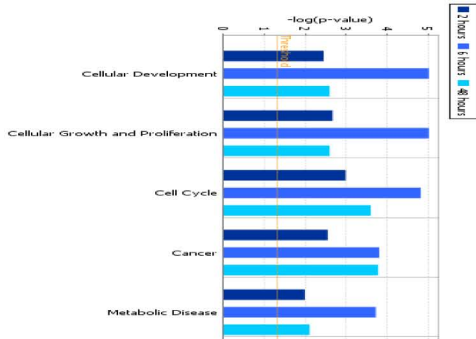
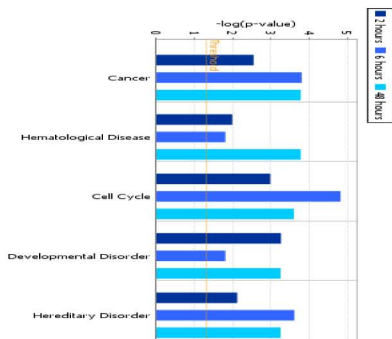
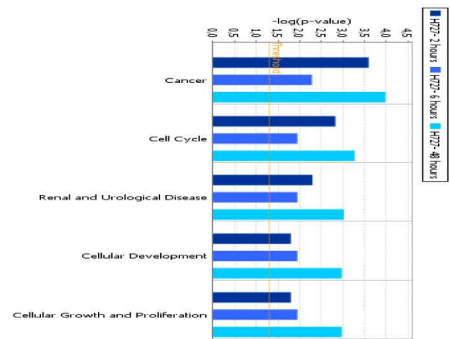
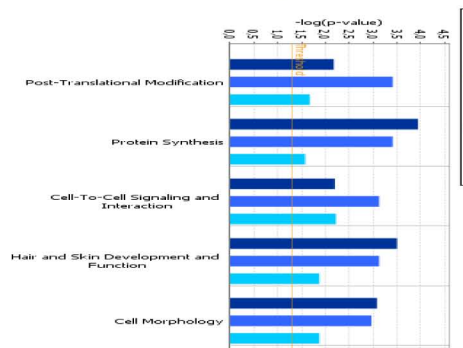
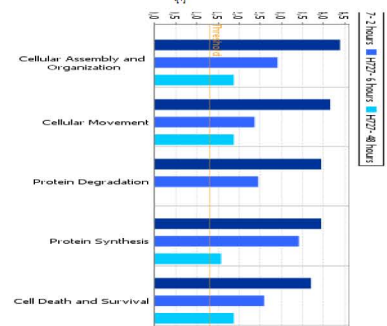


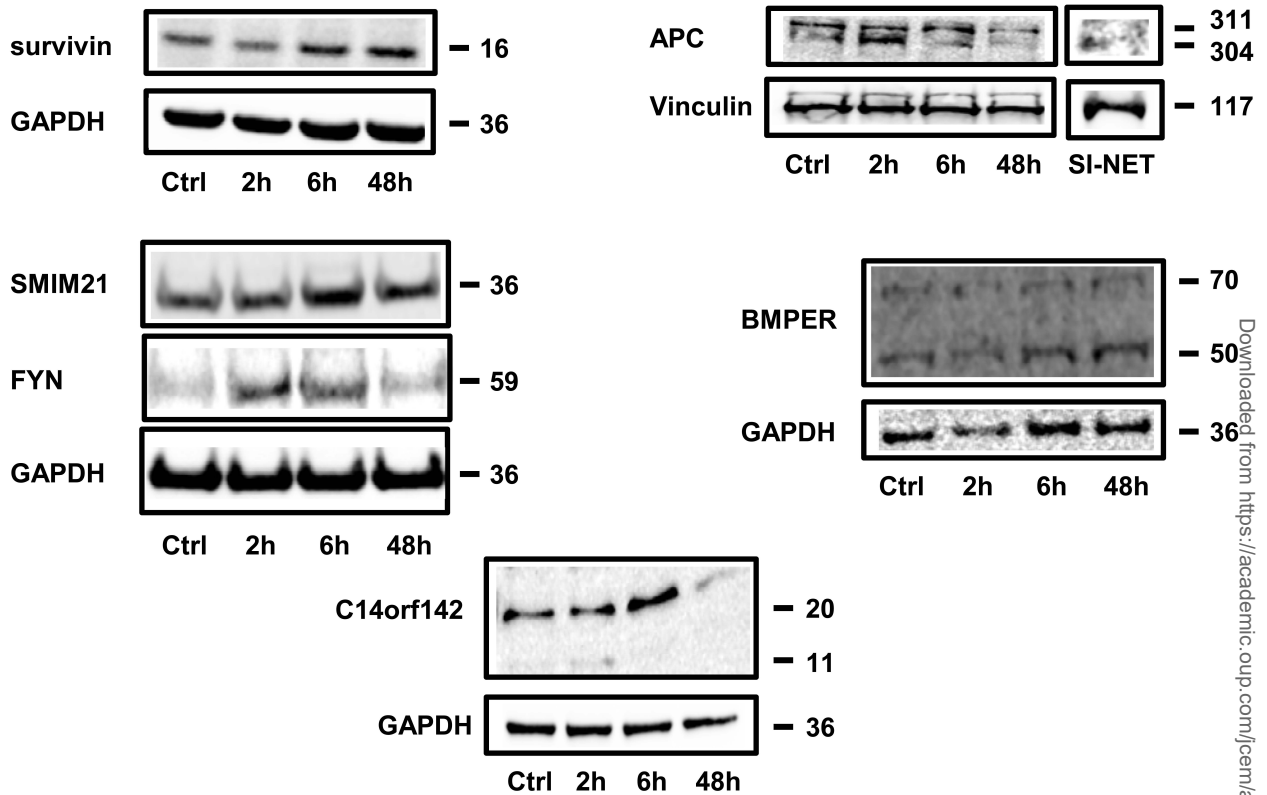
Supplementary Figure S1. Expression of the SSTR2 and SSTR5 proteins and the SSTR1-5 genes. (A) Photomicrographs showing immunocytochemistry analysis of NET cell lines and the primary SI-NET culture. Immunohistochemistry of normal pancreas shows expression in Langerhans islets as a positive control. Magnification: 160x in HC45 and primary SI-NET or 400x in the other cases. **(B)** Relative mRNA expression levels were determined by qRT-PCR and normalized to the endogenous control *ACTB*.

HC45**H727****2 hours****6 hours****48 hours**

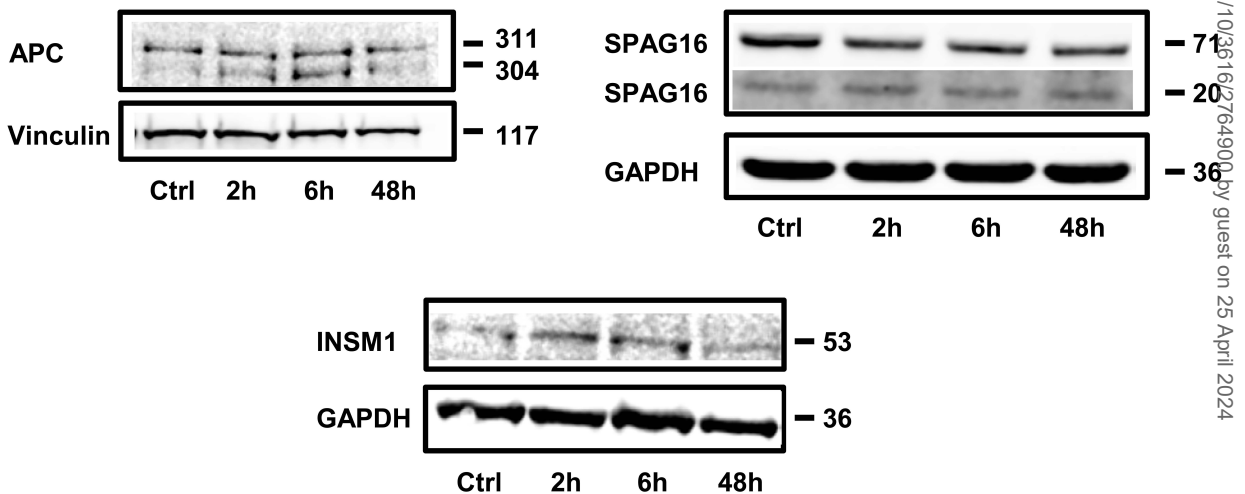
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Supplementary Figure S2. Principal component analysis (PCA) and pathways identified by Ingenuity analyses after lanreotide treatment. Top: PCA was performed for all proteins quantified in controls and after lanreotide treatment at 2, 6 and 48 hours for the HC45 and H727 cell lines. For each condition two replicates are shown. Below: The top five pathways are shown for altered proteins at 2, 6 and 48 hours lanreotide treatment of HC45 and H727 cells.

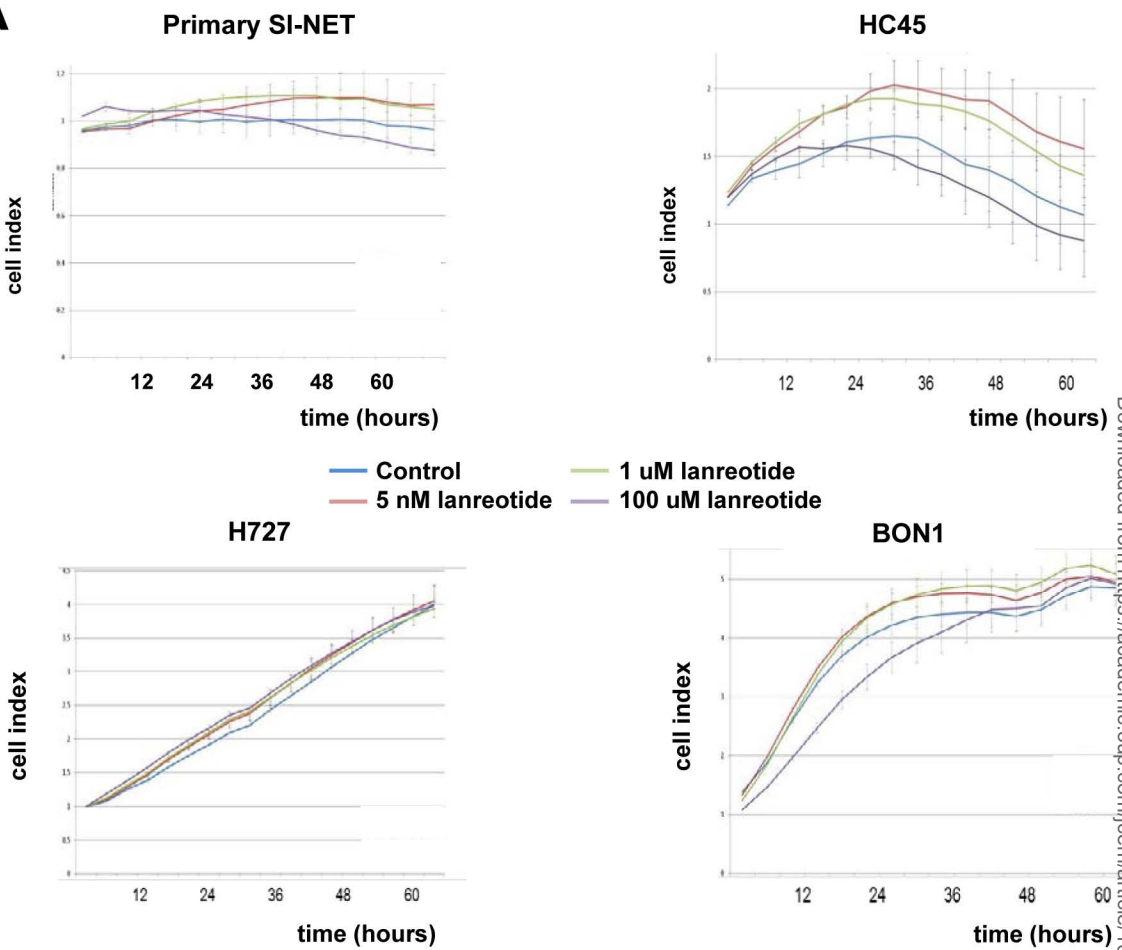
A HC45



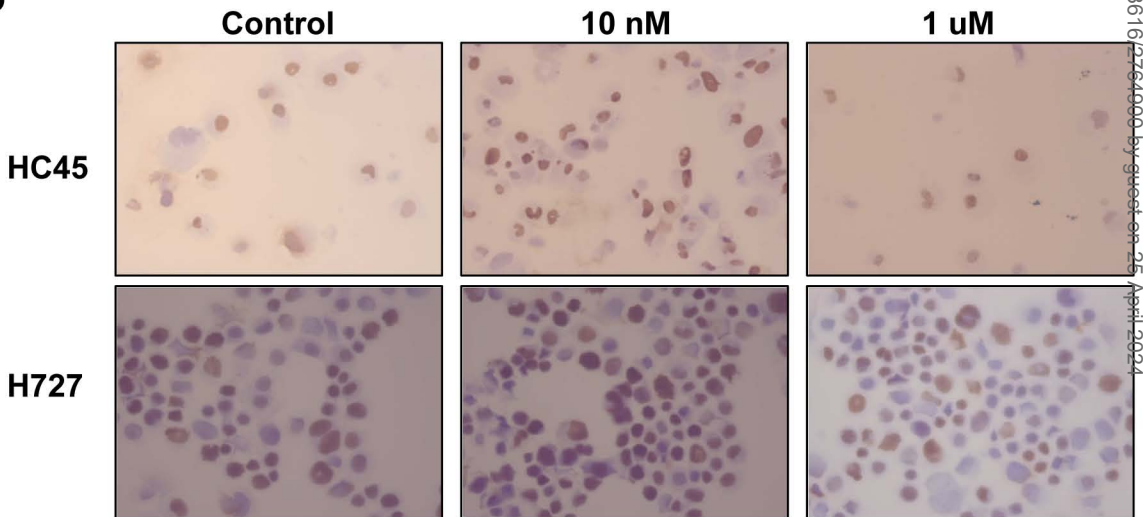
B H727



Supplementary Figure S3. Western blot analysis of selected deregulated proteins identified by proteomics experiments. Detection of APC, SMIM21, survivin, C14orf142 and BMPER in HC45 (A) and of APC, SPAG16 and INSM in H727 (B) cell lines. Protein sizes are given to the right in kDa. Results are shown for untreated control cells and after 2, 6, and 48 hours of 10 nM lanreotide treatment. Analyses of survivin in HC45 at 48 hours treatment with lanreotide at different concentrations are also shown in (A). The antibody EP701Y was used for detection of APC. GAPDH and Vinculin were used as loading controls.

A

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B

Supplementary Figure S4. Analysis of proliferation and MIB-1 staining of cell lines after lanreotide treatment. (A) Time resolution xCELLigence was used to assess possible effects on proliferation in lanreotide treated cells and untreated controls.

(B) MIB1 immunocytochemistry was performed for HC45 and H727 in control cells (without treatment) and after 9 days of treatment with lanreotide at different concentrations.

Magnification: 160× and 400× for HC45 or H727

Supplementary Table S1. Summary of clinical characteristics for NET cases (n=112) included in the TMA.

Parameter	observations	(%)
Gender		
female	55	(49.1)
male	57	(50.9)
Age		
median years	55	
range years	12 - 91	
Primary tumor site		
pancreas	28	(25.0)
small bowel	28	(25.0)
stomach	1	(0.9)
appendix	43	(38.4)
colon and rectum	7	(6.3)
unknown origin	5	(4.5)
Functionality		
functional	7	(6.3)
non-functional	53	(47.3)
unknown	52	(46.4)
Grade		
Grade 1	74	(66.1)
Grade 2	11	(9.8)
Grade 3	8	(7.1)
unknown	19	(17.0)
Stage		
Stage I	34	(30.4)
Stage II	17	(15.2)
Stage III	17	(15.2)
Stage IV	17	(15.2)
unknown	27	(24.1)
T classification		
T1	20	(17.9)
T2	23	(20.5)
T3	21	(18.8)
T4	18	(16.1)
unknown	30	(26.8)
Proliferation index (Ki-67)		
<2%	67	(59.8)
2-20%	13	(11.6)
>20%	7	(6.3)
unknown	25	(22.3)
Lymph node involvement		

present	23	(20.5)
not identified	58	(51.8)
unknown	31	(27.7)
<i>Distant metastases</i>		
present	17	(15.2)
not identified	95	(84.8)
<i>Relapse</i>		
no	84	(75.0)
yes	24	(21.4)
unknown	4	(3.6)
<i>Death</i>		
no	87	(77.7)
yes	23	(20.5)
unknown	2	(1.8)
<i>Follow up</i>		
median years	12	
range years	1-34	

Supplementary Table S3. Differentially expressed proteins after lanreotide treatment.

Accession number	Gene symbol	Expression vs. Control	P-value
HC45 cells - 2 hours lanreotide (178 proteins)			
Q9H147	Deoxynucleotidyltransferase terminal-interacting protein 1 OS=Homo sapiens GN=DNTTIP1 PE=1 SV=2 - [TDIF1_HUMAN]	0.574	0.042
P62714	Serine/threonine-protein phosphatase 2A catalytic subunit beta isoform OS=Homo sapiens GN=PPP2CB PE=1 SV=1 - [PP2AB_HUMAN]	0.590	0.002
P01258	Calcitonin OS=Homo sapiens GN=CALCA PE=1 SV=2 - [CALC_HUMAN]	0.687	0.022
O60573	Eukaryotic translation initiation factor 4E type 2 OS=Homo sapiens GN=EIF4E2 PE=1 SV=1 - [IF4E2_HUMAN]	0.764	0.013
Q9UPW8	Protein unc-13 homolog A OS=Homo sapiens GN=UNC13A PE=2 SV=4 - [UN13A_HUMAN]	0.785	0.013
O43852	Calumenin OS=Homo sapiens GN=CALU PE=1 SV=2 - [CALU_HUMAN]	0.808	0.002
Q96HR8	H/ACA ribonucleoprotein complex non-core subunit NAF1 OS=Homo sapiens GN=NAF1 PE=1 SV=2 - [NAF1_HUMAN]	0.817	0.029
P10244	Myb-related protein B OS=Homo sapiens GN=MYBL2 PE=1 SV=1 - [MYBB_HUMAN]	0.819	0.036
P10114	Ras-related protein Rap-2a OS=Homo sapiens GN=RAP2A PE=1 SV=1 - [RAP2A_HUMAN]	0.824	0.020
Q13557-12	Isoform Delta 12 of Calcium/calmodulin-dependent protein kinase type II subunit delta OS=Homo sapiens GN=CAMK2D - [KCC2D_HUMAN]	0.826	0.030
P62633-3	Isoform 3 of Cellular nucleic acid-binding protein OS=Homo sapiens GN=CNBP - [CNBP_HUMAN]	0.829	0.039
P0C0S5	Histone H2A.Z OS=Homo sapiens GN=H2AFZ PE=1 SV=2 - [H2AZ_HUMAN]	0.832	0.025
Q15814	Tubulin-specific chaperone C OS=Homo sapiens GN=TBCC PE=1 SV=2 - [TBCC_HUMAN]	0.836	0.027
Q9UBL3-2	Isoform 2 of Set1/Ash2 histone methyltransferase complex subunit ASH2 OS=Homo sapiens GN=ASH2L - [ASH2L_HUMAN]	0.839	0.033
Q8N3Y1	F-box/WD repeat-containing protein 8 OS=Homo sapiens GN=FBXW8 PE=1 SV=2 - [FBXW8_HUMAN]	0.845	0.011
P15408-3	Isoform 3 of Fos-related antigen 2 OS=Homo sapiens GN=FOSL2 - [FOSL2_HUMAN]	0.846	0.008
Q9GZY4	Uncharacterized protein C7orf44 OS=Homo sapiens GN=C7orf44 PE=1 SV=1 - [CG044_HUMAN]	0.846	0.023
P15336	Cyclic AMP-dependent transcription factor ATF-2 OS=Homo sapiens GN=ATF2 PE=1 SV=4 - [ATF2_HUMAN]	0.849	0.044
O43181	NADH dehydrogenase [ubiquinone] iron-sulfur protein 4, mitochondrial OS=Homo sapiens GN=NDUFS4 PE=1 SV=1 - [NDUS4_HUMAN]	0.852	0.037
O75695	Protein XRP2 OS=Homo sapiens GN=RP2 PE=1 SV=4 - [XRP2_HUMAN]	0.853	0.037
Q96BP2	Coiled-coil-helix-coiled-coil-helix domain-containing protein 1 OS=Homo sapiens GN=CHCHD1 PE=1 SV=1 - [CHCH1_HUMAN]	0.854	0.040
P13726	Tissue factor OS=Homo sapiens GN=F3 PE=1 SV=1 - [TF_HUMAN]	0.859	0.021
Q6ZUT1	Uncharacterized protein C11orf57 OS=Homo sapiens GN=C11orf57 PE=1 SV=2 - [CK057_HUMAN]	0.863	0.017
Q9P2K5-2	Isoform 2 of Myelin expression factor 2 OS=Homo sapiens GN=MYEF2 - [MYEF2_HUMAN]	0.875	0.028

P35749	Myosin-11 OS=Homo sapiens GN=MYH11 PE=1 SV=3 - [MYH11_HUMAN]	0.877	0.039
P05455	Lupus La protein OS=Homo sapiens GN=SSB PE=1 SV=2 - [LA_HUMAN]	0.877	0.026
Q12933-4	Isoform 4 of TNF receptor-associated factor 2 OS=Homo sapiens GN=TRAF2 - [TRAF2_HUMAN]	0.881	0.008
Q6PI78	Transmembrane protein 65 OS=Homo sapiens GN=TMEM65 PE=1 SV=2 - [TMM65_HUMAN]	0.882	0.019
O75367-3	Isoform 3 of Core histone macro-H2A.1 OS=Homo sapiens GN=H2AFY - [H2AY_HUMAN]	0.883	0.037
Q9HBD1-6	Isoform 6 of RING finger and CCCH-type zinc finger domain-containing protein 2 OS=Homo sapiens GN=RC3H2 - [RC3H2_HUMAN]	0.883	0.009
Q68EM7-2	Isoform 2 of Rho GTPase-activating protein 17 OS=Homo sapiens GN=ARHGAP17 - [RHG17_HUMAN]	0.884	0.024
Q9Y673	Dolichyl-phosphate beta-glucosyltransferase OS=Homo sapiens GN=ALG5 PE=1 SV=1 - [ALG5_HUMAN]	0.890	0.044
Q8N4V1	Membrane magnesium transporter 1 OS=Homo sapiens GN=MMGT1 PE=1 SV=1 - [MMGT1_HUMAN]	0.890	0.017
P01111	GTPase NRas OS=Homo sapiens GN=NRAS PE=1 SV=1 - [RASN_HUMAN]	0.892	0.019
Q96GZ6-6	Isoform 6 of Solute carrier family 41 member 3 OS=Homo sapiens GN=SLC41A3 - [S41A3_HUMAN]	0.893	0.039
Q86Y91-2	Isoform 2 of Kinesin-like protein KIF18B OS=Homo sapiens GN=KIF18B - [KI18B_HUMAN]	0.895	0.035
Q9BZK7	F-box-like/WD repeat-containing protein TBL1XR1 OS=Homo sapiens GN=TBL1XR1 PE=1 SV=1 - [TBL1R_HUMAN]	0.899	0.025
O43719	HIV Tat-specific factor 1 OS=Homo sapiens GN=HTATSF1 PE=1 SV=1 - [HTSF1_HUMAN]	0.899	0.016
Q9H0L4	Cleavage stimulation factor subunit 2 tau variant OS=Homo sapiens GN=CSTF2T PE=1 SV=1 - [CSTFT_HUMAN]	0.900	0.001
Q9Y2R0	Coiled-coil domain-containing protein 56 OS=Homo sapiens GN=CCDC56 PE=1 SV=1 - [CCD56_HUMAN]	0.902	0.013
Q15019	Septin-2 OS=Homo sapiens GN=SEPT2 PE=1 SV=1 - [SEPT2_HUMAN]	0.903	0.041
Q9H5Q4	Dimethyladenosine transferase 2, mitochondrial OS=Homo sapiens GN=TFB2M PE=1 SV=1 - [TFB2M_HUMAN]	0.903	0.016
Q96KQ7-2	Isoform 2 of Histone-lysine N-methyltransferase EHMT2 OS=Homo sapiens GN=EHMT2 - [EHMT2_HUMAN]	0.903	0.042
Q8NBN3-3	Isoform 3 of Transmembrane protein 87A OS=Homo sapiens GN=TMEM87A - [TM87A_HUMAN]	0.904	0.044
P26440	Isovaleryl-CoA dehydrogenase, mitochondrial OS=Homo sapiens GN=IVD PE=1 SV=1 - [IVD_HUMAN]	0.904	0.014
Q9NVP1	ATP-dependent RNA helicase DDX18 OS=Homo sapiens GN=DDX18 PE=1 SV=2 - [DDX18_HUMAN]	0.904	0.045
Q9P0L0	Vesicle-associated membrane protein-associated protein A OS=Homo sapiens GN=VAPA PE=1 SV=3 - [VAPA_HUMAN]	0.904	0.035
P41227	N-alpha-acetyltransferase 10 OS=Homo sapiens GN=NAA10 PE=1 SV=1 - [NAA10_HUMAN]	0.904	0.004
P52895	Aldo-keto reductase family 1 member C2 OS=Homo sapiens GN=AKR1C2 PE=1 SV=3 - [AK1C2_HUMAN]	0.905	0.042
Q9H2J4	Phosducin-like protein 3 OS=Homo sapiens GN=PDCL3 PE=1 SV=1 - [PDCL3_HUMAN]	0.906	0.048
P08195	4F2 cell-surface antigen heavy chain OS=Homo sapiens GN=SLC3A2 PE=1 SV=3 - [4F2_HUMAN]	0.906	0.029
Q9H082	Ras-related protein Rab-33B OS=Homo sapiens GN=RAB33B PE=1 SV=1 - [RB33B_HUMAN]	0.908	0.032
Q3KQV9	UDP-N-acetylhexosamine pyrophosphorylase-like protein 1 OS=Homo sapiens GN=UAP1L1 PE=2 SV=2 - [UAP1L_HUMAN]	0.910	0.027

Q9Y314	Nitric oxide synthase-interacting protein OS=Homo sapiens GN=NOSIP PE=1 SV=1 - [NOSIP_HUMAN]	0.914	0.042
P50402	Emerin OS=Homo sapiens GN=EMD PE=1 SV=1 - [EMD_HUMAN]	0.919	0.014
P27824	Calnexin OS=Homo sapiens GN=CANX PE=1 SV=2 - [CALX_HUMAN]	0.920	0.031
P52565	Rho GDP-dissociation inhibitor 1 OS=Homo sapiens GN=ARHGDI1 PE=1 SV=3 - [GDIR1_HUMAN]	0.923	0.027
Q5H9R7-3	Isoform 3 of Serine/threonine-protein phosphatase 6 regulatory subunit 3 OS=Homo sapiens GN=PPP6R3 - [PP6R3_HUMAN]	0.924	0.028
P37235	Hippocalcin-like protein 1 OS=Homo sapiens GN=HPCAL1 PE=1 SV=3 - [HPCL1_HUMAN]	0.924	0.039
Q99653	Calcium-binding protein p22 OS=Homo sapiens GN=CHP PE=1 SV=3 - [CHP1_HUMAN]	0.924	0.016
O15270	Serine palmitoyltransferase 2 OS=Homo sapiens GN=SPTLC2 PE=1 SV=1 - [SPTC2_HUMAN]	0.925	0.043
Q9NYP7	Elongation of very long chain fatty acids protein 5 OS=Homo sapiens GN=ELOVL5 PE=1 SV=1 - [ELOV5_HUMAN]	0.926	0.039
Q14681	BTB/POZ domain-containing protein KCTD2 OS=Homo sapiens GN=KCTD2 PE=1 SV=3 - [KCTD2_HUMAN]	0.928	0.022
Q0ZGT2-2	Isoform 2 of Nexilin OS=Homo sapiens GN=NEXN - [NEXN_HUMAN]	0.929	0.028
O14776-2	Isoform 2 of Transcription elongation regulator 1 OS=Homo sapiens GN=TCERG1 - [TCRG1_HUMAN]	0.930	0.044
O14828	Secretory carrier-associated membrane protein 3 OS=Homo sapiens GN=SCAMP3 PE=1 SV=3 - [SCAM3_HUMAN]	0.930	0.032
Q8NEY8-6	Isoform 6 of Periphilin-1 OS=Homo sapiens GN=PPHLN1 - [PPHLN_HUMAN]	0.931	0.044
P35580	Myosin-10 OS=Homo sapiens GN=MYH10 PE=1 SV=3 - [MYH10_HUMAN]	0.932	0.010
P68431	Histone H3.1 OS=Homo sapiens GN=HIST1H3A PE=1 SV=2 - [H31_HUMAN]	0.934	0.012
Q14669	Probable E3 ubiquitin-protein ligase TRIP12 OS=Homo sapiens GN=TRIP12 PE=1 SV=1 - [TRIPC_HUMAN]	0.934	0.005
P13995	Bifunctional methylenetetrahydrofolate dehydrogenase/cyclohydrolase, mitochondrial OS=Homo sapiens GN=MTHFD2 PE=1 SV=2 - [MTDC_HUMAN]	0.935	0.035
Q9UGP8	Translocation protein SEC63 homolog OS=Homo sapiens GN=SEC63 PE=1 SV=2 - [SEC63_HUMAN]	0.940	0.029
Q93009	Ubiquitin carboxyl-terminal hydrolase 7 OS=Homo sapiens GN=USP7 PE=1 SV=2 - [UBP7_HUMAN]	0.949	0.018
P06396-2	Isoform 2 of Gelsolin OS=Homo sapiens GN=GSN - [GELS_HUMAN]	0.949	0.038
Q9H4A4	Aminopeptidase B OS=Homo sapiens GN=RNPEP PE=1 SV=2 - [AMPB_HUMAN]	0.954	0.045
P43034	Platelet-activating factor acetylhydrolase IB subunit alpha OS=Homo sapiens GN=PFAH1B1 PE=1 SV=2 - [LIS1_HUMAN]	0.954	0.020
Q96KB5	Lymphokine-activated killer T-cell-originated protein kinase OS=Homo sapiens GN=PBK PE=1 SV=3 - [TOPK_HUMAN]	0.954	0.029
Q13310-2	Isoform 2 of Polyadenylate-binding protein 4 OS=Homo sapiens GN=PABPC4 - [PABP4_HUMAN]	0.957	0.008
P55884	Eukaryotic translation initiation factor 3 subunit B OS=Homo sapiens GN=EIF3B PE=1 SV=3 - [EIF3B_HUMAN]	0.957	0.044
Q15435	Protein phosphatase 1 regulatory subunit 7 OS=Homo sapiens GN=PPP1R7 PE=1 SV=1 - [PP1R7_HUMAN]	0.957	0.049
O14745	Na(+)/H(+) exchange regulatory cofactor NHE-RF1 OS=Homo sapiens GN=SLC9A3R1 PE=1 SV=4 - [NHRF1_HUMAN]	0.960	0.020
P06733	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2 - [ENOA_HUMAN]	0.963	0.005

O15371	Eukaryotic translation initiation factor 3 subunit D OS=Homo sapiens GN=EIF3D PE=1 SV=1 - [EIF3D_HUMAN]	0.964	0.038
P31751	RAC-beta serine/threonine-protein kinase OS=Homo sapiens GN=AKT2 PE=1 SV=2 - [AKT2_HUMAN]	0.965	0.044
P01137	Transforming growth factor beta-1 OS=Homo sapiens GN=TGFB1 PE=1 SV=2 - [TGFB1_HUMAN]	0.967	0.037
Q9Y2L9-2	Isoform 2 of Leucine-rich repeat and calponin homology domain-containing protein 1 OS=Homo sapiens GN=LRCH1 - [LRCH1_HUMAN]	0.968	0.048
Q96EU7	C1GALT1-specific chaperone 1 OS=Homo sapiens GN=C1GALT1C1 PE=1 SV=1 - [C1GLC_HUMAN]	0.970	0.007
P14866	Heterogeneous nuclear ribonucleoprotein L OS=Homo sapiens GN=HNRNPL PE=1 SV=2 - [HNRPL_HUMAN]	0.972	0.023
Q00839-2	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU - [HNRPU_HUMAN]	0.972	0.023
Q13459-2	Isoform Short of Myosin-IXb OS=Homo sapiens GN=MYO9B - [MYO9B_HUMAN]	0.973	0.022
Q92609	TBC1 domain family member 5 OS=Homo sapiens GN=TBC1D5 PE=1 SV=1 - [TBCD5_HUMAN]	0.974	0.019
O43707	Alpha-actinin-4 OS=Homo sapiens GN=ACTN4 PE=1 SV=2 - [ACTN4_HUMAN]	0.976	0.023
Q9NZN5-2	Isoform 2 of Rho guanine nucleotide exchange factor 12 OS=Homo sapiens GN=ARHGEF12 - [ARHGC_HUMAN]	0.977	0.035
O00541-2	Isoform 2 of Pescadillo homolog OS=Homo sapiens GN=PES1 - [PESC_HUMAN]	0.979	0.015
P22314	Ubiquitin-like modifier-activating enzyme 1 OS=Homo sapiens GN=UBA1 PE=1 SV=3 - [UBA1_HUMAN]	0.987	0.045
Q99575	Ribonucleases P/MRP protein subunit POP1 OS=Homo sapiens GN=POP1 PE=1 SV=2 - [POP1_HUMAN]	1.017	0.022
P04632	Calpain small subunit 1 OS=Homo sapiens GN=CAPNS1 PE=1 SV=1 - [CPNS1_HUMAN]	1.019	0.033
P42345	Serine/threonine-protein kinase mTOR OS=Homo sapiens GN=MTOR PE=1 SV=1 - [MTOR_HUMAN]	1.025	0.006
P17987	T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1 - [TCPA_HUMAN]	1.026	0.023
P30046	D-dopachrome decarboxylase OS=Homo sapiens GN=DDT PE=1 SV=3 - [DOPD_HUMAN]	1.029	0.045
P30153	Serine/threonine-protein phosphatase 2A 65 kDa regulatory subunit A alpha isoform OS=Homo sapiens GN=PPP2R1A PE=1 SV=4 - [2AAA_HUMAN]	1.029	0.025
Q8N556	Actin filament-associated protein 1 OS=Homo sapiens GN=AFAP1 PE=1 SV=2 - [AFAP1_HUMAN]	1.031	0.010
Q6UX07-2	Isoform 2 of Dehydrogenase/reductase SDR family member 13 OS=Homo sapiens GN=DHRS13 - [DHR13_HUMAN]	1.032	0.021
Q9Y5P4-2	Isoform 2 of Collagen type IV alpha-3-binding protein OS=Homo sapiens GN=COL4A3BP - [C43BP_HUMAN]	1.035	0.038
Q8N3R9	MAGUK p55 subfamily member 5 OS=Homo sapiens GN=MPP5 PE=1 SV=3 - [MPP5_HUMAN]	1.037	0.007
Q9NSE4	Isoleucine--tRNA ligase, mitochondrial OS=Homo sapiens GN=IARS2 PE=1 SV=2 - [SYIM_HUMAN]	1.038	0.012
Q03518	Antigen peptide transporter 1 OS=Homo sapiens GN=TAP1 PE=1 SV=2 - [TAP1_HUMAN]	1.046	0.028
P45974-2	Isoform Short of Ubiquitin carboxyl-terminal hydrolase 5 OS=Homo sapiens GN=USP5 - [UBP5_HUMAN]	1.048	0.036
Q00613-2	Isoform Short of Heat shock factor protein 1 OS=Homo sapiens GN=HSF1 - [HSF1_HUMAN]	1.050	0.023
Q08AD1-2	Isoform 2 of Calmodulin-regulated spectrin-associated protein 2 OS=Homo sapiens GN=CAMSAP2 - [CAMP2_HUMAN]	1.051	0.039
Q08431-3	Isoform 3 of Lactadherin OS=Homo sapiens GN=MFGE8 - [MFGM_HUMAN]	1.054	0.046

Q96CN9	GRIP and coiled-coil domain-containing protein 1 OS=Homo sapiens GN=GCC1 PE=1 SV=1 - [GCC1_HUMAN]	1.056	0.015
P62995-3	Isoform 3 of Transformer-2 protein homolog beta OS=Homo sapiens GN=TRA2B - [TRA2B_HUMAN]	1.057	0.040
P08910	Abhydrolase domain-containing protein 2 OS=Homo sapiens GN=ABHD2 PE=2 SV=1 - [ABHD2_HUMAN]	1.060	0.022
Q9NTZ6	RNA-binding protein 12 OS=Homo sapiens GN=RBM12 PE=1 SV=1 - [RBM12_HUMAN]	1.060	0.015
Q13426-2	Isoform 2 of DNA repair protein XRCC4 OS=Homo sapiens GN=XRCC4 - [XRCC4_HUMAN]	1.062	0.005
P27144	Adenylate kinase isoenzyme 4, mitochondrial OS=Homo sapiens GN=AK4 PE=1 SV=1 - [KAD4_HUMAN]	1.063	0.007
Q15185	Prostaglandin E synthase 3 OS=Homo sapiens GN=PTGES3 PE=1 SV=1 - [TEBP_HUMAN]	1.063	0.048
P56962	Syntaxin-17 OS=Homo sapiens GN=STX17 PE=1 SV=2 - [STX17_HUMAN]	1.066	0.036
Q9BVM4	Gamma-glutamylaminocyclotransferase OS=Homo sapiens GN=A2LD1 PE=1 SV=2 - [A2LD1_HUMAN]	1.067	0.028
P62136	Serine/threonine-protein phosphatase PP1-alpha catalytic subunit OS=Homo sapiens GN=PPP1CA PE=1 SV=1 - [PP1A_HUMAN]	1.067	0.044
Q9NW82	WD repeat-containing protein 70 OS=Homo sapiens GN=WDR70 PE=1 SV=1 - [WDR70_HUMAN]	1.072	0.014
Q96CW6	Probable RNA polymerase II nuclear localization protein SLC7A6 OS=Homo sapiens GN=SLC7A6OS PE=1 SV=2 - [S7A6O_HUMAN]	1.074	0.006
P51812	Ribosomal protein S6 kinase alpha-3 OS=Homo sapiens GN=RPS6KA3 PE=1 SV=1 - [KS6A3_HUMAN]	1.074	0.012
Q96QZ7-3	Isoform 3 of Membrane-associated guanylate kinase, WW and PDZ domain-containing protein 1 OS=Homo sapiens GN=MAGI1 - [MAGI1_HUMAN]	1.076	0.035
Q969F9-2	Isoform 2 of Hermansky-Pudlak syndrome 3 protein OS=Homo sapiens GN=HPS3 - [HPS3_HUMAN]	1.080	0.022
O96033	Molybdopterine synthase sulfur carrier subunit OS=Homo sapiens GN=MOCS2 PE=1 SV=1 - [MOC2A_HUMAN]	1.081	0.039
Q8N653	Leucine-zipper-like transcriptional regulator 1 OS=Homo sapiens GN=LZTR1 PE=2 SV=2 - [LZTR1_HUMAN]	1.082	0.005
O15111	Inhibitor of nuclear factor kappa-B kinase subunit alpha OS=Homo sapiens GN=CHUK PE=1 SV=2 - [IKKA_HUMAN]	1.086	0.020
Q13098-5	Isoform 4 of COP9 signalosome complex subunit 1 OS=Homo sapiens GN=GPS1 - [CSN1_HUMAN]	1.089	0.020
Q8TDY2-2	Isoform 2 of RB1-inducible coiled-coil protein 1 OS=Homo sapiens GN=RB1CC1 - [RBCC1_HUMAN]	1.094	0.010
P48553	Trafficking protein particle complex subunit 10 OS=Homo sapiens GN=TRAPPC10 PE=1 SV=2 - [TPC10_HUMAN]	1.095	0.034
P47914	60S ribosomal protein L29 OS=Homo sapiens GN=RPL29 PE=1 SV=2 - [RL29_HUMAN]	1.096	0.031
P14678-2	Isoform SM-B of Small nuclear ribonucleoprotein-associated proteins B and B' OS=Homo sapiens GN=SNRNP - [RSMB_HUMAN]	1.099	0.022
Q96ED9-2	Isoform 2 of Protein Hook homolog 2 OS=Homo sapiens GN=HOOK2 - [HOOK2_HUMAN]	1.104	0.033
P61803	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit DAD1 OS=Homo sapiens GN=DAD1 PE=1 SV=3 - [DAD1_HUMAN]	1.110	0.040
P10646-2	Isoform Beta of Tissue factor pathway inhibitor OS=Homo sapiens GN=TFPI - [TFPI1_HUMAN]	1.111	0.033
A2RUC4-2	Isoform 2 of tRNA tryptophan-synthesizing protein 5 OS=Homo sapiens GN=TYW5 - [TYW5_HUMAN]	1.113	0.005
Q9NWB6	Arginine and glutamate-rich protein 1 OS=Homo sapiens GN=ARGLU1 PE=1 SV=1 - [ARGL1_HUMAN]	1.116	0.041
Q9NRH1	Uncharacterized protein C7orf36 OS=Homo sapiens GN=C7orf36 PE=1 SV=1 - [CG036_HUMAN]	1.117	0.027

O75879	Glutamyl-tRNA(Gln) amidotransferase subunit B, mitochondrial OS=Homo sapiens GN=PET112 PE=1 SV=1 - [GATB_HUMAN]	1.119	0.027
Q86WQ0	Nuclear receptor 2C2-associated protein OS=Homo sapiens GN=NR2C2AP PE=1 SV=1 - [NR2CA_HUMAN]	1.122	0.005
Q96LT7-2	Isoform 2 of Uncharacterized protein C9orf72 OS=Homo sapiens GN=C9orf72 - [CI072_HUMAN]	1.123	0.020
Q5TEU4-2	Isoform 2 of Probable methyltransferase C20orf7, mitochondrial OS=Homo sapiens GN=C20orf7 - [CT007_HUMAN]	1.124	0.017
Q9UI26	Importin-11 OS=Homo sapiens GN=IPO11 PE=1 SV=1 - [IPO11_HUMAN]	1.129	0.004
P19256-2	Isoform 2 of Lymphocyte function-associated antigen 3 OS=Homo sapiens GN=CD58 - [LFA3_HUMAN]	1.129	0.048
Q9Y672	Dolichyl pyrophosphate Man9GlcNAc2 alpha-1,3-glucosyltransferase OS=Homo sapiens GN=ALG6 PE=1 SV=1 - [ALG6_HUMAN]	1.131	0.014
Q7Z7G8-2	Isoform 2 of Vacuolar protein sorting-associated protein 13B OS=Homo sapiens GN=VPS13B - [VP13B_HUMAN]	1.140	0.032
Q9Y2D4	Exocyst complex component 6B OS=Homo sapiens GN=EXOC6B PE=1 SV=3 - [EXC6B_HUMAN]	1.144	0.037
Q9Y6B6	GTP-binding protein SAR1b OS=Homo sapiens GN=SAR1B PE=1 SV=1 - [SAR1B_HUMAN]	1.144	0.030
Q9P0U4	CpG-binding protein OS=Homo sapiens GN=CXXC1 PE=1 SV=2 - [CXXC1_HUMAN]	1.146	0.039
Q9UGU0-2	Isoform 2 of Transcription factor 20 OS=Homo sapiens GN=TCF20 - [TCF20_HUMAN]	1.151	0.011
Q96EU6-2	Isoform 2 of Ribosomal RNA processing protein 36 homolog OS=Homo sapiens GN=RRP36 - [RRP36_HUMAN]	1.152	0.033
Q99470	Stromal cell-derived factor 2 OS=Homo sapiens GN=SDF2 PE=1 SV=2 - [SDF2_HUMAN]	1.152	0.025
P49419-2	Isoform 2 of Alpha-aminoadipic semialdehyde dehydrogenase OS=Homo sapiens GN=ALDH7A1 - [AL7A1_HUMAN]	1.161	0.016
Q8NB15-2	Isoform 2 of Zinc finger protein 511 OS=Homo sapiens GN=ZNF511 - [ZN511_HUMAN]	1.165	0.039
Q9P1Z2-4	Isoform 4 of Calcium-binding and coiled-coil domain-containing protein 1 OS=Homo sapiens GN=CALCOCO1 - [CACO1_HUMAN]	1.166	0.025
Q6BDS2	UHRF1-binding protein 1 OS=Homo sapiens GN=UHRF1BP1 PE=1 SV=1 - [URFB1_HUMAN]	1.166	0.044
Q9UKV5	E3 ubiquitin-protein ligase AMFR OS=Homo sapiens GN=AMFR PE=1 SV=2 - [AMFR2_HUMAN]	1.177	0.021
Q13261-8	Isoform 7 of Interleukin-15 receptor subunit alpha OS=Homo sapiens GN=IL15RA - [I15RA_HUMAN]	1.179	0.003
O00522-2	Isoform 2 of Krev interaction trapped protein 1 OS=Homo sapiens GN=KRIT1 - [KRIT1_HUMAN]	1.187	0.045
Q8N7R7-3	Isoform 3 of Cyclin-Y-like protein 1 OS=Homo sapiens GN=CCNYL1 - [CCYL1_HUMAN]	1.192	0.033
Q96P48-2	Isoform 2 of Arf-GAP with Rho-GAP domain, ANK repeat and PH domain-containing protein 1 OS=Homo sapiens GN=ARAP1 - [ARAP1_HUMAN]	1.193	0.037
Q15058	Kinesin-like protein KIF14 OS=Homo sapiens GN=KIF14 PE=1 SV=1 - [KIF14_HUMAN]	1.196	0.027
Q9NYB0	Telomeric repeat-binding factor 2-interacting protein 1 OS=Homo sapiens GN=TERF2IP PE=1 SV=1 - [TE2IP_HUMAN]	1.204	0.016
Q15637-6	Isoform 6 of Splicing factor 1 OS=Homo sapiens GN=SF1 - [SF01_HUMAN]	1.210	0.027
Q9HCP0-2	Isoform 1S of Casein kinase I isoform gamma-1 OS=Homo sapiens GN=CSNK1G1 - [KC1G1_HUMAN]	1.218	0.045
Q8N387	Mucin-15 OS=Homo sapiens GN=MUC15 PE=2 SV=2 - [MUC15_HUMAN]	1.226	0.011
O75907	Diacylglycerol O-acyltransferase 1 OS=Homo sapiens GN=DGAT1 PE=1 SV=2 - [DGAT1_HUMAN]	1.248	0.049

Q9UKX7-2	Isoform 2 of Nuclear pore complex protein Nup50 OS=Homo sapiens GN=NUP50 - [NUP50_HUMAN]	1.255	0.049
Q14653	Interferon regulatory factor 3 OS=Homo sapiens GN=IRF3 PE=1 SV=1 - [IRF3_HUMAN]	1.258	0.022
Q969R8	Integrin-alpha FG-GAP repeat-containing protein 2 OS=Homo sapiens GN=ITFG2 PE=2 SV=1 - [ITFG2_HUMAN]	1.263	0.038
P13051-2	Isoform 1 of Uracil-DNA glycosylase OS=Homo sapiens GN=UNG - [UNG_HUMAN]	1.284	0.046
Q96JP5-2	Isoform 2 of E3 ubiquitin-protein ligase ZFP91 OS=Homo sapiens GN=ZFP91 - [ZFP91_HUMAN]	1.297	0.050
Q9P2W1	Homologous-pairing protein 2 homolog OS=Homo sapiens GN=PSMC3IP PE=1 SV=1 - [HOP2_HUMAN]	1.309	0.029
P25054-2	Isoform Short of Adenomatous polyposis coli protein OS=Homo sapiens GN=APC - [APC_HUMAN]	1.365	0.029
O15145	Actin-related protein 2/3 complex subunit 3 OS=Homo sapiens GN=ARPC3 PE=1 SV=3 - [ARPC3_HUMAN]	1.471	0.013
Q8N8U9	BMP-binding endothelial regulator protein OS=Homo sapiens GN=BMPER PE=1 SV=3 - [BMPER_HUMAN]	1.527	0.020

HC45 cells - 6 hours lanreotide (268 proteins)

P13645	Keratin, type I cytoskeletal 10 OS=Homo sapiens GN=KRT10 PE=1 SV=6 - [K1C10_HUMAN]	0.594	0.035
Q15051-3	Isoform 3 of IQ calmodulin-binding motif-containing protein 1 OS=Homo sapiens GN=IQCB1 - [IQCB1_HUMAN]	0.610	0.019
P13647	Keratin, type II cytoskeletal 5 OS=Homo sapiens GN=KRT5 PE=1 SV=3 - [K2C5_HUMAN]	0.629	0.007
Q9BXV9	Uncharacterized protein C14orf142 OS=Homo sapiens GN=C14orf142 PE=2 SV=2 - [CN142_HUMAN]	0.668	0.013
Q4ZIN3-2	Isoform 2 of Membralin OS=Homo sapiens GN=C19orf6 - [MBRL_HUMAN]	0.679	0.027
O96020-2	Isoform Short of G1/S-specific cyclin-E2 OS=Homo sapiens GN=CCNE2 - [CCNE2_HUMAN]	0.734	0.004
O75607	Nucleoplasmin-3 OS=Homo sapiens GN=NPM3 PE=1 SV=3 - [NPM3_HUMAN]	0.738	0.014
P51795-2	Isoform 2 of H(+)/Cl(-) exchange transporter 5 OS=Homo sapiens GN=CLCN5 - [CLCN5_HUMAN]	0.739	0.050
O95199-2	Isoform 2 of RCC1 and BTB domain-containing protein 2 OS=Homo sapiens GN=RCBT2 - [RCBT2_HUMAN]	0.751	0.042
P02675	Fibrinogen beta chain OS=Homo sapiens GN=FGB PE=1 SV=2 - [FIBB_HUMAN]	0.751	0.029
Q9NRG4	N-lysine methyltransferase SMYD2 OS=Homo sapiens GN=SMYD2 PE=1 SV=2 - [SMYD2_HUMAN]	0.758	0.023
Q96AX1	Vacuolar protein sorting-associated protein 33A OS=Homo sapiens GN=VPS33A PE=1 SV=1 - [VP33A_HUMAN]	0.763	0.027
Q9ULH0-3	Isoform 3 of Kinase D-interacting substrate of 220 kDa OS=Homo sapiens GN=KIDINS220 - [KDIS_HUMAN]	0.773	0.043
Q12982	BCL2/adenovirus E1B 19 kDa protein-interacting protein 2 OS=Homo sapiens GN=BNIP2 PE=1 SV=1 - [BNIP2_HUMAN]	0.776	0.009
Q9NQT8	Kinesin-like protein KIF13B OS=Homo sapiens GN=KIF13B PE=1 SV=1 - [K113B_HUMAN]	0.777	0.042
O95243-3	Isoform 3 of Methyl-CpG-binding domain protein 4 OS=Homo sapiens GN=MBD4 - [MBD4_HUMAN]	0.781	0.048
P38936	Cyclin-dependent kinase inhibitor 1 OS=Homo sapiens GN=CDKN1A PE=1 SV=3 - [CDN1A_HUMAN]	0.792	0.026
Q8N6R0-3	Isoform 3 of Methyltransferase-like protein 13 OS=Homo sapiens GN=METTL13 - [MTL13_HUMAN]	0.793	0.035

Q16678	Cytochrome P450 1B1 OS=Homo sapiens GN=CYP1B1 PE=1 SV=2 - [CP1B1_HUMAN]	0.794	0.047
P16860	Natriuretic peptides B OS=Homo sapiens GN=NPPB PE=1 SV=1 - [ANFB_HUMAN]	0.801	0.010
Q9C035-5	Isoform Epsilon of Tripartite motif-containing protein 5 OS=Homo sapiens GN=TRIM5 - [TRIM5_HUMAN]	0.803	0.006
Q9UK76	Hematological and neurological expressed 1 protein OS=Homo sapiens GN=HN1 PE=1 SV=3 - [HN1_HUMAN]	0.805	0.046
P10909-3	Isoform 3 of Clusterin OS=Homo sapiens GN=CLU - [CLUS_HUMAN]	0.809	0.033
Q12805-2	Isoform 2 of EGF-containing fibulin-like extracellular matrix protein 1 OS=Homo sapiens GN=EFEMP1 - [FBLN3_HUMAN]	0.813	0.042
Q9NPD8	Ubiquitin-conjugating enzyme E2 T OS=Homo sapiens GN=UBE2T PE=1 SV=1 - [UBE2T_HUMAN]	0.817	0.038
Q7Z5L2-2	Isoform 2 of Growth inhibition and differentiation-related protein 88 OS=Homo sapiens GN=GIDRP88 - [GIDRP_HUMAN]	0.819	0.002
Q14168-2	Isoform 2 of MAGUK p55 subfamily member 2 OS=Homo sapiens GN=MPP2 - [MPP2_HUMAN]	0.832	0.010
O60678-2	Isoform 2 of Protein arginine N-methyltransferase 3 OS=Homo sapiens GN=PRMT3 - [ANM3_HUMAN]	0.833	0.031
P27701	CD82 antigen OS=Homo sapiens GN=CD82 PE=1 SV=1 - [CD82_HUMAN]	0.837	0.027
Q99988	Growth/differentiation factor 15 OS=Homo sapiens GN=GDF15 PE=1 SV=3 - [GDF15_HUMAN]	0.837	0.017
Q02447-4	Isoform 4 of Transcription factor Sp3 OS=Homo sapiens GN=SP3 - [SP3_HUMAN]	0.839	0.019
O43633	Charged multivesicular body protein 2a OS=Homo sapiens GN=CHMP2A PE=1 SV=1 - [CHM2A_HUMAN]	0.847	0.010
Q96EU7	C1GALT1-specific chaperone 1 OS=Homo sapiens GN=C1GALT1C1 PE=1 SV=1 - [C1GLC_HUMAN]	0.850	0.042
Q8WX19	Transcriptional repressor p66-beta OS=Homo sapiens GN=GATAD2B PE=1 SV=1 - [P66B_HUMAN]	0.852	0.048
Q8NC51-2	Isoform 2 of Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERBP1 - [PAIRB_HUMAN]	0.854	0.034
O94953	Lysine-specific demethylase 4B OS=Homo sapiens GN=KDM4B PE=1 SV=4 - [KDM4B_HUMAN]	0.857	0.016
Q9NRW3	Probable DNA dC->dU-editing enzyme APOBEC-3C OS=Homo sapiens GN=APOBEC3C PE=1 SV=2 - [ABC3C_HUMAN]	0.860	0.024
P22694-8	Isoform 8 of cAMP-dependent protein kinase catalytic subunit beta OS=Homo sapiens GN=PRKACB - [KAPCB_HUMAN]	0.861	0.042
Q13685	Angio-associated migratory cell protein OS=Homo sapiens GN=AAMP PE=1 SV=2 - [AAMP_HUMAN]	0.863	0.010
Q14696	LDLR chaperone MESD OS=Homo sapiens GN=MESDC2 PE=1 SV=2 - [MESD_HUMAN]	0.864	0.047
Q9Y546	Leucine-rich repeat-containing protein 42 OS=Homo sapiens GN=LRRC42 PE=1 SV=1 - [LRC42_HUMAN]	0.866	0.041
Q16831	Uridine phosphorylase 1 OS=Homo sapiens GN=UPP1 PE=1 SV=1 - [UPP1_HUMAN]	0.866	0.029
P42677	40S ribosomal protein S27 OS=Homo sapiens GN=RPS27 PE=1 SV=3 - [RS27_HUMAN]	0.867	0.022
P56589	Peroxisomal biogenesis factor 3 OS=Homo sapiens GN=PEX3 PE=1 SV=1 - [PEX3_HUMAN]	0.868	0.021
Q96KQ7-2	Isoform 2 of Histone-lysine N-methyltransferase EHMT2 OS=Homo sapiens GN=EHMT2 - [EHMT2_HUMAN]	0.869	0.038
Q9NQX7-2	Isoform 2 of Integral membrane protein 2C OS=Homo sapiens GN=ITM2C - [ITM2C_HUMAN]	0.870	0.048
Q8N4A0	Polypeptide N-acetylgalactosaminyltransferase 4 OS=Homo sapiens GN=GALNT4 PE=1 SV=2 - [GALT4_HUMAN]	0.878	0.001

P50402	Emerin OS=Homo sapiens GN=EMD PE=1 SV=1 - [EMD_HUMAN]	0.879	0.017
P0C0S5	Histone H2A.Z OS=Homo sapiens GN=H2AFZ PE=1 SV=2 - [H2AZ_HUMAN]	0.880	0.035
O43324	Eukaryotic translation elongation factor 1 epsilon-1 OS=Homo sapiens GN=EEF1E1 PE=1 SV=1 - [MCA3_HUMAN]	0.881	0.043
Q8IXQ4-2	Isoform 2 of Uncharacterized protein KIAA1704 OS=Homo sapiens GN=KIAA1704 - [K1704_HUMAN]	0.882	0.038
Q9H0E3-2	Isoform 2 of Histone deacetylase complex subunit SAP130 OS=Homo sapiens GN=SAP130 - [SP130_HUMAN]	0.883	0.030
Q96JG6	Coiled-coil domain-containing protein 132 OS=Homo sapiens GN=CCDC132 PE=1 SV=3 - [CC132_HUMAN]	0.884	0.023
Q9H2W6	39S ribosomal protein L46, mitochondrial OS=Homo sapiens GN=MRPL46 PE=1 SV=1 - [RM46_HUMAN]	0.884	0.021
Q06609-2	Isoform 2 of DNA repair protein RAD51 homolog 1 OS=Homo sapiens GN=RAD51 - [RAD51_HUMAN]	0.885	0.018
Q14191	Werner syndrome ATP-dependent helicase OS=Homo sapiens GN=WRN PE=1 SV=2 - [WRN_HUMAN]	0.886	0.006
Q5HYK3	2-methoxy-6-polyprenyl-1,4-benzoquinol methylase, mitochondrial OS=Homo sapiens GN=COQ5 PE=1 SV=2 - [COQ5_HUMAN]	0.891	0.007
P27695	DNA-(apurinic or apyrimidinic site) lyase OS=Homo sapiens GN=APEX1 PE=1 SV=2 - [APEX1_HUMAN]	0.891	0.026
Q9NUE0	Palmitoyltransferase ZDHHC18 OS=Homo sapiens GN=ZDHHC18 PE=2 SV=2 - [ZDH18_HUMAN]	0.892	0.016
Q9HD23-2	Isoform 2 of Magnesium transporter MRS2 homolog, mitochondrial OS=Homo sapiens GN=MRS2 - [MRS2_HUMAN]	0.892	0.018
P48506	Glutamate--cysteine ligase catalytic subunit OS=Homo sapiens GN=GCLC PE=1 SV=2 - [GSH1_HUMAN]	0.894	0.033
P34949	Mannose-6-phosphate isomerase OS=Homo sapiens GN=MPI PE=1 SV=2 - [MPI_HUMAN]	0.894	0.025
Q9UJY4	ADP-ribosylation factor-binding protein GGA2 OS=Homo sapiens GN=GGA2 PE=1 SV=3 - [GGA2_HUMAN]	0.897	0.037
Q96Q89-4	Isoform 4 of Kinesin-like protein KIF20B OS=Homo sapiens GN=KIF20B - [KI20B_HUMAN]	0.899	0.001
P18859	ATP synthase-coupling factor 6, mitochondrial OS=Homo sapiens GN=ATP5J PE=1 SV=1 - [ATP5J_HUMAN]	0.900	0.010
O15212	Prefoldin subunit 6 OS=Homo sapiens GN=PFDN6 PE=1 SV=1 - [PFD6_HUMAN]	0.900	0.047
P51636-3	Isoform C of Caveolin-2 OS=Homo sapiens GN=CAV2 - [CAV2_HUMAN]	0.901	0.000
Q9HBR0	Putative sodium-coupled neutral amino acid transporter 10 OS=Homo sapiens GN=SLC38A10 PE=1 SV=2 - [S38AA_HUMAN]	0.903	0.016
Q92966	snRNA-activating protein complex subunit 3 OS=Homo sapiens GN=SNAPC3 PE=1 SV=1 - [SNPC3_HUMAN]	0.903	0.028
Q9UBP0-4	Isoform 4 of Spastin OS=Homo sapiens GN=SPAST - [SPAST_HUMAN]	0.904	0.017
Q8IZM8	Zinc finger protein 654 OS=Homo sapiens GN=ZNF654 PE=1 SV=3 - [ZN654_HUMAN]	0.904	0.002
P08195	4F2 cell-surface antigen heavy chain OS=Homo sapiens GN=SLC3A2 PE=1 SV=3 - [4F2_HUMAN]	0.905	0.021
Q9Y3B9	RRP15-like protein OS=Homo sapiens GN=RRP15 PE=1 SV=2 - [RRP15_HUMAN]	0.906	0.025
P07108	Acyl-CoA-binding protein OS=Homo sapiens GN=DBI PE=1 SV=2 - [ACBP_HUMAN]	0.907	0.008
Q9UKR5	Probable ergosterol biosynthetic protein 28 OS=Homo sapiens GN=C14orf1 PE=1 SV=1 - [ERG28_HUMAN]	0.908	0.000
O60826	Coiled-coil domain-containing protein 22 OS=Homo sapiens GN=CCDC22 PE=1 SV=1 - [CCD22_HUMAN]	0.908	0.048

Q12931	Heat shock protein 75 kDa, mitochondrial OS=Homo sapiens GN=TRAP1 PE=1 SV=3 - [TRAP1_HUMAN]	0.910	0.034
Q8TD19	Serine/threonine-protein kinase Nek9 OS=Homo sapiens GN=NEK9 PE=1 SV=2 - [NEK9_HUMAN]	0.911	0.044
P24385	G1/S-specific cyclin-D1 OS=Homo sapiens GN=CCND1 PE=1 SV=1 - [CCND1_HUMAN]	0.913	0.032
Q12986-2	Isoform 2 of Transcriptional repressor NF-X1 OS=Homo sapiens GN=NFX1 - [NFX1_HUMAN]	0.915	0.009
Q96KP1	Exocyst complex component 2 OS=Homo sapiens GN=EXOC2 PE=1 SV=1 - [EXOC2_HUMAN]	0.917	0.003
Q13049	E3 ubiquitin-protein ligase TRIM32 OS=Homo sapiens GN=TRIM32 PE=1 SV=2 - [TRIM32_HUMAN]	0.919	0.010
Q9UDT6-2	Isoform 2 of CAP-Gly domain-containing linker protein 2 OS=Homo sapiens GN=CLIP2 - [CLIP2_HUMAN]	0.920	0.032
Q16563-2	Isoform 2 of Synaptophysin-like protein 1 OS=Homo sapiens GN=SYPL1 - [SYPL1_HUMAN]	0.920	0.008
Q3KQV9	UDP-N-acetylhexosamine pyrophosphorylase-like protein 1 OS=Homo sapiens GN=UAP1L1 PE=2 SV=2 - [UAP1L_HUMAN]	0.923	0.032
O00429-4	Isoform 3 of Dynamin-1-like protein OS=Homo sapiens GN=DNM1L - [DNM1L_HUMAN]	0.925	0.028
Q9H0G5	Nuclear speckle splicing regulatory protein 1 OS=Homo sapiens GN=NSRP1 PE=1 SV=1 - [NSRP1_HUMAN]	0.927	0.022
Q9BX63-2	Isoform 2 of Fanconi anemia group J protein OS=Homo sapiens GN=BRIP1 - [FANCI_HUMAN]	0.930	0.008
Q92685	Dol-P-Man:Man(5)GlcNAc(2)-PP-Dol alpha-1,3-mannosyltransferase OS=Homo sapiens GN=ALG3 PE=1 SV=1 - [ALG3_HUMAN]	0.931	0.038
Q8N9N7	Leucine-rich repeat-containing protein 57 OS=Homo sapiens GN=LRRC57 PE=1 SV=1 - [LRC57_HUMAN]	0.932	0.025
P61204	ADP-ribosylation factor 3 OS=Homo sapiens GN=ARF3 PE=1 SV=2 - [ARF3_HUMAN]	0.932	0.032
P49257	Protein ERGIC-53 OS=Homo sapiens GN=LMAN1 PE=1 SV=2 - [LMAN1_HUMAN]	0.932	0.012
Q9P2X3	Protein IMPACT OS=Homo sapiens GN=IMPACT PE=1 SV=2 - [IMPACT_HUMAN]	0.933	0.018
P06730	Eukaryotic translation initiation factor 4E OS=Homo sapiens GN=EIF4E PE=1 SV=2 - [IF4E_HUMAN]	0.934	0.029
P15408-3	Isoform 3 of Fos-related antigen 2 OS=Homo sapiens GN=FOSL2 - [FOSL2_HUMAN]	0.934	0.022
Q9BV79-2	Isoform 2 of Trans-2-enoyl-CoA reductase, mitochondrial OS=Homo sapiens GN=MECR - [MECR_HUMAN]	0.937	0.034
O95613	Pericentrin OS=Homo sapiens GN=PCNT PE=1 SV=4 - [PCNT_HUMAN]	0.937	0.025
O43314-2	Isoform 2 of Inositol hexakisphosphate and diphosphoinositol-pentakisphosphate kinase 2 OS=Homo sapiens GN=PPIP5K2 - [VIP2_HUMAN]	0.937	0.043
Q14457	Beclin-1 OS=Homo sapiens GN=BECN1 PE=1 SV=2 - [BECN1_HUMAN]	0.939	0.033
Q96AE7	Tetratricopeptide repeat protein 17 OS=Homo sapiens GN=TTC17 PE=1 SV=1 - [TTC17_HUMAN]	0.939	0.021
O94761	ATP-dependent DNA helicase Q4 OS=Homo sapiens GN=RECQL4 PE=1 SV=1 - [RECQ4_HUMAN]	0.939	0.034
Q9NY27-3	Isoform 3 of Serine/threonine-protein phosphatase 4 regulatory subunit 2 OS=Homo sapiens GN=PPP4R2 - [PP4R2_HUMAN]	0.940	0.032
P15260	Interferon gamma receptor 1 OS=Homo sapiens GN=IFNGR1 PE=1 SV=1 - [INGR1_HUMAN]	0.940	0.039
Q6NUK1	Calcium-binding mitochondrial carrier protein SCaMC-1 OS=Homo sapiens GN=SLC25A24 PE=1 SV=2 - [SCMC1_HUMAN]	0.941	0.035
O94826	Mitochondrial import receptor subunit TOM70 OS=Homo sapiens GN=TOMM70A PE=1 SV=1 - [TOM70_HUMAN]	0.942	0.018

Q96BJ3	Axin interactor, dorsalization-associated protein OS=Homo sapiens GN=AIDA PE=1 SV=1 - [AIDA_HUMAN]	0.943	0.025
Q8N138-4	Isoform 2 of ORM1-like protein 3 OS=Homo sapiens GN=ORMDL3 - [ORML3_HUMAN]	0.946	0.027
P24928	DNA-directed RNA polymerase II subunit RPB1 OS=Homo sapiens GN=POLR2A PE=1 SV=2 - [RPB1_HUMAN]	0.947	0.027
O14979-3	Isoform 3 of Heterogeneous nuclear ribonucleoprotein D-like OS=Homo sapiens GN=HNRPDL - [HNRDL_HUMAN]	0.948	0.044
Q71RC2-3	Isoform 3 of La-related protein 4 OS=Homo sapiens GN=LARP4 - [LARP4_HUMAN]	0.948	0.029
Q9H082	Ras-related protein Rab-33B OS=Homo sapiens GN=RAB33B PE=1 SV=1 - [RB33B_HUMAN]	0.949	0.048
Q75QN2-2	Isoform 2 of Integrator complex subunit 8 OS=Homo sapiens GN=INTS8 - [INT8_HUMAN]	0.949	0.046
Q9UPP1-3	Isoform 3 of Histone lysine demethylase PHF8 OS=Homo sapiens GN=PHF8 - [PHF8_HUMAN]	0.949	0.041
Q8N653	Leucine-zipper-like transcriptional regulator 1 OS=Homo sapiens GN=LZTR1 PE=2 SV=2 - [LZTR1_HUMAN]	0.950	0.041
Q5GLZ8-6	Isoform 6 of Probable E3 ubiquitin-protein ligase HERC4 OS=Homo sapiens GN=HERC4 - [HERC4_HUMAN]	0.951	0.028
Q9Y4W6	AFG3-like protein 2 OS=Homo sapiens GN=AFG3L2 PE=1 SV=2 - [AFG32_HUMAN]	0.952	0.016
Q96DE0	U8 snoRNA-decapping enzyme OS=Homo sapiens GN=NUDT16 PE=1 SV=2 - [NUD16_HUMAN]	0.955	0.035
O00151	PDZ and LIM domain protein 1 OS=Homo sapiens GN=PDLIM1 PE=1 SV=4 - [PDL1_HUMAN]	0.955	0.035
Q5VYS8-4	Isoform 4 of Terminal uridylyltransferase 7 OS=Homo sapiens GN=ZCCHC6 - [TUT7_HUMAN]	0.955	0.014
Q9BYW2	Histone-lysine N-methyltransferase SETD2 OS=Homo sapiens GN=SETD2 PE=1 SV=3 - [SETD2_HUMAN]	0.957	0.027
P09496-2	Isoform Non-brain of Clathrin light chain A OS=Homo sapiens GN=CLTA - [CLCA_HUMAN]	0.957	0.040
P52788	Spermine synthase OS=Homo sapiens GN=SMS PE=1 SV=2 - [SPSY_HUMAN]	0.957	0.047
P62424	60S ribosomal protein L7a OS=Homo sapiens GN=RPL7A PE=1 SV=2 - [RL7A_HUMAN]	0.957	0.018
P31937	3-hydroxyisobutyrate dehydrogenase, mitochondrial OS=Homo sapiens GN=HIBADH PE=1 SV=2 - [3HIDH_HUMAN]	0.959	0.033
P51151	Ras-related protein Rab-9A OS=Homo sapiens GN=RAB9A PE=1 SV=1 - [RAB9A_HUMAN]	0.960	0.019
P14921-2	Isoform c-ETS-1B of Protein C-ets-1 OS=Homo sapiens GN=ETS1 - [ETS1_HUMAN]	0.960	0.045
Q86YV9	Hermansky-Pudlak syndrome 6 protein OS=Homo sapiens GN=HPS6 PE=1 SV=1 - [HPS6_HUMAN]	0.961	0.034
P31040	Succinate dehydrogenase [ubiquinone] flavoprotein subunit, mitochondrial OS=Homo sapiens GN=SDHA PE=1 SV=2 - [DHSA_HUMAN]	0.963	0.016
Q9Y618-5	Isoform 5 of Nuclear receptor corepressor 2 OS=Homo sapiens GN=NCOR2 - [NCOR2_HUMAN]	0.964	0.015
Q9HCD5	Nuclear receptor coactivator 5 OS=Homo sapiens GN=NCOA5 PE=1 SV=2 - [NCOA5_HUMAN]	0.967	0.016
Q9HCG8	Pre-mRNA-splicing factor CWC22 homolog OS=Homo sapiens GN=CWC22 PE=1 SV=3 - [CWC22_HUMAN]	0.972	0.004
Q9BXW9-2	Isoform 2 of Fanconi anemia group D2 protein OS=Homo sapiens GN=FANCD2 - [FACD2_HUMAN]	0.973	0.028
Q99543	DnaJ homolog subfamily C member 2 OS=Homo sapiens GN=DNAJC2 PE=1 SV=4 - [DNJC2_HUMAN]	0.982	0.025
Q16512	Serine/threonine-protein kinase N1 OS=Homo sapiens GN=PKN1 PE=1 SV=2 - [PKN1_HUMAN]	0.985	0.049

P51858	Hepatoma-derived growth factor OS=Homo sapiens GN=HDGF PE=1 SV=1 - [HDGF_HUMAN]	0.986	0.033
Q16643	Drebrin OS=Homo sapiens GN=DBN1 PE=1 SV=4 - [DREB_HUMAN]	0.997	0.014
P53992	Protein transport protein Sec24C OS=Homo sapiens GN=SEC24C PE=1 SV=3 - [SC24C_HUMAN]	1.015	0.045
O15031	Plexin-B2 OS=Homo sapiens GN=PLXNB2 PE=1 SV=3 - [PLXB2_HUMAN]	1.016	0.017
P41252	Isoleucine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=IARS PE=1 SV=2 - [SYIC_HUMAN]	1.025	0.048
P42574	Caspase-3 OS=Homo sapiens GN=CASP3 PE=1 SV=2 - [CASP3_HUMAN]	1.025	0.049
Q9NSE4	Isoleucine--tRNA ligase, mitochondrial OS=Homo sapiens GN=IARS2 PE=1 SV=2 - [SYIM_HUMAN]	1.025	0.029
Q8NFW8	N-acylneuraminate cytidyltransferase OS=Homo sapiens GN=CMAS PE=1 SV=2 - [NEUA_HUMAN]	1.026	0.043
Q14012	Calcium/calmodulin-dependent protein kinase type 1 OS=Homo sapiens GN=CAMK1 PE=1 SV=1 - [KCC1A_HUMAN]	1.031	0.038
P06400	Retinoblastoma-associated protein OS=Homo sapiens GN=RB1 PE=1 SV=2 - [RB_HUMAN]	1.032	0.011
Q99613	Eukaryotic translation initiation factor 3 subunit C OS=Homo sapiens GN=EIF3C PE=1 SV=1 - [EIF3C_HUMAN]	1.034	0.030
P11441	Ubiquitin-like protein 4A OS=Homo sapiens GN=UBL4A PE=1 SV=1 - [UBL4A_HUMAN]	1.035	0.041
Q02543	60S ribosomal protein L18a OS=Homo sapiens GN=RPL18A PE=1 SV=2 - [RL18A_HUMAN]	1.035	0.004
Q13765	Nascent polypeptide-associated complex subunit alpha OS=Homo sapiens GN=NACA PE=1 SV=1 - [NACA_HUMAN]	1.037	0.039
P17706-2	Isoform PTPA of Tyrosine-protein phosphatase non-receptor type 2 OS=Homo sapiens GN=PTPN2 - [PTN2_HUMAN]	1.037	0.034
Q9UJX5	Anaphase-promoting complex subunit 4 OS=Homo sapiens GN=ANAPC4 PE=1 SV=2 - [APC4_HUMAN]	1.038	0.045
P36551	Coproporphyrinogen-III oxidase, mitochondrial OS=Homo sapiens GN=CPOX PE=1 SV=3 - [HEM6_HUMAN]	1.040	0.040
Q9P2J5	Leucine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=LARS PE=1 SV=2 - [SYLC_HUMAN]	1.041	0.012
Q96KA5-2	Isoform 2 of Cleft lip and palate transmembrane protein 1-like protein OS=Homo sapiens GN=CLPTM1L - [CLP1L_HUMAN]	1.044	0.043
O95376	E3 ubiquitin-protein ligase ARIH2 OS=Homo sapiens GN=ARIH2 PE=1 SV=1 - [ARI2_HUMAN]	1.045	0.045
P27448-3	Isoform 3 of MAP/microtubule affinity-regulating kinase 3 OS=Homo sapiens GN=MARK3 - [MARK3_HUMAN]	1.046	0.047
O15269	Serine palmitoyltransferase 1 OS=Homo sapiens GN=SPTLC1 PE=1 SV=1 - [SPTC1_HUMAN]	1.046	0.007
Q6ZUT1	Uncharacterized protein C11orf57 OS=Homo sapiens GN=C11orf57 PE=1 SV=2 - [CK057_HUMAN]	1.048	0.022
Q9Y6A5	Transforming acidic coiled-coil-containing protein 3 OS=Homo sapiens GN=TACC3 PE=1 SV=1 - [TACC3_HUMAN]	1.048	0.009
P48723	Heat shock 70 kDa protein 13 OS=Homo sapiens GN=HSPA13 PE=1 SV=1 - [HSP13_HUMAN]	1.049	0.036
O75694-2	Isoform 2 of Nuclear pore complex protein Nup155 OS=Homo sapiens GN=NUP155 - [NU155_HUMAN]	1.050	0.037
Q7RTV5	UPF0308 protein C9orf21 OS=Homo sapiens GN=C9orf21 PE=2 SV=1 - [CI021_HUMAN]	1.050	0.014
Q7Z3B4	Nucleoporin p54 OS=Homo sapiens GN=NUP54 PE=1 SV=2 - [NUP54_HUMAN]	1.051	0.045
P00491	Purine nucleoside phosphorylase OS=Homo sapiens GN=PNP PE=1 SV=2 - [PNPH_HUMAN]	1.051	0.018

Q04837	Single-stranded DNA-binding protein, mitochondrial OS=Homo sapiens GN=SSBP1 PE=1 SV=1 - [SSBP_HUMAN]	1.051	0.029
Q96RL7-4	Isoform 4 of Vacuolar protein sorting-associated protein 13A OS=Homo sapiens GN=VPS13A - [VP13A_HUMAN]	1.052	0.044
P54136	Arginine--tRNA ligase, cytoplasmic OS=Homo sapiens GN=RARS PE=1 SV=2 - [SYRC_HUMAN]	1.052	0.026
Q8N5M1	ATP synthase mitochondrial F1 complex assembly factor 2 OS=Homo sapiens GN=ATPAF2 PE=1 SV=1 - [ATPF2_HUMAN]	1.053	0.014
Q9UKB1-2	Isoform A of F-box/WD repeat-containing protein 11 OS=Homo sapiens GN=FBXW11 - [FBW1B_HUMAN]	1.054	0.049
Q9Y2L9-2	Isoform 2 of Leucine-rich repeat and calponin homology domain-containing protein 1 OS=Homo sapiens GN=LRCH1 - [LRCH1_HUMAN]	1.055	0.033
P62841	40S ribosomal protein S15 OS=Homo sapiens GN=RPS15 PE=1 SV=2 - [RS15_HUMAN]	1.055	0.046
Q9H0H5	Rac GTPase-activating protein 1 OS=Homo sapiens GN=RACGAP1 PE=1 SV=1 - [RGAP1_HUMAN]	1.055	0.033
Q93008-1	Isoform 2 of Probable ubiquitin carboxyl-terminal hydrolase FAF-X OS=Homo sapiens GN=USP9X - [USP9X_HUMAN]	1.056	0.033
Q8TDW4-8	Isoform 8 of Suppressor of tumorigenicity 7 protein-like OS=Homo sapiens GN=ST7L - [ST7L_HUMAN]	1.057	0.048
P25786	Proteasome subunit alpha type-1 OS=Homo sapiens GN=PSMA1 PE=1 SV=1 - [PSA1_HUMAN]	1.057	0.018
P08238	Heat shock protein HSP 90-beta OS=Homo sapiens GN=HSP90AB1 PE=1 SV=4 - [HS90B_HUMAN]	1.058	0.029
O60524-4	Isoform 4 of Nuclear export mediator factor NEMF OS=Homo sapiens GN=NEMF - [NEMF_HUMAN]	1.058	0.010
Q13613-2	Isoform 1A of Myotubularin-related protein 1 OS=Homo sapiens GN=MTMR1 - [MTMR1_HUMAN]	1.058	0.001
P43034	Platelet-activating factor acetylhydrolase IB subunit alpha OS=Homo sapiens GN=PFAFH1B1 PE=1 SV=2 - [LIS1_HUMAN]	1.065	0.011
P39748	Flap endonuclease 1 OS=Homo sapiens GN=FEN1 PE=1 SV=1 - [FEN1_HUMAN]	1.065	0.005
P62266	40S ribosomal protein S23 OS=Homo sapiens GN=RPS23 PE=1 SV=3 - [RS23_HUMAN]	1.067	0.035
Q15006	Tetratricopeptide repeat protein 35 OS=Homo sapiens GN=TTC35 PE=1 SV=1 - [TTC35_HUMAN]	1.067	0.017
P37802	Transgelin-2 OS=Homo sapiens GN=TAGLN2 PE=1 SV=3 - [TAGL2_HUMAN]	1.068	0.036
O60493	Sorting nexin-3 OS=Homo sapiens GN=SNX3 PE=1 SV=3 - [SNX3_HUMAN]	1.068	0.010
Q53GS9	U4/U6.U5 tri-snRNP-associated protein 2 OS=Homo sapiens GN=USP39 PE=1 SV=2 - [SNUT2_HUMAN]	1.069	0.041
Q9NP79	Vacuolar protein sorting-associated protein VTA1 homolog OS=Homo sapiens GN=VTA1 PE=1 SV=1 - [VTA1_HUMAN]	1.069	0.038
O60812	Heterogeneous nuclear ribonucleoprotein C-like 1 OS=Homo sapiens GN=HNRNPCL1 PE=1 SV=1 - [HNRCL_HUMAN]	1.070	0.001
Q99797	Mitochondrial intermediate peptidase OS=Homo sapiens GN=MIPEP PE=1 SV=2 - [MIPEP_HUMAN]	1.071	0.050
Q99661-2	Isoform 2 of Kinesin-like protein KIF2C OS=Homo sapiens GN=KIF2C - [KIF2C_HUMAN]	1.072	0.034
P40189	Interleukin-6 receptor subunit beta OS=Homo sapiens GN=IL6ST PE=1 SV=2 - [IL6RB_HUMAN]	1.073	0.039
Q5SXM8	DNL-type zinc finger protein OS=Homo sapiens GN=DNLZ PE=2 SV=1 - [DNLZ_HUMAN]	1.073	0.021
Q9NVG8	TBC1 domain family member 13 OS=Homo sapiens GN=TBC1D13 PE=1 SV=3 - [TBC13_HUMAN]	1.074	0.015
Q9BYG3	MKI67 FHA domain-interacting nucleolar phosphoprotein OS=Homo sapiens GN=MKI67IP PE=1 SV=1 - [MK67I_HUMAN]	1.075	0.011

Q8N6Y2-2	Isoform 2 of Leucine-rich repeat-containing protein 17 OS=Homo sapiens GN=LRR17 - [LRR17_HUMAN]	1.076	0.014
P21281	V-type proton ATPase subunit B, brain isoform OS=Homo sapiens GN=ATP6V1B2 PE=1 SV=3 - [VATB2_HUMAN]	1.077	0.039
P52565	Rho GDP-dissociation inhibitor 1 OS=Homo sapiens GN=ARHGDI1 PE=1 SV=3 - [GDIR1_HUMAN]	1.077	0.049
Q9BWD1	Acetyl-CoA acetyltransferase, cytosolic OS=Homo sapiens GN=ACAT2 PE=1 SV=2 - [THIC_HUMAN]	1.077	0.037
P42285	Superkiller viralicidic activity 2-like 2 OS=Homo sapiens GN=SKIV2L2 PE=1 SV=3 - [SK2L2_HUMAN]	1.079	0.037
P48147	Prolyl endopeptidase OS=Homo sapiens GN=PREP PE=1 SV=2 - [PPCE_HUMAN]	1.079	0.039
Q658Y4	Protein FAM91A1 OS=Homo sapiens GN=FAM91A1 PE=1 SV=3 - [F91A1_HUMAN]	1.080	0.029
Q9BYT8	Neurolysin, mitochondrial OS=Homo sapiens GN=NLN PE=1 SV=1 - [NEUL_HUMAN]	1.081	0.044
Q96GM8	Target of EGR1 protein 1 OS=Homo sapiens GN=TOE1 PE=1 SV=1 - [TOE1_HUMAN]	1.083	0.032
P28066	Proteasome subunit alpha type-5 OS=Homo sapiens GN=PSMA5 PE=1 SV=3 - [PSA5_HUMAN]	1.083	0.008
Q9H0U4	Ras-related protein Rab-1B OS=Homo sapiens GN=RAB1B PE=1 SV=1 - [RAB1B_HUMAN]	1.083	0.031
P62070	Ras-related protein R-Ras2 OS=Homo sapiens GN=RRAS2 PE=1 SV=1 - [RRAS2_HUMAN]	1.083	0.019
Q15648	Mediator of RNA polymerase II transcription subunit 1 OS=Homo sapiens GN=MED1 PE=1 SV=4 - [MED1_HUMAN]	1.084	0.021
P61970	Nuclear transport factor 2 OS=Homo sapiens GN=NUTF2 PE=1 SV=1 - [NTF2_HUMAN]	1.086	0.022
Q8WUA2	Peptidyl-prolyl cis-trans isomerase-like 4 OS=Homo sapiens GN=PPIL4 PE=1 SV=1 - [PPIL4_HUMAN]	1.087	0.003
Q9NYK5	39S ribosomal protein L39, mitochondrial OS=Homo sapiens GN=MRPL39 PE=1 SV=3 - [RM39_HUMAN]	1.087	0.044
O95067	G2/mitotic-specific cyclin-B2 OS=Homo sapiens GN=CCNB2 PE=1 SV=1 - [CCNB2_HUMAN]	1.087	0.017
Q9BRS2	Serine/threonine-protein kinase RIO1 OS=Homo sapiens GN=RIOK1 PE=1 SV=2 - [RIOK1_HUMAN]	1.088	0.027
Q96P11	Putative methyltransferase NSUN5 OS=Homo sapiens GN=NSUN5 PE=1 SV=2 - [NSUN5_HUMAN]	1.088	0.023
O95164	Ubiquitin-like protein 3 OS=Homo sapiens GN=UBL3 PE=1 SV=1 - [UBL3_HUMAN]	1.089	0.040
P82914	28S ribosomal protein S15, mitochondrial OS=Homo sapiens GN=MRPS15 PE=1 SV=1 - [RT15_HUMAN]	1.089	0.041
Q9Y2L5-2	Isoform 2 of Trafficking protein particle complex subunit 8 OS=Homo sapiens GN=TRAPPC8 - [TPPC8_HUMAN]	1.091	0.003
P00505	Aspartate aminotransferase, mitochondrial OS=Homo sapiens GN=GOT2 PE=1 SV=3 - [AATM_HUMAN]	1.092	0.009
Q9UIL8-2	Isoform 2 of PHD finger protein 11 OS=Homo sapiens GN=PHF11 - [PHF11_HUMAN]	1.092	0.031
Q96T37-2	Isoform 2 of Putative RNA-binding protein 15 OS=Homo sapiens GN=RBM15 - [RBM15_HUMAN]	1.093	0.029
P35659	Protein DEK OS=Homo sapiens GN=DEK PE=1 SV=1 - [DEK_HUMAN]	1.096	0.034
P11177-2	Isoform 2 of Pyruvate dehydrogenase E1 component subunit beta, mitochondrial OS=Homo sapiens GN=PDHB - [ODPB_HUMAN]	1.101	0.040
P09104	Gamma-enolase OS=Homo sapiens GN=ENO2 PE=1 SV=3 - [ENOG_HUMAN]	1.104	0.046
Q9NV35	Probable 7,8-dihydro-8-oxoguanine triphosphatase NUDT15 OS=Homo sapiens GN=NUDT15 PE=1 SV=1 - [NUDT15_HUMAN]	1.106	0.029

O95163	Elongator complex protein 1 OS=Homo sapiens GN=IKBKAP PE=1 SV=3 - [ELP1_HUMAN]	1.112	0.021
Q00534	Cyclin-dependent kinase 6 OS=Homo sapiens GN=CDK6 PE=1 SV=1 - [CDK6_HUMAN]	1.117	0.006
P52594-2	Isoform 2 of Arf-GAP domain and FG repeats-containing protein 1 OS=Homo sapiens GN=AGFG1 - [AGFG1_HUMAN]	1.120	0.046
O76021	Ribosomal L1 domain-containing protein 1 OS=Homo sapiens GN=RSL1D1 PE=1 SV=3 - [RL1D1_HUMAN]	1.122	0.005
Q8IWB9	Testis-expressed sequence 2 protein OS=Homo sapiens GN=TEX2 PE=1 SV=2 - [TEX2_HUMAN]	1.123	0.019
Q96NW4	Ankyrin repeat domain-containing protein 27 OS=Homo sapiens GN=ANKRD27 PE=1 SV=2 - [ANR27_HUMAN]	1.123	0.012
Q2TAY7	WD40 repeat-containing protein SMU1 OS=Homo sapiens GN=SMU1 PE=1 SV=2 - [SMU1_HUMAN]	1.125	0.041
Q96S19-3	Isoform 3 of UPF0585 protein C16orf13 OS=Homo sapiens GN=C16orf13 - [CP013_HUMAN]	1.125	0.029
P49720	Proteasome subunit beta type-3 OS=Homo sapiens GN=PSMB3 PE=1 SV=2 - [PSB3_HUMAN]	1.128	0.036
Q15185	Prostaglandin E synthase 3 OS=Homo sapiens GN=PTGES3 PE=1 SV=1 - [TEBP_HUMAN]	1.128	0.030
Q04721	Neurogenic locus notch homolog protein 2 OS=Homo sapiens GN=NOTCH2 PE=1 SV=3 - [NOTC2_HUMAN]	1.130	0.046
Q9P2E9	Ribosome-binding protein 1 OS=Homo sapiens GN=RRBP1 PE=1 SV=4 - [RRBP1_HUMAN]	1.131	0.001
Q8I WV7	E3 ubiquitin-protein ligase UBR1 OS=Homo sapiens GN=UBR1 PE=1 SV=1 - [UBR1_HUMAN]	1.132	0.038
Q9H9J2	39S ribosomal protein L44, mitochondrial OS=Homo sapiens GN=MRPL44 PE=1 SV=1 - [RM44_HUMAN]	1.137	0.016
P55287-2	Isoform 2 of Cadherin-11 OS=Homo sapiens GN=CDH11 - [CAD11_HUMAN]	1.140	0.021
Q8NFI3-2	Isoform 2 of Cytosolic endo-beta-N-acetylglucosaminidase OS=Homo sapiens GN=ENGASE - [ENASE_HUMAN]	1.140	0.006
Q96IF1	LIM domain-containing protein ajuba OS=Homo sapiens GN=AJUBA PE=1 SV=1 - [AJUBA_HUMAN]	1.142	0.001
Q02318	Sterol 26-hydroxylase, mitochondrial OS=Homo sapiens GN=CYP27A1 PE=1 SV=1 - [CP27A_HUMAN]	1.144	0.015
O75947	ATP synthase subunit d, mitochondrial OS=Homo sapiens GN=ATP5H PE=1 SV=3 - [ATP5H_HUMAN]	1.148	0.043
Q8IWA4-3	Isoform 3 of Mitofusin-1 OS=Homo sapiens GN=MFN1 - [MFN1_HUMAN]	1.148	0.016
Q9P0P0	E3 ubiquitin-protein ligase RNF181 OS=Homo sapiens GN=RNF181 PE=1 SV=1 - [RN181_HUMAN]	1.152	0.049
P49757-4	Isoform 4 of Protein numb homolog OS=Homo sapiens GN=NUMB - [NUMB_HUMAN]	1.152	0.034
Q9H8G2	Uncharacterized protein C9orf82 OS=Homo sapiens GN=C9orf82 PE=1 SV=2 - [CI082_HUMAN]	1.162	0.047
Q96GG9	DCN1-like protein 1 OS=Homo sapiens GN=DCUN1D1 PE=1 SV=1 - [DCNL1_HUMAN]	1.166	0.017
P61599	N-alpha-acetyltransferase 20 OS=Homo sapiens GN=NAA20 PE=1 SV=1 - [NAA20_HUMAN]	1.168	0.016
O96033	Molybdopterin synthase sulfur carrier subunit OS=Homo sapiens GN=MOCS2 PE=1 SV=1 - [MOC2A_HUMAN]	1.185	0.036
Q9UIM3	FK506-binding protein-like OS=Homo sapiens GN=FKBPL PE=1 SV=1 - [FKBPL_HUMAN]	1.185	0.008
Q00535-2	Isoform 2 of Cyclin-dependent kinase 5 OS=Homo sapiens GN=CDK5 - [CDK5_HUMAN]	1.191	0.026
Q2TB10	Zinc finger protein 800 OS=Homo sapiens GN=ZNF800 PE=1 SV=1 - [ZN800_HUMAN]	1.191	0.025

Q9BYI3	Hyccin OS=Homo sapiens GN=FAM126A PE=1 SV=2 - [HYCCI_HUMAN]	1.206	0.027
Q96BA8-2	Isoform 2 of Cyclic AMP-responsive element-binding protein 3-like protein 1 OS=Homo sapiens GN=CREB3L1 - [CR3L1_HUMAN]	1.212	0.045
P23469-3	Isoform 3 of Receptor-type tyrosine-protein phosphatase epsilon OS=Homo sapiens GN=PTPRE - [PTPRE_HUMAN]	1.221	0.047
Q8IVF5	T-lymphoma invasion and metastasis-inducing protein 2 OS=Homo sapiens GN=TIAM2 PE=2 SV=4 - [TIAM2_HUMAN]	1.222	0.048
Q15181	Inorganic pyrophosphatase OS=Homo sapiens GN=PPA1 PE=1 SV=2 - [IPYR_HUMAN]	1.237	0.025
Q9H7D0	Dedicator of cytokinesis protein 5 OS=Homo sapiens GN=DOCK5 PE=1 SV=3 - [DOCK5_HUMAN]	1.254	0.005
Q8IXH7-4	Isoform NELF-D of Negative elongation factor C/D OS=Homo sapiens GN=TH1L - [NELFD_HUMAN]	1.289	0.017
Q9BYD2	39S ribosomal protein L9, mitochondrial OS=Homo sapiens GN=MRPL9 PE=1 SV=2 - [RM09_HUMAN]	1.310	0.012
P01889	HLA class I histocompatibility antigen, B-7 alpha chain OS=Homo sapiens GN=HLA-B PE=1 SV=3 - [1B07_HUMAN]	1.318	0.029
Q9H3K2	Growth hormone-inducible transmembrane protein OS=Homo sapiens GN=GHITM PE=1 SV=2 - [GHITM_HUMAN]	1.392	0.046
Q9Y2C4	Nuclease EXOG, mitochondrial OS=Homo sapiens GN=EXOG PE=1 SV=2 - [EXOG_HUMAN]	1.435	0.025
O75880	Protein SCO1 homolog, mitochondrial OS=Homo sapiens GN=SCO1 PE=1 SV=1 - [SCO1_HUMAN]	1.465	0.034
O15235	28S ribosomal protein S12, mitochondrial OS=Homo sapiens GN=MRPS12 PE=1 SV=1 - [RT12_HUMAN]	1.556	0.028
O75323	Protein NipSnap homolog 2 OS=Homo sapiens GN=GBAS PE=1 SV=1 - [NIPS2_HUMAN]	1.586	0.036
P62877	E3 ubiquitin-protein ligase RBX1 OS=Homo sapiens GN=RBX1 PE=1 SV=1 - [RBX1_HUMAN]	1.590	0.039
Q86VY9	Transmembrane protein 200A OS=Homo sapiens GN=TMEM200A PE=2 SV=1 - [T200A_HUMAN]	1.640	0.041
Q3B7S5	Putative uncharacterized protein C18orf62 OS=Homo sapiens GN=C18orf62 PE=1 SV=1 - [CR062_HUMAN]	2.078	0.012
Q9BYX7	Putative beta-actin-like protein 3 OS=Homo sapiens GN=POTEKP PE=5 SV=1 - [ACTBM_HUMAN]	2.298	0.048

HC45 cells - 48 hours lanreotide (397 proteins)

P13473	Lysosome-associated membrane glycoprotein 2 OS=Homo sapiens GN=LAMP2 PE=1 SV=2 - [LAMP2_HUMAN]	0.396	0.035
P53539	Protein fosB OS=Homo sapiens GN=FOSB PE=1 SV=1 - [FOSB_HUMAN]	0.457	0.002
P60903	Protein S100-A10 OS=Homo sapiens GN=S100A10 PE=1 SV=2 - [S10AA_HUMAN]	0.529	0.043
Q04695	Keratin, type I cytoskeletal 17 OS=Homo sapiens GN=KRT17 PE=1 SV=2 - [K1C17_HUMAN]	0.542	0.009
Q96J01	THO complex subunit 3 OS=Homo sapiens GN=THOC3 PE=1 SV=1 - [THOC3_HUMAN]	0.545	0.023
P0CG34	Thymosin beta-15A OS=Homo sapiens GN=TMSB15A PE=1 SV=1 - [TB15A_HUMAN]	0.571	0.001
O95243-3	Isoform 3 of Methyl-CpG-binding domain protein 4 OS=Homo sapiens GN=MBD4 - [MBD4_HUMAN]	0.595	0.014
Q16678	Cytochrome P450 1B1 OS=Homo sapiens GN=CYP1B1 PE=1 SV=2 - [CP1B1_HUMAN]	0.626	0.033
Q9H8M7	Protein FAM188A OS=Homo sapiens GN=FAM188A PE=1 SV=1 - [F188A_HUMAN]	0.628	0.031

P05114	Non-histone chromosomal protein HMG-14 OS=Homo sapiens GN=HMGN1 PE=1 SV=3 - [HMGN1_HUMAN]	0.635	0.042
O95218-2	Isoform 2 of Zinc finger Ran-binding domain-containing protein 2 OS=Homo sapiens GN=ZRANB2 - [ZRAB2_HUMAN]	0.636	0.042
Q16831	Uridine phosphorylase 1 OS=Homo sapiens GN=UPP1 PE=1 SV=1 - [UPP1_HUMAN]	0.638	0.006
Q15005	Signal peptidase complex subunit 2 OS=Homo sapiens GN=SPCS2 PE=1 SV=3 - [SPCS2_HUMAN]	0.649	0.046
Q9Y4P8-3	Isoform 3 of WD repeat domain phosphoinositide-interacting protein 2 OS=Homo sapiens GN=WIP12 - [WIP12_HUMAN]	0.657	0.050
Q6UWJ1-3	Isoform 3 of Transmembrane and coiled-coil domain-containing protein 3 OS=Homo sapiens GN=TMCO3 - [TMCO3_HUMAN]	0.667	0.036
P09493-3	Isoform 3 of Tropomyosin alpha-1 chain OS=Homo sapiens GN=TPM1 - [TPM1_HUMAN]	0.681	0.048
O95210	Starch-binding domain-containing protein 1 OS=Homo sapiens GN=STBD1 PE=1 SV=1 - [STBD1_HUMAN]	0.686	0.015
Q9Y314	Nitric oxide synthase-interacting protein OS=Homo sapiens GN=NOSIP PE=1 SV=1 - [NOSIP_HUMAN]	0.687	0.040
Q12982	BCL2/adenovirus E1B 19 kDa protein-interacting protein 2 OS=Homo sapiens GN=BNIP2 PE=1 SV=1 - [BNIP2_HUMAN]	0.690	0.015
P42677	40S ribosomal protein S27 OS=Homo sapiens GN=RPS27 PE=1 SV=3 - [RS27_HUMAN]	0.691	0.044
P68036	Ubiquitin-conjugating enzyme E2 L3 OS=Homo sapiens GN=UBE2L3 PE=1 SV=1 - [UB2L3_HUMAN]	0.692	0.047
Q6JQN1-4	Isoform 4 of Acyl-CoA dehydrogenase family member 10 OS=Homo sapiens GN=ACAD10 - [ACD10_HUMAN]	0.697	0.022
Q96S52-2	Isoform 2 of GPI transamidase component PIG-S OS=Homo sapiens GN=PIGS - [PIGS_HUMAN]	0.698	0.042
Q15847	Adipose most abundant gene transcript 2 protein OS=Homo sapiens GN=APM2 PE=1 SV=1 - [APM2_HUMAN]	0.699	0.029
O75607	Nucleoplasmin-3 OS=Homo sapiens GN=NPM3 PE=1 SV=3 - [NPM3_HUMAN]	0.700	0.039
P10620	Microsomal glutathione S-transferase 1 OS=Homo sapiens GN=MGST1 PE=1 SV=1 - [MGST1_HUMAN]	0.703	0.030
Q9UPW8	Protein unc-13 homolog A OS=Homo sapiens GN=UNC13A PE=2 SV=4 - [UN13A_HUMAN]	0.707	0.011
P09429	High mobility group protein B1 OS=Homo sapiens GN=HMGB1 PE=1 SV=3 - [HMGB1_HUMAN]	0.715	0.033
Q9UK76	Hematological and neurological expressed 1 protein OS=Homo sapiens GN=HN1 PE=1 SV=3 - [HN1_HUMAN]	0.720	0.042
O60333-3	Isoform 3 of Kinesin-like protein KIF1B OS=Homo sapiens GN=KIF1B - [KIF1B_HUMAN]	0.724	0.011
Q5TAQ9	DDB1- and CUL4-associated factor 8 OS=Homo sapiens GN=DCAF8 PE=1 SV=1 - [DCAF8_HUMAN]	0.728	0.012
Q8WUM9	Sodium-dependent phosphate transporter 1 OS=Homo sapiens GN=SLC20A1 PE=1 SV=1 - [S20A1_HUMAN]	0.729	0.013
P10619	Lysosomal protective protein OS=Homo sapiens GN=CTSA PE=1 SV=2 - [PPGB_HUMAN]	0.731	0.013
Q13601	KRR1 small subunit processome component homolog OS=Homo sapiens GN=KRR1 PE=1 SV=4 - [KRR1_HUMAN]	0.736	0.003
Q15370	Transcription elongation factor B polypeptide 2 OS=Homo sapiens GN=TCEB2 PE=1 SV=1 - [ELOB_HUMAN]	0.744	0.026
P55040	GTP-binding protein GEM OS=Homo sapiens GN=GEM PE=1 SV=1 - [GEM_HUMAN]	0.748	0.039
Q76N89	E3 ubiquitin-protein ligase HECW1 OS=Homo sapiens GN=HECW1 PE=1 SV=3 - [HECW1_HUMAN]	0.750	0.031
Q8NC51-2	Isoform 2 of Plasminogen activator inhibitor 1 RNA-binding protein OS=Homo sapiens GN=SERP1 - [PAIRB_HUMAN]	0.754	0.015

Q504U0	Uncharacterized protein C4orf46 OS=Homo sapiens GN=C4orf46 PE=2 SV=1 - [CD046_HUMAN]	0.755	0.041
Q96II8	Leucine-rich repeat and calponin homology domain-containing protein 3 OS=Homo sapiens GN=LRCH3 PE=1 SV=2 - [LRCH3_HUMAN]	0.756	0.024
Q8NI22	Multiple coagulation factor deficiency protein 2 OS=Homo sapiens GN=MCFD2 PE=1 SV=1 - [MCFD2_HUMAN]	0.762	0.003
Q9H147	Deoxynucleotidyltransferase terminal-interacting protein 1 OS=Homo sapiens GN=DNTTIP1 PE=1 SV=2 - [TDIF1_HUMAN]	0.763	0.008
O43759-2	Isoform 1B of Synaptogyrin-1 OS=Homo sapiens GN=SYNGR1 - [SNG1_HUMAN]	0.766	0.035
P0C0S5	Histone H2A.Z OS=Homo sapiens GN=H2AFZ PE=1 SV=2 - [H2AZ_HUMAN]	0.766	0.016
Q96DG6	Carboxymethylenebutenolidase homolog OS=Homo sapiens GN=CMBL PE=1 SV=1 - [CMBL_HUMAN]	0.766	0.006
O43865	Putative adenosylhomocysteinase 2 OS=Homo sapiens GN=AHCYL1 PE=1 SV=2 - [SAHH2_HUMAN]	0.768	0.019
Q9H1C7	UPF0467 protein C5orf32 OS=Homo sapiens GN=C5orf32 PE=2 SV=1 - [CE032_HUMAN]	0.770	0.035
Q6PCE3	Glucose 1,6-bisphosphate synthase OS=Homo sapiens GN=PGM2L1 PE=1 SV=3 - [PGM2L_HUMAN]	0.771	0.043
Q05707-2	Isoform 2 of Collagen alpha-1(XIV) chain OS=Homo sapiens GN=COL14A1 - [COEA1_HUMAN]	0.771	0.046
Q68CZ2	Tensin-3 OS=Homo sapiens GN=TNS3 PE=1 SV=2 - [TENS3_HUMAN]	0.774	0.029
P53701	Cytochrome c-type heme lyase OS=Homo sapiens GN=HCCS PE=1 SV=1 - [CCHL_HUMAN]	0.775	0.044
P63241	Eukaryotic translation initiation factor 5A-1 OS=Homo sapiens GN=EIF5A PE=1 SV=2 - [IF5A1_HUMAN]	0.778	0.031
P41227	N-alpha-acetyltransferase 10 OS=Homo sapiens GN=NAA10 PE=1 SV=1 - [NAA10_HUMAN]	0.778	0.037
Q16585	Beta-sarcoglycan OS=Homo sapiens GN=SGCB PE=1 SV=1 - [SGCB_HUMAN]	0.779	0.007
A8MXV4	Nucleoside diphosphate-linked moiety X motif 19, mitochondrial OS=Homo sapiens GN=NUDT19 PE=1 SV=1 - [NUDT19_HUMAN]	0.781	0.016
O43493-2	Isoform TGN46 of Trans-Golgi network integral membrane protein 2 OS=Homo sapiens GN=TGOLN2 - [TGON2_HUMAN]	0.783	0.020
Q8N4V1	Membrane magnesium transporter 1 OS=Homo sapiens GN=MMGT1 PE=1 SV=1 - [MMGT1_HUMAN]	0.783	0.008
O43676	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 3 OS=Homo sapiens GN=NDUFB3 PE=1 SV=3 - [NDUB3_HUMAN]	0.784	0.030
P49069	Calcium signal-modulating cyclophilin ligand OS=Homo sapiens GN=CAMLG PE=1 SV=1 - [CAMLG_HUMAN]	0.784	0.046
P61326	Protein mago nashi homolog OS=Homo sapiens GN=MAGOH PE=1 SV=1 - [MGN_HUMAN]	0.784	0.032
P14921-2	Isoform c-ETS-1B of Protein C-ets-1 OS=Homo sapiens GN=ETS1 - [ETS1_HUMAN]	0.787	0.002
Q99959-2	Isoform 1 of Plakophilin-2 OS=Homo sapiens GN=PKP2 - [PKP2_HUMAN]	0.788	0.026
O75695	Protein XRP2 OS=Homo sapiens GN=RP2 PE=1 SV=4 - [XRP2_HUMAN]	0.788	0.001
Q15365	Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2 - [PCBP1_HUMAN]	0.790	0.029
P62873	Guanine nucleotide-binding protein G(I)/G(S)/G(T) subunit beta-1 OS=Homo sapiens GN=GNB1 PE=1 SV=3 - [GBB1_HUMAN]	0.790	0.020
Q99536	Synaptic vesicle membrane protein VAT-1 homolog OS=Homo sapiens GN=VAT1 PE=1 SV=2 - [VAT1_HUMAN]	0.790	0.048
P60953	Cell division control protein 42 homolog OS=Homo sapiens GN=CDC42 PE=1 SV=2 - [CDC42_HUMAN]	0.792	0.010

Q9UKD2	mRNA turnover protein 4 homolog OS=Homo sapiens GN=MRTO4 PE=1 SV=2 - [MRT4_HUMAN]	0.795	0.014
Q92551	Inositol hexakisphosphate kinase 1 OS=Homo sapiens GN=IP6K1 PE=1 SV=3 - [IP6K1_HUMAN]	0.796	0.038
Q9H814	Phosphorylated adapter RNA export protein OS=Homo sapiens GN=PHAX PE=1 SV=1 - [PHAX_HUMAN]	0.802	0.025
Q02952	A-kinase anchor protein 12 OS=Homo sapiens GN=AKAP12 PE=1 SV=4 - [AKA12_HUMAN]	0.804	0.042
Q86XL3	Ankyrin repeat and LEM domain-containing protein 2 OS=Homo sapiens GN=ANKLE2 PE=1 SV=4 - [ANKL2_HUMAN]	0.807	0.006
P04818	Thymidylate synthase OS=Homo sapiens GN=TYMS PE=1 SV=3 - [TYSY_HUMAN]	0.810	0.040
Q9UL26	Ras-related protein Rab-22A OS=Homo sapiens GN=RAB22A PE=1 SV=2 - [RB22A_HUMAN]	0.813	0.048
O43324	Eukaryotic translation elongation factor 1 epsilon-1 OS=Homo sapiens GN=EEF1E1 PE=1 SV=1 - [MCA3_HUMAN]	0.817	0.017
O43752	Syntaxin-6 OS=Homo sapiens GN=STX6 PE=1 SV=1 - [STX6_HUMAN]	0.819	0.018
P00736	Complement C1r subcomponent OS=Homo sapiens GN=C1R PE=1 SV=2 - [C1R_HUMAN]	0.819	0.015
O75347	Tubulin-specific chaperone A OS=Homo sapiens GN=TBCA PE=1 SV=3 - [TBCA_HUMAN]	0.821	0.037
Q9BX40	Protein LSM14 homolog B OS=Homo sapiens GN=LSM14B PE=1 SV=1 - [LS14B_HUMAN]	0.826	0.049
Q9ULJ7	Ankyrin repeat domain-containing protein 50 OS=Homo sapiens GN=ANKRD50 PE=1 SV=4 - [ANR50_HUMAN]	0.826	0.017
P27695	DNA-(apurinic or apyrimidinic site) lyase OS=Homo sapiens GN=APEX1 PE=1 SV=2 - [APEX1_HUMAN]	0.827	0.002
Q5M775-5	Isoform 5 of Cytospin-B OS=Homo sapiens GN=SPECC1 - [CYTSB_HUMAN]	0.831	0.048
Q5T3F8	Transmembrane protein 63B OS=Homo sapiens GN=TMEM63B PE=1 SV=1 - [TM63B_HUMAN]	0.831	0.007
Q9NRF8	CTP synthase 2 OS=Homo sapiens GN=CTPS2 PE=1 SV=1 - [PYRG2_HUMAN]	0.832	0.030
Q674X7-3	Isoform 3 of Kazrin OS=Homo sapiens GN=KAZN - [KAZRN_HUMAN]	0.836	0.047
Q9UKL0	REST corepressor 1 OS=Homo sapiens GN=RCOR1 PE=1 SV=1 - [RCOR1_HUMAN]	0.836	0.007
Q14696	LDLR chaperone MESD OS=Homo sapiens GN=MESDC2 PE=1 SV=2 - [MESD_HUMAN]	0.837	0.027
Q01543-2	Isoform 2 of Friend leukemia integration 1 transcription factor OS=Homo sapiens GN=FLI1 - [FLI1_HUMAN]	0.840	0.049
Q16342	Programmed cell death protein 2 OS=Homo sapiens GN=PDCD2 PE=1 SV=2 - [PDCD2_HUMAN]	0.842	0.012
Q9NZI7-4	Isoform 2 of Upstream-binding protein 1 OS=Homo sapiens GN=UBP1 - [UBIP1_HUMAN]	0.842	0.040
Q8ND56	Protein LSM14 homolog A OS=Homo sapiens GN=LSM14A PE=1 SV=3 - [LS14A_HUMAN]	0.845	0.040
Q96PK6	RNA-binding protein 14 OS=Homo sapiens GN=RBM14 PE=1 SV=2 - [RBM14_HUMAN]	0.845	0.011
Q96RE7	Nucleus accumbens-associated protein 1 OS=Homo sapiens GN=NACC1 PE=1 SV=1 - [NACC1_HUMAN]	0.847	0.041
Q5SWX8-3	Isoform 3 of Protein odr-4 homolog OS=Homo sapiens GN=ODR4 - [ODR4_HUMAN]	0.848	0.007
Q9NWM3	CUE domain-containing protein 1 OS=Homo sapiens GN=CUEDC1 PE=1 SV=1 - [CUED1_HUMAN]	0.849	0.044
Q8WUW1	Protein BRICK1 OS=Homo sapiens GN=BRK1 PE=1 SV=1 - [BRK1_HUMAN]	0.850	0.049

P15408-3	Isoform 3 of Fos-related antigen 2 OS=Homo sapiens GN=FOSL2 - [FOSL2_HUMAN]	0.851	0.004
Q96B26	Exosome complex component RRP43 OS=Homo sapiens GN=EXOSC8 PE=1 SV=1 - [EXOSC8_HUMAN]	0.852	0.021
P46783	40S ribosomal protein S10 OS=Homo sapiens GN=RPS10 PE=1 SV=1 - [RPS10_HUMAN]	0.853	0.024
Q9NX58	Cell growth-regulating nucleolar protein OS=Homo sapiens GN=LYAR PE=1 SV=2 - [LYAR_HUMAN]	0.854	0.038
Q13490	Baculoviral IAP repeat-containing protein 2 OS=Homo sapiens GN=BIRC2 PE=1 SV=2 - [BIRC2_HUMAN]	0.855	0.039
P51114	Fragile X mental retardation syndrome-related protein 1 OS=Homo sapiens GN=FXR1 PE=1 SV=3 - [FXR1_HUMAN]	0.855	0.017
Q9UBL3-2	Isoform 2 of Set1/Ash2 histone methyltransferase complex subunit ASH2 OS=Homo sapiens GN=ASH2L - [ASH2L_HUMAN]	0.856	0.034
Q16543	Hsp90 co-chaperone Cdc37 OS=Homo sapiens GN=CDC37 PE=1 SV=1 - [CDC37_HUMAN]	0.861	0.032
Q9HBL0	Tensin-1 OS=Homo sapiens GN=TNS1 PE=1 SV=2 - [TNS1_HUMAN]	0.861	0.039
P50750	Cyclin-dependent kinase 9 OS=Homo sapiens GN=CDK9 PE=1 SV=3 - [CDK9_HUMAN]	0.862	0.013
P61244-3	Isoform 3 of Protein max OS=Homo sapiens GN=MAX - [MAX_HUMAN]	0.862	0.034
Q03111	Protein ENL OS=Homo sapiens GN=MLLT1 PE=1 SV=2 - [ENL_HUMAN]	0.862	0.048
Q96CP2	FLYWCH family member 2 OS=Homo sapiens GN=FLYWCH2 PE=1 SV=1 - [FWCH2_HUMAN]	0.863	0.033
Q14161-2	Isoform 2 of ARF GTPase-activating protein GIT2 OS=Homo sapiens GN=GIT2 - [GIT2_HUMAN]	0.863	0.049
Q12874	Splicing factor 3A subunit 3 OS=Homo sapiens GN=SF3A3 PE=1 SV=1 - [SF3A3_HUMAN]	0.863	0.032
P00387-2	Isoform 2 of NADH-cytochrome b5 reductase 3 OS=Homo sapiens GN=CYB5R3 - [NB5R3_HUMAN]	0.864	0.011
Q96PN7-2	Isoform 2 of Transcriptional-regulating factor 1 OS=Homo sapiens GN=TRERF1 - [TREF1_HUMAN]	0.868	0.037
Q9BVM4	Gamma-glutamylaminocyclotransferase OS=Homo sapiens GN=A2LD1 PE=1 SV=2 - [A2LD1_HUMAN]	0.869	0.038
O43913-2	Isoform 2 of Origin recognition complex subunit 5 OS=Homo sapiens GN=ORC5 - [ORC5_HUMAN]	0.870	0.040
Q96GZ6-6	Isoform 6 of Solute carrier family 41 member 3 OS=Homo sapiens GN=SLC41A3 - [S41A3_HUMAN]	0.870	0.047
O14818	Proteasome subunit alpha type-7 OS=Homo sapiens GN=PSMA7 PE=1 SV=1 - [PSA7_HUMAN]	0.873	0.038
Q96DI7	U5 small nuclear ribonucleoprotein 40 kDa protein OS=Homo sapiens GN=SNRNP40 PE=1 SV=1 - [SNR40_HUMAN]	0.874	0.035
Q14393-3	Isoform 3 of Growth arrest-specific protein 6 OS=Homo sapiens GN=GAS6 - [GAS6_HUMAN]	0.875	0.040
Q8N138-4	Isoform 2 of ORM1-like protein 3 OS=Homo sapiens GN=ORMDL3 - [ORML3_HUMAN]	0.875	0.001
Q9Y448	Putative TRAF4-associated factor 1 OS=Homo sapiens GN=TRAF4AF1 PE=1 SV=2 - [T4AF1_HUMAN]	0.876	0.009
Q9UPT8	Zinc finger CCCH domain-containing protein 4 OS=Homo sapiens GN=ZC3H4 PE=1 SV=3 - [ZC3H4_HUMAN]	0.878	0.035
O14641	Segment polarity protein dishevelled homolog DVL-2 OS=Homo sapiens GN=DVL2 PE=1 SV=1 - [DVL2_HUMAN]	0.880	0.049
P08107	Heat shock 70 kDa protein 1A/1B OS=Homo sapiens GN=HSPA1A PE=1 SV=5 - [HSP71_HUMAN]	0.880	0.013
O00567	Nucleolar protein 56 OS=Homo sapiens GN=NOP56 PE=1 SV=4 - [NOP56_HUMAN]	0.881	0.025

Q00423	Echinoderm microtubule-associated protein-like 1 OS=Homo sapiens GN=EML1 PE=1 SV=3 - [EMAL1_HUMAN]	0.882	0.049
Q9Y282	Endoplasmic reticulum-Golgi intermediate compartment protein 3 OS=Homo sapiens GN=ERGIC3 PE=1 SV=1 - [ERGI3_HUMAN]	0.882	0.033
Q5PRF9	Protein Smaug homolog 2 OS=Homo sapiens GN=SAMD4B PE=1 SV=1 - [SMAG2_HUMAN]	0.883	0.034
P02786	Transferrin receptor protein 1 OS=Homo sapiens GN=TFRC PE=1 SV=2 - [TFR1_HUMAN]	0.884	0.045
Q92845	Kinesin-associated protein 3 OS=Homo sapiens GN=KIFAP3 PE=1 SV=2 - [KIFA3_HUMAN]	0.884	0.027
P54652	Heat shock-related 70 kDa protein 2 OS=Homo sapiens GN=HSPA2 PE=1 SV=1 - [HSP72_HUMAN]	0.886	0.005
Q13905-2	Isoform Short of Rap guanine nucleotide exchange factor 1 OS=Homo sapiens GN=RAPGEF1 - [RPGF1_HUMAN]	0.887	0.042
Q5T6V5	UPF0553 protein C9orf64 OS=Homo sapiens GN=C9orf64 PE=1 SV=1 - [CI064_HUMAN]	0.887	0.007
Q94992	Protein HEXIM1 OS=Homo sapiens GN=HEXIM1 PE=1 SV=1 - [HEXI1_HUMAN]	0.887	0.019
P61916	Epididymal secretory protein E1 OS=Homo sapiens GN=NPC2 PE=1 SV=1 - [NPC2_HUMAN]	0.892	0.020
P25398	40S ribosomal protein S12 OS=Homo sapiens GN=RPS12 PE=1 SV=3 - [RS12_HUMAN]	0.892	0.028
Q96ST2-2	Isoform 2 of Protein IWS1 homolog OS=Homo sapiens GN=IWS1 - [IWS1_HUMAN]	0.893	0.042
O43402	Neighbor of COX4 OS=Homo sapiens GN=COX4NB PE=1 SV=1 - [CX4NB_HUMAN]	0.893	0.046
Q9BV79-2	Isoform 2 of Trans-2-enoyl-CoA reductase, mitochondrial OS=Homo sapiens GN=MECR - [MECR_HUMAN]	0.893	0.020
P26641	Elongation factor 1-gamma OS=Homo sapiens GN=EEF1G PE=1 SV=3 - [EF1G_HUMAN]	0.894	0.016
Q92615	La-related protein 4B OS=Homo sapiens GN=LARP4B PE=1 SV=3 - [LAR4B_HUMAN]	0.894	0.048
Q9P2K5-2	Isoform 2 of Myelin expression factor 2 OS=Homo sapiens GN=MYEF2 - [MYEF2_HUMAN]	0.895	0.047
Q9NVA1-2	Isoform 2 of Ubiquinol-cytochrome c reductase complex chaperone CBP3 homolog OS=Homo sapiens GN=UQCC - [UQCC_HUMAN]	0.895	0.044
Q9H490-2	Isoform 2 of Phosphatidylinositol glycan anchor biosynthesis class U protein OS=Homo sapiens GN=PIGU - [PIGU_HUMAN]	0.896	0.031
Q9NVC6	Mediator of RNA polymerase II transcription subunit 17 OS=Homo sapiens GN=MED17 PE=1 SV=2 - [MED17_HUMAN]	0.896	0.046
P22626	Heterogeneous nuclear ribonucleoproteins A2/B1 OS=Homo sapiens GN=HNRNPA2B1 PE=1 SV=2 - [ROA2_HUMAN]	0.896	0.010
P62937	Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens GN=PPIA PE=1 SV=2 - [PPIA_HUMAN]	0.897	0.019
Q5HYK3	2-methoxy-6-polyprenyl-1,4-benzoquinol methylase, mitochondrial OS=Homo sapiens GN=COQ5 PE=1 SV=2 - [COQ5_HUMAN]	0.897	0.027
Q9Y2D5-6	Isoform 4 of A-kinase anchor protein 2 OS=Homo sapiens GN=AKAP2 - [AKAP2_HUMAN]	0.897	0.046
Q9NSV4	Protein diaphanous homolog 3 OS=Homo sapiens GN=DIAPH3 PE=1 SV=4 - [DIAP3_HUMAN]	0.901	0.048
Q13976	cGMP-dependent protein kinase 1 OS=Homo sapiens GN=PRKG1 PE=1 SV=3 - [KGP1_HUMAN]	0.902	0.026
P30046	D-dopachrome decarboxylase OS=Homo sapiens GN=DDT PE=1 SV=3 - [DOPD_HUMAN]	0.903	0.019
P51151	Ras-related protein Rab-9A OS=Homo sapiens GN=RAB9A PE=1 SV=1 - [RAB9A_HUMAN]	0.905	0.002
Q92581-2	Isoform 2 of Sodium/hydrogen exchanger 6 OS=Homo sapiens GN=SLC9A6 - [SL9A6_HUMAN]	0.906	0.039

P08754	Guanine nucleotide-binding protein G(k) subunit alpha OS=Homo sapiens GN=GNAI3 PE=1 SV=3 - [GNAI3_HUMAN]	0.906	0.043
Q14061	Cytochrome c oxidase copper chaperone OS=Homo sapiens GN=COX17 PE=1 SV=2 - [COX17_HUMAN]	0.907	0.012
P09012	U1 small nuclear ribonucleoprotein A OS=Homo sapiens GN=SNRPA PE=1 SV=3 - [SNRPA_HUMAN]	0.907	0.026
P18621	60S ribosomal protein L17 OS=Homo sapiens GN=RPL17 PE=1 SV=3 - [RPL17_HUMAN]	0.909	0.018
O14908	PDZ domain-containing protein GIPC1 OS=Homo sapiens GN=GIPC1 PE=1 SV=2 - [GIPC1_HUMAN]	0.909	0.044
P54727	UV excision repair protein RAD23 homolog B OS=Homo sapiens GN=RAD23B PE=1 SV=1 - [RAD23B_HUMAN]	0.911	0.035
Q9Y3F4	Serine-threonine kinase receptor-associated protein OS=Homo sapiens GN=STRAP PE=1 SV=1 - [STRAP_HUMAN]	0.911	0.034
P07237	Protein disulfide-isomerase OS=Homo sapiens GN=P4HB PE=1 SV=3 - [PDIA1_HUMAN]	0.912	0.010
Q5H9R7-3	Isoform 3 of Serine/threonine-protein phosphatase 6 regulatory subunit 3 OS=Homo sapiens GN=PPP6R3 - [PPP6R3_HUMAN]	0.912	0.038
P11279	Lysosome-associated membrane glycoprotein 1 OS=Homo sapiens GN=LAMP1 PE=1 SV=3 - [LAMP1_HUMAN]	0.914	0.008
O15042-2	Isoform 2 of U2 snRNP-associated SURP motif-containing protein OS=Homo sapiens GN=U2SURP - [SR140_HUMAN]	0.915	0.042
Q2NKX8	DNA excision repair protein ERCC-6-like OS=Homo sapiens GN=ERCC6L PE=1 SV=1 - [ERC6L_HUMAN]	0.915	0.044
Q13443	Disintegrin and metalloproteinase domain-containing protein 9 OS=Homo sapiens GN=ADAM9 PE=1 SV=1 - [ADAM9_HUMAN]	0.916	0.015
Q9ULR0	Pre-mRNA-splicing factor ISY1 homolog OS=Homo sapiens GN=ISY1 PE=1 SV=3 - [ISY1_HUMAN]	0.916	0.020
Q14680	Maternal embryonic leucine zipper kinase OS=Homo sapiens GN=MELK PE=1 SV=3 - [MELK_HUMAN]	0.918	0.000
Q9NVP1	ATP-dependent RNA helicase DDX18 OS=Homo sapiens GN=DDX18 PE=1 SV=2 - [DDX18_HUMAN]	0.918	0.041
Q9NXW2	DnaJ homolog subfamily B member 12 OS=Homo sapiens GN=DNAJB12 PE=1 SV=4 - [DJB12_HUMAN]	0.919	0.023
Q8N3Z3-3	Isoform 3 of GTP-binding protein 8 OS=Homo sapiens GN=GTPBP8 - [GTPBP8_HUMAN]	0.920	0.024
P36551	Coproporphyrinogen-III oxidase, mitochondrial OS=Homo sapiens GN=CPOX PE=1 SV=3 - [HEM6_HUMAN]	0.920	0.043
P08397-2	Isoform 2 of Porphobilinogen deaminase OS=Homo sapiens GN=HMBS - [HEM3_HUMAN]	0.920	0.048
Q14C86-2	Isoform 2 of GTPase-activating protein and VPS9 domain-containing protein 1 OS=Homo sapiens GN=GAPVD1 - [GAPD1_HUMAN]	0.921	0.033
Q9BW62	Katanin p60 ATPase-containing subunit A-like 1 OS=Homo sapiens GN=KATNAL1 PE=2 SV=1 - [KATL1_HUMAN]	0.921	0.037
Q86UP2	Kinectin OS=Homo sapiens GN=KTN1 PE=1 SV=1 - [KTN1_HUMAN]	0.922	0.039
A6NDG6	Phosphoglycolate phosphatase OS=Homo sapiens GN=PGP PE=1 SV=1 - [PGP_HUMAN]	0.923	0.004
O43719	HIV Tat-specific factor 1 OS=Homo sapiens GN=HTATSF1 PE=1 SV=1 - [HTSF1_HUMAN]	0.924	0.033
P17844	Probable ATP-dependent RNA helicase DDX5 OS=Homo sapiens GN=DDX5 PE=1 SV=1 - [DDX5_HUMAN]	0.925	0.025
Q71RC2-3	Isoform 3 of La-related protein 4 OS=Homo sapiens GN=LARP4 - [LARP4_HUMAN]	0.926	0.020
Q8IV38	Ankyrin repeat and MYND domain-containing protein 2 OS=Homo sapiens GN=ANKMY2 PE=1 SV=1 - [ANKY2_HUMAN]	0.928	0.049
Q13564	NEDD8-activating enzyme E1 regulatory subunit OS=Homo sapiens GN=NAE1 PE=1 SV=1 - [ULA1_HUMAN]	0.929	0.010

Q8NBX0	Saccharopine dehydrogenase-like oxidoreductase OS=Homo sapiens GN=SCCPDH PE=1 SV=1 - [SCPDH_HUMAN]	0.929	0.024
Q93050-1	Isoform 2 of V-type proton ATPase 116 kDa subunit a isoform 1 OS=Homo sapiens GN=ATP6V0A1 - [VPP1_HUMAN]	0.929	0.008
Q92930	Ras-related protein Rab-8B OS=Homo sapiens GN=RAB8B PE=1 SV=2 - [RAB8B_HUMAN]	0.930	0.031
Q9BQG0	Myb-binding protein 1A OS=Homo sapiens GN=MYBBP1A PE=1 SV=2 - [MBB1A_HUMAN]	0.930	0.047
P14625	Endoplasmic reticulum chaperone protein OS=Homo sapiens GN=HSP90B1 PE=1 SV=1 - [ENPL_HUMAN]	0.931	0.037
Q9Y266	Nuclear migration protein nudC OS=Homo sapiens GN=NUDC PE=1 SV=1 - [NUDC_HUMAN]	0.932	0.047
Q92966	snRNA-activating protein complex subunit 3 OS=Homo sapiens GN=SNAPC3 PE=1 SV=1 - [SNPC3_HUMAN]	0.932	0.037
Q9UGP8	Translocation protein SEC63 homolog OS=Homo sapiens GN=SEC63 PE=1 SV=2 - [SEC63_HUMAN]	0.933	0.017
Q7L523	Ras-related GTP-binding protein A OS=Homo sapiens GN=RRAGA PE=1 SV=1 - [RRAGA_HUMAN]	0.933	0.028
Q5JSH3-2	Isoform 2 of WD repeat-containing protein 44 OS=Homo sapiens GN=WDR44 - [WDR44_HUMAN]	0.935	0.023
Q63HQ0	AP-1 complex-associated regulatory protein OS=Homo sapiens GN=AP1AR PE=1 SV=1 - [AP1AR_HUMAN]	0.938	0.013
Q7Z6E9	E3 ubiquitin-protein ligase RBBP6 OS=Homo sapiens GN=RBBP6 PE=1 SV=1 - [RBBP6_HUMAN]	0.939	0.043
O60346	PH domain leucine-rich repeat-containing protein phosphatase 1 OS=Homo sapiens GN=PHLPP1 PE=1 SV=3 - [PHLP1_HUMAN]	0.940	0.035
P61923	Coatamer subunit zeta-1 OS=Homo sapiens GN=COPZ1 PE=1 SV=1 - [COPZ1_HUMAN]	0.941	0.044
P08758	Annexin A5 OS=Homo sapiens GN=ANXA5 PE=1 SV=2 - [ANXA5_HUMAN]	0.941	0.048
Q93009	Ubiquitin carboxyl-terminal hydrolase 7 OS=Homo sapiens GN=USP7 PE=1 SV=2 - [UBP7_HUMAN]	0.942	0.008
Q96TC7	Regulator of microtubule dynamics protein 3 OS=Homo sapiens GN=FAM82A2 PE=1 SV=2 - [RMD3_HUMAN]	0.942	0.035
P49662	Caspase-4 OS=Homo sapiens GN=CASP4 PE=1 SV=1 - [CASP4_HUMAN]	0.945	0.040
Q06323	Proteasome activator complex subunit 1 OS=Homo sapiens GN=PSME1 PE=1 SV=1 - [PSME1_HUMAN]	0.945	0.039
O15305	Phosphomannomutase 2 OS=Homo sapiens GN=PMM2 PE=1 SV=1 - [PMM2_HUMAN]	0.945	0.028
P46821	Microtubule-associated protein 1B OS=Homo sapiens GN=MAP1B PE=1 SV=2 - [MAP1B_HUMAN]	0.946	0.017
Q9P2W9	Syntaxin-18 OS=Homo sapiens GN=STX18 PE=1 SV=1 - [STX18_HUMAN]	0.946	0.009
Q92685	Dol-P-Man:Man(5)GlcNAc(2)-PP-Dol alpha-1,3-mannosyltransferase OS=Homo sapiens GN=ALG3 PE=1 SV=1 - [ALG3_HUMAN]	0.946	0.029
Q0VDG4	Secernin-3 OS=Homo sapiens GN=SCRN3 PE=1 SV=1 - [SCRN3_HUMAN]	0.948	0.043
P09651	Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNPA1 PE=1 SV=5 - [ROA1_HUMAN]	0.948	0.037
P18206-2	Isoform 1 of Vinculin OS=Homo sapiens GN=VCL - [VINC_HUMAN]	0.949	0.050
P43034	Platelet-activating factor acetylhydrolase 1B subunit alpha OS=Homo sapiens GN=PAFAH1B1 PE=1 SV=2 - [LIS1_HUMAN]	0.949	0.020
Q13459-2	Isoform Short of Myosin-IXb OS=Homo sapiens GN=MYO9B - [MYO9B_HUMAN]	0.950	0.010
O00151	PDZ and LIM domain protein 1 OS=Homo sapiens GN=PDLIM1 PE=1 SV=4 - [PDL1_HUMAN]	0.950	0.029

O94760	N(G),N(G)-dimethylarginine dimethylaminohydrolase 1 OS=Homo sapiens GN=DDAH1 PE=1 SV=3 - [DDAH1_HUMAN]	0.950	0.042
Q99653	Calcium-binding protein p22 OS=Homo sapiens GN=CHP PE=1 SV=3 - [CHP1_HUMAN]	0.952	0.045
O75396	Vesicle-trafficking protein SEC22b OS=Homo sapiens GN=SEC22B PE=1 SV=4 - [SC22B_HUMAN]	0.952	0.042
P39023	60S ribosomal protein L3 OS=Homo sapiens GN=RPL3 PE=1 SV=2 - [RL3_HUMAN]	0.953	0.046
P13797	Plastin-3 OS=Homo sapiens GN=PLS3 PE=1 SV=4 - [PLST_HUMAN]	0.953	0.026
P13639	Elongation factor 2 OS=Homo sapiens GN=EEF2 PE=1 SV=4 - [EF2_HUMAN]	0.953	0.046
Q99848	Probable rRNA-processing protein EBP2 OS=Homo sapiens GN=EBNA1BP2 PE=1 SV=2 - [EBP2_HUMAN]	0.953	0.035
Q9P265	Disco-interacting protein 2 homolog B OS=Homo sapiens GN=DIP2B PE=1 SV=3 - [DIP2B_HUMAN]	0.954	0.003
Q00341	Vigilin OS=Homo sapiens GN=HDLBP PE=1 SV=2 - [VIGLN_HUMAN]	0.955	0.023
Q70UQ0-4	Isoform 4 of Inhibitor of nuclear factor kappa-B kinase-interacting protein OS=Homo sapiens GN=IKBIP - [IKIP_HUMAN]	0.958	0.018
P62195	26S protease regulatory subunit 8 OS=Homo sapiens GN=PSMC5 PE=1 SV=1 - [PRS8_HUMAN]	0.960	0.031
Q9Y4W6	AFG3-like protein 2 OS=Homo sapiens GN=AFG3L2 PE=1 SV=2 - [AFG32_HUMAN]	0.961	0.023
P11413	Glucose-6-phosphate 1-dehydrogenase OS=Homo sapiens GN=G6PD PE=1 SV=4 - [G6PD_HUMAN]	0.961	0.011
Q14677	Clathrin interactor 1 OS=Homo sapiens GN=CLINT1 PE=1 SV=1 - [EPN4_HUMAN]	0.964	0.014
P28340	DNA polymerase delta catalytic subunit OS=Homo sapiens GN=POLD1 PE=1 SV=2 - [DPOD1_HUMAN]	0.977	0.040
P62258	14-3-3 protein epsilon OS=Homo sapiens GN=YWHAE PE=1 SV=1 - [1433E_HUMAN]	0.979	0.032
Q86W56-2	Isoform 2 of Poly(ADP-ribose) glycohydrolase OS=Homo sapiens GN=PARG - [PARG_HUMAN]	1.015	0.041
O43491	Band 4.1-like protein 2 OS=Homo sapiens GN=EPB41L2 PE=1 SV=1 - [E41L2_HUMAN]	1.025	0.046
Q7KZ85	Transcription elongation factor SPT6 OS=Homo sapiens GN=SUPT6H PE=1 SV=2 - [SPT6H_HUMAN]	1.029	0.040
Q9Y262	Eukaryotic translation initiation factor 3 subunit L OS=Homo sapiens GN=EIF3L PE=1 SV=1 - [EIF3L_HUMAN]	1.030	0.014
Q86UV5-2	Isoform 2 of Ubiquitin carboxyl-terminal hydrolase 48 OS=Homo sapiens GN=USP48 - [UBP48_HUMAN]	1.030	0.025
Q14669	Probable E3 ubiquitin-protein ligase TRIP12 OS=Homo sapiens GN=TRIP12 PE=1 SV=1 - [TRIPC_HUMAN]	1.031	0.047
Q96AE4	Far upstream element-binding protein 1 OS=Homo sapiens GN=FUBP1 PE=1 SV=3 - [FUBP1_HUMAN]	1.031	0.015
O43617	Trafficking protein particle complex subunit 3 OS=Homo sapiens GN=TRAPPC3 PE=1 SV=1 - [TPPC3_HUMAN]	1.042	0.045
Q5VIR6-4	Isoform 4 of Vacuolar protein sorting-associated protein 53 homolog OS=Homo sapiens GN=VPS53 - [VPS53_HUMAN]	1.042	0.020
Q3L8U1-2	Isoform 2 of Chromodomain-helicase-DNA-binding protein 9 OS=Homo sapiens GN=CHD9 - [CHD9_HUMAN]	1.042	0.036
P16435	NADPH--cytochrome P450 reductase OS=Homo sapiens GN=POR PE=1 SV=2 - [NCPR_HUMAN]	1.044	0.011
P68366	Tubulin alpha-4A chain OS=Homo sapiens GN=TUBA4A PE=1 SV=1 - [TBA4A_HUMAN]	1.053	0.034
Q9H0R6	Glutamyl-tRNA(Gln) amidotransferase subunit A, mitochondrial OS=Homo sapiens GN=QRSL1 PE=1 SV=2 - [GATA_HUMAN]	1.054	0.024

P17987	T-complex protein 1 subunit alpha OS=Homo sapiens GN=TCP1 PE=1 SV=1 - [TCPA_HUMAN]	1.057	0.004
Q0IIM8-2	Isoform 2 of TBC1 domain family member 8B OS=Homo sapiens GN=TBC1D8B - [TBC8B_HUMAN]	1.058	0.021
Q01970	1-phosphatidylinositol-4,5-bisphosphate phosphodiesterase beta-3 OS=Homo sapiens GN=PLCB3 PE=1 SV=2 - [PLCB3_HUMAN]	1.061	0.028
Q9BW60	Elongation of very long chain fatty acids protein 1 OS=Homo sapiens GN=ELOVL1 PE=1 SV=1 - [ELOV1_HUMAN]	1.061	0.013
Q5JVF3-3	Isoform 3 of PCI domain-containing protein 2 OS=Homo sapiens GN=PCID2 - [PCID2_HUMAN]	1.065	0.009
Q13596-2	Isoform 1A of Sorting nexin-1 OS=Homo sapiens GN=SNX1 - [SNX1_HUMAN]	1.065	0.040
Q6P2E9	Enhancer of mRNA-decapping protein 4 OS=Homo sapiens GN=EDC4 PE=1 SV=1 - [EDC4_HUMAN]	1.065	0.028
P51003	Poly(A) polymerase alpha OS=Homo sapiens GN=PAPOLA PE=1 SV=4 - [PAPOA_HUMAN]	1.068	0.020
Q99661-2	Isoform 2 of Kinesin-like protein KIF2C OS=Homo sapiens GN=KIF2C - [KIF2C_HUMAN]	1.068	0.035
P23258	Tubulin gamma-1 chain OS=Homo sapiens GN=TUBG1 PE=1 SV=2 - [TBG1_HUMAN]	1.068	0.029
Q93008-1	Isoform 2 of Probable ubiquitin carboxyl-terminal hydrolase FAF-X OS=Homo sapiens GN=USP9X - [USP9X_HUMAN]	1.069	0.039
Q92896	Golgi apparatus protein 1 OS=Homo sapiens GN=GLG1 PE=1 SV=2 - [GSLG1_HUMAN]	1.069	0.047
Q96GM8	Target of EGR1 protein 1 OS=Homo sapiens GN=TOE1 PE=1 SV=1 - [TOE1_HUMAN]	1.069	0.040
P45880-2	Isoform 2 of Voltage-dependent anion-selective channel protein 2 OS=Homo sapiens GN=VDAC2 - [VDAC2_HUMAN]	1.071	0.029
O15031	Plexin-B2 OS=Homo sapiens GN=PLXNB2 PE=1 SV=3 - [PLXB2_HUMAN]	1.071	0.024
Q147X3	N-alpha-acetyltransferase 30 OS=Homo sapiens GN=NAA30 PE=1 SV=1 - [NAA30_HUMAN]	1.073	0.049
O75879	Glutamyl-tRNA(Gln) amidotransferase subunit B, mitochondrial OS=Homo sapiens GN=PET112 PE=1 SV=1 - [GATB_HUMAN]	1.073	0.049
Q9NY61	Protein AATF OS=Homo sapiens GN=AATF PE=1 SV=1 - [AATF_HUMAN]	1.074	0.021
Q9ULW0	Targeting protein for Xklp2 OS=Homo sapiens GN=TPX2 PE=1 SV=2 - [TPX2_HUMAN]	1.074	0.046
Q92734	Protein TFG OS=Homo sapiens GN=TFG PE=1 SV=2 - [TFG_HUMAN]	1.077	0.014
Q96H20-2	Isoform 2 of Vacuolar-sorting protein SNF8 OS=Homo sapiens GN=SNF8 - [SNF8_HUMAN]	1.082	0.032
Q9BW19	Kinesin-like protein KIFC1 OS=Homo sapiens GN=KIFC1 PE=1 SV=2 - [KIFC1_HUMAN]	1.085	0.007
P62136	Serine/threonine-protein phosphatase PP1-alpha catalytic subunit OS=Homo sapiens GN=PPP1CA PE=1 SV=1 - [PP1A_HUMAN]	1.085	0.038
Q5MIZ7-3	Isoform 3 of Serine/threonine-protein phosphatase 4 regulatory subunit 3B OS=Homo sapiens GN=SMEK2 - [P4R3B_HUMAN]	1.090	0.004
Q96CW6	Probable RNA polymerase II nuclear localization protein SLC7A6OS OS=Homo sapiens GN=SLC7A6OS PE=1 SV=2 - [S7A6O_HUMAN]	1.091	0.005
Q9UMY1-2	Isoform 2 of Nucleolar protein 7 OS=Homo sapiens GN=NOL7 - [NOL7_HUMAN]	1.092	0.042
Q13155	Aminoacyl tRNA synthase complex-interacting multifunctional protein 2 OS=Homo sapiens GN=AIMP2 PE=1 SV=2 - [AIMP2_HUMAN]	1.093	0.044
Q96BJ3	Axin interactor, dorsalization-associated protein OS=Homo sapiens GN=AIDA PE=1 SV=1 - [AIDA_HUMAN]	1.093	0.021
O95163	Elongator complex protein 1 OS=Homo sapiens GN=IKBKAP PE=1 SV=3 - [ELP1_HUMAN]	1.095	0.044

Q9UGV2-3	Isoform 3 of Protein NDRG3 OS=Homo sapiens GN=NDRG3 - [NDRG3_HUMAN]	1.095	0.003
Q9Y4E5-2	Isoform 2 of Zinc finger protein 451 OS=Homo sapiens GN=ZNF451 - [ZN451_HUMAN]	1.097	0.017
Q6SJ93	Protein FAM111B OS=Homo sapiens GN=FAM111B PE=2 SV=1 - [F111B_HUMAN]	1.101	0.018
P50148	Guanine nucleotide-binding protein G(q) subunit alpha OS=Homo sapiens GN=GNAQ PE=1 SV=4 - [GNAQ_HUMAN]	1.102	0.006
Q15006	Tetratricopeptide repeat protein 35 OS=Homo sapiens GN=TTC35 PE=1 SV=1 - [TTC35_HUMAN]	1.105	0.006
Q5JTD0-2	Isoform 2 of Tight junction-associated protein 1 OS=Homo sapiens GN=TJAP1 - [TJAP1_HUMAN]	1.105	0.040
P61457	Pterin-4-alpha-carbinolamine dehydratase OS=Homo sapiens GN=PCBD1 PE=1 SV=2 - [PHS_HUMAN]	1.106	0.016
Q6IPX3	Transcription elongation factor A protein-like 6 OS=Homo sapiens GN=TCEAL6 PE=1 SV=1 - [TCAL6_HUMAN]	1.107	0.046
Q6IA86-4	Isoform 4 of Elongator complex protein 2 OS=Homo sapiens GN=ELP2 - [ELP2_HUMAN]	1.110	0.017
P12111-2	Isoform 2 of Collagen alpha-3(VI) chain OS=Homo sapiens GN=COL6A3 - [CO6A3_HUMAN]	1.111	0.009
Q9H8V3-2	Isoform 2 of Protein ECT2 OS=Homo sapiens GN=ECT2 - [ECT2_HUMAN]	1.111	0.034
P49641-1	Isoform 1 of Alpha-mannosidase 2x OS=Homo sapiens GN=MAN2A2 - [MA2A2_HUMAN]	1.111	0.049
Q32P28	Prolyl 3-hydroxylase 1 OS=Homo sapiens GN=LEPRE1 PE=1 SV=2 - [P3H1_HUMAN]	1.112	0.036
Q9NRR5	Ubiquilin-4 OS=Homo sapiens GN=UBQLN4 PE=1 SV=2 - [UBQL4_HUMAN]	1.115	0.010
Q9UN37	Vacuolar protein sorting-associated protein 4A OS=Homo sapiens GN=VPS4A PE=1 SV=1 - [VPS4A_HUMAN]	1.116	0.020
O95864-3	Isoform 3 of Fatty acid desaturase 2 OS=Homo sapiens GN=FADS2 - [FADS2_HUMAN]	1.118	0.042
P0DJ07	Uncharacterized protein C19orf79 OS=Homo sapiens GN=C19orf79 PE=4 SV=1 - [CS079_HUMAN]	1.120	0.001
Q6NUQ1	RAD50-interacting protein 1 OS=Homo sapiens GN=RINT1 PE=1 SV=1 - [RINT1_HUMAN]	1.122	0.043
P21399	Cytoplasmic aconitate hydratase OS=Homo sapiens GN=ACO1 PE=1 SV=3 - [ACOC_HUMAN]	1.123	0.008
Q8WYP3	Ras and Rab interactor 2 OS=Homo sapiens GN=RIN2 PE=1 SV=1 - [RIN2_HUMAN]	1.125	0.030
Q8TAF3-4	Isoform 4 of WD repeat-containing protein 48 OS=Homo sapiens GN=WDR48 - [WDR48_HUMAN]	1.126	0.035
P57678	Gem-associated protein 4 OS=Homo sapiens GN=GEMIN4 PE=1 SV=2 - [GEMI4_HUMAN]	1.126	0.027
Q9Y614	Ubiquitin carboxyl-terminal hydrolase 3 OS=Homo sapiens GN=USP3 PE=1 SV=2 - [UBP3_HUMAN]	1.127	0.036
Q8IX04	Ubiquitin-conjugating enzyme E2 variant 3 OS=Homo sapiens GN=UEVLD PE=1 SV=2 - [UEVLD_HUMAN]	1.127	0.027
Q9BY32	Inosine triphosphate pyrophosphatase OS=Homo sapiens GN=ITPA PE=1 SV=2 - [ITPA_HUMAN]	1.129	0.016
O95391	Pre-mRNA-splicing factor SLU7 OS=Homo sapiens GN=SLU7 PE=1 SV=2 - [SLU7_HUMAN]	1.129	0.045
Q9BYT8	Neurolysin, mitochondrial OS=Homo sapiens GN=NLN PE=1 SV=1 - [NEUL_HUMAN]	1.130	0.017
P18887	DNA repair protein XRCC1 OS=Homo sapiens GN=XRCC1 PE=1 SV=2 - [XRCC1_HUMAN]	1.130	0.032
P31930	Cytochrome b-c1 complex subunit 1, mitochondrial OS=Homo sapiens GN=UQCRC1 PE=1 SV=3 - [QCR1_HUMAN]	1.130	0.034

P28290	Sperm-specific antigen 2 OS=Homo sapiens GN=SSFA2 PE=1 SV=3 - [SSFA2_HUMAN]	1.131	0.014
Q9UPP1-3	Isoform 3 of Histone lysine demethylase PHF8 OS=Homo sapiens GN=PHF8 - [PHF8_HUMAN]	1.132	0.043
Q8WWN8	Arf-GAP with Rho-GAP domain, ANK repeat and PH domain-containing protein 3 OS=Homo sapiens GN=ARAP3 PE=1 SV=1 - [ARAP3_HUMAN]	1.132	0.003
O94855	Protein transport protein Sec24D OS=Homo sapiens GN=SEC24D PE=1 SV=2 - [SC24D_HUMAN]	1.132	0.042
O00743	Serine/threonine-protein phosphatase 6 catalytic subunit OS=Homo sapiens GN=PPP6C PE=1 SV=1 - [PPP6_HUMAN]	1.134	0.017
O60869-2	Isoform 2 of Endothelial differentiation-related factor 1 OS=Homo sapiens GN=EDF1 - [EDF1_HUMAN]	1.135	0.048
Q9NZW5	MAGUK p55 subfamily member 6 OS=Homo sapiens GN=MPP6 PE=1 SV=2 - [MPP6_HUMAN]	1.136	0.014
O95983-2	Isoform 2 of Methyl-CpG-binding domain protein 3 OS=Homo sapiens GN=MBD3 - [MBD3_HUMAN]	1.137	0.026
P21266	Glutathione S-transferase Mu 3 OS=Homo sapiens GN=GSTM3 PE=1 SV=3 - [GSTM3_HUMAN]	1.139	0.037
Q658Y4	Protein FAM91A1 OS=Homo sapiens GN=FAM91A1 PE=1 SV=3 - [F91A1_HUMAN]	1.141	0.033
Q9P270	SLAIN motif-containing protein 2 OS=Homo sapiens GN=SLAIN2 PE=1 SV=2 - [SLAI2_HUMAN]	1.147	0.002
Q8TDY2-2	Isoform 2 of RB1-inducible coiled-coil protein 1 OS=Homo sapiens GN=RB1CC1 - [RBCC1_HUMAN]	1.149	0.025
Q05048	Cleavage stimulation factor subunit 1 OS=Homo sapiens GN=CSTF1 PE=1 SV=1 - [CSTF1_HUMAN]	1.154	0.013
Q9Y697-2	Isoform Cytoplasmic of Cysteine desulfurase, mitochondrial OS=Homo sapiens GN=NFS1 - [NFS1_HUMAN]	1.155	0.026
Q96I23	Protein preY, mitochondrial OS=Homo sapiens GN=PREY PE=1 SV=1 - [PREY_HUMAN]	1.155	0.005
Q04721	Neurogenic locus notch homolog protein 2 OS=Homo sapiens GN=NOTCH2 PE=1 SV=3 - [NOTC2_HUMAN]	1.156	0.008
O75947	ATP synthase subunit d, mitochondrial OS=Homo sapiens GN=ATP5H PE=1 SV=3 - [ATP5H_HUMAN]	1.156	0.011
Q96CG8	Collagen triple helix repeat-containing protein 1 OS=Homo sapiens GN=CTHRC1 PE=1 SV=1 - [CTHR1_HUMAN]	1.159	0.041
P11177-2	Isoform 2 of Pyruvate dehydrogenase E1 component subunit beta, mitochondrial OS=Homo sapiens GN=PDHB - [ODPB_HUMAN]	1.160	0.002
Q07864	DNA polymerase epsilon catalytic subunit A OS=Homo sapiens GN=POLE PE=1 SV=5 - [DPOE1_HUMAN]	1.162	0.008
Q9H5V9	UPF0428 protein CXorf56 OS=Homo sapiens GN=CXorf56 PE=1 SV=1 - [CX056_HUMAN]	1.162	0.046
Q9UKA9	Polypyrimidine tract-binding protein 2 OS=Homo sapiens GN=PTBP2 PE=1 SV=1 - [PTBP2_HUMAN]	1.163	0.004
Q8NHG7	Small VCP/p97-interacting protein OS=Homo sapiens GN=SVIP PE=2 SV=1 - [SVIP_HUMAN]	1.163	0.041
P35268	60S ribosomal protein L22 OS=Homo sapiens GN=RPL22 PE=1 SV=2 - [RL22_HUMAN]	1.164	0.021
Q9Y2L5-2	Isoform 2 of Trafficking protein particle complex subunit 8 OS=Homo sapiens GN=TRAPPC8 - [TPPC8_HUMAN]	1.164	0.047
P24468	COUP transcription factor 2 OS=Homo sapiens GN=NR2F2 PE=1 SV=1 - [COT2_HUMAN]	1.166	0.025
P78406	mRNA export factor OS=Homo sapiens GN=RAE1 PE=1 SV=1 - [RAE1L_HUMAN]	1.172	0.008
Q99715-4	Isoform 4 of Collagen alpha-1(XII) chain OS=Homo sapiens GN=COL12A1 - [COCA1_HUMAN]	1.172	0.050
Q9Y6A4	UPF0468 protein C16orf80 OS=Homo sapiens GN=C16orf80 PE=1 SV=1 - [CP080_HUMAN]	1.172	0.035

Q9Y3B3	Transmembrane emp24 domain-containing protein 7 OS=Homo sapiens GN=TMED7 PE=1 SV=2 - [TMED7_HUMAN]	1.177	0.049
Q13261-8	Isoform 7 of Interleukin-15 receptor subunit alpha OS=Homo sapiens GN=IL15RA - [IL15RA_HUMAN]	1.180	0.014
Q9NRS6-2	Isoform 2 of Sorting nexin-15 OS=Homo sapiens GN=SNX15 - [SNX15_HUMAN]	1.182	0.039
Q4J6C6-4	Isoform 4 of Prolyl endopeptidase-like OS=Homo sapiens GN=PREPL - [PPCEL_HUMAN]	1.183	0.002
P62070	Ras-related protein R-Ras2 OS=Homo sapiens GN=RRAS2 PE=1 SV=1 - [RRAS2_HUMAN]	1.185	0.028
Q86VW0	SEC14 domain and spectrin repeat-containing protein 1 OS=Homo sapiens GN=SESTD1 PE=1 SV=2 - [SESD1_HUMAN]	1.185	0.049
P16383-2	Isoform 2 of GC-rich sequence DNA-binding factor 2 OS=Homo sapiens GN=GCFC2 - [GCFC2_HUMAN]	1.187	0.021
P25490	Transcriptional repressor protein YY1 OS=Homo sapiens GN=YY1 PE=1 SV=2 - [TTY1_HUMAN]	1.187	0.024
P62995-3	Isoform 3 of Transformer-2 protein homolog beta OS=Homo sapiens GN=TRA2B - [TRA2B_HUMAN]	1.187	0.020
P04181	Ornithine aminotransferase, mitochondrial OS=Homo sapiens GN=OAT PE=1 SV=1 - [OAT_HUMAN]	1.187	0.019
Q8IZL8	Proline-, glutamic acid- and leucine-rich protein 1 OS=Homo sapiens GN=PELP1 PE=1 SV=2 - [PELP1_HUMAN]	1.189	0.043
Q9H974	Queuine tRNA-ribosyltransferase subunit QTRTD1 OS=Homo sapiens GN=QTRTD1 PE=1 SV=1 - [QTRD1_HUMAN]	1.192	0.044
Q7L5N1	COP9 signalosome complex subunit 6 OS=Homo sapiens GN=COPS6 PE=1 SV=1 - [CSN6_HUMAN]	1.196	0.014
O60493	Sorting nexin-3 OS=Homo sapiens GN=SNX3 PE=1 SV=3 - [SNX3_HUMAN]	1.196	0.045
Q9UBG0	C-type mannose receptor 2 OS=Homo sapiens GN=MRC2 PE=1 SV=2 - [MRC2_HUMAN]	1.198	0.030
Q9NQZ5	StAR-related lipid transfer protein 7, mitochondrial OS=Homo sapiens GN=STARD7 PE=1 SV=2 - [STAR7_HUMAN]	1.198	0.024
P23469-3	Isoform 3 of Receptor-type tyrosine-protein phosphatase epsilon OS=Homo sapiens GN=PTPRE - [PTPRE_HUMAN]	1.201	0.042
P43121	Cell surface glycoprotein MUC18 OS=Homo sapiens GN=MCAM PE=1 SV=2 - [MUC18_HUMAN]	1.202	0.032
P02656	Apolipoprotein C-III OS=Homo sapiens GN=APOC3 PE=1 SV=1 - [APOC3_HUMAN]	1.204	0.047
Q9NW82	WD repeat-containing protein 70 OS=Homo sapiens GN=WDR70 PE=1 SV=1 - [WDR70_HUMAN]	1.208	0.016
Q9Y2S7	Polymerase delta-interacting protein 2 OS=Homo sapiens GN=POLDIP2 PE=1 SV=1 - [PDIP2_HUMAN]	1.214	0.042
O43861-2	Isoform 2 of Probable phospholipid-transporting ATPase IIB OS=Homo sapiens GN=ATP9B - [ATP9B_HUMAN]	1.214	0.040
Q96L92-2	Isoform 3 of Sorting nexin-27 OS=Homo sapiens GN=SNX27 - [SNX27_HUMAN]	1.216	0.003
O43708-2	Isoform 2 of Maleylacetoacetate isomerase OS=Homo sapiens GN=GSTZ1 - [MAAI_HUMAN]	1.218	0.039
Q9UL40	Zinc finger protein 346 OS=Homo sapiens GN=ZNF346 PE=1 SV=1 - [ZN346_HUMAN]	1.221	0.004
Q14353	Guanidinoacetate N-methyltransferase OS=Homo sapiens GN=GAMT PE=1 SV=1 - [GAMT_HUMAN]	1.221	0.018
Q9BQ24	Zinc finger FYVE domain-containing protein 21 OS=Homo sapiens GN=ZFYVE21 PE=1 SV=1 - [ZFY21_HUMAN]	1.225	0.020
P02452	Collagen alpha-1(I) chain OS=Homo sapiens GN=COL1A1 PE=1 SV=5 - [CO1A1_HUMAN]	1.231	0.040
Q9BYI3	Hyccin OS=Homo sapiens GN=FAM126A PE=1 SV=2 - [HYCCI_HUMAN]	1.232	0.038

Q15181	Inorganic pyrophosphatase OS=Homo sapiens GN=PPA1 PE=1 SV=2 - [IPYR_HUMAN]	1.235	0.027
Q8IXH7-4	Isoform NELF-D of Negative elongation factor C/D OS=Homo sapiens GN=TH1L - [NELFD_HUMAN]	1.240	0.046
P57768	Sorting nexin-16 OS=Homo sapiens GN=SNX16 PE=1 SV=2 - [SNX16_HUMAN]	1.241	0.047
O75419	Cell division control protein 45 homolog OS=Homo sapiens GN=CDC45 PE=1 SV=1 - [CDC45_HUMAN]	1.244	0.044
Q9HCP0-2	Isoform 1S of Casein kinase I isoform gamma-1 OS=Homo sapiens GN=CSNK1G1 - [KC1G1_HUMAN]	1.244	0.044
Q96GG9	DCN1-like protein 1 OS=Homo sapiens GN=DCUN1D1 PE=1 SV=1 - [DCNL1_HUMAN]	1.247	0.021
O00522-2	Isoform 2 of Krev interaction trapped protein 1 OS=Homo sapiens GN=KRIT1 - [KRIT1_HUMAN]	1.255	0.040
O75911	Short-chain dehydrogenase/reductase 3 OS=Homo sapiens GN=DHRS3 PE=1 SV=2 - [DHRS3_HUMAN]	1.255	0.015
Q9UDX5	Mitochondrial fission process protein 1 OS=Homo sapiens GN=MTFP1 PE=1 SV=1 - [MTFP1_HUMAN]	1.262	0.016
P49757-4	Isoform 4 of Protein numb homolog OS=Homo sapiens GN=NUMB - [NUMB_HUMAN]	1.264	0.013
Q8TBM8	DnaJ homolog subfamily B member 14 OS=Homo sapiens GN=DNAJB14 PE=2 SV=1 - [DJB14_HUMAN]	1.265	0.025
Q9HC52	Chromobox protein homolog 8 OS=Homo sapiens GN=CBX8 PE=1 SV=3 - [CBX8_HUMAN]	1.268	0.033
Q8IVL1-4	Isoform 4 of Neuron navigator 2 OS=Homo sapiens GN=NAV2 - [NAV2_HUMAN]	1.270	0.022
Q8NB15-2	Isoform 2 of Zinc finger protein 511 OS=Homo sapiens GN=ZNF511 - [ZN511_HUMAN]	1.275	0.017
Q92879-2	Isoform 2 of CUGBP Elav-like family member 1 OS=Homo sapiens GN=CELF1 - [CELF1_HUMAN]	1.277	0.036
P49716	CCAAT/enhancer-binding protein delta OS=Homo sapiens GN=CEBPD PE=1 SV=2 - [CEBPD_HUMAN]	1.288	0.038
P62158	Calmodulin OS=Homo sapiens GN=CALM1 PE=1 SV=2 - [CALM_HUMAN]	1.295	0.029
P35520	Cystathionine beta-synthase OS=Homo sapiens GN=CBS PE=1 SV=2 - [CBS_HUMAN]	1.296	0.041
Q9Y6A1-4	Isoform 4 of Protein O-mannosyl-transferase 1 OS=Homo sapiens GN=POMT1 - [POMT1_HUMAN]	1.309	0.039
Q9UIL1-4	Isoform 4 of Short coiled-coil protein OS=Homo sapiens GN=SCOC - [SCOC_HUMAN]	1.334	0.020
P98172	Ephrin-B1 OS=Homo sapiens GN=EFNB1 PE=1 SV=1 - [EFNB1_HUMAN]	1.337	0.047
Q86U90	YrdC domain-containing protein, mitochondrial OS=Homo sapiens GN=YRDC PE=1 SV=1 - [YRDC_HUMAN]	1.346	0.044
O15392-7	Isoform 7 of Baculoviral IAP repeat-containing protein 5 OS=Homo sapiens GN=BIRC5 - [BIRC5_HUMAN]	1.352	0.012
Q9NR33	DNA polymerase epsilon subunit 4 OS=Homo sapiens GN=POLE4 PE=1 SV=2 - [DPOE4_HUMAN]	1.365	0.044
Q9H8G2	Uncharacterized protein C9orf82 OS=Homo sapiens GN=C9orf82 PE=1 SV=2 - [CI082_HUMAN]	1.371	0.011
P14316	Interferon regulatory factor 2 OS=Homo sapiens GN=IRF2 PE=1 SV=2 - [IRF2_HUMAN]	1.379	0.036
Q92529-2	Isoform p52 of SHC-transforming protein 3 OS=Homo sapiens GN=SHC3 - [SHC3_HUMAN]	1.390	0.031
Q5FWF5-2	Isoform 2 of N-acetyltransferase ESCO1 OS=Homo sapiens GN=ESCO1 - [ESCO1_HUMAN]	1.399	0.044
O00635	Tripartite motif-containing protein 38 OS=Homo sapiens GN=TRIM38 PE=2 SV=1 - [TRI38_HUMAN]	1.434	0.045

Q8NFH3	Nucleoporin Nup43 OS=Homo sapiens GN=NUP43 PE=1 SV=1 - [NUP43_HUMAN]	1.468	0.035
P11166	Solute carrier family 2, facilitated glucose transporter member 1 OS=Homo sapiens GN=SLC2A1 PE=1 SV=2 - [GTR1_HUMAN]	1.495	0.003
Q99943	1-acyl-sn-glycerol-3-phosphate acyltransferase alpha OS=Homo sapiens GN=AGPAT1 PE=2 SV=2 - [PLCA_HUMAN]	1.501	0.049
O14735	CDP-diacylglycerol--inositol 3-phosphatidyltransferase OS=Homo sapiens GN=CDIPT PE=1 SV=1 - [CDIPT_HUMAN]	1.516	0.034
O15235	28S ribosomal protein S12, mitochondrial OS=Homo sapiens GN=MRPS12 PE=1 SV=1 - [RT12_HUMAN]	1.584	0.027
P06241-3	Isoform 3 of Tyrosine-protein kinase Fyn OS=Homo sapiens GN=FYN - [FYN_HUMAN]	1.663	0.048
P42766	60S ribosomal protein L35 OS=Homo sapiens GN=RPL35 PE=1 SV=2 - [RL35_HUMAN]	1.689	0.019
Q9BZM5	NKG2D ligand 2 OS=Homo sapiens GN=ULBP2 PE=1 SV=1 - [N2DL2_HUMAN]	1.705	0.003
Q9Y3E7-4	Isoform 4 of Charged multivesicular body protein 3 OS=Homo sapiens GN=CHMP3 - [CHMP3_HUMAN]	1.760	0.043
Q15819	Ubiquitin-conjugating enzyme E2 variant 2 OS=Homo sapiens GN=UBE2V2 PE=1 SV=4 - [UB2V2_HUMAN]	1.911	0.046
O15145	Actin-related protein 2/3 complex subunit 3 OS=Homo sapiens GN=ARPC3 PE=1 SV=3 - [ARPC3_HUMAN]	2.512	0.045

H727 cells - 2 hours lanreotide (397 proteins)

P84243	Histone H3.3 OS=Homo sapiens GN=H3F3A PE=1 SV=2 - [H33_HUMAN]	0.523	0.013
Q6DD87	Zinc finger protein 787 OS=Homo sapiens GN=ZNF787 PE=1 SV=2 - [ZN787_HUMAN]	0.641	0.038
P53671-2	Isoform LIMK2b of LIM domain kinase 2 OS=Homo sapiens GN=LIMK2 - [LIMK2_HUMAN]	0.719	0.042
Q9UJH8	Meteorin OS=Homo sapiens GN=METR_N PE=2 SV=2 - [METRN_HUMAN]	0.758	0.030
Q8IZQ1-2	Isoform 2 of WD repeat and FYVE domain-containing protein 3 OS=Homo sapiens GN=WDFY3 - [WDFY3_HUMAN]	0.783	0.048
Q9UPQ8	Dolichol kinase OS=Homo sapiens GN=DOLK PE=1 SV=1 - [DOLK_HUMAN]	0.788	0.042
O94766	Galactosylgalactosylxylosylprotein 3-beta-glucuronosyltransferase 3 OS=Homo sapiens GN=B3GAT3 PE=1 SV=2 - [B3GA3_HUMAN]	0.798	0.035
P05981	Serine protease hepsin OS=Homo sapiens GN=HPN PE=1 SV=1 - [HEPS_HUMAN]	0.798	0.001
Q9BVS4	Serine/threonine-protein kinase RIO2 OS=Homo sapiens GN=RIOK2 PE=1 SV=2 - [RIOK2_HUMAN]	0.811	0.016
Q2TAA2	Isoamyl acetate-hydrolyzing esterase 1 homolog OS=Homo sapiens GN=IAH1 PE=1 SV=1 - [IAH1_HUMAN]	0.822	0.004
O00220	Tumor necrosis factor receptor superfamily member 10A OS=Homo sapiens GN=TNFRSF10A PE=1 SV=3 - [TR10A_HUMAN]	0.825	0.015
Q96DZ1	Endoplasmic reticulum lectin 1 OS=Homo sapiens GN=ERLEC1 PE=1 SV=1 - [ERLEC_HUMAN]	0.827	0.042
Q8N183	Mimitin, mitochondrial OS=Homo sapiens GN=NDUF2 PE=1 SV=1 - [MIMIT_HUMAN]	0.828	0.030
O75356	Ectonucleoside triphosphate diphosphohydrolase 5 OS=Homo sapiens GN=ENTPD5 PE=1 SV=1 - [ENTP5_HUMAN]	0.828	0.049
Q96AB3	Isochorismatase domain-containing protein 2, mitochondrial OS=Homo sapiens GN=ISOC2 PE=1 SV=1 - [ISOC2_HUMAN]	0.831	0.014
Q8NCR9	Clarín-3 OS=Homo sapiens GN=CLRN3 PE=2 SV=1 - [CLRN3_HUMAN]	0.851	0.034

O75838	Calcium and integrin-binding family member 2 OS=Homo sapiens GN=CIB2 PE=2 SV=1 - [CIB2_HUMAN]	0.852	0.042
P62854	40S ribosomal protein S26 OS=Homo sapiens GN=RPS26 PE=1 SV=3 - [RS26_HUMAN]	0.856	0.003
Q8WW12-2	Isoform 2 of PEST proteolytic signal-containing nuclear protein OS=Homo sapiens GN=PCNP - [PCNP_HUMAN]	0.857	0.043
Q7Z5U6	WD repeat-containing protein 53 OS=Homo sapiens GN=WDR53 PE=2 SV=1 - [WDR53_HUMAN]	0.858	0.029
P0CG38	POTE ankyrin domain family member I OS=Homo sapiens GN=POTEI PE=3 SV=1 - [POTEI_HUMAN]	0.858	0.047
O60858	E3 ubiquitin-protein ligase TRIM13 OS=Homo sapiens GN=TRIM13 PE=1 SV=2 - [TRIM13_HUMAN]	0.859	0.020
P60484	Phosphatidylinositol-3,4,5-trisphosphate 3-phosphatase and dual-specificity protein phosphatase PTEN OS=Homo sapiens GN=PTEN PE=1 SV=1 - [PTEN_HUMAN]	0.863	0.034
Q9BSA9	Transmembrane protein 175 OS=Homo sapiens GN=TMEM175 PE=1 SV=1 - [TM175_HUMAN]	0.865	0.029
Q5TC12	ATP synthase mitochondrial F1 complex assembly factor 1 OS=Homo sapiens GN=ATPAF1 PE=1 SV=1 - [ATPF1_HUMAN]	0.865	0.016
O43581	Synaptotagmin-7 OS=Homo sapiens GN=SYT7 PE=1 SV=3 - [SYT7_HUMAN]	0.867	0.041
P45880-2	Isoform 2 of Voltage-dependent anion-selective channel protein 2 OS=Homo sapiens GN=VDAC2 - [VDAC2_HUMAN]	0.869	0.034
Q9BT43	DNA-directed RNA polymerase III subunit RPC7-like OS=Homo sapiens GN=POLR3GL PE=2 SV=1 - [RPC7L_HUMAN]	0.873	0.041
Q9HBL7	Transmembrane protein C9orf46 OS=Homo sapiens GN=C9orf46 PE=2 SV=1 - [CI046_HUMAN]	0.873	0.038
Q5VWZ2-2	Isoform 2 of Lysophospholipase-like protein 1 OS=Homo sapiens GN=LYPLAL1 - [LYPL1_HUMAN]	0.874	0.001
O95400	CD2 antigen cytoplasmic tail-binding protein 2 OS=Homo sapiens GN=CD2BP2 PE=1 SV=1 - [CD2B2_HUMAN]	0.874	0.001
Q86WQ0	Nuclear receptor 2C2-associated protein OS=Homo sapiens GN=NR2C2AP PE=1 SV=1 - [NR2CA_HUMAN]	0.878	0.023
P05204	Non-histone chromosomal protein HMG-17 OS=Homo sapiens GN=HMGN2 PE=1 SV=3 - [HMGN2_HUMAN]	0.879	0.019
Q96SI9	Spermatid perinuclear RNA-binding protein OS=Homo sapiens GN=STRBP PE=1 SV=1 - [STRBP_HUMAN]	0.879	0.047
Q9BXT8-4	Isoform 4 of RING finger protein 17 OS=Homo sapiens GN=RNF17 - [RNF17_HUMAN]	0.881	0.018
P15121	Aldose reductase OS=Homo sapiens GN=AKR1B1 PE=1 SV=3 - [ALDR_HUMAN]	0.885	0.050
Q14508-2	Isoform 2 of WAP four-disulfide core domain protein 2 OS=Homo sapiens GN=WFDC2 - [WFDC2_HUMAN]	0.886	0.021
Q9BS40	Latexin OS=Homo sapiens GN=LXN PE=1 SV=2 - [LXN_HUMAN]	0.886	0.014
O60613	15 kDa selenoprotein OS=Homo sapiens GN=SEP15 PE=1 SV=3 - [SEP15_HUMAN]	0.886	0.036
Q9NQ88	Probable fructose-2,6-bisphosphatase TIGAR OS=Homo sapiens GN=TIGAR PE=1 SV=1 - [TIGAR_HUMAN]	0.888	0.018
Q99622	Protein C10 OS=Homo sapiens GN=C12orf57 PE=1 SV=1 - [C10_HUMAN]	0.889	0.001
P30405	Peptidyl-prolyl cis-trans isomerase F, mitochondrial OS=Homo sapiens GN=PPIF PE=1 SV=1 - [PPIF_HUMAN]	0.891	0.038
Q969H8	UPF0556 protein C19orf10 OS=Homo sapiens GN=C19orf10 PE=1 SV=1 - [CS010_HUMAN]	0.892	0.032
Q9BY50	Signal peptidase complex catalytic subunit SEC11C OS=Homo sapiens GN=SEC11C PE=1 SV=3 - [SC11C_HUMAN]	0.892	0.012

Q12797	Aspartyl/asparaginyl beta-hydroxylase OS=Homo sapiens GN=ASPH PE=1 SV=3 - [ASPH_HUMAN]	0.892	0.045
P11387	DNA topoisomerase 1 OS=Homo sapiens GN=TOP1 PE=1 SV=2 - [TOP1_HUMAN]	0.893	0.043
Q8N4P3	Guanosine-3',5'-bis(diphosphate) 3'-pyrophosphohydrolase MESH1 OS=Homo sapiens GN=HDDC3 PE=1 SV=3 - [MESH1_HUMAN]	0.895	0.002
Q6NUQ1	RAD50-interacting protein 1 OS=Homo sapiens GN=RINT1 PE=1 SV=1 - [RINT1_HUMAN]	0.896	0.035
Q15717	ELAV-like protein 1 OS=Homo sapiens GN=ELAVL1 PE=1 SV=2 - [ELAV1_HUMAN]	0.896	0.040
Q05682-5	Isoform 5 of Caldesmon OS=Homo sapiens GN=CALD1 - [CALD1_HUMAN]	0.896	0.047
Q9NWM0-2	Isoform 2 of Spermine oxidase OS=Homo sapiens GN=SMOX - [SMOX_HUMAN]	0.896	0.048
Q99547	M-phase phosphoprotein 6 OS=Homo sapiens GN=MPHOSPH6 PE=1 SV=2 - [MPH6_HUMAN]	0.897	0.019
O00217	NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial OS=Homo sapiens GN=NDUFS8 PE=1 SV=1 - [NDUS8_HUMAN]	0.898	0.047
P82664	28S ribosomal protein S10, mitochondrial OS=Homo sapiens GN=MRPS10 PE=1 SV=2 - [RT10_HUMAN]	0.898	0.026
Q9NZD2	Glycolipid transfer protein OS=Homo sapiens GN=GLTP PE=1 SV=3 - [GLTP_HUMAN]	0.898	0.029
Q9BVK6	Transmembrane emp24 domain-containing protein 9 OS=Homo sapiens GN=TMED9 PE=1 SV=2 - [TMED9_HUMAN]	0.900	0.025
P00505	Aspartate aminotransferase, mitochondrial OS=Homo sapiens GN=GOT2 PE=1 SV=3 - [AATM_HUMAN]	0.900	0.000
P82914	28S ribosomal protein S15, mitochondrial OS=Homo sapiens GN=MRPS15 PE=1 SV=1 - [RT15_HUMAN]	0.900	0.024
P62820	Ras-related protein Rab-1A OS=Homo sapiens GN=RAB1A PE=1 SV=3 - [RAB1A_HUMAN]	0.902	0.013
Q9NUE0	Palmitoyltransferase ZDHHC18 OS=Homo sapiens GN=ZDHHC18 PE=2 SV=2 - [ZDH18_HUMAN]	0.903	0.013
O94874-2	Isoform 2 of E3 UFM1-protein ligase 1 OS=Homo sapiens GN=UFL1 - [UFL1_HUMAN]	0.904	0.032
O95363	Phenylalanine--tRNA ligase, mitochondrial OS=Homo sapiens GN=FARS2 PE=1 SV=1 - [SYFM_HUMAN]	0.906	0.027
Q8N1G0	Zinc finger protein 687 OS=Homo sapiens GN=ZNF687 PE=1 SV=1 - [ZN687_HUMAN]	0.906	0.001
Q04837	Single-stranded DNA-binding protein, mitochondrial OS=Homo sapiens GN=SSBP1 PE=1 SV=1 - [SSBP_HUMAN]	0.907	0.016
Q13405	39S ribosomal protein L49, mitochondrial OS=Homo sapiens GN=MRPL49 PE=1 SV=1 - [RM49_HUMAN]	0.907	0.028
Q15390	Mitochondrial fission regulator 1 OS=Homo sapiens GN=MTFR1 PE=1 SV=2 - [MTFR1_HUMAN]	0.908	0.020
P42765	3-ketoacyl-CoA thiolase, mitochondrial OS=Homo sapiens GN=ACAA2 PE=1 SV=2 - [THIM_HUMAN]	0.908	0.039
P07711	Cathepsin L1 OS=Homo sapiens GN=CTSL1 PE=1 SV=2 - [CATL1_HUMAN]	0.910	0.034
P10515	Dihydrolipoyllysine-residue acetyltransferase component of pyruvate dehydrogenase complex, mitochondrial OS=Homo sapiens GN=DLAT PE=1 SV=3 - [ODP2_HUMAN]	0.912	0.045
O75489	NADH dehydrogenase [ubiquinone] iron-sulfur protein 3, mitochondrial OS=Homo sapiens GN=NDUFS3 PE=1 SV=1 - [NDUS3_HUMAN]	0.913	0.034
Q13740-2	Isoform 2 of CD166 antigen OS=Homo sapiens GN=ALCAM - [CD166_HUMAN]	0.914	0.042
P49458	Signal recognition particle 9 kDa protein OS=Homo sapiens GN=SRP9 PE=1 SV=2 - [SRP09_HUMAN]	0.914	0.044

Q9UNA1-2	Isoform 2 of Rho GTPase-activating protein 26 OS=Homo sapiens GN=ARHGAP26 - [RHG26_HUMAN]	0.914	0.022
P43007	Neutral amino acid transporter A OS=Homo sapiens GN=SLC1A4 PE=1 SV=1 - [SATT_HUMAN]	0.915	0.039
Q99459	Cell division cycle 5-like protein OS=Homo sapiens GN=CDC5L PE=1 SV=2 - [CDC5L_HUMAN]	0.917	0.027
P49802-2	Isoform 2 of Regulator of G-protein signaling 7 OS=Homo sapiens GN=RGS7 - [RGS7_HUMAN]	0.917	0.036
Q6NZY4	Zinc finger CCHC domain-containing protein 8 OS=Homo sapiens GN=ZCCHC8 PE=1 SV=2 - [ZCHC8_HUMAN]	0.917	0.033
Q96G03	Phosphoglucomutase-2 OS=Homo sapiens GN=PGM2 PE=1 SV=4 - [PGM2_HUMAN]	0.917	0.047
P37287	Phosphatidylinositol N-acetylglucosaminyltransferase subunit A OS=Homo sapiens GN=PIGA PE=1 SV=1 - [PIGA_HUMAN]	0.918	0.019
Q9H9H4	Vacuolar protein sorting-associated protein 37B OS=Homo sapiens GN=VPS37B PE=1 SV=1 - [VP37B_HUMAN]	0.919	0.034
O95372	Acyl-protein thioesterase 2 OS=Homo sapiens GN=LYPLA2 PE=1 SV=1 - [LYPA2_HUMAN]	0.920	0.032
Q9NV06	DDB1- and CUL4-associated factor 13 OS=Homo sapiens GN=DCAF13 PE=1 SV=2 - [DCA13_HUMAN]	0.920	0.009
Q92759	General transcription factor IIH subunit 4 OS=Homo sapiens GN=GTF2H4 PE=2 SV=1 - [TF2H4_HUMAN]	0.920	0.005
Q8NBX0	Saccharopine dehydrogenase-like oxidoreductase OS=Homo sapiens GN=SCCPDH PE=1 SV=1 - [SCPDL_HUMAN]	0.921	0.029
Q02880-2	Isoform Beta-1 of DNA topoisomerase 2-beta OS=Homo sapiens GN=TOP2B - [TOP2B_HUMAN]	0.922	0.009
Q9C040	Tripartite motif-containing protein 2 OS=Homo sapiens GN=TRIM2 PE=1 SV=1 - [TRIM2_HUMAN]	0.922	0.012
P27797	Calreticulin OS=Homo sapiens GN=CALR PE=1 SV=1 - [CALR_HUMAN]	0.922	0.006
Q9NQE9	Histidine triad nucleotide-binding protein 3 OS=Homo sapiens GN=HINT3 PE=1 SV=1 - [HINT3_HUMAN]	0.923	0.013
P11177-2	Isoform 2 of Pyruvate dehydrogenase E1 component subunit beta, mitochondrial OS=Homo sapiens GN=PDHB - [ODPB_HUMAN]	0.923	0.015
P13010	X-ray repair cross-complementing protein 5 OS=Homo sapiens GN=XRCC5 PE=1 SV=3 - [XRCC5_HUMAN]	0.925	0.026
P37108	Signal recognition particle 14 kDa protein OS=Homo sapiens GN=SRP14 PE=1 SV=2 - [SRP14_HUMAN]	0.928	0.019
Q99623	Prohibitin-2 OS=Homo sapiens GN=PHB2 PE=1 SV=2 - [PHB2_HUMAN]	0.929	0.042
O00559	Receptor-binding cancer antigen expressed on SiSo cells OS=Homo sapiens GN=EBAG9 PE=1 SV=1 - [RCAS1_HUMAN]	0.930	0.038
O15014	Zinc finger protein 609 OS=Homo sapiens GN=ZNF609 PE=1 SV=2 - [ZN609_HUMAN]	0.931	0.048
O75506	Heat shock factor-binding protein 1 OS=Homo sapiens GN=HSBP1 PE=1 SV=1 - [HSBP1_HUMAN]	0.931	0.042
Q92484	Acid sphingomyelinase-like phosphodiesterase 3a OS=Homo sapiens GN=SMPDL3A PE=1 SV=2 - [ASM3A_HUMAN]	0.932	0.025
P83916	Chromobox protein homolog 1 OS=Homo sapiens GN=CBX1 PE=1 SV=1 - [CBX1_HUMAN]	0.933	0.020
P05455	Lupus La protein OS=Homo sapiens GN=SSB PE=1 SV=2 - [LA_HUMAN]	0.933	0.010
Q92552	28S ribosomal protein S27, mitochondrial OS=Homo sapiens GN=MRPS27 PE=1 SV=3 - [RT27_HUMAN]	0.934	0.042
Q00839-2	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU - [HNRPU_HUMAN]	0.935	0.015
P21127-8	Isoform 8 of Cyclin-dependent kinase 11B OS=Homo sapiens GN=CDK11B - [CD11B_HUMAN]	0.936	0.018

Q9Y3A3-2	Isoform 2 of MOB-like protein phocein OS=Homo sapiens GN=MOB4 - [PHOCN_HUMAN]	0.936	0.022
Q9H0W9-2	Isoform 2 of Ester hydrolase C11orf54 OS=Homo sapiens GN=C11orf54 - [CK054_HUMAN]	0.936	0.031
Q9UNM6	26S proteasome non-ATPase regulatory subunit 13 OS=Homo sapiens GN=PSMD13 PE=1 SV=2 - [PSD13_HUMAN]	0.936	0.001
P12532	Creatine kinase U-type, mitochondrial OS=Homo sapiens GN=CKMT1A PE=1 SV=1 - [KCRU_HUMAN]	0.937	0.023
P10636-4	Isoform Tau-B of Microtubule-associated protein tau OS=Homo sapiens GN=MAPT - [TAU_HUMAN]	0.937	0.030
Q12904	Aminoacyl tRNA synthase complex-interacting multifunctional protein 1 OS=Homo sapiens GN=AIMP1 PE=1 SV=2 - [AIMP1_HUMAN]	0.938	0.033
Q75QN2-2	Isoform 2 of Integrator complex subunit 8 OS=Homo sapiens GN=INTS8 - [INT8_HUMAN]	0.939	0.004
P39656	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit OS=Homo sapiens GN=DDOST PE=1 SV=4 - [OST48_HUMAN]	0.940	0.041
Q32P41	tRNA (guanine(37)-N1)-methyltransferase OS=Homo sapiens GN=TRMT5 PE=1 SV=2 - [TRM5_HUMAN]	0.940	0.023
Q9H490-2	Isoform 2 of Phosphatidylinositol glycan anchor biosynthesis class U protein OS=Homo sapiens GN=PIGU - [PIGU_HUMAN]	0.945	0.024
P26368-2	Isoform 2 of Splicing factor U2AF 65 kDa subunit OS=Homo sapiens GN=U2AF2 - [U2AF2_HUMAN]	0.945	0.010
Q8NA19-2	Isoform 2 of Lethal(3)malignant brain tumor-like protein 4 OS=Homo sapiens GN=L3MBTL4 - [LMBL4_HUMAN]	0.946	0.024
O15511	Actin-related protein 2/3 complex subunit 5 OS=Homo sapiens GN=ARPC5 PE=1 SV=3 - [ARPC5_HUMAN]	0.947	0.032
Q9H0U4	Ras-related protein Rab-1B OS=Homo sapiens GN=RAB1B PE=1 SV=1 - [RAB1B_HUMAN]	0.947	0.032
Q13561	Dynactin subunit 2 OS=Homo sapiens GN=DCTN2 PE=1 SV=4 - [DCTN2_HUMAN]	0.947	0.048
Q96AQ8	Coiled-coil domain-containing protein 90A, mitochondrial OS=Homo sapiens GN=CCDC90A PE=2 SV=1 - [CC90A_HUMAN]	0.948	0.002
Q969Q5	Ras-related protein Rab-24 OS=Homo sapiens GN=RAB24 PE=1 SV=1 - [RAB24_HUMAN]	0.949	0.040
P41732	Tetraspanin-7 OS=Homo sapiens GN=TSPAN7 PE=1 SV=2 - [TSN7_HUMAN]	0.950	0.036
Q06481	Amyloid-like protein 2 OS=Homo sapiens GN=APLP2 PE=1 SV=2 - [APLP2_HUMAN]	0.951	0.030
P05783	Keratin, type I cytoskeletal 18 OS=Homo sapiens GN=KRT18 PE=1 SV=2 - [K1C18_HUMAN]	0.951	0.038
Q99797	Mitochondrial intermediate peptidase OS=Homo sapiens GN=MIPEP PE=1 SV=2 - [MIPEP_HUMAN]	0.951	0.002
P43304	Glycerol-3-phosphate dehydrogenase, mitochondrial OS=Homo sapiens GN=GPD2 PE=1 SV=3 - [GPDM_HUMAN]	0.954	0.008
O14524	Transmembrane protein 194A OS=Homo sapiens GN=TMEM194A PE=1 SV=2 - [T194A_HUMAN]	0.954	0.046
P35232	Prohibitin OS=Homo sapiens GN=PHB PE=1 SV=1 - [PHB_HUMAN]	0.955	0.037
O00165-5	Isoform 5 of HCLS1-associated protein X-1 OS=Homo sapiens GN=HAX1 - [HAX1_HUMAN]	0.955	0.042
Q96C90	Protein phosphatase 1 regulatory subunit 14B OS=Homo sapiens GN=PPP1R14B PE=1 SV=3 - [PP14B_HUMAN]	0.956	0.048
Q9BVS5	Potential tRNA (adenine(58)-N(1))-methyltransferase catalytic subunit TRMT61B OS=Homo sapiens GN=TRMT61B PE=1 SV=2 - [TR61B_HUMAN]	0.958	0.036
Q6PL24	Protein TMED8 OS=Homo sapiens GN=TMED8 PE=1 SV=1 - [TMED8_HUMAN]	0.959	0.006
P41091	Eukaryotic translation initiation factor 2 subunit 3 OS=Homo sapiens GN=EIF2S3 PE=1 SV=3 - [IF2G_HUMAN]	0.960	0.047

Q8WXF1-2	Isoform 2 of Paraspeckle component 1 OS=Homo sapiens GN=PSPC1 - [PSPC1_HUMAN]	0.962	0.009
Q5JRX3	Presequence protease, mitochondrial OS=Homo sapiens GN=PITRM1 PE=1 SV=2 - [PREP_HUMAN]	0.962	0.035
P10809	60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2 - [CH60_HUMAN]	0.965	0.024
P09104	Gamma-enolase OS=Homo sapiens GN=ENO2 PE=1 SV=3 - [ENOG_HUMAN]	0.966	0.020
P42858	Huntingtin OS=Homo sapiens GN=HTT PE=1 SV=2 - [HD_HUMAN]	0.966	0.023
Q12962	Transcription initiation factor TFIID subunit 10 OS=Homo sapiens GN=TAF10 PE=1 SV=1 - [TAF10_HUMAN]	0.967	0.000
Q9NZB2-4	Isoform D of Constitutive coactivator of PPAR-gamma-like protein 1 OS=Homo sapiens GN=FAM120A - [F120A_HUMAN]	0.968	0.016
Q9BXP5-4	Isoform 4 of Serrate RNA effector molecule homolog OS=Homo sapiens GN=SRRT - [SRRT_HUMAN]	0.968	0.034
P61421	V-type proton ATPase subunit d 1 OS=Homo sapiens GN=ATP6V0D1 PE=1 SV=1 - [VA0D1_HUMAN]	0.969	0.039
P07910-2	Isoform C1 of Heterogeneous nuclear ribonucleoproteins C1/C2 OS=Homo sapiens GN=HNRNPC - [HNRPC_HUMAN]	0.970	0.043
Q96H20-2	Isoform 2 of Vacuolar-sorting protein SNF8 OS=Homo sapiens GN=SNF8 - [SNF8_HUMAN]	0.972	0.011
P07437	Tubulin beta chain OS=Homo sapiens GN=TUBB PE=1 SV=2 - [TBB5_HUMAN]	0.973	0.040
Q08211	ATP-dependent RNA helicase A OS=Homo sapiens GN=DHX9 PE=1 SV=4 - [DHX9_HUMAN]	0.977	0.047
Q8N392-2	Isoform 2 of Rho GTPase-activating protein 18 OS=Homo sapiens GN=ARHGAP18 - [RHG18_HUMAN]	0.981	0.018
Q8WUM4	Programmed cell death 6-interacting protein OS=Homo sapiens GN=PDCD6IP PE=1 SV=1 - [PDC6I_HUMAN]	0.984	0.022
P06733	Alpha-enolase OS=Homo sapiens GN=ENO1 PE=1 SV=2 - [ENOA_HUMAN]	0.986	0.044
Q96S19-3	Isoform 3 of UPF0585 protein C16orf13 OS=Homo sapiens GN=C16orf13 - [CP013_HUMAN]	1.011	0.005
P42345	Serine/threonine-protein kinase mTOR OS=Homo sapiens GN=MTOR PE=1 SV=1 - [MTOR_HUMAN]	1.021	0.034
Q9NVU7	Protein SDA1 homolog OS=Homo sapiens GN=SDAD1 PE=1 SV=3 - [SDA1_HUMAN]	1.022	0.029
P52735-3	Isoform 3 of Guanine nucleotide exchange factor VAV2 OS=Homo sapiens GN=VAV2 - [VAV2_HUMAN]	1.024	0.009
Q15796-2	Isoform Short of Mothers against decapentaplegic homolog 2 OS=Homo sapiens GN=SMAD2 - [SMAD2_HUMAN]	1.026	0.026
O60292	Signal-induced proliferation-associated 1-like protein 3 OS=Homo sapiens GN=SIPA1L3 PE=1 SV=3 - [SI1L3_HUMAN]	1.028	0.002
Q9H0N5	Pterin-4-alpha-carbinolamine dehydratase 2 OS=Homo sapiens GN=PCBD2 PE=1 SV=4 - [PHS2_HUMAN]	1.032	0.026
Q6PI98	INO80 complex subunit C OS=Homo sapiens GN=INO80C PE=1 SV=1 - [IN80C_HUMAN]	1.042	0.037
Q99576-4	Isoform 3 of TSC22 domain family protein 3 OS=Homo sapiens GN=TSC22D3 - [T22D3_HUMAN]	1.042	0.032
O15379	Histone deacetylase 3 OS=Homo sapiens GN=HDAC3 PE=1 SV=2 - [HDAC3_HUMAN]	1.042	0.026
Q8IY18	Structural maintenance of chromosomes protein 5 OS=Homo sapiens GN=SMC5 PE=1 SV=2 - [SMC5_HUMAN]	1.043	0.011
Q96A65	Exocyst complex component 4 OS=Homo sapiens GN=EXOC4 PE=1 SV=1 - [EXOC4_HUMAN]	1.043	0.043
Q9BQ04	RNA-binding protein 4B OS=Homo sapiens GN=RBM4B PE=1 SV=1 - [RBM4B_HUMAN]	1.044	0.017

Q96T51	RUN and FYVE domain-containing protein 1 OS=Homo sapiens GN=RUFY1 PE=1 SV=2 - [RUFY1_HUMAN]	1.046	0.022
O14976	Cyclin-G-associated kinase OS=Homo sapiens GN=GAK PE=1 SV=2 - [GAK_HUMAN]	1.050	0.049
Q9H2P0	Activity-dependent neuroprotector homeobox protein OS=Homo sapiens GN=ADNP PE=1 SV=1 - [ADNP_HUMAN]	1.050	0.024
Q99470	Stromal cell-derived factor 2 OS=Homo sapiens GN=SDF2 PE=1 SV=2 - [SDF2_HUMAN]	1.053	0.044
O00291	Huntingtin-interacting protein 1 OS=Homo sapiens GN=HIP1 PE=1 SV=5 - [HIP1_HUMAN]	1.054	0.019
O96017-12	Isoform 12 of Serine/threonine-protein kinase Chk2 OS=Homo sapiens GN=CHK2 - [CHK2_HUMAN]	1.055	0.033
Q9NTK5	Obg-like ATPase 1 OS=Homo sapiens GN=OLA1 PE=1 SV=2 - [OLA1_HUMAN]	1.058	0.017
Q9NWT6	Hypoxia-inducible factor 1-alpha inhibitor OS=Homo sapiens GN=HIF1AN PE=1 SV=2 - [HIF1N_HUMAN]	1.060	0.005
Q96S55-2	Isoform 2 of ATPase WRNIP1 OS=Homo sapiens GN=WRNIP1 - [WRIP1_HUMAN]	1.061	0.036
Q53RY4-2	Isoform 2 of Keratinocyte-associated protein 3 OS=Homo sapiens GN=KRTCAP3 - [KCP3_HUMAN]	1.062	0.050
Q00537	Cyclin-dependent kinase 17 OS=Homo sapiens GN=CDK17 PE=1 SV=2 - [CDK17_HUMAN]	1.063	0.024
Q7Z7G8-2	Isoform 2 of Vacuolar protein sorting-associated protein 13B OS=Homo sapiens GN=VPS13B - [VP13B_HUMAN]	1.064	0.045
P61962	DDB1- and CUL4-associated factor 7 OS=Homo sapiens GN=DCAF7 PE=1 SV=1 - [DCAF7_HUMAN]	1.064	0.037
Q99543	DnaJ homolog subfamily C member 2 OS=Homo sapiens GN=DNAJC2 PE=1 SV=4 - [DNJC2_HUMAN]	1.064	0.043
P34896-3	Isoform 3 of Serine hydroxymethyltransferase, cytosolic OS=Homo sapiens GN=SHMT1 - [GLYC_HUMAN]	1.066	0.017
P35250	Replication factor C subunit 2 OS=Homo sapiens GN=RFC2 PE=1 SV=3 - [RFC2_HUMAN]	1.067	0.006
Q15911	Zinc finger homeobox protein 3 OS=Homo sapiens GN=ZFH3 PE=1 SV=2 - [ZFH3_HUMAN]	1.069	0.028
Q7L4I2	Arginine/serine-rich coiled-coil protein 2 OS=Homo sapiens GN=RSRC2 PE=1 SV=1 - [RSRC2_HUMAN]	1.070	0.020
Q15365	Poly(rC)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2 - [PCBP1_HUMAN]	1.072	0.010
Q8NFD5	AT-rich interactive domain-containing protein 1B OS=Homo sapiens GN=ARID1B PE=1 SV=2 - [ARI1B_HUMAN]	1.072	0.039
Q16222-2	Isoform AGX1 of UDP-N-acetylhexosamine pyrophosphorylase OS=Homo sapiens GN=UAP1 - [UAP1_HUMAN]	1.074	0.015
Q92541	RNA polymerase-associated protein RTF1 homolog OS=Homo sapiens GN=RTF1 PE=1 SV=4 - [RTF1_HUMAN]	1.075	0.018
Q96DX4	RING finger and SPRY domain-containing protein 1 OS=Homo sapiens GN=RSPRY1 PE=2 SV=1 - [RSPRY_HUMAN]	1.076	0.024
Q9H497	Torsin-3A OS=Homo sapiens GN=TOR3A PE=1 SV=1 - [TOR3A_HUMAN]	1.076	0.045
Q13574-2	Isoform 2 of Diacylglycerol kinase zeta OS=Homo sapiens GN=DGKZ - [DGKZ_HUMAN]	1.076	0.000
Q96N67-4	Isoform 4 of Dedicator of cytokinesis protein 7 OS=Homo sapiens GN=DOCK7 - [DOCK7_HUMAN]	1.078	0.044
Q7Z3T8	Zinc finger FYVE domain-containing protein 16 OS=Homo sapiens GN=ZFYVE16 PE=1 SV=3 - [ZFY16_HUMAN]	1.078	0.050
Q14654	ATP-sensitive inward rectifier potassium channel 11 OS=Homo sapiens GN=KCNJ11 PE=1 SV=2 - [IRK11_HUMAN]	1.079	0.015
Q9P2G1	Ankyrin repeat and IBR domain-containing protein 1 OS=Homo sapiens GN=ANKIB1 PE=1 SV=3 - [AKIB1_HUMAN]	1.079	0.019

Q69YN2	CWF19-like protein 1 OS=Homo sapiens GN=CWF19L1 PE=1 SV=2 - [C19L1_HUMAN]	1.081	0.022
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1 - [AMBP_HUMAN]	1.081	0.026
O00151	PDZ and LIM domain protein 1 OS=Homo sapiens GN=PDLIM1 PE=1 SV=4 - [PDL1_HUMAN]	1.082	0.027
P27701	CD82 antigen OS=Homo sapiens GN=CD82 PE=1 SV=1 - [CD82_HUMAN]	1.083	0.023
P62314	Small nuclear ribonucleoprotein Sm D1 OS=Homo sapiens GN=SNRPD1 PE=1 SV=1 - [SMD1_HUMAN]	1.085	0.023
O75113	NEDD4-binding protein 1 OS=Homo sapiens GN=N4BP1 PE=1 SV=4 - [N4BP1_HUMAN]	1.086	0.046
P49760-3	Isoform 3 of Dual specificity protein kinase CLK2 OS=Homo sapiens GN=CLK2 - [CLK2_HUMAN]	1.089	0.024
Q5TA45-2	Isoform 2 of Integrator complex subunit 11 OS=Homo sapiens GN=CPSF3L - [INT11_HUMAN]	1.090	0.046
Q6AZY7-2	Isoform 2 of Scavenger receptor class A member 3 OS=Homo sapiens GN=SCARA3 - [SCAR3_HUMAN]	1.090	0.022
Q9H3H1-4	Isoform 4 of tRNA dimethylallyltransferase, mitochondrial OS=Homo sapiens GN=TRIT1 - [MOD5_HUMAN]	1.091	0.039
Q9H0G5	Nuclear speckle splicing regulatory protein 1 OS=Homo sapiens GN=NSRP1 PE=1 SV=1 - [NSRP1_HUMAN]	1.092	0.029
Q96F10	Diamine acetyltransferase 2 OS=Homo sapiens GN=SAT2 PE=1 SV=1 - [SAT2_HUMAN]	1.092	0.037
O94880-2	Isoform 2 of PHD finger protein 14 OS=Homo sapiens GN=PHF14 - [PHF14_HUMAN]	1.093	0.047
Q6NTF9	Rhomboid domain-containing protein 2 OS=Homo sapiens GN=RHBDD2 PE=2 SV=2 - [RHBD2_HUMAN]	1.094	0.038
Q9BVJ7	Dual specificity protein phosphatase 23 OS=Homo sapiens GN=DUSP23 PE=1 SV=1 - [DUS23_HUMAN]	1.094	0.046
P46779	60S ribosomal protein L28 OS=Homo sapiens GN=RPL28 PE=1 SV=3 - [RL28_HUMAN]	1.095	0.039
Q9Y2G2-2	Isoform 2 of Caspase recruitment domain-containing protein 8 OS=Homo sapiens GN=CARD8 - [CARD8_HUMAN]	1.095	0.047
O14972	Down syndrome critical region protein 3 OS=Homo sapiens GN=DSCR3 PE=1 SV=1 - [DSCR3_HUMAN]	1.095	0.041
Q92968	Peroxisomal membrane protein PEX13 OS=Homo sapiens GN=PEX13 PE=1 SV=2 - [PEX13_HUMAN]	1.096	0.034
Q9ULV0	Myosin-Vb OS=Homo sapiens GN=MYO5B PE=1 SV=3 - [MYO5B_HUMAN]	1.096	0.033
P30419	Glycylpeptide N-tetradecanoyltransferase 1 OS=Homo sapiens GN=NMT1 PE=1 SV=2 - [NMT1_HUMAN]	1.097	0.046
O75467	Zinc finger protein 324A OS=Homo sapiens GN=ZNF324 PE=1 SV=1 - [Z324A_HUMAN]	1.098	0.029
O60294	Leucine carboxyl methyltransferase 2 OS=Homo sapiens GN=LCMT2 PE=1 SV=3 - [LCMT2_HUMAN]	1.100	0.044
Q659A1	NMDA receptor-regulated protein 2 OS=Homo sapiens GN=NARG2 PE=1 SV=2 - [NARG2_HUMAN]	1.100	0.038
Q9H061	Transmembrane protein 126A OS=Homo sapiens GN=TMEM126A PE=1 SV=1 - [T126A_HUMAN]	1.101	0.030
Q96SZ5	2-aminoethanethiol dioxygenase OS=Homo sapiens GN=ADO PE=1 SV=2 - [AEDO_HUMAN]	1.101	0.040
Q9BRS2	Serine/threonine-protein kinase RIO1 OS=Homo sapiens GN=RIOK1 PE=1 SV=2 - [RIOK1_HUMAN]	1.103	0.024
Q4KMP7	TBC1 domain family member 10B OS=Homo sapiens GN=TBC1D10B PE=1 SV=3 - [TB10B_HUMAN]	1.104	0.030
Q6PJW8-2	Isoform 2 of Consortin OS=Homo sapiens GN=CNST - [CNST_HUMAN]	1.105	0.025

Q9GZNI	Actin-related protein 6 OS=Homo sapiens GN=ACTR6 PE=1 SV=1 - [ARP6_HUMAN]	1.106	0.046
Q9UBN7	Histone deacetylase 6 OS=Homo sapiens GN=HDAC6 PE=1 SV=2 - [HDAC6_HUMAN]	1.107	0.045
Q12986	Transcriptional repressor NF-X1 OS=Homo sapiens GN=NFX1 PE=1 SV=2 - [NFX1_HUMAN]	1.107	0.011
P42680	Tyrosine-protein kinase Tec OS=Homo sapiens GN=TEC PE=1 SV=2 - [TEC_HUMAN]	1.109	0.016
Q9NW75-2	Isoform 2 of G patch domain-containing protein 2 OS=Homo sapiens GN=GPATCH2 - [GPTC2_HUMAN]	1.110	0.041
Q92830	Histone acetyltransferase KAT2A OS=Homo sapiens GN=KAT2A PE=1 SV=3 - [KAT2A_HUMAN]	1.111	0.026
O15194-2	Isoform 2 of CTD small phosphatase-like protein OS=Homo sapiens GN=CTDSPL - [CTDSL_HUMAN]	1.113	0.045
Q15223	Poliovirus receptor-related protein 1 OS=Homo sapiens GN=PVRL1 PE=1 SV=3 - [PVRL1_HUMAN]	1.114	0.005
O43505	N-acetyllactosaminide beta-1,3-N-acetylglucosaminyltransferase OS=Homo sapiens GN=B3GNT1 PE=1 SV=1 - [B3GN1_HUMAN]	1.115	0.043
Q92733	Proline-rich protein PRCC OS=Homo sapiens GN=PRCC PE=1 SV=1 - [PRCC_HUMAN]	1.116	0.025
Q13425	Beta-2-syntrophin OS=Homo sapiens GN=SNTB2 PE=1 SV=1 - [SNTB2_HUMAN]	1.116	0.036
Q9BX95	Sphingosine-1-phosphate phosphatase 1 OS=Homo sapiens GN=SGPP1 PE=1 SV=2 - [SGPP1_HUMAN]	1.117	0.021
P59780	AP-3 complex subunit sigma-2 OS=Homo sapiens GN=AP3S2 PE=2 SV=1 - [AP3S2_HUMAN]	1.121	0.034
Q9BVC5	Ashwin OS=Homo sapiens GN=C2orf49 PE=1 SV=1 - [ASHWN_HUMAN]	1.126	0.038
P61313	60S ribosomal protein L15 OS=Homo sapiens GN=RPL15 PE=1 SV=2 - [RL15_HUMAN]	1.127	0.017
Q9NXN4-2	Isoform 2 of Ganglioside-induced differentiation-associated protein 2 OS=Homo sapiens GN=GDAP2 - [GDAP2_HUMAN]	1.127	0.003
O00268	Transcription initiation factor TFIID subunit 4 OS=Homo sapiens GN=TAF4 PE=1 SV=2 - [TAF4_HUMAN]	1.129	0.029
O00628	Peroxisomal targeting signal 2 receptor OS=Homo sapiens GN=PEX7 PE=1 SV=1 - [PEX7_HUMAN]	1.136	0.027
Q96PQ7	Kelch-like protein 5 OS=Homo sapiens GN=KLHL5 PE=2 SV=3 - [KLHL5_HUMAN]	1.136	0.008
Q86VR2	Protein FAM134C OS=Homo sapiens GN=FAM134C PE=1 SV=1 - [F134C_HUMAN]	1.140	0.020
P32121-2	Isoform 3 of Beta-arrestin-2 OS=Homo sapiens GN=ARRB2 - [ARRB2_HUMAN]	1.141	0.004
Q9NQG7-2	Isoform 2 of Hermansky-Pudlak syndrome 4 protein OS=Homo sapiens GN=HPS4 - [HPS4_HUMAN]	1.142	0.036
Q9H7X7-2	Isoform 2 of Rab-like protein 5 OS=Homo sapiens GN=RABL5 - [RABL5_HUMAN]	1.145	0.039
Q9UQR0	Sex comb on midleg-like protein 2 OS=Homo sapiens GN=SCML2 PE=1 SV=1 - [SCML2_HUMAN]	1.145	0.023
Q9C029-3	Isoform 2 of Tripartite motif-containing protein 7 OS=Homo sapiens GN=TRIM7 - [TRIM7_HUMAN]	1.147	0.041
O94916-2	Isoform A of Nuclear factor of activated T-cells 5 OS=Homo sapiens GN=NFAT5 - [NFAT5_HUMAN]	1.149	0.038
Q5T8P6-6	Isoform 6 of RNA-binding protein 26 OS=Homo sapiens GN=RBM26 - [RBM26_HUMAN]	1.149	0.048
Q14494-2	Isoform 2 of Nuclear factor erythroid 2-related factor 1 OS=Homo sapiens GN=NFE2L1 - [NF2L1_HUMAN]	1.152	0.031
Q9NU19	TBC1 domain family member 22B OS=Homo sapiens GN=TBC1D22B PE=1 SV=3 - [TB22B_HUMAN]	1.153	0.049

Q9BT25-2	Isoform 2 of HAUS augmin-like complex subunit 8 OS=Homo sapiens GN=HAUS8 - [HAUS8_HUMAN]	1.157	0.037
P27930-2	Isoform Short of Interleukin-1 receptor type 2 OS=Homo sapiens GN=IL1R2 - [IL1R2_HUMAN]	1.158	0.044
Q9Y256	CAAX prenyl protease 2 OS=Homo sapiens GN=RCE1 PE=1 SV=1 - [FACE2_HUMAN]	1.158	0.047
Q9P2Q2	FERM domain-containing protein 4A OS=Homo sapiens GN=FRMD4A PE=1 SV=3 - [FRM4A_HUMAN]	1.158	0.010
Q6P444	Protein FAM54A OS=Homo sapiens GN=FAM54A PE=2 SV=2 - [FA54A_HUMAN]	1.162	0.030
Q99758	ATP-binding cassette sub-family A member 3 OS=Homo sapiens GN=ABCA3 PE=1 SV=2 - [ABCA3_HUMAN]	1.164	0.021
O15169-2	Isoform 2 of Axin-1 OS=Homo sapiens GN=AXIN1 - [AXIN1_HUMAN]	1.166	0.040
Q9BUK0	Coiled-coil-helix-coiled-coil-helix domain-containing protein 7 OS=Homo sapiens GN=CHCHD7 PE=2 SV=1 - [CHCH7_HUMAN]	1.174	0.045
O14519	Cyclin-dependent kinase 2-associated protein 1 OS=Homo sapiens GN=CDK2AP1 PE=1 SV=1 - [CDKA1_HUMAN]	1.176	0.027
Q8IWY9-1	Isoform 1 of Codanin-1 OS=Homo sapiens GN=CDAN1 - [CDAN1_HUMAN]	1.177	0.029
Q9NSA3	Beta-catenin-interacting protein 1 OS=Homo sapiens GN=CTNNBIP1 PE=1 SV=1 - [CNBP1_HUMAN]	1.184	0.032
Q9H2G9	Golgin-45 OS=Homo sapiens GN=BLZF1 PE=1 SV=2 - [GO45_HUMAN]	1.190	0.039
Q9NPI8	Fanconi anemia group F protein OS=Homo sapiens GN=FANCF PE=1 SV=1 - [FANCF_HUMAN]	1.191	0.021
Q12805-2	Isoform 2 of EGF-containing fibulin-like extracellular matrix protein 1 OS=Homo sapiens GN=EFEMP1 - [FBLN3_HUMAN]	1.195	0.012
P35527	Keratin, type I cytoskeletal 9 OS=Homo sapiens GN=KRT9 PE=1 SV=3 - [K1C9_HUMAN]	1.197	0.039
Q8TB52	F-box only protein 30 OS=Homo sapiens GN=FBXO30 PE=1 SV=3 - [FBX30_HUMAN]	1.199	0.046
Q96RT8	Gamma-tubulin complex component 5 OS=Homo sapiens GN=TUBGCP5 PE=1 SV=1 - [GCP5_HUMAN]	1.203	0.011
Q12983	BCL2/adenovirus E1B 19 kDa protein-interacting protein 3 OS=Homo sapiens GN=BNIP3 PE=1 SV=2 - [BNIP3_HUMAN]	1.204	0.008
P55197	Protein AF-10 OS=Homo sapiens GN=MLLT10 PE=1 SV=1 - [AF10_HUMAN]	1.215	0.009
Q9H7E9	UPF0488 protein C8orf33 OS=Homo sapiens GN=C8orf33 PE=1 SV=1 - [CH033_HUMAN]	1.219	0.044
P08670	Vimentin OS=Homo sapiens GN=VIM PE=1 SV=4 - [VIME_HUMAN]	1.226	0.027
Q14296	Fas-activated serine/threonine kinase OS=Homo sapiens GN=FASTK PE=1 SV=1 - [FASTK_HUMAN]	1.228	0.023
Q86YD1-3	Isoform 3 of Prostate tumor-overexpressed gene 1 protein OS=Homo sapiens GN=PTOV1 - [PTOV1_HUMAN]	1.229	0.025
Q99985	Semaphorin-3C OS=Homo sapiens GN=SEMA3C PE=1 SV=2 - [SEM3C_HUMAN]	1.230	0.012
Q8N394	Transmembrane and TPR repeat-containing protein 2 OS=Homo sapiens GN=TMTC2 PE=2 SV=1 - [TMTC2_HUMAN]	1.232	0.040
Q15413-3	Isoform 3 of Ryanodine receptor 3 OS=Homo sapiens GN=RYSR3 - [RYSR3_HUMAN]	1.243	0.011
P00973-2	Isoform p41 of 2'-5'-oligoadenylate synthase 1 OS=Homo sapiens GN=OAS1 - [OAS1_HUMAN]	1.245	0.005
Q9Y6B6	GTP-binding protein SAR1b OS=Homo sapiens GN=SAR1B PE=1 SV=1 - [SAR1B_HUMAN]	1.246	0.030
Q13795	ADP-ribosylation factor-related protein 1 OS=Homo sapiens GN=ARFRP1 PE=1 SV=1 - [ARFRP_HUMAN]	1.247	0.047

P13611-2	Isoform V1 of Versican core protein OS=Homo sapiens GN=VCAN - [CSPG2_HUMAN]	1.254	0.044
Q8IYN2	Transcription elongation factor A protein-like 8 OS=Homo sapiens GN=TCEAL8 PE=2 SV=1 - [TCAL8_HUMAN]	1.257	0.043
O00506	Serine/threonine-protein kinase 25 OS=Homo sapiens GN=STK25 PE=1 SV=1 - [STK25_HUMAN]	1.260	0.036
Q5SWA1	Protein phosphatase 1 regulatory subunit 15B OS=Homo sapiens GN=PPP1R15B PE=1 SV=1 - [PR15B_HUMAN]	1.263	0.016
Q13164-2	Isoform 2 of Mitogen-activated protein kinase 7 OS=Homo sapiens GN=MAPK7 - [MK07_HUMAN]	1.272	0.030
O43542	DNA repair protein XRCC3 OS=Homo sapiens GN=XRCC3 PE=1 SV=1 - [XRCC3_HUMAN]	1.272	0.013
O60930	Ribonuclease H1 OS=Homo sapiens GN=RNASEH1 PE=1 SV=2 - [RNH1_HUMAN]	1.274	0.026
O15040	Tectonin beta-propeller repeat-containing protein 2 OS=Homo sapiens GN=TECPR2 PE=1 SV=4 - [TCPR2_HUMAN]	1.276	0.000
O15055	Period circadian protein homolog 2 OS=Homo sapiens GN=PER2 PE=1 SV=2 - [PER2_HUMAN]	1.277	0.006
Q9P2W1	Homologous-pairing protein 2 homolog OS=Homo sapiens GN=PSMC3IP PE=1 SV=1 - [HOP2_HUMAN]	1.289	0.030
Q6DCA0	AMMECR1-like protein OS=Homo sapiens GN=AMMECR1L PE=1 SV=1 - [AMERL_HUMAN]	1.291	0.009
Q9NQ34	Transmembrane protein 9B OS=Homo sapiens GN=TMEM9B PE=1 SV=1 - [TMM9B_HUMAN]	1.317	0.044
Q96S15-2	Isoform 2 of WD repeat-containing protein 24 OS=Homo sapiens GN=WDR24 - [WDR24_HUMAN]	1.318	0.040
Q9Y6M4-3	Isoform 3 of Casein kinase I isoform gamma-3 OS=Homo sapiens GN=CSNK1G3 - [KC1G3_HUMAN]	1.320	0.040
P15923-2	Isoform E47 of Transcription factor E2-alpha OS=Homo sapiens GN=TCF3 - [TFE2_HUMAN]	1.322	0.017
Q562F6-2	Isoform 2 of Shugoshin-like 2 OS=Homo sapiens GN=SGOL2 - [SGOL2_HUMAN]	1.354	0.030
P49447	Cytochrome b561 OS=Homo sapiens GN=CYB561 PE=2 SV=2 - [CY561_HUMAN]	1.359	0.043
Q99715-4	Isoform 4 of Collagen alpha-1(XII) chain OS=Homo sapiens GN=COL12A1 - [COCA1_HUMAN]	1.368	0.042
O14628-4	Isoform 4 of Zinc finger protein 195 OS=Homo sapiens GN=ZNF195 - [ZN195_HUMAN]	1.440	0.040
Q08043	Alpha-actinin-3 OS=Homo sapiens GN=ACTN3 PE=1 SV=2 - [ACTN3_HUMAN]	1.453	0.009
P36894	Bone morphogenetic protein receptor type-1A OS=Homo sapiens GN=BMPR1A PE=1 SV=2 - [BMR1A_HUMAN]	1.533	0.023
P53667-2	Isoform 2 of LIM domain kinase 1 OS=Homo sapiens GN=LIMK1 - [LIMK1_HUMAN]	1.581	0.014
Q9UBT7-2	Isoform 2 of Alpha-catulin OS=Homo sapiens GN=CTNNAL1 - [CTNL1_HUMAN]	1.610	0.013
Q8IYH5-2	Isoform 2 of ZZ-type zinc finger-containing protein 3 OS=Homo sapiens GN=ZZZ3 - [ZZZ3_HUMAN]	1.942	0.037
H727 - 6 hours lanreotide (205 proteins)			
Q8N0X2-4	Isoform 4 of Sperm-associated antigen 16 protein OS=Homo sapiens GN=SPAG16 - [SPG16_HUMAN]	0.106	0.000
O76027	Annexin A9 OS=Homo sapiens GN=ANXA9 PE=1 SV=3 - [ANXA9_HUMAN]	0.419	0.039
P11926	Ornithine decarboxylase OS=Homo sapiens GN=ODC1 PE=1 SV=2 - [DCOR_HUMAN]	0.527	0.005

Q92802	NEDD4-binding protein 2-like 2 OS=Homo sapiens GN=N4BP2L2 PE=1 SV=1 - [N42L2_HUMAN]	0.551	0.045
Q9H8M1	Coenzyme Q-binding protein COQ10 homolog B, mitochondrial OS=Homo sapiens GN=COQ10B PE=2 SV=1 - [CQ10B_HUMAN]	0.625	0.021
P18847	Cyclic AMP-dependent transcription factor ATF-3 OS=Homo sapiens GN=ATF3 PE=1 SV=2 - [ATF3_HUMAN]	0.628	0.049
O75382-3	Isoform Gamma of Tripartite motif-containing protein 3 OS=Homo sapiens GN=TRIM3 - [TRIM3_HUMAN]	0.653	0.019
Q15011-3	Isoform 3 of Homocysteine-responsive endoplasmic reticulum-resident ubiquitin-like domain member 1 protein OS=Homo sapiens GN=HERPUD1 - [HERP1_HUMAN]	0.682	0.042
Q96DN5-2	Isoform 2 of WD repeat-containing protein 67 OS=Homo sapiens GN=WDR67 - [WDR67_HUMAN]	0.723	0.024
Q99576-4	Isoform 3 of TSC22 domain family protein 3 OS=Homo sapiens GN=TSC22D3 - [T22D3_HUMAN]	0.728	0.001
Q71DI3	Histone H3.2 OS=Homo sapiens GN=HIST2H3A PE=1 SV=3 - [H32_HUMAN]	0.728	0.049
Q9H0X9	Oxysterol-binding protein-related protein 5 OS=Homo sapiens GN=OSBPL5 PE=1 SV=1 - [OSBL5_HUMAN]	0.740	0.011
O60907-2	Isoform 2 of F-box-like/WD repeat-containing protein TBL1X OS=Homo sapiens GN=TBL1X - [TBL1X_HUMAN]	0.753	0.042
P53667-2	Isoform 2 of LIM domain kinase 1 OS=Homo sapiens GN=LIMK1 - [LIMK1_HUMAN]	0.772	0.034
Q9H3U5-2	Isoform 2 of Major facilitator superfamily domain-containing protein 1 OS=Homo sapiens GN=MFSD1 - [MFSD1_HUMAN]	0.784	0.049
O43246	Cationic amino acid transporter 4 OS=Homo sapiens GN=SLC7A4 PE=2 SV=3 - [CTR4_HUMAN]	0.791	0.008
Q08AN1	Zinc finger protein 616 OS=Homo sapiens GN=ZNF616 PE=2 SV=2 - [ZN616_HUMAN]	0.827	0.048
P19532	Transcription factor E3 OS=Homo sapiens GN=TFE3 PE=1 SV=4 - [TFE3_HUMAN]	0.829	0.042
Q5T7W0-3	Isoform 3 of Zinc finger protein 618 OS=Homo sapiens GN=ZNF618 - [ZN618_HUMAN]	0.830	0.027
Q9NUE0	Palmitoyltransferase ZDHHC18 OS=Homo sapiens GN=ZDHHC18 PE=2 SV=2 - [ZDH18_HUMAN]	0.832	0.035
P42685	Tyrosine-protein kinase FRK OS=Homo sapiens GN=FRK PE=1 SV=1 - [FRK_HUMAN]	0.839	0.017
P17275	Transcription factor jun-B OS=Homo sapiens GN=JUNB PE=1 SV=1 - [JUNB_HUMAN]	0.845	0.039
Q3MIT2	Putative tRNA pseudouridine synthase Pus10 OS=Homo sapiens GN=PUS10 PE=1 SV=1 - [PUS10_HUMAN]	0.845	0.041
Q8IYU4-2	Isoform 2 of Ubiquilin-like protein OS=Homo sapiens GN=UBQLNL - [UBQLN_HUMAN]	0.854	0.037
Q08495	Dematin OS=Homo sapiens GN=EPB49 PE=1 SV=3 - [DEMA_HUMAN]	0.855	0.023
P43007	Neutral amino acid transporter A OS=Homo sapiens GN=SLC1A4 PE=1 SV=1 - [SATT_HUMAN]	0.859	0.014
Q07866-3	Isoform G of Kinesin light chain 1 OS=Homo sapiens GN=KLC1 - [KLC1_HUMAN]	0.859	0.041
Q2M3G4-2	Isoform 2 of Protein Shroom1 OS=Homo sapiens GN=SHROOM1 - [SHRM1_HUMAN]	0.860	0.043
Q969S8-2	Isoform 2 of Histone deacetylase 10 OS=Homo sapiens GN=HDAC10 - [HDA10_HUMAN]	0.861	0.040
Q8N1P7	Absent in melanoma 1-like protein OS=Homo sapiens GN=AIM1L PE=2 SV=1 - [AIM1L_HUMAN]	0.867	0.013
O75054	Immunoglobulin superfamily member 3 OS=Homo sapiens GN=IGSF3 PE=1 SV=3 - [IGSF3_HUMAN]	0.870	0.008
Q9Y6K5	2'-5'-oligoadenylate synthase 3 OS=Homo sapiens GN=OAS3 PE=1 SV=3 - [OAS3_HUMAN]	0.873	0.044

Q9C0D6	FH2 domain-containing protein 1 OS=Homo sapiens GN=FHDC1 PE=2 SV=2 - [FHDC1_HUMAN]	0.875	0.019
Q96EN8	Molybdenum cofactor sulfurase OS=Homo sapiens GN=MOCOS PE=1 SV=2 - [MOCOS_HUMAN]	0.877	0.012
P31146	Coronin-1A OS=Homo sapiens GN=CORO1A PE=1 SV=4 - [COR1A_HUMAN]	0.880	0.014
P55210-2	Isoform Beta of Caspase-7 OS=Homo sapiens GN=CASP7 - [CASP7_HUMAN]	0.883	0.042
P15121	Aldose reductase OS=Homo sapiens GN=AKR1B1 PE=1 SV=3 - [ALDR_HUMAN]	0.883	0.023
Q9UIK4	Death-associated protein kinase 2 OS=Homo sapiens GN=DAPK2 PE=1 SV=1 - [DAPK2_HUMAN]	0.886	0.049
Q8TEY7-2	Isoform 2 of Ubiquitin carboxyl-terminal hydrolase 33 OS=Homo sapiens GN=USP33 - [UBP33_HUMAN]	0.893	0.002
Q4VC44	FLYWCH-type zinc finger-containing protein 1 OS=Homo sapiens GN=FLYWCH1 PE=1 SV=2 - [FWCH1_HUMAN]	0.899	0.018
Q9BR77	Coiled-coil domain-containing protein 77 OS=Homo sapiens GN=CCDC77 PE=2 SV=1 - [CCD77_HUMAN]	0.899	0.046
Q9UNA1-2	Isoform 2 of Rho GTPase-activating protein 26 OS=Homo sapiens GN=ARHGAP26 - [RHG26_HUMAN]	0.899	0.000
P07711	Cathepsin L1 OS=Homo sapiens GN=CTSL1 PE=1 SV=2 - [CATL1_HUMAN]	0.901	0.019
P21953	2-oxoisovalerate dehydrogenase subunit beta, mitochondrial OS=Homo sapiens GN=BCKDHB PE=1 SV=2 - [ODBB_HUMAN]	0.901	0.030
Q5VWZ2-2	Isoform 2 of Lysophospholipase-like protein 1 OS=Homo sapiens GN=LYPLAL1 - [LYPL1_HUMAN]	0.902	0.003
O60858	E3 ubiquitin-protein ligase TRIM13 OS=Homo sapiens GN=TRIM13 PE=1 SV=2 - [TRI13_HUMAN]	0.903	0.043
P55789	FAD-linked sulfhydryl oxidase ALR OS=Homo sapiens GN=GFER PE=1 SV=2 - [ALR_HUMAN]	0.906	0.035
P16144-4	Isoform Beta-4D of Integrin beta-4 OS=Homo sapiens GN=ITGB4 - [ITB4_HUMAN]	0.907	0.037
Q9P0J1	[Pyruvate dehydrogenase [acetyl-transferring]]-phosphatase 1, mitochondrial OS=Homo sapiens GN=PDP1 PE=1 SV=3 - [PDP1_HUMAN]	0.908	0.049
Q86SQ7-3	Isoform 3 of Serologically defined colon cancer antigen 8 OS=Homo sapiens GN=SDCCAG8 - [SDCG8_HUMAN]	0.910	0.047
Q9UI14	Prenylated Rab acceptor protein 1 OS=Homo sapiens GN=RABAC1 PE=1 SV=1 - [PRAF1_HUMAN]	0.911	0.035
Q969E8	Pre-rRNA-processing protein TSR2 homolog OS=Homo sapiens GN=TSR2 PE=1 SV=1 - [TSR2_HUMAN]	0.912	0.027
Q9HBU6	Ethanolamine kinase 1 OS=Homo sapiens GN=ETNK1 PE=1 SV=1 - [EKI1_HUMAN]	0.914	0.031
P53634	Dipeptidyl peptidase 1 OS=Homo sapiens GN=CTSC PE=1 SV=2 - [CATC_HUMAN]	0.915	0.031
Q2TAA2	Isoamyl acetate-hydrolyzing esterase 1 homolog OS=Homo sapiens GN=IAH1 PE=1 SV=1 - [IAH1_HUMAN]	0.917	0.012
O14727-3	Isoform 3 of Apoptotic protease-activating factor 1 OS=Homo sapiens GN=APAF1 - [APAF_HUMAN]	0.917	0.022
O95372	Acyl-protein thioesterase 2 OS=Homo sapiens GN=LYPLA2 PE=1 SV=1 - [LYPA2_HUMAN]	0.918	0.026
Q08AG7	Mitotic-spindle organizing protein 1 OS=Homo sapiens GN=MZT1 PE=1 SV=2 - [MZT1_HUMAN]	0.919	0.033
P82914	28S ribosomal protein S15, mitochondrial OS=Homo sapiens GN=MRPS15 PE=1 SV=1 - [RT15_HUMAN]	0.919	0.010
A8MPS7	UPF0249 protein ydjC homolog OS=Homo sapiens GN=YDJC PE=2 SV=1 - [YDJC_HUMAN]	0.923	0.025
A6NJ78	Probable methyltransferase-like protein 15 OS=Homo sapiens GN=METT15 PE=2 SV=1 - [MET15_HUMAN]	0.923	0.020

Q9UBN7	Histone deacetylase 6 OS=Homo sapiens GN=HDAC6 PE=1 SV=2 - [HDAC6_HUMAN]	0.924	0.044
Q9BWM7	Sideroflexin-3 OS=Homo sapiens GN=SFXN3 PE=2 SV=2 - [SFXN3_HUMAN]	0.925	0.003
Q9UJS0	Calcium-binding mitochondrial carrier protein Aralar2 OS=Homo sapiens GN=SLC25A13 PE=1 SV=2 - [CMC2_HUMAN]	0.925	0.023
Q7Z3D6-4	Isoform 4 of UPF0317 protein C14orf159, mitochondrial OS=Homo sapiens GN=C14orf159 - [CN159_HUMAN]	0.926	0.037
P35237	Serpin B6 OS=Homo sapiens GN=SERPINB6 PE=1 SV=3 - [SPB6_HUMAN]	0.927	0.032
P09467	Fructose-1,6-bisphosphatase 1 OS=Homo sapiens GN=FBP1 PE=1 SV=5 - [F16P1_HUMAN]	0.927	0.035
Q6ZS17-2	Isoform 2 of Protein FAM65A OS=Homo sapiens GN=FAM65A - [FA65A_HUMAN]	0.927	0.039
O15511	Actin-related protein 2/3 complex subunit 5 OS=Homo sapiens GN=ARPC5 PE=1 SV=3 - [ARPC5_HUMAN]	0.928	0.020
O00165-5	Isoform 5 of HCLS1-associated protein X-1 OS=Homo sapiens GN=HAX1 - [HAX1_HUMAN]	0.929	0.003
O75396	Vesicle-trafficking protein SEC22b OS=Homo sapiens GN=SEC22B PE=1 SV=4 - [SC22B_HUMAN]	0.929	0.030
P35270	Sepiapterin reductase OS=Homo sapiens GN=SPR PE=1 SV=1 - [SPRE_HUMAN]	0.930	0.042
Q8IWF6	Protein FAM116A OS=Homo sapiens GN=FAM116A PE=2 SV=1 - [F116A_HUMAN]	0.931	0.049
Q9HBH1	Peptide deformylase, mitochondrial OS=Homo sapiens GN=PDF PE=1 SV=1 - [DEFM_HUMAN]	0.931	0.045
Q16706	Alpha-mannosidase 2 OS=Homo sapiens GN=MAN2A1 PE=1 SV=2 - [MA2A1_HUMAN]	0.932	0.016
P13807	Glycogen [starch] synthase, muscle OS=Homo sapiens GN=GYS1 PE=1 SV=2 - [GYS1_HUMAN]	0.932	0.034
O75427	Leucine-rich repeat and calponin homology domain-containing protein 4 OS=Homo sapiens GN=LRCH4 PE=1 SV=2 - [LRCH4_HUMAN]	0.935	0.016
Q8WXS8-2	Isoform B of A disintegrin and metalloproteinase with thrombospondin motifs 14 OS=Homo sapiens GN=ADAMTS14 - [ATS14_HUMAN]	0.936	0.041
Q9UFC0	Leucine-rich repeat and WD repeat-containing protein 1 OS=Homo sapiens GN=LRWD1 PE=1 SV=2 - [LRWD1_HUMAN]	0.936	0.023
P51649	Succinate-semialdehyde dehydrogenase, mitochondrial OS=Homo sapiens GN=ALDH5A1 PE=1 SV=2 - [SSDH_HUMAN]	0.936	0.035
Q8N7R7	Cyclin-Y-like protein 1 OS=Homo sapiens GN=CCNYL1 PE=1 SV=2 - [CCYL1_HUMAN]	0.938	0.042
P04040	Catalase OS=Homo sapiens GN=CAT PE=1 SV=3 - [CATA_HUMAN]	0.938	0.031
O96000	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=Homo sapiens GN=NDUFB10 PE=1 SV=3 - [NDUBA_HUMAN]	0.938	0.007
O14795	Protein unc-13 homolog B OS=Homo sapiens GN=UNC13B PE=1 SV=2 - [UN13B_HUMAN]	0.939	0.043
Q05209	Tyrosine-protein phosphatase non-receptor type 12 OS=Homo sapiens GN=PTPN12 PE=1 SV=3 - [PTN12_HUMAN]	0.940	0.044
Q9UHG3	Prenylcysteine oxidase 1 OS=Homo sapiens GN=PCYOX1 PE=1 SV=3 - [PCYOX_HUMAN]	0.941	0.004
Q9BPY3	Protein FAM118B OS=Homo sapiens GN=FAM118B PE=1 SV=1 - [F118B_HUMAN]	0.947	0.024
P49281-2	Isoform 1 of Natural resistance-associated macrophage protein 2 OS=Homo sapiens GN=SLC11A2 - [NRAM2_HUMAN]	0.949	0.018
O94973-2	Isoform 2 of AP-2 complex subunit alpha-2 OS=Homo sapiens GN=AP2A2 - [AP2A2_HUMAN]	0.950	0.047
P42765	3-ketoacyl-CoA thiolase, mitochondrial OS=Homo sapiens GN=ACAA2 PE=1 SV=2 - [THIM_HUMAN]	0.950	0.026

Q32P41	tRNA (guanine(37)-N1)-methyltransferase OS=Homo sapiens GN=TRMT5 PE=1 SV=2 - [TRM5_HUMAN]	0.951	0.050
O15056-2	Isoform 2B1 of Synaptojanin-2 OS=Homo sapiens GN=SYNJ2 - [SYNJ2_HUMAN]	0.951	0.028
Q96AQ8	Coiled-coil domain-containing protein 90A, mitochondrial OS=Homo sapiens GN=CCDC90A PE=2 SV=1 - [CC90A_HUMAN]	0.951	0.002
Q9BV57	1,2-dihydroxy-3-keto-5-methylthiopentene dioxygenase OS=Homo sapiens GN=ADI1 PE=1 SV=1 - [MTND_HUMAN]	0.954	0.033
P29350	Tyrosine-protein phosphatase non-receptor type 6 OS=Homo sapiens GN=PTPN6 PE=1 SV=1 - [PTN6_HUMAN]	0.956	0.028
O43609	Protein sprouty homolog 1 OS=Homo sapiens GN=SPRY1 PE=1 SV=2 - [SPY1_HUMAN]	0.957	0.000
P43304	Glycerol-3-phosphate dehydrogenase, mitochondrial OS=Homo sapiens GN=GPD2 PE=1 SV=3 - [GPDM_HUMAN]	0.960	0.038
Q9NRF8	CTP synthase 2 OS=Homo sapiens GN=CTPS2 PE=1 SV=1 - [PYRG2_HUMAN]	0.961	0.038
Q9P2R7-2	Isoform 2 of Succinyl-CoA ligase [ADP-forming] subunit beta, mitochondrial OS=Homo sapiens GN=SUCLA2 - [SUCB1_HUMAN]	0.962	0.050
Q9Y230	RuvB-like 2 OS=Homo sapiens GN=RUVBL2 PE=1 SV=3 - [RUVB2_HUMAN]	0.963	0.004
P48634	Protein PRRC2A OS=Homo sapiens GN=PRRC2A PE=1 SV=3 - [PRC2A_HUMAN]	0.963	0.037
P31150	Rab GDP dissociation inhibitor alpha OS=Homo sapiens GN=GDI1 PE=1 SV=2 - [GDIA_HUMAN]	0.964	0.031
Q5ZPR3	CD276 antigen OS=Homo sapiens GN=CD276 PE=1 SV=1 - [CD276_HUMAN]	0.964	0.025
Q9Y5K5-2	Isoform 2 of Ubiquitin carboxyl-terminal hydrolase isozyme L5 OS=Homo sapiens GN=UCHL5 - [UCHL5_HUMAN]	0.964	0.049
Q9Y296	Trafficking protein particle complex subunit 4 OS=Homo sapiens GN=TRAPPC4 PE=1 SV=1 - [TPPC4_HUMAN]	0.966	0.028
Q13948	Protein CASP OS=Homo sapiens GN=CUX1 PE=1 SV=2 - [CASP_HUMAN]	0.970	0.005
Q7Z3E2	Uncharacterized protein C10orf118 OS=Homo sapiens GN=C10orf118 PE=2 SV=2 - [CJ118_HUMAN]	0.973	0.029
P28715	DNA repair protein complementing XP-G cells OS=Homo sapiens GN=ERCC5 PE=1 SV=3 - [ERCC5_HUMAN]	0.974	0.015
Q8N5M1	ATP synthase mitochondrial F1 complex assembly factor 2 OS=Homo sapiens GN=ATPAF2 PE=1 SV=1 - [ATPF2_HUMAN]	0.982	0.023
Q9P265	Disco-interacting protein 2 homolog B OS=Homo sapiens GN=DIP2B PE=1 SV=3 - [DIP2B_HUMAN]	0.983	0.043
Q8IXN7	N-acetylaspartyl-glutamate synthetase A OS=Homo sapiens GN=RIMKLA PE=2 SV=2 - [RIMKA_HUMAN]	0.984	0.030
P07384	Calpain-1 catalytic subunit OS=Homo sapiens GN=CAPN1 PE=1 SV=1 - [CAN1_HUMAN]	0.985	0.030
O00268	Transcription initiation factor TFIID subunit 4 OS=Homo sapiens GN=TAF4 PE=1 SV=2 - [TAF4_HUMAN]	1.019	0.048
Q09161	Nuclear cap-binding protein subunit 1 OS=Homo sapiens GN=NCBP1 PE=1 SV=1 - [NCBP1_HUMAN]	1.019	0.019
Q9Y608-4	Isoform 4 of Leucine-rich repeat flightless-interacting protein 2 OS=Homo sapiens GN=LRRFIP2 - [LRRF2_HUMAN]	1.024	0.036
Q8TC07-2	Isoform 2 of TBC1 domain family member 15 OS=Homo sapiens GN=TBC1D15 - [TBC15_HUMAN]	1.026	0.045
P62330	ADP-ribosylation factor 6 OS=Homo sapiens GN=ARF6 PE=1 SV=2 - [ARF6_HUMAN]	1.027	0.027
P05556	Integrin beta-1 OS=Homo sapiens GN=ITGB1 PE=1 SV=2 - [ITB1_HUMAN]	1.029	0.026
Q9BVC6	Transmembrane protein 109 OS=Homo sapiens GN=TMEM109 PE=1 SV=1 - [TM109_HUMAN]	1.032	0.017

Q6Y7W6	PERQ amino acid-rich with GYF domain-containing protein 2 OS=Homo sapiens GN=GIGYF2 PE=1 SV=1 - [PERQ2_HUMAN]	1.037	0.021
Q9Y2S0	DNA-directed RNA polymerases I and III subunit RPAC2 OS=Homo sapiens GN=POLR1D PE=1 SV=1 - [RPAC2_HUMAN]	1.038	0.027
O43617	Trafficking protein particle complex subunit 3 OS=Homo sapiens GN=TRAPPC3 PE=1 SV=1 - [TPPC3_HUMAN]	1.039	0.045
Q9H3L0	Methylmalonic aciduria and homocystinuria type D protein, mitochondrial OS=Homo sapiens GN=MMADHC PE=1 SV=2 - [MMAD_HUMAN]	1.040	0.022
Q9NXV6	CDKN2A-interacting protein OS=Homo sapiens GN=CDKN2AIP PE=1 SV=3 - [CARF_HUMAN]	1.041	0.037
Q96B96	Promethin OS=Homo sapiens GN=TMEM159 PE=2 SV=2 - [TM159_HUMAN]	1.043	0.021
Q99613	Eukaryotic translation initiation factor 3 subunit C OS=Homo sapiens GN=EIF3C PE=1 SV=1 - [EIF3C_HUMAN]	1.044	0.045
O43681	ATPase ASNA1 OS=Homo sapiens GN=ASNA1 PE=1 SV=2 - [ASNA_HUMAN]	1.047	0.016
Q9BRX2	Protein pelota homolog OS=Homo sapiens GN=PELO PE=1 SV=2 - [PELO_HUMAN]	1.050	0.009
O60292	Signal-induced proliferation-associated 1-like protein 3 OS=Homo sapiens GN=SIPA1L3 PE=1 SV=3 - [SI1L3_HUMAN]	1.050	0.015
O15164-2	Isoform Short of Transcription intermediary factor 1-alpha OS=Homo sapiens GN=TRIM24 - [TIF1A_HUMAN]	1.051	0.040
Q8NHY2-3	Isoform 3 of E3 ubiquitin-protein ligase RFWD2 OS=Homo sapiens GN=RFWD2 - [RFWD2_HUMAN]	1.051	0.005
Q7Z401	C-myc promoter-binding protein OS=Homo sapiens GN=DENND4A PE=1 SV=2 - [MYCPP_HUMAN]	1.052	0.030
Q9UPA5	Protein bassoon OS=Homo sapiens GN=BSN PE=1 SV=4 - [BSN_HUMAN]	1.053	0.008
O95486	Protein transport protein Sec24A OS=Homo sapiens GN=SEC24A PE=1 SV=2 - [SC24A_HUMAN]	1.053	0.041
Q12824-2	Isoform B of SWI/SNF-related matrix-associated actin-dependent regulator of chromatin subfamily B member 1 OS=Homo sapiens GN=SMARCB1 - [SNF5_HUMAN]	1.055	0.005
Q15633-2	Isoform 2 of RISC-loading complex subunit TARBP2 OS=Homo sapiens GN=TARBP2 - [TRBP2_HUMAN]	1.057	0.017
Q9GZS1-2	Isoform 2 of DNA-directed RNA polymerase I subunit RPA49 OS=Homo sapiens GN=POLR1E - [RPA49_HUMAN]	1.058	0.010
Q13283	Ras GTPase-activating protein-binding protein 1 OS=Homo sapiens GN=G3BP1 PE=1 SV=1 - [G3BP1_HUMAN]	1.060	0.044
Q13356	Peptidyl-prolyl cis-trans isomerase-like 2 OS=Homo sapiens GN=PPIL2 PE=1 SV=1 - [PPIL2_HUMAN]	1.063	0.036
Q4LE39-2	Isoform 2 of AT-rich interactive domain-containing protein 4B OS=Homo sapiens GN=ARID4B - [ARI4B_HUMAN]	1.063	0.049
Q8NEZ2-2	Isoform 2 of Vacuolar protein sorting-associated protein 37A OS=Homo sapiens GN=VPS37A - [VP37A_HUMAN]	1.065	0.011
Q92786	Prospero homeobox protein 1 OS=Homo sapiens GN=PROX1 PE=1 SV=2 - [PROX1_HUMAN]	1.067	0.046
Q9NRX2	39S ribosomal protein L17, mitochondrial OS=Homo sapiens GN=MRPL17 PE=1 SV=1 - [RM17_HUMAN]	1.067	0.028
Q6ZSJ8	Uncharacterized protein C1orf122 OS=Homo sapiens GN=C1orf122 PE=2 SV=2 - [CA122_HUMAN]	1.071	0.012
Q9NR28-2	Isoform 2 of Diablo homolog, mitochondrial OS=Homo sapiens GN=DIABLO - [DBLOH_HUMAN]	1.071	0.003
Q9BX95	Sphingosine-1-phosphate phosphatase 1 OS=Homo sapiens GN=SGPP1 PE=1 SV=2 - [SGPP1_HUMAN]	1.072	0.042
Q6GQQ9	OTU domain-containing protein 7B OS=Homo sapiens GN=OTUD7B PE=1 SV=1 - [OTU7B_HUMAN]	1.073	0.002
Q96KQ7-2	Isoform 2 of Histone-lysine N-methyltransferase EHMT2 OS=Homo sapiens GN=EHMT2 - [EHMT2_HUMAN]	1.074	0.014

Q12772	Sterol regulatory element-binding protein 2 OS=Homo sapiens GN=SREBF2 PE=1 SV=2 - [SRBP2_HUMAN]	1.075	0.045
Q15527	Surfeit locus protein 2 OS=Homo sapiens GN=SURF2 PE=1 SV=3 - [SURF2_HUMAN]	1.075	0.006
P06748-2	Isoform 2 of Nucleophosmin OS=Homo sapiens GN=NPM1 - [NPM_HUMAN]	1.075	0.020
Q96LA8-2	Isoform 2 of Protein arginine N-methyltransferase 6 OS=Homo sapiens GN=PRMT6 - [ANM6_HUMAN]	1.077	0.015
O76003	Glutaredoxin-3 OS=Homo sapiens GN=GLRX3 PE=1 SV=2 - [GLRX3_HUMAN]	1.080	0.024
Q9UBP6	tRNA (guanine-N(7)-)-methyltransferase OS=Homo sapiens GN=METTTL1 PE=1 SV=1 - [TRMB_HUMAN]	1.082	0.010
Q96SZ5	2-aminoethanethiol dioxygenase OS=Homo sapiens GN=ADO PE=1 SV=2 - [AEDO_HUMAN]	1.084	0.022
P25686-2	Isoform 2 of DnaJ homolog subfamily B member 2 OS=Homo sapiens GN=DNAJB2 - [DNJB2_HUMAN]	1.085	0.030
Q9Y5F0	Protocadherin beta-13 OS=Homo sapiens GN=PCDHB13 PE=2 SV=1 - [PCDBD_HUMAN]	1.090	0.033
Q12805-2	Isoform 2 of EGF-containing fibulin-like extracellular matrix protein 1 OS=Homo sapiens GN=EFEMP1 - [FBLN3_HUMAN]	1.091	0.018
P27701	CD82 antigen OS=Homo sapiens GN=CD82 PE=1 SV=1 - [CD82_HUMAN]	1.092	0.015
Q13325	Interferon-induced protein with tetratricopeptide repeats 5 OS=Homo sapiens GN=IFIT5 PE=1 SV=1 - [IFIT5_HUMAN]	1.094	0.010
Q8IXW5-2	Isoform 2 of Putative RNA polymerase II subunit B1 CTD phosphatase RPAP2 OS=Homo sapiens GN=RPAP2 - [RPAP2_HUMAN]	1.098	0.048
Q01850	Cerebellar degeneration-related protein 2 OS=Homo sapiens GN=CDR2 PE=1 SV=2 - [CDR2_HUMAN]	1.099	0.036
Q96ES7	SAGA-associated factor 29 homolog OS=Homo sapiens GN=CCDC101 PE=1 SV=1 - [SGF29_HUMAN]	1.099	0.043
Q86VR2	Protein FAM134C OS=Homo sapiens GN=FAM134C PE=1 SV=1 - [F134C_HUMAN]	1.102	0.040
Q9NUP7	tRNA guanosine-2'-O-methyltransferase TRM13 homolog OS=Homo sapiens GN=CCDC76 PE=1 SV=2 - [TRM13_HUMAN]	1.104	0.031
Q9NRG0	Chromatin accessibility complex protein 1 OS=Homo sapiens GN=CHRC1 PE=1 SV=1 - [CHRC1_HUMAN]	1.104	0.022
Q92979	Ribosomal RNA small subunit methyltransferase NEP1 OS=Homo sapiens GN=EMG1 PE=1 SV=4 - [NEP1_HUMAN]	1.107	0.029
Q9UN86-2	Isoform B of Ras GTPase-activating protein-binding protein 2 OS=Homo sapiens GN=G3BP2 - [G3BP2_HUMAN]	1.107	0.015
P53804-3	Isoform TPRDIII of E3 ubiquitin-protein ligase TTC3 OS=Homo sapiens GN=TTC3 - [TTC3_HUMAN]	1.107	0.003
O43865	Putative adenosylhomocysteinase 2 OS=Homo sapiens GN=AHCYL1 PE=1 SV=2 - [SAHH2_HUMAN]	1.108	0.046
Q9UHW5	GPN-loop GTPase 3 OS=Homo sapiens GN=GPN3 PE=1 SV=2 - [GPN3_HUMAN]	1.108	0.003
P50542-3	Isoform 3 of Peroxisomal targeting signal 1 receptor OS=Homo sapiens GN=PEX5 - [PEX5_HUMAN]	1.109	0.048
Q96BJ3	Axin interactor, dorsalization-associated protein OS=Homo sapiens GN=AIDA PE=1 SV=1 - [AIDA_HUMAN]	1.111	0.037
Q93063	Exostosin-2 OS=Homo sapiens GN=EXT2 PE=1 SV=1 - [EXT2_HUMAN]	1.111	0.013
Q9NS00-2	Isoform 2 of Glycoprotein-N-acetylgalactosamine 3-beta-galactosyltransferase 1 OS=Homo sapiens GN=C1GALT1 - [C1GLT_HUMAN]	1.118	0.043
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1 - [AMBP_HUMAN]	1.120	0.026
Q86Y91-2	Isoform 2 of Kinesin-like protein KIF18B OS=Homo sapiens GN=KIF18B - [KI18B_HUMAN]	1.122	0.046

Q95801	Tetratricopeptide repeat protein 4 OS=Homo sapiens GN=TTC4 PE=1 SV=3 - [TTC4_HUMAN]	1.130	0.035
Q8TF30	WASP homolog-associated protein with actin, membranes and microtubules OS=Homo sapiens GN=WHAMM PE=1 SV=2 - [WHAMM_HUMAN]	1.133	0.021
Q9H6E5	Speckle targeted PIP5K1A-regulated poly(A) polymerase OS=Homo sapiens GN=TUT1 PE=1 SV=2 - [STPAP_HUMAN]	1.137	0.020
Q14181	DNA polymerase alpha subunit B OS=Homo sapiens GN=POLA2 PE=1 SV=2 - [DPOA2_HUMAN]	1.147	0.011
Q96FX2	DPH3 homolog OS=Homo sapiens GN=DPH3 PE=1 SV=1 - [DPH3_HUMAN]	1.154	0.022
Q8WUH6	UPF0444 transmembrane protein C12orf23 OS=Homo sapiens GN=C12orf23 PE=1 SV=1 - [CL023_HUMAN]	1.162	0.047
Q96K49-2	Isoform 2 of Transmembrane protein 87B OS=Homo sapiens GN=TMEM87B - [TM87B_HUMAN]	1.162	0.009
Q9BPX7	UPF0415 protein C7orf25 OS=Homo sapiens GN=C7orf25 PE=1 SV=1 - [CG025_HUMAN]	1.165	0.036
Q99697-3	Isoform PTX2A of Pituitary homeobox 2 OS=Homo sapiens GN=PITX2 - [PITX2_HUMAN]	1.165	0.026
Q9BWS9-3	Isoform 3 of Chitinase domain-containing protein 1 OS=Homo sapiens GN=CHID1 - [CHID1_HUMAN]	1.165	0.035
Q95466	Formin-like protein 1 OS=Homo sapiens GN=FMNL1 PE=1 SV=3 - [FMNL_HUMAN]	1.165	0.022
Q9UGY1	Nucleolar protein 12 OS=Homo sapiens GN=NOL12 PE=1 SV=1 - [NOL12_HUMAN]	1.170	0.010
P49427	Ubiquitin-conjugating enzyme E2 R1 OS=Homo sapiens GN=CDC34 PE=1 SV=2 - [UB2R1_HUMAN]	1.173	0.050
Q9UL54-3	Isoform 3 of Serine/threonine-protein kinase TAO2 OS=Homo sapiens GN=TAOK2 - [TAOK2_HUMAN]	1.176	0.006
Q6UXM1-2	Isoform 2 of Leucine-rich repeats and immunoglobulin-like domains protein 3 OS=Homo sapiens GN=LRIG3 - [LRIG3_HUMAN]	1.176	0.048
Q06547-4	Isoform 4 of GA-binding protein subunit beta-1 OS=Homo sapiens GN=GABPB1 - [GABP1_HUMAN]	1.179	0.041
O43734-4	Isoform 4 of Adapter protein CIKS OS=Homo sapiens GN=TRAF3IP2 - [CIKS_HUMAN]	1.181	0.042
Q12983	BCL2/adenovirus E1B 19 kDa protein-interacting protein 3 OS=Homo sapiens GN=BNIP3 PE=1 SV=2 - [BNIP3_HUMAN]	1.183	0.044
Q9Y653-2	Isoform 2 of G-protein coupled receptor 56 OS=Homo sapiens GN=GPR56 - [GPR56_HUMAN]	1.191	0.013
Q8TCY9-3	Isoform 3 of Up-regulator of cell proliferation OS=Homo sapiens GN=URGCP - [URGCP_HUMAN]	1.199	0.026
Q9NXV2	BTB/POZ domain-containing protein KCTD5 OS=Homo sapiens GN=KCTD5 PE=1 SV=1 - [KCTD5_HUMAN]	1.208	0.016
Q16787-1	Isoform 1 of Laminin subunit alpha-3 OS=Homo sapiens GN=LAMA3 - [LAMA3_HUMAN]	1.212	0.017
Q96A19	Coiled-coil domain-containing protein 102A OS=Homo sapiens GN=CCDC102A PE=1 SV=2 - [C102A_HUMAN]	1.218	0.007
Q92823-3	Isoform 3 of Neuronal cell adhesion molecule OS=Homo sapiens GN=NRCAM - [NRCAM_HUMAN]	1.224	0.010
Q5T2R2-3	Isoform 3 of Decaprenyl-diphosphate synthase subunit 1 OS=Homo sapiens GN=PDSS1 - [DPS1_HUMAN]	1.242	0.038
P17707	S-adenosylmethionine decarboxylase proenzyme OS=Homo sapiens GN=AMD1 PE=1 SV=2 - [DCAM_HUMAN]	1.259	0.046
Q6ZRI6-2	Isoform 2 of Uncharacterized protein C15orf39 OS=Homo sapiens GN=C15orf39 - [CO039_HUMAN]	1.305	0.005
O75081-2	Isoform 2 of Protein CBFA2T3 OS=Homo sapiens GN=CBFA2T3 - [MTG16_HUMAN]	1.383	0.013

H727 - 48 hours lanreotide (233 proteins)

Q8N0X2-4	Isoform 4 of Sperm-associated antigen 16 protein OS=Homo sapiens GN=SPAG16 - [SPG16_HUMAN]	0.094	0.003
P11926	Ornithine decarboxylase OS=Homo sapiens GN=ODC1 PE=1 SV=2 - [DCOR_HUMAN]	0.357	0.011
A0FGR8-2	Isoform 2 of Extended synaptotagmin-2 OS=Homo sapiens GN=ESYT2 - [ESYT2_HUMAN]	0.510	0.029
O76027	Annexin A9 OS=Homo sapiens GN=ANXA9 PE=1 SV=3 - [ANXA9_HUMAN]	0.613	0.017
P38936	Cyclin-dependent kinase inhibitor 1 OS=Homo sapiens GN=CDKN1A PE=1 SV=3 - [CDN1A_HUMAN]	0.621	0.021
P30825	High affinity cationic amino acid transporter 1 OS=Homo sapiens GN=SLC7A1 PE=1 SV=1 - [CTR1_HUMAN]	0.635	0.049
Q01101	Insulinoma-associated protein 1 OS=Homo sapiens GN=INSM1 PE=2 SV=1 - [INSM1_HUMAN]	0.670	0.046
P04798	Cytochrome P450 1A1 OS=Homo sapiens GN=CYP1A1 PE=1 SV=1 - [CP1A1_HUMAN]	0.688	0.046
Q15011-3	Isoform 3 of Homocysteine-responsive endoplasmic reticulum-resident ubiquitin-like domain member 1 protein OS=Homo sapiens GN=HERPUD1 - [HERP1_HUMAN]	0.706	0.038
P06732	Creatine kinase M-type OS=Homo sapiens GN=CKM PE=1 SV=2 - [KCRM_HUMAN]	0.738	0.040
P17275	Transcription factor jun-B OS=Homo sapiens GN=JUNB PE=1 SV=1 - [JUNB_HUMAN]	0.744	0.013
Q8N594-2	Isoform 2 of MPN domain-containing protein OS=Homo sapiens GN=MPND - [MPND_HUMAN]	0.744	0.033
Q6P9G4	Transmembrane protein 154 OS=Homo sapiens GN=TMEM154 PE=2 SV=2 - [TM154_HUMAN]	0.758	0.040
Q53GA4	Pleckstrin homology-like domain family A member 2 OS=Homo sapiens GN=PHLDA2 PE=1 SV=2 - [PHLA2_HUMAN]	0.761	0.008
O43474-4	Isoform 3 of Krueppel-like factor 4 OS=Homo sapiens GN=KLF4 - [KLF4_HUMAN]	0.785	0.044
Q9NPA3	Mid1-interacting protein 1 OS=Homo sapiens GN=MID1IP1 PE=1 SV=1 - [M1IP1_HUMAN]	0.792	0.031
P17707	S-adenosylmethionine decarboxylase proenzyme OS=Homo sapiens GN=AMD1 PE=1 SV=2 - [DCAM_HUMAN]	0.800	0.049
Q5T7W0-3	Isoform 3 of Zinc finger protein 618 OS=Homo sapiens GN=ZNF618 - [ZN618_HUMAN]	0.805	0.014
P48739-2	Isoform 2 of Phosphatidylinositol transfer protein beta isoform OS=Homo sapiens GN=PITPNB - [PIPNB_HUMAN]	0.805	0.034
P08247	Synaptophysin OS=Homo sapiens GN=SYP PE=1 SV=3 - [SYPH_HUMAN]	0.811	0.046
P53355-2	Isoform 2 of Death-associated protein kinase 1 OS=Homo sapiens GN=DAPK1 - [DAPK1_HUMAN]	0.819	0.006
P43007	Neutral amino acid transporter A OS=Homo sapiens GN=SLC1A4 PE=1 SV=1 - [SATT_HUMAN]	0.819	0.015
O43581	Synaptotagmin-7 OS=Homo sapiens GN=SYT7 PE=1 SV=3 - [SYT7_HUMAN]	0.826	0.022
Q7RTP0-2	Isoform 2 of Magnesium transporter NIPA1 OS=Homo sapiens GN=NIPA1 - [NIPA1_HUMAN]	0.831	0.034
Q8NET8-3	Isoform 3 of Transient receptor potential cation channel subfamily V member 3 OS=Homo sapiens GN=TRPV3 - [TRPV3_HUMAN]	0.835	0.024
P40121	Macrophage-capping protein OS=Homo sapiens GN=CAPG PE=1 SV=2 - [CAPG_HUMAN]	0.837	0.041
Q9Y394-2	Isoform 2 of Dehydrogenase/reductase SDR family member 7 OS=Homo sapiens GN=DHRS7 - [DHRS7_HUMAN]	0.839	0.039
P10253	Lysosomal alpha-glucosidase OS=Homo sapiens GN=GAA PE=1 SV=4 - [LYAG_HUMAN]	0.853	0.023

P23634-7	Isoform ZB of Plasma membrane calcium-transporting ATPase 4 OS=Homo sapiens GN=ATP2B4 - [AT2B4_HUMAN]	0.861	0.050
P30042-2 Q9NNW7-2	Isoform Short of ES1 protein homolog, mitochondrial OS=Homo sapiens GN=C21orf33 - [ES1_HUMAN]	0.866	0.046
Q02218	Isoform 2 of Thioredoxin reductase 2, mitochondrial OS=Homo sapiens GN=TXNRD2 - [TRXR2_HUMAN]	0.870	0.019
P01116	2-oxoglutarate dehydrogenase, mitochondrial OS=Homo sapiens GN=OGDH PE=1 SV=3 - [ODO1_HUMAN]	0.875	0.024
P40939	GTPase KRas OS=Homo sapiens GN=KRAS PE=1 SV=1 - [RASK_HUMAN]	0.877	0.036
Q06546	Trifunctional enzyme subunit alpha, mitochondrial OS=Homo sapiens GN=HADHA PE=1 SV=2 - [ECHA_HUMAN]	0.878	0.029
Q15651-2	GA-binding protein alpha chain OS=Homo sapiens GN=GABPA PE=1 SV=1 - [GABPA_HUMAN]	0.879	0.026
Q08722	Isoform 2 of High mobility group nucleosome-binding domain-containing protein 3 OS=Homo sapiens GN=HMGN3 - [HMGN3_HUMAN]	0.881	0.007
P17301	Leukocyte surface antigen CD47 OS=Homo sapiens GN=CD47 PE=1 SV=1 - [CD47_HUMAN]	0.882	0.035
Q8NFH8-3	Integrin alpha-2 OS=Homo sapiens GN=ITGA2 PE=1 SV=1 - [ITA2_HUMAN]	0.886	0.014
P11117	Isoform 3 of RalBP1-associated Eps domain-containing protein 2 OS=Homo sapiens GN=REPS2 - [REPS2_HUMAN]	0.887	0.039
Q96AB3	Lysosomal acid phosphatase OS=Homo sapiens GN=ACP2 PE=1 SV=3 - [PPAL_HUMAN]	0.888	0.017
O95372	Isochorismatase domain-containing protein 2, mitochondrial OS=Homo sapiens GN=ISOC2 PE=1 SV=1 - [ISOC2_HUMAN]	0.891	0.006
Q7Z3D6-4	Acyl-protein thioesterase 2 OS=Homo sapiens GN=LYPLA2 PE=1 SV=1 - [LYPA2_HUMAN]	0.892	0.033
P38159	Isoform 4 of UPF0317 protein C14orf159, mitochondrial OS=Homo sapiens GN=C14orf159 - [CN159_HUMAN]	0.895	0.021
P41732	RNA-binding motif protein, X chromosome OS=Homo sapiens GN=RBMX PE=1 SV=3 - [RBMX_HUMAN]	0.898	0.025
Q9BQ95	Tetraspanin-7 OS=Homo sapiens GN=TSPAN7 PE=1 SV=2 - [TSN7_HUMAN]	0.898	0.031
P40925	Evolutionarily conserved signaling intermediate in Toll pathway, mitochondrial OS=Homo sapiens GN=ECSIT PE=1 SV=1 - [ECSIT_HUMAN]	0.898	0.047
Q6P161	Malate dehydrogenase, cytoplasmic OS=Homo sapiens GN=MDH1 PE=1 SV=4 - [MDHC_HUMAN]	0.900	0.050
Q96HE7	39S ribosomal protein L54, mitochondrial OS=Homo sapiens GN=MRPL54 PE=1 SV=1 - [RM54_HUMAN]	0.901	0.015
O43246	ERO1-like protein alpha OS=Homo sapiens GN=ERO1L PE=1 SV=2 - [ERO1A_HUMAN]	0.903	0.024
Q96D53	Cationic amino acid transporter 4 OS=Homo sapiens GN=SLC7A4 PE=2 SV=3 - [CTR4_HUMAN]	0.904	0.012
P42765	Uncharacterized aarF domain-containing protein kinase 4 OS=Homo sapiens GN=ADCK4 PE=1 SV=2 - [ADCK4_HUMAN]	0.904	0.026
P47985	3-ketoacyl-CoA thiolase, mitochondrial OS=Homo sapiens GN=ACAA2 PE=1 SV=2 - [THIM_HUMAN]	0.904	0.014
P13688-2	Cytochrome b-c1 complex subunit Rieske, mitochondrial OS=Homo sapiens GN=UQCRFS1 PE=1 SV=2 - [UCRI_HUMAN]	0.905	0.025
A8MXV4	Isoform 2 of Carcinoembryonic antigen-related cell adhesion molecule 1 OS=Homo sapiens GN=CEACAM1 - [CEAM1_HUMAN]	0.909	0.046
P28347	Nucleoside diphosphate-linked moiety X motif 19, mitochondrial OS=Homo sapiens GN=NUDT19 PE=1 SV=1 - [NUD19_HUMAN]	0.909	0.037
Q9UNA1-2	Transcriptional enhancer factor TEF-1 OS=Homo sapiens GN=TEAD1 PE=1 SV=2 - [TEAD1_HUMAN]	0.910	0.011
	Isoform 2 of Rho GTPase-activating protein 26 OS=Homo sapiens GN=ARHGAP26 - [RHG26_HUMAN]	0.912	0.002

Q9Y3D3	28S ribosomal protein S16, mitochondrial OS=Homo sapiens GN=MRPS16 PE=1 SV=1 - [RT16_HUMAN]	0.912	0.035
Q6UB35	Monofunctional C1-tetrahydrofolate synthase, mitochondrial OS=Homo sapiens GN=MTHFD1L PE=1 SV=1 - [C1TM_HUMAN]	0.913	0.037
P17509	Homeobox protein Hox-B6 OS=Homo sapiens GN=HOXB6 PE=1 SV=4 - [HXB6_HUMAN]	0.913	0.021
O96000	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 10 OS=Homo sapiens GN=NDUFB10 PE=1 SV=3 - [NDUBA_HUMAN]	0.914	0.026
O00220	Tumor necrosis factor receptor superfamily member 10A OS=Homo sapiens GN=TNFRSF10A PE=1 SV=3 - [TR10A_HUMAN]	0.914	0.018
Q9HCU5	Prolactin regulatory element-binding protein OS=Homo sapiens GN=PREB PE=1 SV=2 - [PREB_HUMAN]	0.915	0.020
P62820	Ras-related protein Rab-1A OS=Homo sapiens GN=RAB1A PE=1 SV=3 - [RAB1A_HUMAN]	0.918	0.032
P51148	Ras-related protein Rab-5C OS=Homo sapiens GN=RAB5C PE=1 SV=2 - [RAB5C_HUMAN]	0.918	0.004
P61026	Ras-related protein Rab-10 OS=Homo sapiens GN=RAB10 PE=1 SV=1 - [RAB10_HUMAN]	0.920	0.048
O15347	High mobility group protein B3 OS=Homo sapiens GN=HMGB3 PE=1 SV=4 - [HMGB3_HUMAN]	0.920	0.023
Q13153	Serine/threonine-protein kinase PAK 1 OS=Homo sapiens GN=PAK1 PE=1 SV=2 - [PAK1_HUMAN]	0.920	0.050
Q00577	Transcriptional activator protein Pur-alpha OS=Homo sapiens GN=PURA PE=1 SV=2 - [PURA_HUMAN]	0.921	0.001
O60885	Bromodomain-containing protein 4 OS=Homo sapiens GN=BRD4 PE=1 SV=2 - [BRD4_HUMAN]	0.921	0.026
Q13405	39S ribosomal protein L49, mitochondrial OS=Homo sapiens GN=MRPL49 PE=1 SV=1 - [RM49_HUMAN]	0.921	0.039
P25705	ATP synthase subunit alpha, mitochondrial OS=Homo sapiens GN=ATP5A1 PE=1 SV=1 - [ATPA_HUMAN]	0.921	0.043
P49411	Elongation factor Tu, mitochondrial OS=Homo sapiens GN=TUFM PE=1 SV=2 - [EFTU_HUMAN]	0.922	0.025
Q93099	Homogentisate 1,2-dioxygenase OS=Homo sapiens GN=HGD PE=1 SV=2 - [HGD_HUMAN]	0.923	0.005
P84103	Serine/arginine-rich splicing factor 3 OS=Homo sapiens GN=SRSF3 PE=1 SV=1 - [SRSF3_HUMAN]	0.923	0.045
Q9Y276	Mitochondrial chaperone BCS1 OS=Homo sapiens GN=BCS1L PE=1 SV=1 - [BCS1_HUMAN]	0.925	0.011
Q96EK6	Glucosamine 6-phosphate N-acetyltransferase OS=Homo sapiens GN=GNPNAT1 PE=1 SV=1 - [GNA1_HUMAN]	0.926	0.043
Q14671	Pumilio homolog 1 OS=Homo sapiens GN=PUM1 PE=1 SV=3 - [PUM1_HUMAN]	0.927	0.048
Q460N5-4	Isoform 4 of Poly [ADP-ribose] polymerase 14 OS=Homo sapiens GN=PARP14 - [PAR14_HUMAN]	0.927	0.019
O95573	Long-chain-fatty-acid--CoA ligase 3 OS=Homo sapiens GN=ACSL3 PE=1 SV=3 - [ACSL3_HUMAN]	0.928	0.008
Q9NRF8	CTP synthase 2 OS=Homo sapiens GN=CTPS2 PE=1 SV=1 - [PYRG2_HUMAN]	0.929	0.011
P26447	Protein S100-A4 OS=Homo sapiens GN=S100A4 PE=1 SV=1 - [S10A4_HUMAN]	0.929	0.034
Q9H9P8	L-2-hydroxyglutarate dehydrogenase, mitochondrial OS=Homo sapiens GN=L2HGDH PE=1 SV=3 - [L2HDH_HUMAN]	0.929	0.013
Q16706	Alpha-mannosidase 2 OS=Homo sapiens GN=MAN2A1 PE=1 SV=2 - [MA2A1_HUMAN]	0.930	0.044
Q8TDJ6	DmX-like protein 2 OS=Homo sapiens GN=DMXL2 PE=1 SV=2 - [DMXL2_HUMAN]	0.932	0.041
P12532	Creatine kinase U-type, mitochondrial OS=Homo sapiens GN=CKMT1A PE=1 SV=1 - [KCRU_HUMAN]	0.932	0.031

P55199	RNA polymerase II elongation factor ELL OS=Homo sapiens GN=ELL PE=1 SV=1 - [ELL_HUMAN]	0.933	0.028
P57737-2	Isoform 2 of Coronin-7 OS=Homo sapiens GN=CORO7 - [CORO7_HUMAN]	0.933	0.023
Q96EP5-2	Isoform 2 of DAZ-associated protein 1 OS=Homo sapiens GN=DAZAP1 - [DAZP1_HUMAN]	0.934	0.033
Q9H0V9	VIP36-like protein OS=Homo sapiens GN=LMAN2L PE=1 SV=1 - [LMA2L_HUMAN]	0.935	0.046
Q9UHG3	Prenylcysteine oxidase 1 OS=Homo sapiens GN=PCYOX1 PE=1 SV=3 - [PCYOX_HUMAN]	0.936	0.022
P10746	Uroporphyrinogen-III synthase OS=Homo sapiens GN=UROS PE=1 SV=1 - [HEM4_HUMAN]	0.936	0.006
Q8NFW8	N-acylneuraminate cytidyltransferase OS=Homo sapiens GN=CMAS PE=1 SV=2 - [NEUA_HUMAN]	0.939	0.046
P61088	Ubiquitin-conjugating enzyme E2 N OS=Homo sapiens GN=UBE2N PE=1 SV=1 - [UBE2N_HUMAN]	0.939	0.016
Q96BQ5	Coiled-coil domain-containing protein 127 OS=Homo sapiens GN=CCDC127 PE=1 SV=1 - [CC127_HUMAN]	0.940	0.012
Q92766	Ras-responsive element-binding protein 1 OS=Homo sapiens GN=RREB1 PE=1 SV=3 - [RREB1_HUMAN]	0.941	0.039
Q9NXG2	THUMP domain-containing protein 1 OS=Homo sapiens GN=THUMPD1 PE=1 SV=2 - [THUM1_HUMAN]	0.941	0.020
O95785	Protein Wiz OS=Homo sapiens GN=WIZ PE=1 SV=2 - [WIZ_HUMAN]	0.942	0.014
P00505	Aspartate aminotransferase, mitochondrial OS=Homo sapiens GN=GOT2 PE=1 SV=3 - [AATM_HUMAN]	0.943	0.008
P10809	60 kDa heat shock protein, mitochondrial OS=Homo sapiens GN=HSPD1 PE=1 SV=2 - [CH60_HUMAN]	0.944	0.010
Q00839-2	Isoform Short of Heterogeneous nuclear ribonucleoprotein U OS=Homo sapiens GN=HNRNPU - [HNRPU_HUMAN]	0.944	0.011
P61160	Actin-related protein 2 OS=Homo sapiens GN=ACTR2 PE=1 SV=1 - [ARP2_HUMAN]	0.944	0.026
P15153	Ras-related C3 botulinum toxin substrate 2 OS=Homo sapiens GN=RAC2 PE=1 SV=1 - [RAC2_HUMAN]	0.946	0.033
P36955	Pigment epithelium-derived factor OS=Homo sapiens GN=SERPINF1 PE=1 SV=4 - [PEDF_HUMAN]	0.946	0.040
Q96EN8	Molybdenum cofactor sulfurase OS=Homo sapiens GN=MOCOS PE=1 SV=2 - [MOCOS_HUMAN]	0.947	0.025
Q8N9N7	Leucine-rich repeat-containing protein 57 OS=Homo sapiens GN=LRRC57 PE=1 SV=1 - [LRC57_HUMAN]	0.948	0.044
P82914	28S ribosomal protein S15, mitochondrial OS=Homo sapiens GN=MRPS15 PE=1 SV=1 - [RT15_HUMAN]	0.949	0.027
Q15059	Bromodomain-containing protein 3 OS=Homo sapiens GN=BRD3 PE=1 SV=1 - [BRD3_HUMAN]	0.949	0.028
Q8TEM1	Nuclear pore membrane glycoprotein 210 OS=Homo sapiens GN=NUP210 PE=1 SV=3 - [PO210_HUMAN]	0.949	0.016
O43674	NADH dehydrogenase [ubiquinone] 1 beta subcomplex subunit 5, mitochondrial OS=Homo sapiens GN=NDUFB5 PE=1 SV=1 - [NDUB5_HUMAN]	0.950	0.014
Q15061	WD repeat-containing protein 43 OS=Homo sapiens GN=WDR43 PE=1 SV=3 - [WDR43_HUMAN]	0.950	0.040
P35270	Sepiapterin reductase OS=Homo sapiens GN=SPR PE=1 SV=1 - [SPRE_HUMAN]	0.951	0.037
P23786	Carnitine O-palmitoyltransferase 2, mitochondrial OS=Homo sapiens GN=CPT2 PE=1 SV=2 - [CPT2_HUMAN]	0.954	0.027
Q9UNM6	26S proteasome non-ATPase regulatory subunit 13 OS=Homo sapiens GN=PSMD13 PE=1 SV=2 - [PSD13_HUMAN]	0.956	0.002
O75934	Pre-mRNA-splicing factor SPF27 OS=Homo sapiens GN=BCAS2 PE=1 SV=1 - [SPF27_HUMAN]	0.961	0.046

Q13232	Nucleoside diphosphate kinase 3 OS=Homo sapiens GN=NME3 PE=1 SV=2 - [NDK3_HUMAN]	0.962	0.002
Q8IW92	Beta-galactosidase-1-like protein 2 OS=Homo sapiens GN=GLB1L2 PE=2 SV=1 - [GLBL2_HUMAN]	0.962	0.035
Q5JTZ9	Alanine--tRNA ligase, mitochondrial OS=Homo sapiens GN=AARS2 PE=1 SV=1 - [SYAM_HUMAN]	0.964	0.004
Q12904	Aminoacyl tRNA synthase complex-interacting multifunctional protein 1 OS=Homo sapiens GN=AIMP1 PE=1 SV=2 - [AIMP1_HUMAN]	0.969	0.011
Q9H4B7	Tubulin beta-1 chain OS=Homo sapiens GN=TUBB1 PE=1 SV=1 - [TBB1_HUMAN]	0.971	0.011
Q8N1G0	Zinc finger protein 687 OS=Homo sapiens GN=ZNF687 PE=1 SV=1 - [ZN687_HUMAN]	0.974	0.012
O00139-2	Isoform 2 of Kinesin-like protein KIF2A OS=Homo sapiens GN=KIF2A - [KIF2A_HUMAN]	0.978	0.025
Q96S66	Chloride channel CLIC-like protein 1 OS=Homo sapiens GN=CLCC1 PE=1 SV=1 - [CLCC1_HUMAN]	0.981	0.046
Q8TB36	Ganglioside-induced differentiation-associated protein 1 OS=Homo sapiens GN=GDAP1 PE=1 SV=3 - [GDAP1_HUMAN]	0.982	0.032
Q9Y618-5	Isoform 5 of Nuclear receptor corepressor 2 OS=Homo sapiens GN=NCOR2 - [NCOR2_HUMAN]	0.986	0.016
Q15075	Early endosome antigen 1 OS=Homo sapiens GN=EEA1 PE=1 SV=2 - [EEA1_HUMAN]	0.994	0.046
O94822	E3 ubiquitin-protein ligase listerin OS=Homo sapiens GN=LTN1 PE=1 SV=6 - [LTN1_HUMAN]	1.013	0.031
Q9Y2J2-2	Isoform B of Band 4.1-like protein 3 OS=Homo sapiens GN=EPB41L3 - [E41L3_HUMAN]	1.014	0.020
Q15006	Tetratricopeptide repeat protein 35 OS=Homo sapiens GN=TTC35 PE=1 SV=1 - [TTC35_HUMAN]	1.018	0.029
O95376	E3 ubiquitin-protein ligase ARIH2 OS=Homo sapiens GN=ARIH2 PE=1 SV=1 - [ARI2_HUMAN]	1.024	0.002
Q03468	DNA excision repair protein ERCC-6 OS=Homo sapiens GN=ERCC6 PE=1 SV=1 - [ERCC6_HUMAN]	1.024	0.020
P54289-4	Isoform 4 of Voltage-dependent calcium channel subunit alpha-2/delta-1 OS=Homo sapiens GN=CACNA2D1 - [CA2D1_HUMAN]	1.030	0.032
O00462	Beta-mannosidase OS=Homo sapiens GN=MANBA PE=1 SV=3 - [MANBA_HUMAN]	1.031	0.047
Q96CW1-2	Isoform 2 of AP-2 complex subunit mu OS=Homo sapiens GN=AP2M1 - [AP2M1_HUMAN]	1.033	0.050
O00233-2	Isoform p27-S of 26S proteasome non-ATPase regulatory subunit 9 OS=Homo sapiens GN=PSMD9 - [PSMD9_HUMAN]	1.037	0.010
O00232	26S proteasome non-ATPase regulatory subunit 12 OS=Homo sapiens GN=PSMD12 PE=1 SV=3 - [PSD12_HUMAN]	1.038	0.009
Q9H773	dCTP pyrophosphatase 1 OS=Homo sapiens GN=DCTPP1 PE=1 SV=1 - [DCTP1_HUMAN]	1.040	0.026
Q96CN9	GRIP and coiled-coil domain-containing protein 1 OS=Homo sapiens GN=GCC1 PE=1 SV=1 - [GCC1_HUMAN]	1.044	0.020
Q9BV73-2	Isoform 2 of Centrosome-associated protein CEP250 OS=Homo sapiens GN=CEP250 - [CP250_HUMAN]	1.048	0.024
O60343-2	Isoform 2 of TBC1 domain family member 4 OS=Homo sapiens GN=TBC1D4 - [TBCD4_HUMAN]	1.062	0.030
Q9Y294	Histone chaperone ASF1A OS=Homo sapiens GN=ASF1A PE=1 SV=1 - [ASF1A_HUMAN]	1.066	0.035
Q53H54	tRNA wybutosine-synthesizing protein 2 homolog OS=Homo sapiens GN=TRMT12 PE=1 SV=1 - [TYW2_HUMAN]	1.067	0.024
Q96QU8	Exportin-6 OS=Homo sapiens GN=XPO6 PE=1 SV=1 - [XPO6_HUMAN]	1.069	0.008
Q6UX53	Methyltransferase-like protein 7B OS=Homo sapiens GN=METTTL7B PE=2 SV=2 - [MET7B_HUMAN]	1.069	0.049

Q13325	Interferon-induced protein with tetratricopeptide repeats 5 OS=Homo sapiens GN=IFIT5 PE=1 SV=1 - [IFIT5_HUMAN]	1.073	0.036
Q9H0F6-2	Isoform 2 of Sharpin OS=Homo sapiens GN=SHARPIN - [SHRPN_HUMAN]	1.073	0.025
Q5T3U5-2	Isoform 2 of Multidrug resistance-associated protein 7 OS=Homo sapiens GN=ABCC10 - [MRP7_HUMAN]	1.073	0.030
Q9BRX2	Protein pelota homolog OS=Homo sapiens GN=PELO PE=1 SV=2 - [PELO_HUMAN]	1.075	0.035
P19784	Casein kinase II subunit alpha' OS=Homo sapiens GN=CSNK2A2 PE=1 SV=1 - [CSK22_HUMAN]	1.077	0.044
Q7L1T6	Cytochrome b5 reductase 4 OS=Homo sapiens GN=CYB5R4 PE=1 SV=1 - [NB5R4_HUMAN]	1.077	0.031
Q96LA8-2	Isoform 2 of Protein arginine N-methyltransferase 6 OS=Homo sapiens GN=PRMT6 - [ANM6_HUMAN]	1.079	0.031
Q8TCY9-3	Isoform 3 of Up-regulator of cell proliferation OS=Homo sapiens GN=URGCP - [URGCP_HUMAN]	1.079	0.012
O00151	PDZ and LIM domain protein 1 OS=Homo sapiens GN=PDLIM1 PE=1 SV=4 - [PDL1_HUMAN]	1.079	0.018
Q99543	DnaJ homolog subfamily C member 2 OS=Homo sapiens GN=DNAJC2 PE=1 SV=4 - [DNJC2_HUMAN]	1.079	0.031
Q8N5F7	NF-kappa-B-activating protein OS=Homo sapiens GN=NKAP PE=1 SV=1 - [NKAP_HUMAN]	1.082	0.029
Q7L5Y1	Mitochondrial enolase superfamily member 1 OS=Homo sapiens GN=ENOSF1 PE=1 SV=1 - [ENOF1_HUMAN]	1.082	0.035
Q8N9N5	Protein BANP OS=Homo sapiens GN=BANP PE=1 SV=3 - [BANP_HUMAN]	1.084	0.050
Q9ULX9	Transcription factor MafF OS=Homo sapiens GN=MAFF PE=1 SV=2 - [MAFF_HUMAN]	1.084	0.047
Q9BTW9	Tubulin-specific chaperone D OS=Homo sapiens GN=TBCD PE=1 SV=2 - [TBCD_HUMAN]	1.085	0.040
Q9UPQ9-1	Isoform 2 of Trinucleotide repeat-containing gene 6B protein OS=Homo sapiens GN=TNRC6B - [TNR6B_HUMAN]	1.086	0.039
Q8TF01	Arginine/serine-rich protein PNISR OS=Homo sapiens GN=PNISR PE=1 SV=2 - [PNISR_HUMAN]	1.087	0.012
Q9UPM8	AP-4 complex subunit epsilon-1 OS=Homo sapiens GN=AP4E1 PE=1 SV=2 - [AP4E1_HUMAN]	1.090	0.022
Q86WR0	Coiled-coil domain-containing protein 25 OS=Homo sapiens GN=CCDC25 PE=1 SV=2 - [CCD25_HUMAN]	1.090	0.026
O15294-3	Isoform 1 of UDP-N-acetylglucosamine--peptide N-acetylglucosaminyltransferase 110 kDa subunit OS=Homo sapiens GN=OGT - [OGT1_HUMAN]	1.092	0.050
Q9BUF5	Tubulin beta-6 chain OS=Homo sapiens GN=TUBB6 PE=1 SV=1 - [TBB6_HUMAN]	1.098	0.030
Q8WXW3	Progesterone-induced-blocking factor 1 OS=Homo sapiens GN=PIBF1 PE=2 SV=2 - [PIBF1_HUMAN]	1.098	0.049
P15923-2	Isoform E47 of Transcription factor E2-alpha OS=Homo sapiens GN=TCF3 - [TFE2_HUMAN]	1.099	0.048
Q8TAC2	Josephin-2 OS=Homo sapiens GN=JOSD2 PE=1 SV=1 - [JOS2_HUMAN]	1.102	0.025
O95619	YEATS domain-containing protein 4 OS=Homo sapiens GN=YEATS4 PE=1 SV=1 - [YETS4_HUMAN]	1.104	0.012
Q9P2G3-2	Isoform 2 of Kelch-like protein 14 OS=Homo sapiens GN=KLHL14 - [KLH14_HUMAN]	1.104	0.042
Q9H5V9	UPF0428 protein CXorf56 OS=Homo sapiens GN=CXorf56 PE=1 SV=1 - [CX056_HUMAN]	1.110	0.019
Q9BSK4	Protein fem-1 homolog A OS=Homo sapiens GN=FEM1A PE=1 SV=1 - [FEM1A_HUMAN]	1.113	0.011
Q6NXT6	Transmembrane anterior posterior transformation protein 1 homolog OS=Homo sapiens GN=TAPT1 PE=1 SV=1 - [TAPT1_HUMAN]	1.113	0.047

O43353-2	Isoform 2 of Receptor-interacting serine/threonine-protein kinase 2 OS=Homo sapiens GN=RIPK2 - [RIPK2_HUMAN]	1.117	0.033
Q96IK1	Biorientation of chromosomes in cell division protein 1 OS=Homo sapiens GN=BOD1 PE=1 SV=2 - [BOD1_HUMAN]	1.118	0.034
Q8IXJ9	Putative Polycomb group protein ASXL1 OS=Homo sapiens GN=ASXL1 PE=1 SV=3 - [ASXL1_HUMAN]	1.121	0.004
Q6NTF9	Rhomboid domain-containing protein 2 OS=Homo sapiens GN=RHBDD2 PE=2 SV=2 - [RHBDD2_HUMAN]	1.124	0.017
O94964-2	Isoform 2 of Uncharacterized protein KIAA0889 OS=Homo sapiens GN=KIAA0889 - [K0889_HUMAN]	1.126	0.029
P41214	Eukaryotic translation initiation factor 2D OS=Homo sapiens GN=EIF2D PE=1 SV=3 - [EIF2D_HUMAN]	1.126	0.019
Q06547-4	Isoform 4 of GA-binding protein subunit beta-1 OS=Homo sapiens GN=GABPB1 - [GABP1_HUMAN]	1.127	0.024
Q9Y508-2	Isoform 2 of RING finger protein 114 OS=Homo sapiens GN=RNF114 - [RN114_HUMAN]	1.128	0.049
Q9UK76-3	Isoform 3 of Hematological and neurological expressed 1 protein OS=Homo sapiens GN=HN1 - [HN1_HUMAN]	1.129	0.020
Q5VTB9	E3 ubiquitin-protein ligase RNF220 OS=Homo sapiens GN=RNF220 PE=1 SV=1 - [RN220_HUMAN]	1.130	0.047
Q9NQX7-2	Isoform 2 of Integral membrane protein 2C OS=Homo sapiens GN=ITM2C - [ITM2C_HUMAN]	1.138	0.010
Q92793	CREB-binding protein OS=Homo sapiens GN=CREBBP PE=1 SV=3 - [CBP_HUMAN]	1.139	0.019
Q9Y411-2	Isoform 2 of Myosin-Va OS=Homo sapiens GN=MYO5A - [MYO5A_HUMAN]	1.140	0.028
Q9NRG0	Chromatin accessibility complex protein 1 OS=Homo sapiens GN=CHRAC1 PE=1 SV=1 - [CHRC1_HUMAN]	1.141	0.045
Q1MSJ5-1	Isoform 1 of Centrosome and spindle pole-associated protein 1 OS=Homo sapiens GN=CSPP1 - [CSPP1_HUMAN]	1.142	0.044
Q49AR2	UPF0489 protein C5orf22 OS=Homo sapiens GN=C5orf22 PE=1 SV=2 - [CE022_HUMAN]	1.146	0.035
Q96PE2	Rho guanine nucleotide exchange factor 17 OS=Homo sapiens GN=ARHGEF17 PE=1 SV=1 - [ARHGH_HUMAN]	1.147	0.012
Q8NC54	Keratinocyte-associated transmembrane protein 2 OS=Homo sapiens GN=KCT2 PE=2 SV=2 - [KCT2_HUMAN]	1.147	0.010
Q6NXT1-2	Isoform 2 of Ankyrin repeat domain-containing protein 54 OS=Homo sapiens GN=ANKRD54 - [ANR54_HUMAN]	1.147	0.047
Q9BY67	Cell adhesion molecule 1 OS=Homo sapiens GN=CADM1 PE=1 SV=2 - [CADM1_HUMAN]	1.150	0.004
Q06481	Amyloid-like protein 2 OS=Homo sapiens GN=APLP2 PE=1 SV=2 - [APLP2_HUMAN]	1.154	0.027
O15078	Centrosomal protein of 290 kDa OS=Homo sapiens GN=CEP290 PE=1 SV=2 - [CE290_HUMAN]	1.158	0.033
Q6UWI4	Protein shisa-2 homolog OS=Homo sapiens GN=SHISA2 PE=1 SV=1 - [SHSA2_HUMAN]	1.159	0.031
Q6P6C2-2	Isoform 2 of Probable alpha-ketoglutarate-dependent dioxygenase ABH5 OS=Homo sapiens GN=ALKBH5 - [ALKB5_HUMAN]	1.159	0.034
Q15025-2	Isoform 2 of TNFAIP3-interacting protein 1 OS=Homo sapiens GN=TNIP1 - [TNIP1_HUMAN]	1.159	0.006
Q8TA86	Retinitis pigmentosa 9 protein OS=Homo sapiens GN=RP9 PE=1 SV=2 - [RP9_HUMAN]	1.168	0.027
Q562F6-2	Isoform 2 of Shugoshin-like 2 OS=Homo sapiens GN=SGOL2 - [SGOL2_HUMAN]	1.168	0.028
O43164-2	Isoform 2 of E3 ubiquitin-protein ligase Praja-2 OS=Homo sapiens GN=PJA2 - [PJA2_HUMAN]	1.172	0.015
O43422-2	Isoform Short of 52 kDa repressor of the inhibitor of the protein kinase OS=Homo sapiens GN=PRKRIR - [P52K_HUMAN]	1.174	0.006

Q9Y679-3	Isoform 3 of Ancient ubiquitous protein 1 OS=Homo sapiens GN=AUP1 - [AUP1_HUMAN]	1.177	0.006
Q8NCU7	C2 calcium-dependent domain-containing protein 4A OS=Homo sapiens GN=C2CD4A PE=2 SV=2 - [C2C4A_HUMAN]	1.178	0.022
Q14679	Tubulin polyglutamylase TTLL4 OS=Homo sapiens GN=TTLL4 PE=1 SV=2 - [TTLL4_HUMAN]	1.181	0.007
Q9BQ24	Zinc finger FYVE domain-containing protein 21 OS=Homo sapiens GN=ZFYVE21 PE=1 SV=1 - [ZFY21_HUMAN]	1.184	0.048
O43865	Putative adenosylhomocysteinase 2 OS=Homo sapiens GN=AHCYL1 PE=1 SV=2 - [SAHH2_HUMAN]	1.189	0.007
Q92786	Prospero homeobox protein 1 OS=Homo sapiens GN=PROX1 PE=1 SV=2 - [PROX1_HUMAN]	1.190	0.034
P02760	Protein AMBP OS=Homo sapiens GN=AMBP PE=1 SV=1 - [AMBP_HUMAN]	1.196	0.029
Q9H1A3-2	Isoform 2 of Methyltransferase-like protein 9 OS=Homo sapiens GN=METTLL9 - [METL9_HUMAN]	1.198	0.006
Q9Y6M4-3	Isoform 3 of Casein kinase I isoform gamma-3 OS=Homo sapiens GN=CSNK1G3 - [KC1G3_HUMAN]	1.199	0.032
P12259	Coagulation factor V OS=Homo sapiens GN=F5 PE=1 SV=4 - [FA5_HUMAN]	1.203	0.033
Q9NQS1	Cell death regulator Aven OS=Homo sapiens GN=AVEN PE=1 SV=1 - [AVEN_HUMAN]	1.204	0.018
Q9H8W3	Protein FAM204A OS=Homo sapiens GN=FAM204A PE=2 SV=1 - [F204A_HUMAN]	1.205	0.021
Q9BXS4	Transmembrane protein 59 OS=Homo sapiens GN=TMEM59 PE=1 SV=1 - [TMM59_HUMAN]	1.211	0.020
Q9H9L7	Akirin-1 OS=Homo sapiens GN=AKIRIN1 PE=1 SV=1 - [AKIR1_HUMAN]	1.212	0.043
Q13137	Calcium-binding and coiled-coil domain-containing protein 2 OS=Homo sapiens GN=CALCOCO2 PE=1 SV=1 - [CACO2_HUMAN]	1.215	0.040
O96020-2	Isoform Short of G1/S-specific cyclin-E2 OS=Homo sapiens GN=CCNE2 - [CCNE2_HUMAN]	1.217	0.025
Q9BVV6-2	Isoform 2 of Uncharacterized protein KIAA0586 OS=Homo sapiens GN=KIAA0586 - [K0586_HUMAN]	1.230	0.040
Q15345	Leucine-rich repeat-containing protein 41 OS=Homo sapiens GN=LRRC41 PE=1 SV=3 - [LRC41_HUMAN]	1.231	0.042
Q96S15-2	Isoform 2 of WD repeat-containing protein 24 OS=Homo sapiens GN=WDR24 - [WDR24_HUMAN]	1.236	0.004
Q96RV3-2	Isoform 2 of Pecanex-like protein 1 OS=Homo sapiens GN=PCNX - [PCX1_HUMAN]	1.242	0.042
Q9UHQ1-3	Isoform 3 of Nuclear prelamin A recognition factor OS=Homo sapiens GN=NARF - [NARF_HUMAN]	1.244	0.036
Q86VP1-3	Isoform 3 of Tax1-binding protein 1 OS=Homo sapiens GN=TAX1BP1 - [TAXB1_HUMAN]	1.247	0.026
Q92823-3	Isoform 3 of Neuronal cell adhesion molecule OS=Homo sapiens GN=NRCAM - [NRCAM_HUMAN]	1.253	0.043
P28356	Homeobox protein Hox-D9 OS=Homo sapiens GN=HOXD9 PE=2 SV=5 - [HXD9_HUMAN]	1.278	0.024
Q9BQ16-2	Isoform 2 of Testican-3 OS=Homo sapiens GN=SPOCK3 - [TICN3_HUMAN]	1.294	0.036
Q9NR48-2	Isoform 2 of Histone-lysine N-methyltransferase ASH1L OS=Homo sapiens GN=ASH1L - [ASH1L_HUMAN]	1.300	0.031
Q9NWL6	Asparagine synthetase domain-containing protein 1 OS=Homo sapiens GN=ASNSD1 PE=2 SV=2 - [ASND1_HUMAN]	1.342	0.044
O60930	Ribonuclease H1 OS=Homo sapiens GN=RNASEH1 PE=1 SV=2 - [RNH1_HUMAN]	1.347	0.027
Q9Y625	Glypican-6 OS=Homo sapiens GN=GPC6 PE=2 SV=1 - [GPC6_HUMAN]	1.378	0.020

P07949-2	Isoform 2 of Proto-oncogene tyrosine-protein kinase receptor Ret OS=Homo sapiens GN=RET - [RET_HUMAN]	1.465	0.009
Q08043	Alpha-actinin-3 OS=Homo sapiens GN=ACTN3 PE=1 SV=2 - [ACTN3_HUMAN]	1.735	0.032

Each protein expression ratio was normalized to its non-treated control which was assigned arbitrary value 1

Supplementary Table S4. Significantly up- or down-regulated proteins after lanreotide treatment.

Accession number	Gene symbol	Protein description	Expression vs. Control	P - value
HC45 cells - 2 hours lanreotide				
Q9H147	<i>DNTTIP1</i>	Deoxynucleotidyltransferase terminal-interacting protein 1	0.58	0.042
P62714	<i>PPP2CB</i>	Serine/threonine-protein phosphatase 2A catalytic subunit beta isoform	0.59	0.002
P25054-2	<i>APC</i> *	Isoform Short of Adenomatous polyposis coli protein	1.36	0.029
O15145	<i>ARPC3</i>	Actin-related protein 2/3 complex subunit 3	1.47	0.013
Q8N8U9	<i>BMPER</i> *	BMP-binding endothelial regulator protein	1.52	0.020
HC45 cells - 6 hours lanreotide				
Q13015	<i>MLLT11</i>	Protein AF1q	0.54	0.053
P13645	<i>KRT10</i>	Keratin, type I cytoskeletal 10	0.60	0.035
Q15051-3	<i>IQCB1</i>	Isoform 3 of IQ calmodulin-binding motif-containing protein 1	0.61	0.019
P13647	<i>KRT5</i>	Keratin, type II cytoskeletal 5	0.63	0.007
Q9H147	<i>DNTTIP1</i>	Deoxynucleotidyltransferase terminal-interacting protein 1	0.67	0.051
Q9BXV9	<i>C14orf142</i> *	Uncharacterized protein C14orf142	0.67	0.013
Q4ZIN3-2	<i>C19orf6</i>	Isoform 2 of Membralin	0.68	0.027
Q9H3K2	<i>GHITM</i>	Growth hormone-inducible transmembrane protein	1.41	0.046
Q9Y2C4	<i>EXOG</i>	Nuclease EXOG, mitochondrial	1.45	0.025
O75880	<i>SCO1</i>	Protein SCO1 homolog, mitochondrial	1.45	0.034
O15235	<i>MRPS12</i>	28S ribosomal protein S12, mitochondrial	1.53	0.028
O75323	<i>GBAS</i>	Protein NipSnap homolog 2	1.57	0.036
P62877	<i>RBX1</i>	E3 ubiquitin-protein ligase RBX1	1.60	0.039
P62273	<i>RPS29</i>	40S ribosomal protein S29	1.58	0.054
Q86VY9	<i>TMEM200A</i>	Transmembrane protein 200A	1.61	0.041
Q9Y3E0	<i>GOLT1B</i>	Vesicle transport protein GOT1B	2.00	0.051
Q3B7S5	<i>C18orf62</i> *	<i>SMIM21</i> , Small integral membrane protein 21	2.11	0.012
Q9BYX7	<i>POTEKP</i>	Putative beta-actin-like protein 3	2.21	0.048

HC45 cells - 48 hours lanreotide

P13473	<i>LAMP2</i>	Lysosome-associated membrane glycoprotein 2	0.40	0.035
P53539	<i>FOSB</i>	Protein fosB	0.46	0.002
P60903	<i>S100A10</i>	Protein S100-A10	0.54	0.043
Q96SL4	<i>GPX7</i>	Glutathione peroxidase 7	0.53	0.051
Q04695	<i>KRT17</i>	Keratin, type I cytoskeletal 17	0.54	0.009
Q96J01	<i>THOC3</i>	THO complex subunit 3	0.55	0.023
P17096-2	<i>HMGA1</i>	Isoform HMG-Y of High mobility group protein HMG-I/HMG-Y	0.57	0.054
P0CG34	<i>TMSB15A</i>	Thymosin beta-15A	0.57	0.001
O95243-3	<i>MBD4</i>	Isoform 3 of Methyl-CpG-binding domain protein 4	0.59	0.014
O43852-2	<i>CALU</i>	Isoform 2 of Calumenin	0.62	0.054
Q16678	<i>CYP1B1</i>	Cytochrome P450 1B1	0.63	0.033
Q9H8M7	<i>FAM188A</i>	Protein FAM188A	0.63	0.031
O60573	<i>EIF4E2</i>	Eukaryotic translation initiation factor 4E type 2	0.63	0.053
P05114	<i>HMG1</i>	Non-histone chromosomal protein HMG-14	0.64	0.042
O95218-2	<i>ZRANB2</i>	Isoform 2 of Zinc finger Ran-binding domain-containing protein 2	0.64	0.042
Q16831	<i>UPP1</i>	Uridine phosphorylase 1	0.64	0.006
Q15005	<i>SPCS2</i>	Signal peptidase complex subunit 2	0.66	0.046
Q9Y4P8-3	<i>WIPI2</i>	Isoform 3 of WD repeat domain phosphoinositide-interacting protein 2	0.66	0.050
Q6UWJ1-3	<i>TMCO3</i>	Isoform 3 of Transmembrane and coiled-coil domain-containing protein 3	0.67	0.036
P09493-3	<i>TPM1</i>	Isoform 3 of Tropomyosin alpha-1 chain	0.67	0.048
Q9UIL1-4	<i>SCOC</i>	Isoform 4 of Short coiled-coil protein	1.32	0.020
P98172	<i>EFNB1</i>	Ephrin-B1	1.32	0.047
Q86U90	<i>YRDC</i>	YrdC domain-containing protein, mitochondrial	1.34	0.044
O15392-7	<i>BIRC5</i> *	Survivin, Isoform 7 of Baculoviral IAP repeat-containing protein 5	1.36	0.012
Q9NR33	<i>POLE4</i>	DNA polymerase epsilon subunit 4	1.38	0.044
Q9H8G2	<i>C9orf82</i>	Uncharacterized protein C9orf82	1.38	0.011
P14316	<i>IRF2</i>	Interferon regulatory factor 2	1.40	0.036
Q92529-2	<i>SHC3</i>	Isoform p52 of SHC-transforming protein 3	1.37	0.031
Q5FWF5-2	<i>ESCO1</i>	Isoform 2 of N-acetyltransferase ESCO1	1.40	0.044
O00635	<i>TRIM38</i>	Tripartite motif-containing protein 38	1.42	0.045

Q8NFH3	<i>NUP43</i>	Nucleoporin Nup43	1.45	0.035
P11166	<i>SLC2A1</i>	Solute carrier family 2, facilitated glucose transporter member 1 O	1.50	0.003
Q99943	<i>AGPAT1</i>	1-acyl-sn-glycerol-3-phosphate acyltransferase alpha	1.53	0.049
O14735	<i>CDIPT</i>	CDP-diacylglycerol--inositol 3-phosphatidyltransferase	1.54	0.034
Q7KYR7	<i>BTN2A1</i>	Butyrophilin subfamily 2 member A1	1.54	0.053
O15235	<i>MRPS12</i>	28S ribosomal protein S12, mitochondrial	1.56	0.027
P06241-3	<i>FYN</i> *	Isoform 3 of Tyrosine-protein kinase Fyn	1.63	0.048
P42766	<i>RPL35</i>	60S ribosomal protein L35	1.66	0.019
Q9BZM5	<i>ULBP2</i>	NKG2D ligand 2	1.70	0.003
Q9Y3E7-4	<i>CHMP3</i>	Isoform 4 of Charged multivesicular body protein 3	1.72	0.043
Q15819	<i>UBE2V2</i>	Ubiquitin-conjugating enzyme E2 variant 2	1.86	0.046
O15145	<i>ARPC3</i>	Actin-related protein 2/3 complex subunit 3	2.50	0.045

H727 cells - 2 hours lanreotide

P84243	<i>H3F3A</i>	Histone H3.3	0.52	0.013
Q71DI3	<i>HIST2H3A</i>	Histone H3.2	0.55	0.052
Q9UBC1-2	<i>NFKBIL1</i>	Isoform 2 of NF-kappa-B inhibitor-like protein 1	0.61	0.052
Q6DD87	<i>ZNF787</i>	Zinc finger protein 787	0.63	0.038
P53671-2	<i>LIMK2</i>	Isoform LIMK2b of LIM domain kinase 2	0.72	0.042
Q9UJH8	<i>METRIN</i>	Meteorin	0.76	0.030
Q8IZQ1-2	<i>WDFY3</i>	Isoform 2 of WD repeat and FYVE domain-containing protein 3	0.78	0.048
Q9H7E9	<i>C8orf33</i>	UPF0488 protein C8orf33	1.22	0.044
P08670	<i>VIM</i>	Vimentin	1.23	0.027
Q9UMW8	<i>USP18</i>	Ubl carboxyl-terminal hydrolase 18	1.22	0.053
Q14296	<i>FASTK</i>	Fas-activated serine/threonine kinase	1.23	0.023
Q86YD1-3	<i>PTOV1</i>	Isoform 3 of Prostate tumor-overexpressed gene 1 protein	1.22	0.025
Q99985	<i>SEMA3C</i>	Semaphorin-3C	1.24	0.012
Q8N394	<i>TMTC2</i>	Transmembrane and TPR repeat-containing protein 2	1.22	0.040
Q9NRW4	<i>DUSP22</i>	Dual specificity protein phosphatase 22	1.24	0.054
Q15413-3	<i>RYR3</i>	Isoform 3 of Ryanodine receptor 3	1.24	0.011
P00973-2	<i>OAS1</i>	Isoform p41 of 2'-5'-oligoadenylate synthase 1	1.24	0.005

Q9Y6B6	<i>SAR1B</i>	GTP-binding protein SAR1b	1.25	0.030
Q13795	<i>ARFRP1</i>	ADP-ribosylation factor-related protein 1	1.24	0.047
P13611-2	<i>VCAN</i>	Isoform V1 of Versican core protein	1.27	0.044
Q8IYN2	<i>TCEAL8</i>	Transcription elongation factor A protein-like 8	1.27	0.043
O00506	<i>STK25</i>	Serine/threonine-protein kinase 25	1.27	0.036
P53367-2	<i>ARFIP1</i>	Isoform A of Arfaptin-1]	1.27	0.052
Q5SWA1	<i>PPP1R15B</i>	Protein phosphatase 1 regulatory subunit 15B	1.26	0.016
Q13164-2	<i>MAPK7</i>	Isoform 2 of Mitogen-activated protein kinase 7	1.28	0.030
O43542	<i>XRCC3</i>	DNA repair protein XRCC3	1.27	0.013
O60930	<i>RNASEH1</i>	Ribonuclease H1	1.28	0.026
O15040	<i>TECPR2</i>	Tectonin beta-propeller repeat-containing protein 2	1.28	0.000
O15055	<i>PER2</i>	Period circadian protein homolog 2	1.28	0.006
Q9P2W1	<i>PSMC3IP</i>	Homologous-pairing protein 2 homolog	1.30	0.030
Q6DCA0	<i>AMMECR1L</i>	AMMECR1-like protein	1.29	0.009
Q95807	<i>TMEM50A</i>	Transmembrane protein 50A	1.28	0.055
Q9NQ34	<i>TMEM9B</i>	Transmembrane protein 9B	1.30	0.044
Q96S15-2	<i>WDR24</i>	Isoform 2 of WD repeat-containing protein 24	1.32	0.040
Q9Y6M4-3	<i>CSNK1G3</i>	Isoform 3 of Casein kinase I isoform gamma-3	1.31	0.040
P15923-2	<i>TCF3</i>	Isoform E47 of Transcription factor E2-alpha	1.33	0.017
Q562F6-2	<i>SGOL2</i>	Isoform 2 of Shugoshin-like 2	1.36	0.030
P49447	<i>CYB561</i>	Cytochrome b561	1.34	0.043
Q99715-4	<i>COL12A1</i>	Isoform 4 of Collagen alpha-1(XII) chain	1.38	0.042
O14628-4	<i>ZNF195</i>	Isoform 4 of Zinc finger protein 195	1.43	0.040
Q08043	<i>ACTN3</i>	Alpha-actinin-3	1.46	0.009
P36894	<i>BMPR1A</i>	Bone morphogenetic protein receptor type-1A	1.51	0.023
P53667-2	<i>LIMK1</i>	Isoform 2 of LIM domain kinase 1	1.59	0.014
Q9UBT7-2	<i>CTNNAL1</i>	Isoform 2 of Alpha-catenin	1.59	0.013
Q9C030	<i>TRIM6</i>	Tripartite motif-containing protein 6	1.61	0.053
Q8IYH5-2	<i>ZZZ3</i>	Isoform 2 of ZZ-type zinc finger-containing protein 3	1.89	0.037

HC727 cells - 6 hours lanreotide

Q8N0X2-4	<i>SPAG16</i> *	Isoform 4 of Sperm-associated antigen 16 protein	0.11	0.000
O76027	<i>ANXA9</i>	Annexin A9	0.42	0.039
P11926	<i>ODC1</i>	Ornithine decarboxylase	0.53	0.005
Q92802	<i>N4BP2L2</i>	NEDD4-binding protein 2-like 2	0.55	0.045
Q9H8M1	<i>COQ10B</i>	Coenzyme Q-binding protein COQ10 homolog B, mitochondrial	0.62	0.021
P18847	<i>ATF3</i>	Cyclic AMP-dependent transcription factor ATF-3	0.62	0.049
O75382-3	<i>TRIM3</i>	Isoform Gamma of Tripartite motif-containing protein 3	0.65	0.019
Q15011-3	<i>HERPUD1</i>	Isoform 3 of Homocysteine-responsive endoplasmic reticulum-resident ubiquitin-like domain member 1 protein	0.69	0.042
Q96DN5-2	<i>WDR67</i>	Isoform 2 of WD repeat-containing protein 67	0.73	0.024
Q99576-4	<i>TSC22D3</i>	Isoform 3 of TSC22 domain family protein 3	0.73	0.001
Q71DI3	<i>HIST2H3A</i>	Histone H3.2	0.72	0.049
Q9H0X9	<i>OSBPL5</i>	Oxysterol-binding protein-related protein 5	0.74	0.011
O60907-2	<i>TBL1X</i>	Isoform 2 of F-box-like/WD repeat-containing protein TBL1X	0.75	0.042
P53667-2	<i>LIMK1</i>	Isoform 2 of LIM domain kinase 1	0.78	0.034
Q96A19	<i>CCDC102A</i>	Coiled-coil domain-containing protein 102A	1.22	0.007
Q92823-3	<i>NRCAM</i>	Isoform 3 of Neuronal cell adhesion molecule	1.22	0.010
Q06520	<i>SULT2A1</i>	Bile salt sulfotransferase	1.23	0.053
O14513	<i>NCKAP5</i>	Nck-associated protein 5	1.26	0.053
Q5T2R2-3	<i>PDSS1</i>	Isoform 3 of Decaprenyl-diphosphate synthase subunit 1	1.25	0.038
P17707	<i>AMD1</i>	S-adenosylmethionine decarboxylase proenzyme	1.27	0.046
Q6ZRI6-2	<i>C15orf39</i>	Isoform 2 of Uncharacterized protein C15orf39	1.30	0.005
O75081-2	<i>CBFA2T3</i>	Isoform 2 of Protein CBFA2T3	1.37	0.013

H727 cells - 48 hours lanreotide

Q8N0X2-4	<i>SPAG16</i> *	Isoform 4 of Sperm-associated antigen 16 protein	0.09	0.003
P11926	<i>ODC1</i>	Ornithine decarboxylase	0.36	0.011
A0FGR8-2	<i>ESYT2</i>	Isoform 2 of Extended synaptotagmin-2	0.52	0.029
O76027	<i>ANXA9</i>	Annexin A9	0.61	0.017
P38936	<i>CDKN1A</i>	Cyclin-dependent kinase inhibitor 1	0.62	0.021
P30825	<i>SLC7A1</i>	High affinity cationic amino acid transporter 1	0.63	0.049

Q01101	<i>INSM1</i> *	Insulinoma-associated protein 1	0.66	0.046
P04798	<i>CYP1A1</i>	Cytochrome P450 1A1	0.68	0.046
Q15011-3	<i>HERPUD1</i>	Isoform 3 of Homocysteine-responsive endoplasmic reticulum-resident ubiquitin-like domain member 1 protein	0.71	0.038
P06732	<i>CKM</i>	Creatine kinase M-type	0.73	0.040
P17275	<i>JUNB</i>	Transcription factor jun-B	0.75	0.013
Q8N594-2	<i>MPND</i>	Isoform 2 of MPN domain-containing protein	0.74	0.033
Q9UNP4-2	<i>ST3GAL5</i>	Isoform 2 of Lactosylceramide alpha-2,3-sialyltransferase	0.75	0.054
Q6P9G4	<i>TMEM154</i>	Transmembrane protein 154	0.75	0.040
Q53GA4	<i>PHLDA2</i>	Pleckstrin homology-like domain family A member 2	0.76	0.008
Q53GL0-2	<i>PLEKHO1</i>	Isoform 2 of Pleckstrin homology domain-containing family O member 1	0.77	0.052
O96020-2	<i>CCNE2</i>	Isoform Short of G1/S-specific cyclin-E2	1.22	0.025
Q9BVV6-2	<i>KIAA0586</i>	Isoform 2 of Uncharacterized protein KIAA0586	1.23	0.040
Q15345	<i>LRRC41</i>	Leucine-rich repeat-containing protein 41	1.22	0.042
Q96S15-2	<i>WDR24</i>	Isoform 2 of WD repeat-containing protein 24	1.24	0.004
Q96EN9	<i>C19orf60</i>	Uncharacterized protein C19orf60	1.24	0.053
Q96RV3-2	<i>PCNX</i>	Isoform 2 of Pecanex-like protein 1	1.24	0.042
Q9UHQ1-3	<i>NARF</i>	Isoform 3 of Nuclear prelamin A recognition factor	1.26	0.036
Q86VP1-3	<i>TAX1BP1</i>	Isoform 3 of Tax1-binding protein 1	1.25	0.026
Q92823-3	<i>NRCAM</i>	Isoform 3 of Neuronal cell adhesion molecule	1.25	0.043
P28356	<i>HOXD9</i>	Homeobox protein Hox-D9	1.27	0.024
Q9BQ16-2	<i>SPOCK3</i>	Isoform 2 of Testican-3	1.31	0.036
Q9NR48-2	<i>ASH1L</i>	Isoform 2 of Histone-lysine N-methyltransferase ASH1L	1.32	0.031
Q9NWL6	<i>ASNSD1</i>	Asparagine synthetase domain-containing protein 1	1.35	0.044
O60930	<i>RNASEH1</i>	Ribonuclease H1	1.35	0.027
Q9Y625	<i>GPC6</i>	Glypican-6	1.38	0.020
P07949-2	<i>RET</i>	Isoform 2 of Proto-oncogene tyrosine-protein kinase receptor Ret	1.45	0.009
Q08043	<i>ACTN3</i>	Alpha-actinin-3	1.75	0.032

Each protein expression ratio was normalized to its non-treated control which was assigned arbitrary value 1

Cut-offs for up-regulation and down-regulation were 0.69 and 1.33 for HC45 and 0.78 and 1.22 for H727

* Proteins selected for Western blot analysis and functional studies

Supplementary Table S5. Immunohistochemistry for APC and survivin in SI-NETs.

SI-NET samples	Sample analyzed	SSA treatment	Age at diagnosis	Gender F / M	Ki-67 %	APC				survivin			
						normal		tumor cells		normal		tumor cells	
						N	C	N	C	N	C	N	C
1A	Primary	Yes	75	F	1	0	0	0	0	0	0	0	1
1B	Primary	Yes	75	F	1	0	0	0	0	2	1	2	2
2	Primary	Yes	63	M	1.5	0	1	0	1	1	1	1	1
3	Primary	Yes	74	F	<1	0	2	1	1	0	0	0	0
4A	Primary	Yes	71	F	1	0	0	1	0	2	0	2	0
4B	Primary	Yes	71	F	1	0	3	2	1	0	0	0	0
5A	Primary	Yes	58	F	1.5	0	2	0	2	0	0	2	2
5B	Primary	Yes	58	F	1.5	0	0	0	1	0	0	3	3
6	Primary	Yes	63	F	1	0	0	0	1	0	0	1	1
7A	Metastasis	No	51	M	<1	0	1	0	0	0	0	0	2
7B	Metastasis	No	51	M	<1	0	1	0	0	0	1	2	3
8A	Primary	No	54	M	<1	1	1	0	2	0	0	1	2
8B	Primary	No	54	M	<1	0	0	0	0	0	0	2	3
9	Primary	No	66	M	<1	0	1	0	1	0	0	0	1
10	Primary	No	60	F	<1	0	0	0	1	0	1	0	1
11A	Primary	No	59	M	<1	0	1	0	0	0	0	0	2
11B	Primary	No	59	M	<1	0	0	0	2	0	0	2	2
12A	Primary	No	73	M	1	0	1	0	2	0	0	1	2
12B	Primary	No	73	M	1	0	0	2	1	0	0	1	1
13	Primary	No	47	F	3	0	0	0	0	0	0	0	2

N = nuclear, C = cytoplasmic, normal = normal epithelium

0 = absent, 1 = low, 2 = medium, 3 = high expression

SUPPLEMENTAL MATERIAL

Tissue microarray construction

Following confirmation of the NET diagnosis, representative areas of the tumors were selected on haematoxylin and eosin-stained sections and marked on individual paraffin blocks by expert pathologist in the field. Two tissue cores (1 mm in diameter) were obtained from each specimen. The tissue cores were arrayed into a receptor paraffin block using a tissue microarray workstation (Beecher Instruments, Silver Spring, MD, USA), as described previously (9). A haematoxylin and eosin-stained section of the array was reviewed to confirm the presence of morphologically representative areas of the original lesions.

Genotypes and culturing of NET cell lines

Short tandem repeat (STR) genotyping was performed to assure the authenticity of the BON1 cells (at Biosynthesis Inc. Lewisville, TX) and has been previously published for HC45 and CNDT2 (12). The following genotypes were observed (for HC45 / CNDT2 / BON1) at the genotyped loci including: *D8S1178* (13, 16 / 9, 13, 14 / 10, 12); *D21S11* (29, 32.2 / 27, 31 / 28, 33.2); *D7S820* (8, 8 / 8, 10 / 9, 9); *CSFIPO* (9, 11 / 8, 11 / 10, 11); *D3S1358* (14, 17 / 16, 19 / 17, 17); *THO1* (6, 7 / 6, 10 / 8, 8); *DI3S317* (10, 11 / 8, 11 / 11, 12); *DI6S539* (12, 12 / 12, 13 / 10, 11); *D2S1338* (23, 25 / 16, 16 / 16, 23); *DI9S433* (12, 15 / 14, 14 / 12, 5.2); *vWA* (15, 16 / 17, 18, 25 / 18, 19); *TPOX* (8, 12 / 10, 11 / 9, 9); *DI8S51* (16, 16 / 11, 13 / 12, 12); *AMEL* (X, X / X, X / X, Y); *D5S818* (12, 13 / 11, 14 / 9, 12); and *FGA* (21, 24 / 20, 22 / 23, 23).

HC45 and CNDT2 were cultured in DMEM, BON1 cells were cultured in DMEM/F12 and NCI-H727 in RPMI-1640. All media were supplemented with FBS and the cells were incubated at 37 °C, 5% CO₂ in a humidified incubator.

HiRIEF LC-MS/MS experimental procedures and data analyses

A total of 16 samples were profiled including lanreotide treated HC45 and NCI-H727 cells representing the time points 2, 6 and 48 hours as well as controls in duplicates. For protein preparation, cells were lysed in SDS buffer (4 % SDS, 25 mM HEPES, 1 mM DTT) applying 200 µL buffer/106 cells. The samples were then heated to 95 °C, homogenized by sonication with a probe tip (4 x 10 sec), and centrifuged at 10,000 rpm for 10 minutes at 4 °C. Protein concentrations were measured using the Bio-Rad DC protein assay (Bio-Rad Laboratories, Hercules, CA, USA). Two-hundred µg protein from each sample was digested applying the FASP (Filter Aided Sample Preparation) protocol.¹⁸ One hundred µg of each peptide sample were labeled and pooled using the 8-plex iTRAQ kit (Applied Biosystems, Foster City, CA, USA) according to the manufacturer's instructions. Excess reagent was removed from the pooled sample using an SCX-cartridge (StrataSCX, Phenomenex, Torrence, CA, USA). The eluate was dried in a speed-vac.

The peptide isoelectric focusing procedure has been described before (7). iTRAQ-labeled tryptic peptide samples were dissolved in 300 µL 8 M urea. Narrow range IPG-strips for peptide focusing (pH 3.7 - 4.9) together with dry sample application gels (33×3×2 mm) were supplied by GE Healthcare Bio-Sciences AB, Uppsala, Sweden. After peptide separation samples were freeze-dried in a SpeedVac and kept at -20 °C. Prior to HiRIEF LC-MS/MS analysis, each fraction was re-suspended in 8 µl 3% acetonitrile and 0.1% formic acid.

HiRIEF LC-MS/MS data analysis

In each LC-MS/MS run, the LC auto sampler (HPLC 1,200 system, Agilent Technologies) injected 3 μ l into a C18 guard desalting column (Zorbax 300SB-C18, 5 x 0.3 mm, 5 μ m bead size, Agilent). We then used a 15 cm long C18 picofrit column (100 μ m internal diameter, 5 μ m bead size, Nikkyo Technos Co., Tokyo, Japan) installed onto the nano-electrospray ionization (NSI) source. Solvent and gradient descriptions have been described before (7). Sequest searched all MS/MS spectra under the software platform Proteome Discoverer (v1.3.0.339, Thermo Scientific) against a fasta file containing human tryptic sequences. Settings used were: trypsin with 1 missed cleavage, carbamidomethylation on cysteine and iTRAQ-8 plex on lysine and N-terminal as fixed modifications, and oxidation of methionine as variable modification. Further settings for the MS/MS search have been described before (7). Results were limited to ≥ 1 high-confident unique peptide for quantification using a false discovery rate of <1 % at the peptide level.

The consistency between replicate analysis in HC45 and H727 proteomics profiling data were evaluated by Principal component analysis (PCA) using the SIMCA software.

The software Disease Association Protein-Protein Link Evaluator version 2.0 (DAPPLE, <http://www.broadinstitute.org/mpg/dapple/dapple.php>) (34) was used to identify signaling networks for differentially expressed proteins. The physical connectivity was assessed between differentially expressed proteins with the highest/lowest fold change at different time points. All analyses were performed allowing 20,000 permutations and a common interactor binding degree cut-off of 2. Pathway analyses were carried out using the Ingenuity Pathway

Analysis (IPA) (Qiagen, CA, USA). Expected mass sizes of proteins of interest are according to information at www.uniprot.org.

Confidence interval = 0.95 was calculated as the mean \pm 2 \times (standard deviation) of the isobaric label ratios between non-treated controls and treated samples and was defined as cut-offs. Accordingly 0.69 and 1.33 for HC45 and 0.78 and 1.22 for H727 were considered as cut-offs for down-regulation and up-regulation, respectively. Noisy variables exceeding replicate variation based cut-off were excluded. T-test independent 2-tailed were calculated using Microsoft Excel to compare the expression of any protein in each time point vs. non-treated condition. *P*-values below 0.05 were reported as significant. Expression ratios exceeding the above-mentioned cut-offs were considered as down-regulated or up-regulated. Different conditions in proliferation assays and immunohistochemistry scores were compared applying t-test 2-tailed using Microsoft Excel or analyzed for correlations using Spearman's test IBM SPSS statistics version 20.0.

Immunocytochemistry

Immunocytochemistry was performed essentially as previously described (14). In short, cells were trypsinized and cytopinned onto glass slides and incubated with primary antibodies over-night at 4 °C including: SSTR2 (Sigma-Aldrich, HPA007264) at dilution 1:100 for cell lines and 1:200 for the primary SI-NET culture; SSTR5 (Epitomics, clone UMB4) at 1:250 for cell lines and 1:200 for the primary SI-NET culture; and Chromogranin A (Santa Cruz) at 1:150. MIB1 proliferation index was determined by immunocytochemistry using the MIB1 antibody targeting Ki-67 (Dako) at dilution 1:200. FFPE sections of anonymized pancreatic tissue samples were used as positive controls using previously published methodology (15) and applying the SSTR2 and SSTR5 antibodies at dilution 1:300. Omission of the primary

antibodies served as negative controls, and expression in Langerhans cell islets of normal pancreas as a positive control.

Immunohistochemistry and tissue microarray

Immunohistochemistry (IHC) was performed on the 20 SI-NET tissue samples employing previously published methodology (15) using primary antibodies for survivin (BIRC5 D-8, Santa Cruz Biotechnology) at dilution 1:50 and for APC (EP701Y, Abcam) at 1:100.

Anonymized tissue sections of normal breast and normal thyroid were included as positive controls of survivin and APC, respectively, and as negative controls after omission of the primary antibodies. Nuclear and cytoplasmic staining were observed and evaluated in tumor and normal epithelial cells on the same slides. The result was analyzed using a Zeiss Axioplan 2 (Carl Zeiss Jena GmbH, Jena, Germany) microscope, ProgRes C12 plus camera and ProgRes CapturePro 2.5 software.

IHC analyses of the 112 NETs were performed using previously published methodology (35) on 4 µm sections of the TMAs, obtained by a semi-automated microtome HM3508 (Microm). Briefly, tissue sections were deparaffinized and rehydrated in water, after which antigen retrieval was carried out in a DAKO PT Link in citrate buffer (pH 6). Endogenous peroxidase and non-specific antibody reactivity was blocked with peroxidase blocking reagent (Dako, Glostrup, Denmark) at room temperature for 15 min. The sections were then incubated for 60 min with a rabbit polyclonal survivin antibody (#Ab469, Abcam) diluted at 1:400 and a rabbit monoclonal APC antibody (#Ab40778, Abcam) diluted at 1:100. Detection was performed with EnvisionPlus Detection System (Dako) using an Olympus BX43 microscope with DP72 Olympus camera and LabSense software.

Western blot analysis

After blocking membranes were incubated with primary antibodies overnight for APC (EP701Y, Abcam) at dilution 1:500, APC (op44, MERK, Calbiochem) at 1:1000, BMPER (Sigma-Aldrich) at 1:150, SMIM21 (C18orf62, Sigma-Aldrich) at 1:250, SPAG16 (Aviva Systems Biology) at 1:250, C14orf142 (Aviva Systems Biology) at 1:500, FYN (BD Biosciences) at 1:500, survivin (BIRC5 D-8, Santa Cruz Biotechnology) at 1:200, Chromogranin A (LK2H10, Santa Cruz Biotechnology) at 1:1000 and INSM1 (C-1, Santa Cruz Biotech) at 1:750. Rabbit GAPDH (Cell Signaling Technology) at 1:3,000 or mouse GAPDH (O411, Santa Cruz Biotechnology) at 1:10,000 and Vinculin (7F9, Santa Cruz Biotechnology) at 1:1,000, were used as loading controls. Goat anti-rabbit IgG-HRP (Santa Cruz Biotechnology) at 1:7,500, IRDye 800CW goat anti-mouse IgG (P/N 926-32210, LI-COR Biosciences) at 1:10,000 or IRDye 680RD Goat anti-Rabbit IgG (P/N 926-68071, LI-COR Biosciences) at 1:10,000 served as secondary antibodies and were incubated for one hour at room temperature. Membranes were read in an Odyssey imaging system (LI-COR Biosciences). Expression densities were quantified using Image Studio 2.1 software.

Proliferation assays

For the BrdU proliferation assay, cells were seeded in the 96-well plates with densities of 4,000 cells/well for the primary SI-NET, 4,000 cells /well for HC45, 8,000 cells /well for BON1 and 8,000 cells/well for H727. BrdU labeling reagent was added 48 hours after treatment. Incubation of the cell lines or primary cell culture continued for 4 or 24 hours, respectively. Cells were then fixed and incubated with the BrdU antibody for 90 minutes.

The xCELLigence real time proliferation assays were performed on E-plates following the manufacturer's instructions (Aceabio). Experiments were carried out at 0.5% FBS. Cell

indices were obtained in 1 or 4 hours intervals. Data were normalized to the cell indices at the time of starting of the treatment assigning the arbitrary value of 1. Proliferation index was determined by immunocytochemistry using the MIB1 antibody (Dako) for detection of the Ki-67 antigen.

***In vitro* transfection experiments**

APC (pCMV-Neo-Bam APC Addgene plasmid # 16507) and its empty vector (pCMV-Neo-Bam Addgene plasmid # 16440) were a gift from Bert Vogelstein (36). Since HC45 cells are slow growing and difficult to transfect we used H727 cells for transfection experiments.

Lipofectamine 2000 or RNAiMAX was used for transfection of plasmids or siRNA 24 hours after seeding according to the instructions of the manufacturer (Life Technologies). Forty-eight hours later protein was extracted and used for Western blot analysis. To knock down APC, H727 cells were transfected with shAPC1 and shAPC2 (TRCN0000040093 and TRCN0000040097- Sigma Aldrich) under the same conditions. PLKO.1 eGFP knock-down shRNA (Sigma Aldrich) was used as a control. siRNA control 1 and siSSTR2 (s13500, Life Technologies) were used for *SSTR2* knock-down experiments.

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