

Supplemental Table S-1b. A list of 588 proteins quantified by XPRESS, sorted by protein name.

Acc #	Protein name	TPC	H/L ratio	SD
O00461	130 kD Golgi-localized phosphoprotein	1	0.26	
Q15392	24-dehydrocholesterol reductase precursor (EC 1.3.1.-) (3-beta-hydrox	1	0.36	
Q99460	26S proteasome non-ATPase regulatory subunit 1 (26S proteasome re	5	0.48	0.34
P62280	40S ribosomal protein S11	1	0.90	
P62263	40S ribosomal protein S14	1	0.34	
P62269	40S ribosomal protein S18 (Ke-3) (Ke3)	3	0.35	0.03
P23396	40S ribosomal protein S3	1	0.36	
P62861	40S ribosomal protein S30	1	0.00	
P08195	4F2 cell-surface antigen heavy chain (4F2hc) (Lymphocyte activation a	91	0.50	0.38
Q07020	60S ribosomal protein L18	2	0.92	0.42
Q02543	60S ribosomal protein L18a	1	0.28	
P62829	60S ribosomal protein L23 (Ribosomal protein L17)	3	0.68	0.45
P62899	60S ribosomal protein L31	1	0.37	
P83881	60S ribosomal protein L36a (60S ribosomal protein L44) (Migration-ind	1	1.11	
P36578	60S ribosomal protein L4 (L1)	3	1.21	1.56
P11021	78 kDa glucose-regulated protein precursor (GRP 78) (Immunoglobuli	5	1.26	1.26
Q9UBM7	7-dehydrocholesterol reductase (EC 1.3.1.21) (7-DHC reductase) (Ster	6	0.98	0.79
O00400	Acetyl-coenzyme A transporter 1 (AT-1) (Acetyl-CoA transporter) (Solu	2	0.19	0.12
Q6P7J9	ADAM metalloproteinase with thrombospondin type 1 motif, 7, prepropr	1	1.87	
Q8TB61	Adenosine 3'-phospho 5'-phosphosulfate transporter 1 (PAPS transpor	3	0.91	0.47
P12235	ADP/ATP translocase 1 (Adenine nucleotide translocator 1) (ANT 1) (A	1	0.24	
P05141	ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (ANT 2) (A	10	0.27	0.12
O00116	Alkyldihydroxyacetonephosphate synthase, peroxisomal precursor (EC	1	1.08	
Q11128	Alpha-(1,3)-fucosyltransferase (EC 2.4.1.65) (Galactoside 3-L-fucosyltr	1	1.59	
Q9H6U8	Alpha-1,2-mannosyltransferase ALG9 (EC 2.4.1.-) (Asparagine-linked	1	0.36	
P35609	Alpha-actinin 2 (Alpha actinin skeletal muscle isoform 2) (F-actin cross	1	11.07	
Q15911	Alpha-fetoprotein enhancer binding protein (AT motif-binding factor) (A	1	1.42	
Q86WK6	Amphoterin-induced protein 1 precursor (Alivin-2)	1	0.42	
Q8WWH4	Ankyrin repeat, SAM and basic leucine zipper domain-containing prote	1	0.05	

Acc #	Protein name	TPC	H/L ratio	SD
P04083	Annexin A1 (Annexin I) (Lipocortin I) (Calpactin II) (Chromobindin-9) (p	1	2.53	
Q9UKV3	Apoptotic chromatin condensation inducer in the nucleus (Acinus)	2	2.41	1.16
O14727	Apoptotic protease-activating factor 1 (Apaf-1)	1	0.14	
Q8IU5	ARP5 actin-related protein 5 homolog	1	0.26	
Q9Y4J0	Aspartyl(Asparaginy)l-beta-hydroxylase	2	0.36	0.00
P25705	ATP synthase alpha chain, mitochondrial precursor (EC 3.6.3.14)	26	0.64	0.54
P06576	ATP synthase beta chain, mitochondrial precursor (EC 3.6.3.14)	15	0.43	0.17
P18859	ATP synthase coupling factor 6, mitochondrial precursor (EC 3.6.3.14)	1	0.93	
O75947	ATP synthase D chain, mitochondrial (EC 3.6.3.14)	6	0.85	1.07
P30049	ATP synthase delta chain, mitochondrial precursor (EC 3.6.3.14)	1	1.55	
P56385	ATP synthase e chain, mitochondrial (EC 3.6.3.14)	7	0.20	0.21
O75964	ATP synthase g chain, mitochondrial (EC 3.6.3.14) (ATPase subunit G	4	0.38	0.04
P36542	ATP synthase gamma chain, mitochondrial precursor (EC 3.6.3.14)	4	0.39	0.16
P48047	ATP synthase O subunit, mitochondrial precursor (EC 3.6.3.14) (Oligo	8	0.75	0.47
Q8NBU5	ATPase family AAA domain containing protein 1	4	0.61	1.03
Q86UK0	ATP-binding cassette sub-family A member 12 (ATP-binding cassette	1	2.19	
P33897	ATP-binding cassette sub-family D member 1 (Adrenoleukodystrophy	6	0.74	0.67
P28288	ATP-binding cassette sub-family D member 3 (70 kDa peroxisomal me	10	0.51	0.32
Q9UNQ0	ATP-binding cassette sub-family G member 2 (Placenta-specific ATP-	16	0.56	0.74
Q92499	ATP-dependent RNA helicase DDX1 (EC 3.6.1.-) (DEAD box protein 1	1	2.08	
Q9NT62	Autophagy protein 3-like (APG3-like) (hApg3) (PC3-96 protein)	1	1.82	
Q7Z3C6	Autophagy protein 9-like 1 (APG9-like 1)	1	1.46	
Q8IXQ6	B aggressive lymphoma protein	1	0.87	
O95816	BAG family molecular chaperone regulator 2 (BCL2-associated athano	3	0.65	0.42
Q9BXC9	Bardet-Biedl syndrome 2 protein	3	0.41	0.48
P35613	Basigin precursor (CD147 antigen) (Leukocyte activation antigen M6) (4	0.47	0.25
Q9C0K0	B-cell lymphoma/leukemia 11B (B-cell CLL/lymphoma 11B) (Radiation	1	0.27	
Q9UHQ4	B-cell receptor-associated protein 29 (BCR-associated protein Bap29)	2	0.31	0.04
P51572	B-cell receptor-associated protein 31 (BCR-associated protein Bap31)	17	0.27	0.16
P53674	Beta crystallin B1	1	0.19	

Acc #	Protein name	TPC	H/L ratio	SD
P61769	Beta-2-microglobulin precursor [Contains: Beta-2-microglobulin variant	2	0.50	0.00
Q9Y223	Bifunctional UDP-N-acetylglucosamine 2-epimerase/N-acetylmannosa	1	1.00	
P80723	Brain acid soluble protein 1 (BASP1 protein) (Neuronal axonal membra	8	0.37	0.24
Q9Y6D6	Brefeldin A-inhibited guanine nucleotide-exchange protein 1 (Brefeldin	1	1.66	
Q9NXS3	BTB/POZ domain-containing protein 5	1	0.61	
Q15800	C-4 methylsterol oxidase (EC 1.14.13.72) (Methylsterol monooxygenas	1	1.79	
O75844	CAAX prenyl protease 1 homolog (EC 3.4.24.84) (Prenyl protein-specif	10	0.19	0.18
Q14517	Cadherin-related tumor suppressor homolog precursor (Fat protein ho	1	0.20	
P98194	Calcium-transporting ATPase type 2C member 1 (EC 3.6.3.8) (ATPas	1	0.49	
P27824	Calnexin precursor (Major histocompatibility complex class I antigen-bi	4	0.80	0.79
Q9GZY7	CAMP/cGMP cyclic nucleotide phosphodiesterase 11A3	1	0.50	
Q8NG31	Cancer susceptibility candidate gene 5 protein (ALL1-fused gene from	1	0.07	
P31327	Carbamoyl-phosphate synthase [ammonia], mitochondrial precursor (1	0.35	
P50416	Carnitine O-palmitoyltransferase I, mitochondrial liver isoform (EC 2.3.	3	0.54	0.34
P11717	Cation-independent mannose-6-phosphate receptor precursor (CI Man	1	15.15	
P51636	Caveolin-2	2	0.51	0.00
P13987	CD59 glycoprotein precursor (Membrane attack complex inhibition fact	3	4.33	6.41
Q6ZMV5	CDNA FLJ16643 fis, clone TEST14029528	1	0.24	
Q8NBI5	CDNA PSEC0252 fis, clone NT2RP3003258, highly similar to Likely or	3	0.35	0.21
O75943	Cell cycle checkpoint protein RAD17 (hRad17) (RF-C/activator 1 homo	1	0.15	
Q99459	Cell division cycle 5-like protein (Cdc5-like protein) (Pombe cdc5-relate	2	0.24	0.06
Q15057	Centaurin-beta 2 (Cnt-b2)	1	0.47	
Q66GS9	Centrosomal protein 4 (Centrosomal protein of 135 kDa) (Cep135 prot	1	0.30	
Q70F00	Centrosome spindle pole associated protein	1	0.48	
Q16739	Ceramide glucosyltransferase (EC 2.4.1.80) (Glucosylceramide syntha	1	0.61	
O00590	Chemokine-binding protein 2 (Chemokine-binding protein D6) (C-C ch	2	1.97	2.01
Q8IZ52	Chondroitin sulfate synthase 2 (EC 2.4.1.175) (Glucuronosyl-N-acetylg	1	0.34	
Q5TF85	Chromosome 6 open reading frame 37	2	2.59	0.00
Q6PEV5	CKLFSF1 protein	1	0.40	
Q00610	Clathrin heavy chain 1 (CLH-17)	2	0.67	0.44

Acc #	Protein name	TPC	H/L ratio	SD
Q5W1L8	CMP-sialic acid transporter	1	1.43	
Q9NX63	Coiled-coil-helix-coiled-coil-helix domain containing protein 3	12	0.59	0.33
Q02388	Collagen alpha 1(VII) chain precursor (Long-chain collagen) (LC collag	3	0.50	0.10
Q07021	Complement component 1, Q subcomponent binding protein, mitoch	3	0.48	0.17
Q9NS00	Core1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1,3-galacto	1	0.30	
P57737	Coronin-7 (70 kDa WD repeat tumor rejection antigen homolog)	1	0.04	
P61073	C-X-C chemokine receptor type 4 (CXC-R4) (CXCR-4) (Stromal cell-de	1	0.16	
P20674	Cytochrome c oxidase polypeptide Va, mitochondrial precursor (EC 1.	2	0.61	0.06
P10606	Cytochrome c oxidase polypeptide Vb, mitochondrial precursor (EC 1.	5	0.40	0.37
P09669	Cytochrome c oxidase polypeptide VIc precursor (EC 1.9.3.1)	4	0.40	0.11
P14406	Cytochrome c oxidase polypeptide VIIa-liver/heart, mitochondrial precu	4	7.05	12.57
P13073	Cytochrome c oxidase subunit IV isoform 1, mitochondrial precursor (E	4	0.63	0.26
Q14548	Cytochrome c oxidase subunit VIIa-related protein, mitochondrial prec	2	0.57	0.00
P08574	Cytochrome c1, heme protein, mitochondrial precursor (Cytochrome c-	7	0.60	0.54
P53701	Cytochrome c-type heme lyase (EC 4.4.1.17) (CCHL) (Holocytochrom	1	0.40	
Q16850	Cytochrome P450 51A1 (EC 1.14.13.70) (CYPLI) (P450LI) (Sterol 14-a	1	0.36	
Q6NT55	Cytochrome P450, family 2, subfamily E, polypeptide 2 homolog	1	0.19	
Q96P26	Cytosolic 5'-nucleotidase 1B (EC 3.1.3.5) (Cytosolic 5'-nucleotidase IB	1	0.60	
Q9NYF0	Dapper homolog 1 (hDPR1) (Heptacellular carcinoma novel gene 3 pr	1	0.22	
Q6YHU2	Death receptor interacting protein splice variant	3	157.35	167.37
Q14185	Dedicator of cytokinesis protein 1 (180 kDa protein downstream of CR	1	188.67	
P61803	Defender against cell death 1 (DAD-1)	3	0.42	0.22
Q96HY7	Dehydrogenase E1 and transketolase domain containing protein 1	2	1.63	0.06
Q8NBQ5	Dehydrogenase/reductase SDR family member 8 precursor (EC 1.1.1.	3	0.44	0.33
Q8TB45	DEP domain containing 6	1	0.38	
Q75140	DEP domain containing protein 5	1	0.17	
P81605	Dermcidin precursor (Preproteolysin) [Contains: Survival-promoting pe	2	0.75	0.00
Q4KKU6	Desmoglein 2, preproprotein	3	0.28	0.21
P15924	Desmoplakin (DP) (250/210 kDa paraneoplastic pemphigus antigen)	1	0.50	
Q5FWG1	DGKG protein	1	0.47	

Acc #	Protein name	TPC	H/L ratio	SD
O75907	Diacylglycerol O-acyltransferase 1 (EC 2.3.1.20) (Diglyceride acyltrans	2	1.06	1.21
O95886	Disks large-associated protein 3 (DAP-3) (SAP90/PSD-95-associated	1	0.37	
Q8IY66	DNA ligase IV (Ligase IV, DNA, ATP-dependent)	1	37.85	
P04053	DNA nucleotidylxotransferase (EC 2.7.7.31) (Terminal addition enzy	1	0.18	
P25205	DNA replication licensing factor MCM3 (DNA polymerase alpha holoen	1	27.59	
Q9BUI4	DNA-directed RNA polymerase III 62 kDa polypeptide (EC 2.7.7.6) (R	1	0.66	
O00411	DNA-directed RNA polymerase, mitochondrial precursor (EC 2.7.7.6) (1	6.29	
Q5T219	DnaJ (Hsp40) homolog, subfamily D, member 1 (Growth-inhibiting 22)	1	0.12	
Q86YN1	Dolichyldiphosphatase 1 (EC 3.6.1.43) (Dolichyl pyrophosphate phosp	1	0.58	
Q5VWA5	Dolichyl-diphosphooligosaccharide-protein glycosyltransferase	7	0.41	0.20
P04843	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 67 kDa	45	0.52	0.39
Q92685	Dolichyl-P-Man:Man(5)GlcNAc(2)-PP-dolichyl mannosyltransferase (E	5	0.39	0.17
O75691	Down-regulated in metastasis protein (Key-1A6 protein) (Novel nucleol	2	0.33	0.02
Q6UXB4	DTTR431 (C-type lectin superfamily 4, member G)	1	7.38	
O00429	Dynamin-1-like protein (EC 3.6.5.5) (Dynamin-like protein) (Dnm1p/Vp	2	8.89	0.62
Q13409	Dynein intermediate chain 2, cytosolic (DH IC-2) (Cytoplasmic dynein i	1	0.83	
P11532	Dystrophin	1	0.05	
P13639	Elongation factor 2 (EF-2)	1	0.12	
P49411	Elongation factor Tu, mitochondrial precursor (EF-Tu) (P43)	1	0.58	
Q9BW60	Elongation of very long chain fatty acids protein 1	2	27.80	2.03
Q9BVF7	Elongation protein 3 homolog (Hypothetical protein DKFZp761L098)	1	1.13	
P29317	Ephrin type-A receptor 2 precursor (EC 2.7.1.112) (Tyrosine-protein ki	3	0.17	0.16
Q9UHB6	Epithelial protein lost in neoplasm	1	0.26	
Q14677	Epsin-4 (Epsin-related protein) (EpsinR) (Enthoprotin) (Clathrin interact	1	0.45	
P49257	ERGIC-53 protein precursor (ER-Golgi intermediate compartment 53 k	1	0.09	
Q969X5	ER-Golgi intermediate compartment 32 kDa protein (ERGIC-32)	1	0.30	
P27105	Erythrocyte band 7 integral membrane protein (Stomatin) (Protein 7.2b	5	0.53	0.18
Q99613	Eukaryotic translation initiation factor 3 subunit 8 (eIF3 p110) (eIF3c)	1	2.26	
O60447	EVI-5 homolog	1	0.52	
P52907	F-actin capping protein alpha-1 subunit (CapZ alpha-1)	1	0.23	

Acc #	Protein name	TPC	H/L ratio	SD
Q53EP0	Factor for adipocyte differentiation 104 variant (Fragment)	1	0.07	
Q01469	Fatty acid-binding protein, epidermal (E-FABP) (Psoriasis-associated f	1	1.61	
Q6PCT2	F-box/LRR-repeat protein 19 (F-box and leucine-rich repeat protein 19)	2	0.37	0.05
Q9Y5Y0	Feline leukemia virus subgroup C receptor-related protein 1 (Feline leu	1	0.48	
Q9BS16	FKSG14 protein (Leucine zipper protein AF5alpha) (SoxLZ/Sox6 leucin	1	25.44	
Q9BYX7	FKSG30	1	180.23	
Q9H7J3	FLJ00086 protein (Fragment)	1	0.18	
Q8TEP9	FLJ00144 protein (Fragment)	1	0.08	
Q6IAA8	FLJ20625 protein	1	21.79	
O75955	Flotillin-1	3	0.38	0.16
Q14254	Flotillin-2 (Epidermal surface antigen) (ESA)	3	1.65	1.58
Q92735	FMI protein	1	0.79	
Q14488	Forkhead box protein G1C (Forkhead-related protein FKHL3) (HFK3) (1	0.49	
Q71RG4	FP2653 (ELSD1897) (MGC3123 protein)	2	0.63	0.00
O60353	Frizzled 6 precursor (Frizzled-6) (Fz-6) (hFz6)	2	0.87	0.11
Q6ZV73	FYVE, RhoGEF and PH domain containing protein 6 (Zinc finger FYV	1	0.81	
P41002	G2/mitotic-specific cyclin-F	1	0.85	
P07316	Gamma crystallin B (Gamma crystallin 1-2)	1	0.31	
O95452	Gap junction beta-6 protein (Connexin-30) (Cx30)	1	0.09	
Q6P1A2	Gene rich cluster, C3f gene	3	0.47	0.18
Q7Z6J2	General receptor for phosphoinositides 1-associated scaffold protein (2	2.00	0.77
P39086	Glutamate receptor, ionotropic kainate 1 precursor (Glutamate recepto	1	0.41	
P48506	Glutamate--cysteine ligase catalytic subunit (EC 6.3.2.2) (Gamma-glut	1	0.57	
P48637	Glutathione synthetase (EC 6.3.2.3) (Glutathione synthase) (GSH synt	2	0.06	0.03
Q5JR58	Glyceraldehyde 3-phosphate dehydrogenase (GAPDH)	1	0.17	
P04406	Glyceraldehyde-3-phosphate dehydrogenase, liver (EC 1.2.1.12) (GAP	1	2.09	
P06737	Glycogen phosphorylase, liver form (EC 2.4.1.1)	1	0.42	
Q9NZD2	Glycolipid transfer protein (GLTP)	1	0.48	
Q9UBI6	Guanine nucleotide-binding protein G(I)/G(S)/G(O) gamma-12 subunit	3	0.17	0.07
P62879	Guanine nucleotide-binding protein G(I)/G(S)/G(T) beta subunit 2 (Tra	1	0.80	

Acc #	Protein name	TPC	H/L ratio	SD
P08754	Guanine nucleotide-binding protein G(k), alpha subunit (G(i) alpha-3)	2	0.80	0.34
O95837	Guanine nucleotide-binding protein, alpha-14 subunit (G-protein alpha	1	0.16	
Q6J203	HCV F-transactivated protein 1	1	0.44	
P11142	Heat shock cognate 71 kDa protein (Heat shock 70 kDa protein 8)	9	0.58	0.58
P04792	Heat-shock protein beta-1 (HspB1) (Heat shock 27 kDa protein) (HSP	3	0.87	0.42
Q15034	HECT domain and RCC1-like domain protein 3	1	2.39	
Q969F9	Hermansky-Pudlak syndrome 3 protein	1	1.38	
P16401	Histone H1.5 (Histone H1a)	2	0.25	0.00
P62805	Histone H4	23	0.44	0.24
Q96L73	Histone-lysine N-methyltransferase, H3 lysine-36 and H4 lysine-20 spe	1	0.33	
Q15047	Histone-lysine N-methyltransferase, H3 lysine-9 specific 4 (EC 2.1.1.4	1	3.21	
GAG_Prote	HIV GAG protein	23	8.77	8.72
Q8MH63	HLAT1-3TM	2	0.42	0.12
O14529	Homeobox protein Cux-2 (Cut-like 2)	1	6.43	
Q15011	Homocysteine-responsive endoplasmic reticulum-resident ubiquitin-lik	1	0.41	
P57058	Hormonally up-regulated neu tumor-associated kinase (EC 2.7.1.37) (1	0.60	
Q86YZ3	Hornerin	1	3.58	
Q9Y5U9	HSPC039 protein (Hypothetical transmembrane protein) (Immediate e	1	0.50	
Q9NZY6	HSPC049	1	0.04	
Q9P035	HSPC121	5	0.31	0.04
Q9UJM8	Hydroxyacid oxidase 1 (EC 1.1.3.15) (HAOX1) (Glycolate oxidase) (G	1	0.05	
Q32M75	Hypothetical protein	1	0.29	
Q3B7X4	Hypothetical protein	2	0.43	0.11
Q3KRF4	Hypothetical protein	1	0.50	
Q3MJA4	Hypothetical protein	1	0.12	
Q49AJ3	Hypothetical protein	1	4.58	
Q4G0N0	Hypothetical protein	3	0.60	0.72
Q6PQ33	Hypothetical protein	1	0.53	
Q8NFK8	Hypothetical protein (Novel protein)	1	1.24	
Q8N1H7	Hypothetical protein C14orf39	1	0.53	

Acc #	Protein name	TPC	H/L ratio	SD
Q5H9S5	Hypothetical protein DKFZp313B047	1	0.11	
Q9H0H5	Hypothetical protein DKFZp434C011 (RACGAP1 protein)	1	0.40	
Q8IYM0	Hypothetical protein DKFZP434J0113	2	0.35	0.10
Q8NDA2	Hypothetical protein DKFZp434P0216 (Fragment)	1	1.30	
Q5HY96	Hypothetical protein DKFZp686C0394	1	0.06	
Q6MZH2	Hypothetical protein DKFZp686J07132	1	0.05	
Q68DQ6	Hypothetical protein DKFZp686O22167	1	0.28	
Q6AHZ3	Hypothetical protein DKFZp781B2386	1	32.08	
Q5H9U9	Hypothetical protein DKFZp781D1175	1	0.13	
Q68DK2	Hypothetical protein DKFZp781H1112	1	0.45	
Q6AHZ1	Hypothetical protein DKFZp781O2147	1	0.08	
Q8TC56	Hypothetical protein FAM71B	1	0.41	
Q9NVA4	Hypothetical protein FLJ10846	1	0.24	
Q9H8U9	Hypothetical protein FLJ13213	1	0.57	
Q96A33	Hypothetical protein FLJ14938 (Hypothetical protein NT2RP2003902) (2	0.27	0.02
Q6ZN37	Hypothetical protein FLJ16464	1	0.13	
Q6ZMX2	Hypothetical protein FLJ16617	1	9.13	
Q9NXG7	Hypothetical protein FLJ20261	1	0.43	
Q9NX61	Hypothetical protein FLJ20422 (AVLG582)	1	1.01	
Q9NX40	Hypothetical protein FLJ20455 (TPA018) (Ovarian carcinoma immunor	4	0.87	0.33
Q8TEC2	Hypothetical protein FLJ23658	1	1.90	
Q96N66	Hypothetical protein FLJ31346	1	0.27	
Q96N64	Hypothetical protein FLJ31359	1	5.21	
Q96MH7	Hypothetical protein FLJ32363	1	0.14	
Q96M69	Hypothetical protein FLJ32786	1	0.19	
Q8N1U5	Hypothetical protein FLJ37549	1	0.36	
Q8N8I4	Hypothetical protein FLJ39430	1	0.23	
Q6ZWG9	Hypothetical protein FLJ41103	1	10.50	
Q6ZVH2	Hypothetical protein FLJ42583	1	0.36	
Q6ZV38	Hypothetical protein FLJ43035	1	0.79	

Acc #	Protein name	TPC	H/L ratio	SD
Q6ZTN4	Hypothetical protein FLJ44477	1	7.73	
Q6ZT81	Hypothetical protein FLJ44884	1	9.16	
Q6ZST6	Hypothetical protein FLJ45221	1	0.20	
Q6ZRQ5	Hypothetical protein FLJ46180	1	0.15	
Q6ZQW6	Hypothetical protein FLJ46844	1	0.54	
Q71M32	Hypothetical protein FP14381	1	0.30	
Q96S30	Hypothetical protein gene +108	1	0.31	
Q96RY5	Hypothetical protein KIAA1426	1	0.17	
Q96CP7	Hypothetical protein LOC116238	1	6.23	
Q96BQ5	Hypothetical protein LOC133957 (Hypothetical protein FLJ25701)	1	0.35	
Q9BQ15	Hypothetical protein MGC2731	3	0.33	0.09
Q8NDZ4	Hypothetical protein MGC33365	1	0.65	
Q8IV33	Hypothetical protein MGC34713	1	0.12	
Q96AG4	Hypothetical protein PRO1855	7	0.26	0.14
Q8TC63	IGHG4 protein	1	0.32	
Q3ZCQ8	Import inner membrane translocase subunit TIM50, mitochondrial prec	2	2.06	1.73
Q14164	Inhibitor of nuclear factor kappa-B kinase epsilon subunit (EC 2.7.1.-) (1	0.19	
Q15070	Inner membrane protein OXA1L, mitochondrial precursor (Oxidase ass	2	0.47	0.03
Q14571	Inositol 1,4,5-trisphosphate receptor type 2 (Type 2 inositol 1,4,5-trisph	1	0.80	
Q14573	Inositol 1,4,5-trisphosphate receptor type 3 (Type 3 inositol 1,4,5-trisph	1	0.12	
Q8WYA0	Intraflagellar transport 81 (Carnitine deficiency-associated protein expr	1	3.09	
Q6UXX5	ITI-like protein (Inter-alpha (Globulin) inhibitor H5-like)	1	103.23	
Q8N5M9	Jagunal homolog 1	1	0.32	
Q6A163	Ka35 protein	2	0.71	0.14
Q86Z20	Kenae1	1	0.04	
Q14664	Keratin 10	1	0.17	
P02533	Keratin, type I cytoskeletal 14 (Cytokeratin-14) (CK-14) (Keratin-14) (K	1	0.81	
P08779	Keratin, type I cytoskeletal 16 (Cytokeratin-16) (CK-16) (Keratin-16) (K	1	1.19	
P05783	Keratin, type I cytoskeletal 18 (Cytokeratin-18) (CK-18) (Keratin-18) (K	15	0.35	0.25
P35527	Keratin, type I cytoskeletal 9 (Cytokeratin-9) (CK-9) (Keratin-9) (K9)	37	1.19	1.09

Acc #	Protein name	TPC	H/L ratio	SD
P04264	Keratin, type II cytoskeletal 1 (Cytokeratin-1) (CK-1) (Keratin-1) (K1) (6	77	1.16	0.60
Q7Z794	Keratin, type II cytoskeletal 1b	1	0.42	
P13647	Keratin, type II cytoskeletal 5 (Cytokeratin-5) (CK-5) (Keratin-5) (K5) (5	5	0.97	0.50
P08729	Keratin, type II cytoskeletal 7 (Cytokeratin-7) (CK-7) (Keratin-7) (K7) (S	6	1.44	2.41
P05787	Keratin, type II cytoskeletal 8 (Cytokeratin-8) (CK-8) (Keratin-8) (K8)	5	0.10	0.01
Q6PG45	Keratinocyte associated protein 2	1	0.62	
Q9C0D2	KIAA1731 protein (Fragment)	1	82.56	
Q96JH4	KIAA1853 protein (Fragment)	1	4.30	
Q86UP2	Kinectin (Kinesin receptor) (CG-1 antigen)	1	0.42	
Q14807	Kinesin-like protein KIF22 (Kinesin-like DNA-binding protein) (Kinesin-I	1	0.58	
Q16719	Kynureninase (EC 3.7.1.3) (L-kynurenine hydrolase)	1	0.22	
P48449	Lanosterol synthase (EC 5.4.99.7) (Oxidosqualene--lanosterol cyclase)	1	0.33	
Q01650	Large neutral amino acids transporter small subunit 1 (L-type amino ac	6	0.25	0.16
Q8TC61	Leucine zipper transcription factor-like 1	1	8.93	
O95202	Leucine zipper-EF-hand containing transmembrane protein 1, mitoch	1	0.27	
Q8NHJ6	Leukocyte immunoglobulin-like receptor subfamily B member 4 precu	1	1.03	
O00188	LLGL protein	1	0.52	
Q8N5G0	LOC389203 protein (Fragment)	1	0.17	
Q9BUI5	LOC51234 protein	1	0.54	
Q6PCB7	Long-chain fatty acid transport protein 1 (EC 6.2.1.-) (Fatty acid transp	1	0.10	
Q14108	Lysosome membrane protein II (LIMP II) (Scavenger receptor class B	2	0.25	0.08
P14174	Macrophage migration inhibitory factor (MIF) (Phenylpyruvate tautomer	1	0.41	
Q6UW68	MBC3205 (Hypothetical protein UNQ501)	1	0.47	
O95427	Mcd4p homolog	1	0.34	
O00264	Membrane associated progesterone receptor component 1 (mPR)	5	0.41	0.32
O15173	Membrane associated progesterone receptor component 2 (Progester	4	1.34	1.12
Q9UM00	Membrane protein (STMF151) (LOC54499) (Xenogeneic cross-immun	3	0.37	0.16
Q5BJH2	MGC13159 protein	1	0.28	
P10620	Microsomal glutathione S-transferase 1 (EC 2.5.1.18) (Microsomal GS	5	0.58	0.13
Q5J9I5	Migration-inducing protein 7	1	0.47	

Acc #	Protein name	TPC	H/L ratio	SD
Q8TCT9	Minor histocompatibility antigen H13 (EC 3.4.99.-) (Signal peptide pept	7	0.28	0.06
Q02978	Mitochondrial 2-oxoglutarate/malate carrier protein (OGCP) (Solute car	2	0.20	0.10
Q546F9	Mitochondrial aspartate-glutamate carrier protein	4	0.86	1.12
Q9Y6C9	Mitochondrial carrier homolog 2 (Met-induced mitochondrial protein)	10	0.22	0.14
Q9UBX3	Mitochondrial dicarboxylate carrier	2	0.14	0.15
O60830	Mitochondrial import inner membrane translocase subunit Tim17-B	1	0.31	
Q15388	Mitochondrial import receptor subunit TOM20 homolog (Mitochondrial	1	0.19	
Q9NS69	Mitochondrial import receptor subunit TOM22 homolog (Translocase of	1	0.15	
Q16891	Mitochondrial inner membrane protein (Mitofilin) (p87/89) (Proliferation	33	0.41	0.28
O94826	Mitochondrial precursor proteins import receptor (Translocase of outer	1	0.22	
Q5JPR4	Mitogen-activated protein kinase kinase kinase 15 (Fragment)	1	7.98	
Q9BVC6	Mitsugumin-23 precursor (Mg23)	3	0.50	0.38
P53985	Monocarboxylate transporter 1 (MCT 1)	6	0.69	0.02
O15427	Monocarboxylate transporter 4 (MCT 4) (MCT 3)	12	0.21	0.05
Q9H3T8	MOP-3	1	0.21	
Q8IZD4	mRNA decapping enzyme 1B (EC 3.-.-.-)	1	0.12	
P10244	Myb-related protein B (B-Myb)	1	0.18	
O43795	Myosin Ib (Myosin I alpha) (MMI-alpha) (MMIa) (MYH-1c)	4	2.69	3.65
Q4KMR3	Myosin IE	1	0.41	
P35580	Myosin-10 (Myosin heavy chain, nonmuscle IIb) (Nonmuscle myosin h	3	0.81	1.06
P35579	Myosin-9 (Myosin heavy chain, nonmuscle IIa) (Nonmuscle myosin he	14	0.41	0.17
Q13496	Myotubularin (EC 3.1.3.48)	1	0.56	
Q76KP1	N-acetylgalactosaminyltransferase	2	0.10	0.09
Q86SF2	N-acetylgalactosaminyltransferase 7 (EC 2.4.1.-) (Protein-UDP acetyl	1	0.09	
Q13423	NAD(P) transhydrogenase, mitochondrial precursor (EC 1.6.1.2) (Pyrid	1	0.53	
Q9NXA8	NAD-dependent deacetylase sirtuin-5 (EC 3.5.1.-) (SIR2-like protein 5)	1	0.86	
P00387	NADH-cytochrome b5 reductase (EC 1.6.2.2) (B5R) (Diaphorase-1) (C	3	0.14	0.07
O75489	NADH-ubiquinone oxidoreductase 30 kDa subunit, mitochondrial precu	1	0.53	
P49821	NADH-ubiquinone oxidoreductase 51 kDa subunit, mitochondrial precu	1	1.25	
P28331	NADH-ubiquinone oxidoreductase 75 kDa subunit, mitochondrial precu	1	0.46	

Acc #	Protein name	TPC	H/L ratio	SD
O43676	NADH-ubiquinone oxidoreductase B12 subunit (EC 1.6.5.3) (EC 1.6.99	1	0.49	
O95139	NADH-ubiquinone oxidoreductase B17 subunit (EC 1.6.5.3) (EC 1.6.99	1	0.17	
P03905	NADH-ubiquinone oxidoreductase chain 4 (EC 1.6.5.3) (NADH dehydr	1	0.20	
P03923	NADH-ubiquinone oxidoreductase chain 6 (EC 1.6.5.3) (NADH dehydr	1	0.31	
Q8WU60	NADH-ubiquinone oxidoreductase flavoprotein 3, isoform a	1	0.50	
P16435	NADPH--cytochrome P450 reductase (EC 1.6.2.4) (CPR) (P450R)	22	0.69	0.63
Q13765	Nascent polypeptide-associated complex alpha subunit (NAC-alpha) (1	0.67	
Q86UW6	NEDD4-binding protein 2 (EC 3.-.-) (N4BP2) (BCL-3-binding protein)	1	0.18	
Q9HCC7	NEDD4-like ubiquitin ligase 1	1	0.81	
O43402	Neighbor of COX4	1	0.26	
P32004	Neural cell adhesion molecule L1 precursor (N-CAM L1) (CD171 antig	3	0.37	0.21
P17677	Neuromodulin (Axonal membrane protein GAP-43) (Growth associated	1	2.99	
P30532	Neuronal acetylcholine receptor protein, alpha-5 subunit precursor	1	0.40	
Q14697	Neutral alpha-glucosidase AB precursor (EC 3.2.1.84) (Glucosidase II	1	0.49	
Q15758	Neutral amino acid transporter B(0) (ATB(0)) (Sodium-dependent neutr	6	0.65	0.28
Q9H2Q2	NGT	2	1.47	1.62
P29475	Nitric-oxide synthase, brain (EC 1.14.13.39) (NOS type I) (Neuronal N	1	23.46	
Q5JP08	Novel protein	1	0.22	
Q5SWH9	Novel protein	1	0.33	
Q5VZP5	Novel protein	1	0.24	
Q7LBC6	Nuclear protein 5qNCA	1	1.39	
Q9GZY0	Nuclear RNA export factor 2 (TAP-like protein 2) (TAPL-2)	2	0.01	0.00
P67809	Nuclease sensitive element binding protein 1 (Y-box binding protein 1)	1	0.82	
Q9NR30	Nucleolar RNA helicase 2 (EC 3.6.1.-) (Nucleolar RNA helicase II) (Nu	2	0.92	0.00
Q8NI70	OK/SW-CL.16	1	0.28	
O43869	Olfactory receptor 2T1 (Olfactory receptor 1-25) (OR1-25) (Olfactory re	1	0.05	
Q5TFD7	OTTHUMP00000016784	1	4.20	
Q5JUJ7	OTTHUMP00000017314 (Fragment)	1	0.25	
Q5VYA5	OTTHUMP00000017471	1	0.30	
Q4VXA4	OTTHUMP00000021035	1	0.24	

Acc #	Protein name	TPC	H/L ratio	SD
Q9NQJ4	OTTHUMP00000031372	1	0.53	
P56715	Oxygen-regulated protein 1 (Retinitis pigmentosa RP1 protein) (Retiniti	1	0.49	
P23284	Peptidyl-prolyl cis-trans isomerase B precursor (EC 5.2.1.8) (PPIase) (2	2.86	3.56
Q9Y3E5	Peptidyl-tRNA hydrolase 2, mitochondrial precursor (EC 3.1.1.29) (PT	1	0.29	
O95613	Pericentrin (Pericentrin B) (Kendrin)	1	0.65	
Q00325	Phosphate carrier protein, mitochondrial precursor (PTP)	13	0.35	0.11
O95674	Phosphatidate cytidyltransferase 2 (EC 2.7.7.41) (CDP-diglyceride sy	6	0.83	0.18
O00750	Phosphatidylinositol-4-phosphate 3-kinase C2 domain-containing beta	1	0.86	
P48651	Phosphatidylserine synthase 1 (EC 2.7.8.-) (PtdSer synthase 1) (PSS-	2	0.83	0.29
Q9BVG9	Phosphatidylserine synthase 2 (EC 2.7.8.-) (PtdSer synthase 2) (PSS-	4	0.39	0.04
Q16526	Photolyase (Cryptochrome 1)	6	0.18	0.02
Q4LE51	PIK3CA variant protein (Fragment)	1	0.21	
Q9H307	Pinin (140 kDa nuclear and cell adhesion-related phosphoprotein) (Do	6	0.60	0.39
P20020	Plasma membrane calcium-transporting ATPase 1 (EC 3.6.3.8) (PMC	1	0.24	
Q9Y4D7	Plexin D1 precursor	1	2.68	
Q9HB66	PNAS-117	2	0.37	0.03
Q6NZI2	Polymerase I and transcript release factor (PTRF protein)	1	0.15	
Q10471	Polypeptide N-acetylgalactosaminyltransferase 2 (EC 2.4.1.41) (Protei	2	0.62	0.64
Q96S06	Possible integral membrane	1	10.90	
Q9NS91	Postreplication repair protein RAD18 (hRAD18) (hHR18) (RING finger	1	0.11	
Q719H9	Potassium channel tetramerization domain-containing 1 (KCTD1 protei	1	0.22	
Q96MT3	Prickle-like protein 1 (REST/NRSF-interacting LIM domain protein 1)	1	1.17	
Q96CF8	PRKR interacting protein 1 (IL11 inducible)	1	1.05	
Q9H375	PRO0756	2	65.88	4.32
Q9HD20	Probable cation-transporting ATPase 13A1 (EC 3.6.3.-)	6	0.61	0.20
O94919	Probable endonuclease KIAA0830 precursor (EC 3.1.30.-)	2	0.48	0.00
Q8IWK6	Probable G-protein coupled receptor 125 precursor	1	0.15	
O96008	Probable mitochondrial import receptor subunit TOM40 homolog (Tran	7	0.44	0.48
Q9NPF4	Probable O-sialoglycoprotein endopeptidase (EC 3.4.24.57) (hOSGEP	1	1.62	
Q8IUH4	Probable palmitoyltransferase ZDHHC13 (EC 2.3.1.-) (Zinc finger DHH	1	0.03	

Acc #	Protein name	TPC	H/L ratio	SD
Q9H6R6	Probable palmitoyltransferase ZDHHC6 (EC 2.3.1.-) (Zinc finger DHH	1	0.16	
Q8WUM4	Programmed cell death 6-interacting protein (PDCD6-interacting protei	1	0.27	
P35232	Prohibitin	25	0.42	0.25
Q99623	Prohibitin-2 (B-cell receptor-associated protein BAP37) (Repressor of	44	0.62	0.37
Q9HCU5	Prolactin regulatory element-binding protein (Mammalian guanine nucl	1	1.02	
O14684	Prostaglandin E synthase (EC 5.3.99.3) (Microsomal glutathione S-tra	2	0.28	0.26
Q9NZ45	Protein C10orf70	1	0.78	
Q6P0N0	Protein C14orf106 (P243)	3	0.40	0.45
Q8IVT2	Protein C19orf21	3	0.50	0.16
Q8NC44	Protein C2orf17	1	0.44	
Q9H330	Protein C9orf5 (Protein CG-2)	1	0.08	
Q96CS3	Protein expressed in T-cells and eosinophils in atopic dermatitis (ETE	2	0.21	0.05
Q14165	Protein KIAA0152 precursor	2	0.32	0.01
O43156	Protein KIAA0406	1	0.66	
O94927	Protein KIAA0841	1	3.39	
Q9UPV7	Protein KIAA1045	1	0.10	
Q96JH8	Protein KIAA1849	1	1.34	
Q02156	Protein kinase C, epsilon type (EC 2.7.1.-) (nPKC-epsilon)	1	0.21	
Q15139	Protein kinase, D1 type (EC 2.7.1.-) (nPKC-D1) (Protein kinase D) (Pr	1	10.13	
Q86UE4	Protein LYRIC (Lysine-rich CEACAM1 co-isolated protein) (3D3/lyric) (1	0.79	
P61619	Protein transport protein Sec61 alpha subunit isoform 1 (Sec61 alpha-	3	0.23	0.13
P60468	Protein transport protein Sec61 beta subunit	12	0.46	0.93
Q6Y1H2	Protein tyrosine phosphatase-like member b (Protein tyrosine phospho	1	0.15	
O60725	Protein-S-isoprenylcysteine O-methyltransferase (EC 2.1.1.100) (Isopr	1	0.10	
Q9Y5E6	Protocadherin beta 3 precursor (PCDH-beta3)	1	0.44	
Q8TB72	Pumilio homolog 2 (Pumilio-2)	1	0.53	
Q96QD8	Putative 40-9-1 protein	3	1.40	1.39
Q8IXJ9	Putative Polycomb group protein ASXL1 (Additional sex combs-like pr	2	0.14	0.04
Q9Y4G8	Rap guanine nucleotide exchange factor 2 (PDZ domain containing gu	1	1.42	
P46940	Ras GTPase-activating-like protein IQGAP1 (p195)	1	0.35	

Acc #	Protein name	TPC	H/L ratio	SD
P61026	Ras-related protein Rab-10	1	0.52	
P61106	Ras-related protein Rab-14	2	1.55	1.83
P62820	Ras-related protein Rab-1A (YPT1-related protein)	1	0.44	
Q9H0U4	Ras-related protein Rab-1B	1	0.17	
P61019	Ras-related protein Rab-2A	3	0.54	0.19
Q9BZG1	Ras-related protein Rab-34 (Rab-39) (Ras-related protein Rah)	1	0.18	
P51148	Ras-related protein Rab-5C (RAB5L) (L1880)	4	0.31	0.10
P51149	Ras-related protein Rab-7	17	0.98	1.01
P61006	Ras-related protein Rab-8A (Oncogene c-mel)	2	0.42	0.12
P51151	Ras-related protein Rab-9A (Rab-9)	1	0.23	
Q6YNB5	Relaxin-like factor b	1	0.12	
O95811	Retinoblastoma binding protein 2 homolog 1	1	0.10	
Q8NFJ5	Retinoic acid-induced protein 3 (G-protein coupled receptor family C gr	12	0.78	0.37
Q8IUC4	Rhopilin-2 (GTP-Rho binding protein 2) (76 kDa RhoB effector protein	1	0.04	
P84095	Rho-related GTP-binding protein RhoG	1	0.53	
Q9P2E9	Ribosome-binding protein 1 (Ribosome receptor protein) (180 kDa ribo	1	0.80	
Q9NW13	RNA-binding protein 28 (RNA-binding motif protein 28)	1	0.15	
Q15287	RNA-binding protein with serine-rich domain 1 (SR-related protein LDC	1	0.28	
Q49A11	RPS27L protein	1	0.13	
Q15413	Ryanodine receptor 3 (Brain-type ryanodine receptor) (RyR3) (RYR-3)	2	1.10	1.34
P16615	Sarcoplasmic/endoplasmic reticulum calcium ATPase 2 (EC 3.6.3.8) (18	0.39	0.42
Q8WTV0	Scavenger receptor class B member 1 (SRB1) (SR-BI) (CD36 antigen-	3	0.45	0.10
O75880	SCO1 protein homolog, mitochondrial precursor	1	0.55	
Q4VX17	Secretory carrier membrane protein 3	1	0.30	
Q14674	Separin (EC 3.4.22.49) (Separase) (Caspase-like protein ESPL1) (Extr	1	0.36	
Q13501	Sequestosome-1 (Phosphotyrosine-independent ligand for the Lck SH	2	0.68	0.00
P48595	Serpin B10 (Bomapin) (Protease inhibitor 10)	1	0.44	
Q52WX2	SH3-binding kinase	1	0.18	
Q9H9B4	Sideroflexin-1 (Tricarboxylate carrier protein) (TCC)	6	0.77	0.40
Q9Y6A9	Signal peptidase complex subunit 1 (EC 3.4.-.-) (Microsomal signal pe	1	0.16	

Acc #	Protein name	TPC	H/L ratio	SD
Q15005	Signal peptidase complex subunit 2 (EC 3.4.-.-) (Microsomal signal pe	6	0.45	0.21
P61009	Signal peptidase complex subunit 3 (EC 3.4.-.-) (Microsomal signal pe	5	0.33	0.23
Q9Y5M8	Signal recognition particle receptor beta subunit (SR-beta) (Protein AP	4	0.32	0.10
P40763	Signal transducer and activator of transcription 3 (Acute-phase respon	2	0.18	0.05
Q9BTI6	Similar to flotillin 2	1	0.38	
Q8N5K1	Similar to mouse 1500009M05Rik protein	4	0.42	0.29
Q04837	Single-stranded DNA-binding protein, mitochondrial precursor (Mt-SSB	1	0.68	
Q96EQ0	Small glutamine-rich tetratricopeptide repeat-containing protein B (Sm	1	0.64	
Q96PI1	Small proline-rich protein 4	1	0.16	
P05023	Sodium/potassium-transporting ATPase alpha-1 chain precursor (EC 3	4	0.46	0.22
P14415	Sodium/potassium-transporting ATPase beta-2 chain (Sodium/potassi	1	0.09	
P54709	Sodium/potassium-transporting ATPase beta-3 chain (Sodium/potassi	2	0.26	0.20
Q9Y289	Sodium-dependent multivitamin transporter (Na(+)-dependent multivita	2	0.56	0.28
P11166	Solute carrier family 2, facilitated glucose transporter member 1 (Gluc	2	0.62	0.22
Q6NUK1	Solute carrier family 25 member 24, isoform 1	1	0.23	
Q9Y512	Sorting and assembly machinery component 50 homolog	10	0.41	0.31
Q8TCJ2	Source of immunodominant MHC-associated peptides	6	0.35	0.25
Q13813	Spectrin alpha chain, brain (Spectrin, non-erythroid alpha chain) (Alph	26	0.45	0.47
Q01082	Spectrin beta chain, brain 1 (Spectrin, non-erythroid beta chain 1) (Bet	2	0.29	0.03
O75477	SPFH domain protein 1 precursor (Protein KE04)	1	0.98	
O94905	SPFH domain protein 2 precursor	4	0.41	0.08
Q9NRA0	Sphingosine kinase 2 (EC 2.7.1.-) (SK 2) (SPK 2)	1	0.26	
Q9UHA8	Splicing coactivator subunit SRm300	4	0.54	0.22
Q9NY15	Stabilin-1 precursor (FEEL-1 protein) (MS-1 antigen)	1	149.47	
P52823	Stanniocalcin-1 precursor (STC-1)	1	0.01	
Q53GQ0	Steroid dehydrogenase homolog (Fragment)	1	0.30	
Q9UJZ1	Stomatin-like protein 2 (SLP-2) (EPB72-like 2)	1	0.52	
P38646	Stress-70 protein, mitochondrial precursor (75 kDa glucose regulated	1	0.31	
O95425	Supervillin (Archvillin) (p205/p250)	3	0.39	0.31
Q9NZ01	Synaptic glycoprotein SC2	3	0.41	0.27

Acc #	Protein name	TPC	H/L ratio	SD
Q5UIP0	Telomere-associated protein RIF1 (Rap1-interacting factor 1 homolog)	1	0.07	
Q15006	Tetratricopeptide repeat protein KIAA0103	3	0.55	0.14
P37173	TGF-beta receptor type II precursor (EC 2.7.1.37) (TGFR-2) (TGF-beta	2	0.15	0.07
Q9H0W7	THAP domain-containing protein 2	5	0.30	0.09
Q9H3N1	Thioredoxin domain-containing protein 1 precursor (Transmembrane T	1	0.30	
Q6F5E7	Thioredoxin reductase 2 intronic transcript 1	1	0.22	
Q9Y2X0	Thyroid hormone receptor-associated protein complex 95 kDa compon	1	0.16	
Q15643	Thyroid receptor-interacting protein 11 (TRIP-11) (Golgi-associated mi	1	0.19	
O00463	TNF receptor-associated factor 5 (RING finger protein 84)	1	0.27	
Q969E4	Transcription elongation factor A (SII)-like 3 (MSTP072) (Hypothetical	1	0.37	
Q15369	Transcription elongation factor B polypeptide 1 (RNA polymerase II tra	1	0.46	
Q9UGU0	Transcription factor 20 (Stromelysin 1 PDGF-responsive element-bindi	1	2.37	
Q99551	Transcription termination factor, mitochondrial precursor (mTERF) (Mit	1	0.44	
P02786	Transferrin receptor protein 1 (TfR1) (TR) (TfR) (Trfr) (CD71 antigen) (11	0.78	0.48
Q9Y4A5	Transformation/transcription domain-associated protein (350/400 kDa	1	0.88	
Q15629	Translocation associated membrane protein 1	5	0.50	0.27
Q9UGP8	Translocation protein SEC63 homolog	2	2.08	0.00
P43307	Translocon-associated protein alpha subunit precursor (TRAP-alpha) (1	1.33	
P51571	Translocon-associated protein delta subunit precursor (TRAP-delta) (S	5	0.34	0.08
Q9UNL2	Translocon-associated protein gamma subunit (TRAP-gamma) (Signal	2	0.18	0.06
Q99805	Transmembrane 9 superfamily protein member 2 precursor (p76)	1	0.65	
P84157	Transmembrane anchor protein 1	3	0.36	0.29
P49755	Transmembrane emp24 domain containing protein 10 precursor (Tran	13	0.82	1.13
Q7Z7H5	Transmembrane emp24 domain containing protein 4 precursor	4	0.67	0.02
Q9Y3A6	Transmembrane emp24 domain containing protein 5 precursor	1	0.23	
Q9BVK6	Transmembrane emp24 domain containing protein 9 precursor (Glyco	2	0.48	0.57
P17152	Transmembrane protein 11 (Protein PM1)	2	2.02	1.13
P57088	Transmembrane protein 33 (DB83 protein)	8	0.44	0.18
Q96GC9	Transmembrane protein 49 (TDC1)	1	0.41	
Q9HC07	Transmembrane protein PT27 (TPA regulated locus)	5	0.39	0.18

Acc #	Protein name	TPC	H/L ratio	SD
P53007	Tricarboxylate transport protein, mitochondrial precursor (Citrate trans	4	0.53	0.17
Q8N6D1	Triggering receptor expressed on myeloid cells 5	1	0.23	
Q9UDY6	Tripartite motif protein 10 (RING finger protein 9) (B30-RING finger pro	1	0.74	
Q6UWB4	Tryptophan/serine protease	1	0.68	
P68366	Tubulin alpha-1 chain (Alpha-tubulin 1) (Testis-specific alpha-tubulin) (2	0.22	0.08
Q14679	Tubulin tyrosine ligase-like protein 4	1	0.34	
P23510	Tumor necrosis factor ligand superfamily member 4 (OX40 ligand) (OX	1	1.83	
Q96RJ3	Tumor necrosis factor receptor superfamily member 13C (B cell-activa	1	0.65	
Q561W0	TXNDC14 protein	5	0.33	0.13
Q96PE3	Type I inositol-3,4-bisphosphate 4-phosphatase (EC 3.1.3.66) (Inositol	3	0.19	0.05
Q6RW13	Type-1 angiotensin II receptor-associated protein (AT1 receptor-associ	1	0.94	
P18031	Tyrosine-protein phosphatase, non-receptor type 1 (EC 3.1.3.48) (Prot	2	0.29	0.27
P29074	Tyrosine-protein phosphatase, non-receptor type 4 (EC 3.1.3.48) (Prot	1	0.57	
O75643	U5 small nuclear ribonucleoprotein 200 kDa helicase (EC 3.6.1.-) (U5	1	0.04	
Q9UDW1	Ubiquinol-cytochrome c reductase complex 7.2 kDa protein (EC 1.10.2	1	3.46	
O14949	Ubiquinol-cytochrome c reductase complex ubiquinone-binding protein	3	0.31	0.27
P22695	Ubiquinol-cytochrome-c reductase complex core protein 2, mitochondri	10	0.61	0.16
P31930	Ubiquinol-cytochrome-c reductase complex core protein I, mitochondri	6	0.62	0.56
Q9Y6I4	Ubiquitin carboxyl-terminal hydrolase 3 (EC 3.1.2.15) (Ubiquitin thiolest	1	0.18	
Q86T82	Ubiquitin carboxyl-terminal hydrolase 37 (EC 3.1.2.15) (Ubiquitin thiole	1	15.67	
P45974	Ubiquitin carboxyl-terminal hydrolase 5 (EC 3.1.2.15) (Ubiquitin thiolest	1	1.60	
Q96EP1	Ubiquitin ligase protein CHFR (EC 6.3.2.-) (Checkpoint with forkhead a	1	1.38	
Q86YT6	Ubiquitin ligase protein MIB1 (EC 6.3.2.-) (Mind bomb homolog 1) (DA	1	0.65	
Q9Y385	Ubiquitin-conjugating enzyme E2 J1 (EC 6.3.2.19) (Non-canonical ubiq	1	0.27	
O00124	UBX domain-containing protein 6 (Rep-8 protein) (Reproduction 8 prot	1	0.64	
Q9Y3E0	UPF0198 protein CGI-141	1	0.31	
Q9P0I2	UPF0347 protein LOC55831	2	0.27	0.11
P61421	Vacuolar ATP synthase subunit d (EC 3.6.3.14) (V-ATPase d subunit)	3	0.21	0.14
Q93050	Vacuolar proton translocating ATPase 116 kDa subunit a isoform 1 (V-	2	1.16	0.06
O75396	Vesicle trafficking protein SEC22b (SEC22 vesicle trafficking protein-li	18	0.53	0.31

Acc #	Protein name	TPC	H/L ratio	SD
Q9P0L0	Vesicle-associated membrane protein-associated protein A (VAMP-as	5	0.30	0.13
Q12907	Vesicular integral-membrane protein VIP36 precursor (GP36b glycopr	2	0.39	0.00
P08670	Vimentin	3	0.35	0.06
Q9BQB6	Vitamin K epoxide reductase complex subunit 1 (EC 1.1.4.1) (Vitamin	1	0.37	
P21796	Voltage-dependent anion-selective channel protein 1 (VDAC-1) (hVDA	32	0.60	0.33
Q9Y277	Voltage-dependent anion-selective channel protein 3 (VDAC-3) (hVDA	14	0.54	0.93
Q96DN5	WD-repeat protein 67	1	0.04	
P30291	Wee1-like protein kinase (EC 2.7.1.112) (WEE1hu)	1	0.07	
P56705	Wnt-4 protein precursor	1	0.27	
P49750	YLP motif containing protein 1 (Nuclear protein ZAP3) (ZAP113)	1	105.27	
Q96GI9	ZDHHC8 protein	1	0.08	
Q7Z2W4	Zinc finger CCCH type antiviral protein 1 (Zinc finger CCCH type doma	1	1.06	
Q7Z3T8	Zinc finger FYVE domain-containing protein 16 (Endofin) (Endosome-a	1	0.11	
O14978	Zinc finger protein 263 (Zinc finger protein FPM315)	1	0.47	
Q96RE9	Zinc finger protein 300	1	0.10	
Q96PQ6	Zinc finger protein 317	1	20.40	
Q9Y2H8	Zinc finger protein 510	1	0.02	
Q96ME7	Zinc finger protein 512	1	0.27	
Q9Y6M5	Zinc transporter 1 (ZnT-1) (Solute carrier family 30 member 1)	1	0.30	