

Supplemental Table S-2b. A list of 285 proteins quantified by present algorithm, sorted by protein name.

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
Q99460	26S proteasome non-ATPase regulatory subunit 1 (26S prot	2	0.708	1.195	0.92
P62269	40S ribosomal protein S18 (Ke-3) (Ke3)	2	0.49	0.571	0.529
P08195	4F2 cell-surface antigen heavy chain (4F2hc) (Lymphocyte a	44	0.129	4.31	0.731
Q07020	60S ribosomal protein L18	2	0.926	1.487	1.173
Q02543	60S ribosomal protein L18a	1	0.362	0.362	0.362
P62829	60S ribosomal protein L23 (Ribosomal protein L17)	1	1.105	1.105	1.105
P62899	60S ribosomal protein L31	1	0.933	0.933	0.933
P83881	60S ribosomal protein L36a (60S ribosomal protein L44) (Mig	1	1.149	1.149	1.149
P36578	60S ribosomal protein L4 (L1)	1	0.749	0.749	0.749
O00400	Acetyl-coenzyme A transporter 1 (AT-1) (Acetyl-CoA transpor	1	0.605	0.605	0.605
Q8TB61	Adenosine 3'-phospho 5'-phosphosulfate transporter 1 (PAP	3	0.482	1.76	0.785
P12235	ADP/ATP translocase 1 (Adenine nucleotide translocator 1) (1	0.654	0.654	0.654
P05141	ADP/ATP translocase 2 (Adenine nucleotide translocator 2) (6	0.286	1.105	0.462
O00116	Alkyldihydroxyacetonephosphate synthase, peroxisomal prec	1	1.174	1.174	1.174
Q9H6U8	Alpha-1,2-mannosyltransferase ALG9 (EC 2.4.1.-) (Asparagi	1	0.293	0.293	0.293
Q15911	Alpha-fetoprotein enhancer binding protein (AT motif-binding	1	1.926	1.926	1.926
Q9Y4J0	Aspartyl(Asparaginy)l)beta-hydroxylase	1	0.573	0.573	0.573
P25705	ATP synthase alpha chain, mitochondrial precursor (EC 3.6.3	11	0.385	1.906	0.78
P06576	ATP synthase beta chain, mitochondrial precursor (EC 3.6.3.	5	0.44	1.247	0.795
P18859	ATP synthase coupling factor 6, mitochondrial precursor (EC	1	0.912	0.912	0.912
O75947	ATP synthase D chain, mitochondrial (EC 3.6.3.14)	3	0.11	0.757	0.286
P30049	ATP synthase delta chain, mitochondrial precursor (EC 3.6.3.	1	1.069	1.069	1.069
O75964	ATP synthase g chain, mitochondrial (EC 3.6.3.14) (ATPase	2	0.439	0.7	0.554
P36542	ATP synthase gamma chain, mitochondrial precursor (EC 3.	2	0.122	0.244	0.173
P48047	ATP synthase O subunit, mitochondrial precursor (EC 3.6.3.1	4	0.152	1.328	0.545
Q8NBU5	ATPase family AAA domain containing protein 1	1	0.107	0.107	0.107
P33897	ATP-binding cassette sub-family D member 1 (Adrenoleukod	3	0.311	0.707	0.475
P28288	ATP-binding cassette sub-family D member 3 (70 kDa peroxi	3	0.531	2.046	0.97

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
Q9UNQ0	ATP-binding cassette sub-family G member 2 (Placenta-spec	8	0.129	1.396	0.506
Q9NT62	Autophagy protein 3-like (APG3-like) (hApp3) (PC3-96 protei	1	0.831	0.831	0.831
O95816	BAG family molecular chaperone regulator 2 (BCL2-associat	2	1.303	1.825	1.542
Q9BXC9	Bardet-Biedl syndrome 2 protein	1	0.173	0.173	0.173
P35613	Basigin precursor (CD147 antigen) (Leukocyte activation anti	3	0.506	1.001	0.661
Q9C0K0	B-cell lymphoma/leukemia 11B (B-cell CLL/lymphoma 11B) (1	0.404	0.404	0.404
Q9UHQ4	B-cell receptor-associated protein 29 (BCR-associated protei	2	0.425	0.621	0.514
P51572	B-cell receptor-associated protein 31 (BCR-associated protei	5	0.361	1.583	0.838
P61769	Beta-2-microglobulin precursor [Contains: Beta-2-microglobul	1	0.845	0.845	0.845
P80723	Brain acid soluble protein 1 (BASP1 protein) (Neuronal axon	4	0.122	0.749	0.361
Q9Y6D6	Brefeldin A-inhibited guanine nucleotide-exchange protein 1 (1	1.245	1.245	1.245
Q15800	C-4 methylsterol oxidase (EC 1.14.13.72) (Methylsterol mono	1	0.485	0.485	0.485
O75844	CAAX prenyl protease 1 homolog (EC 3.4.24.84) (Prenyl prot	2	0.654	0.81	0.728
P98194	Calcium-transporting ATPase type 2C member 1 (EC 3.6.3.8	1	1.027	1.027	1.027
P27824	Calnexin precursor (Major histocompatibility complex class I	1	0.509	0.509	0.509
Q9GZY7	CAMP/cGMP cyclic nucleotide phosphodiesterase 11A3	1	0.941	0.941	0.941
P31327	Carbamoyl-phosphate synthase [ammonia], mitochondrial pr	1	0.722	0.722	0.722
P50416	Carnitine O-palmitoyltransferase I, mitochondrial liver isoform	1	0.495	0.495	0.495
P13987	CD59 glycoprotein precursor (Membrane attack complex inhi	1	1.979	1.979	1.979
Q99459	Cell division cycle 5-like protein (Cdc5-like protein) (Pombe c	1	0.431	0.431	0.431
Q15057	Centaurin-beta 2 (Cnt-b2)	1	0.207	0.207	0.207
O00590	Chemokine-binding protein 2 (Chemokine-binding protein D6	1	2.239	2.239	2.239
Q5TF85	Chromosome 6 open reading frame 37	1	5.137	5.137	5.137
Q00610	Clathrin heavy chain 1 (CLH-17)	2	1.088	2.652	1.699
Q5W1L8	CMP-sialic acid transporter	1	1.687	1.687	1.687
Q9NX63	Coiled-coil-helix-coiled-coil-helix domain containing protein 3	4	0.588	1.382	0.865
Q02388	Collagen alpha 1(VII) chain precursor (Long-chain collagen) (1	0.465	0.465	0.465
Q07021	Complement component 1, Q subcomponent binding protein,	3	0.191	0.706	0.313
Q9NS00	Core1 UDP-galactose:N-acetylgalactosamine-alpha-R beta 1	1	0.236	0.236	0.236
P20674	Cytochrome c oxidase polypeptide Va, mitochondrial precurs	1	0.866	0.866	0.866
P10606	Cytochrome c oxidase polypeptide Vb, mitochondrial precurs	3	0.537	1.274	0.733

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
P09669	Cytochrome c oxidase polypeptide VIc precursor (EC 1.9.3.1)	3	0.574	0.716	0.659
P14406	Cytochrome c oxidase polypeptide VIIa-liver/heart, mitochon	2	0.715	1.636	1.082
P13073	Cytochrome c oxidase subunit IV isoform 1, mitochondrial pr	3	0.562	0.974	0.781
O14548	Cytochrome c oxidase subunit VIIa-related protein, mitochon	1	0.271	0.271	0.271
P08574	Cytochrome c1, heme protein, mitochondrial precursor (Cyto	3	0.159	0.959	0.494
P53701	Cytochrome c-type heme lyase (EC 4.4.1.17) (CCHL) (Holoc	1	0.511	0.511	0.511
Q96P26	Cytosolic 5'-nucleotidase 1B (EC 3.1.3.5) (Cytosolic 5'-nucleo	1	0.904	0.904	0.904
P61803	Defender against cell death 1 (DAD-1)	1	0.164	0.164	0.164
Q96HY7	Dehydrogenase E1 and transketolase domain containing prot	1	1.612	1.612	1.612
Q8NBQ5	Dehydrogenase/reductase SDR family member 8 precursor (2	0.276	0.687	0.435
Q8TB45	DEP domain containing 6	1	0.844	0.844	0.844
P81605	Dermcidin precursor (Preproteolysin) [Contains: Survival-pro	1	1.183	1.183	1.183
P15924	Desmoplakin (DP) (250/210 kDa paraneoplastic pemphigus	1	2.116	2.116	2.116
P04053	DNA nucleotidylexotransferase (EC 2.7.7.31) (Terminal additi	1	0.363	0.363	0.363
Q5T219	DnaJ (Hsp40) homolog, subfamily D, member 1 (Growth-inhi	1	0.676	0.676	0.676
Q5VWA5	Dolichyl-diphosphooligosaccharide-protein glycosyltransferas	5	0.471	1.745	0.771
P04843	Dolichyl-diphosphooligosaccharide--protein glycosyltransfera	23	0.18	2.554	0.534
Q92685	Dolichyl-P-Man:Man(5)GlcNAc(2)-PP-dolichyl mannosyltrans	2	0.68	1.521	1.017
O75691	Down-regulated in metastasis protein (Key-1A6 protein) (Nov	1	0.377	0.377	0.377
Q6UXB4	DTTR431 (C-type lectin superfamily 4, member G)	1	48.218	48.218	48.218
Q13409	Dynein intermediate chain 2, cytosolic (DH IC-2) (Cytoplasmic	1	1.212	1.212	1.212
P49411	Elongation factor Tu, mitochondrial precursor (EF-Tu) (P43)	1	0.781	0.781	0.781
Q9UHB6	Epithelial protein lost in neoplasm	1	1.025	1.025	1.025
P27105	Erythrocyte band 7 integral membrane protein (Stomatin) (Pr	1	0.647	0.647	0.647
Q99613	Eukaryotic translation initiation factor 3 subunit 8 (eIF3 p110)	1	7.83	7.83	7.83
Q01469	Fatty acid-binding protein, epidermal (E-FABP) (Psoriasis-as	1	2.447	2.447	2.447
Q9Y5Y0	Feline leukemia virus subgroup C receptor-related protein 1 (1	0.352	0.352	0.352
Q9BYX7	FKSG30	1	181.065	181.065	181.065
Q8TEP9	FLJ00144 protein (Fragment)	1	0.105	0.105	0.105
O75955	Flotillin-1	1	0.418	0.418	0.418
Q14254	Flotillin-2 (Epidermal surface antigen) (ESA)	3	0.796	8.713	2.703

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
Q71RG4	FP2653 (ELSD1897) (MGC3123 protein)	1	0.748	0.748	0.748
Q6ZV73	FYVE, RhoGEF and PH domain containing protein 6 (Zinc fin	1	1.949	1.949	1.949
O95452	Gap junction beta-6 protein (Connexin-30) (Cx30)	1	0.378	0.378	0.378
Q6P1A2	Gene rich cluster, C3f gene	1	0.555	0.555	0.555
P48506	Glutamate--cysteine ligase catalytic subunit (EC 6.3.2.2) (Ga	1	0.786	0.786	0.786
Q5JR58	Glyceraldehyde 3-phosphate dehydrogenase (GAPDH)	1	0.322	0.322	0.322
P04406	Glyceraldehyde-3-phosphate dehydrogenase, liver (EC 1.2.1	1	2.726	2.726	2.726
Q9UBI6	Guanine nucleotide-binding protein G(l)/G(S)/G(O) gamma-1	1	0.169	0.169	0.169
P08754	Guanine nucleotide-binding protein G(k), alpha subunit (G(i)	1	1.08	1.08	1.08
O95837	Guanine nucleotide-binding protein, alpha-14 subunit (G-prot	1	0.292	0.292	0.292
P11142	Heat shock cognate 71 kDa protein (Heat shock 70 kDa prot	2	0.275	0.921	0.503
P04792	Heat-shock protein beta-1 (HspB1) (Heat shock 27 kDa prote	1	1.225	1.225	1.225
P62805	Histone H4	11	0.208	1.305	0.782
GAG_Prote	HIV GAG protein	11	4.06	27.479	10.042
Q8MH63	HLAT1-3TM	1	0.795	0.795	0.795
P57058	Hormonally up-regulated neu tumor-associated kinase (EC 2.	1	1.266	1.266	1.266
Q9P035	HSPC121	3	0.475	0.93	0.6
Q4G0N0	Hypothetical protein	1	1.443	1.443	1.443
Q6PQ33	Hypothetical protein	1	1.18	1.18	1.18
Q8Nfq8	Hypothetical protein (Novel protein)	1	0.894	0.894	0.894
Q8N1H7	Hypothetical protein C14orf39	1	1.165	1.165	1.165
Q5H9S5	Hypothetical protein DKFZp313B047	1	0.202	0.202	0.202
Q9NVA4	Hypothetical protein FLJ10846	1	0.473	0.473	0.473
Q96A33	Hypothetical protein FLJ14938 (Hypothetical protein NT2RP2	1	0.557	0.557	0.557
Q6ZN37	Hypothetical protein FLJ16464	1	0.287	0.287	0.287
Q9NX61	Hypothetical protein FLJ20422 (AVLG582)	1	1.635	1.635	1.635
Q9NX40	Hypothetical protein FLJ20455 (TPA018) (Ovarian carcinoma	1	0.293	0.293	0.293
Q8N8I4	Hypothetical protein FLJ39430	1	0.321	0.321	0.321
Q6ZQW6	Hypothetical protein FLJ46844	1	1.135	1.135	1.135
Q71M32	Hypothetical protein FP14381	1	0.453	0.453	0.453
Q96RY5	Hypothetical protein KIAA1426	1	0.373	0.373	0.373

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
Q96BQ5	Hypothetical protein LOC133957 (Hypothetical protein FLJ25	1	0.506	0.506	0.506
Q9BQ15	Hypothetical protein MGC2731	3	0.32	0.621	0.491
Q96AG4	Hypothetical protein PRO1855	1	0.649	0.649	0.649
Q8TC63	IGHG4 protein	1	1.219	1.219	1.219
Q15070	Inner membrane protein OXA1L, mitochondrial precursor (Ox	1	0.987	0.987	0.987
Q14571	Inositol 1,4,5-trisphosphate receptor type 2 (Type 2 inositol 1,	1	1.013	1.013	1.013
Q8WYA0	Intraflagellar transport 81 (Carnitine deficiency-associated pr	1	5.761	5.761	5.761
Q6A163	Ka35 protein	1	1.109	1.109	1.109
P05783	Keratin, type I cytoskeletal 18 (Cytokeratin-18) (CK-18) (Kera	10	0.262	1.245	0.492
P35527	Keratin, type I cytoskeletal 9 (Cytokeratin-9) (CK-9) (Keratin-	19	0.5	8.506	2.043
P04264	Keratin, type II cytoskeletal 1 (Cytokeratin-1) (CK-1) (Keratin-	41	0.634	4.62	1.88
Q7Z794	Keratin, type II cytoskeletal 1b	1	0.94	0.94	0.94
P13647	Keratin, type II cytoskeletal 5 (Cytokeratin-5) (CK-5) (Keratin-	3	1.492	2.501	1.784
P08729	Keratin, type II cytoskeletal 7 (Cytokeratin-7) (CK-7) (Keratin-	2	0.618	1.228	0.871
P05787	Keratin, type II cytoskeletal 8 (Cytokeratin-8) (CK-8) (Keraton	1	0.218	0.218	0.218
Q6PG45	Keratinocyte associated protein 2	1	0.785	0.785	0.785
P48449	Lanosterol synthase (EC 5.4.99.7) (Oxidosqualene--lanoster	1	0.636	0.636	0.636
Q01650	Large neutral amino acids transporter small subunit 1 (L-type	1	0.283	0.283	0.283
O95202	Leucine zipper-EF-hand containing transmembrane protein 1	1	0.483	0.483	0.483
Q14108	Lysosome membrane protein II (LIMP II) (Scavenger recepto	1	0.205	0.205	0.205
O95427	Mcd4p homolog	1	0.59	0.59	0.59
O00264	Membrane associated progesterone receptor component 1 (2	0.204	0.602	0.35
O15173	Membrane associated progesterone receptor component 2 (3	0.45	1.333	0.746
Q9UM00	Membrane protein (STMF151) (LOC54499) (Xenogeneic cro	2	0.39	0.761	0.545
Q5BJH2	MGC13159 protein	1	0.89	0.89	0.89
P10620	Microsomal glutathione S-transferase 1 (EC 2.5.1.18) (Micro	2	0.837	1.127	0.971
Q8TCT9	Minor histocompatibility antigen H13 (EC 3.4.99.-) (Signal pe	2	0.454	0.531	0.491
Q02978	Mitochondrial 2-oxoglutarate/malate carrier protein (OGCP) (1	0.287	0.287	0.287
Q546F9	Mitochondrial aspartate-glutamate carrier protein	2	0.524	0.643	0.58
Q9Y6C9	Mitochondrial carrier homolog 2 (Met-induced mitochondrial	6	0.161	0.413	0.273
Q9UBX3	Mitochondrial dicarboxylate carrier	1	0.53	0.53	0.53

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
O60830	Mitochondrial import inner membrane translocase subunit Ti	1	1.14	1.14	1.14
Q15388	Mitochondrial import receptor subunit TOM20 homolog (Mito	1	0.524	0.524	0.524
Q16891	Mitochondrial inner membrane protein (Mitofilin) (p87/89) (Pr	15	0.299	2.215	0.76
Q9BVC6	Mitsugumin-23 precursor (Mg23)	1	0.71	0.71	0.71
P53985	Monocarboxylate transporter 1 (MCT 1)	3	0.747	0.882	0.83
O15427	Monocarboxylate transporter 4 (MCT 4) (MCT 3)	7	0.171	0.716	0.361
Q8IZD4	mRNA decapping enzyme 1B (EC 3.-.-.)	1	0.312	0.312	0.312
P10244	Myb-related protein B (B-Myb)	1	0.403	0.403	0.403
O43795	Myosin Ib (Myosin I alpha) (MMI-alpha) (MMIa) (MYH-1c)	2	0.784	0.933	0.855
Q4KMR3	Myosin IE	1	0.582	0.582	0.582
P35580	Myosin-10 (Myosin heavy chain, nonmuscle IIb) (Nonmuscle	2	1.13	1.147	1.138
P35579	Myosin-9 (Myosin heavy chain, nonmuscle IIa) (Nonmuscle	7	0.259	1.779	0.635
Q13496	Myotubularin (EC 3.1.3.48)	1	0.606	0.606	0.606
Q76KP1	N-acetylgalactosaminyltransferase	1	0.228	0.228	0.228
Q13423	NAD(P) transhydrogenase, mitochondrial precursor (EC 1.6.	1	0.408	0.408	0.408
Q9NXA8	NAD-dependent deacetylase sirtuin-5 (EC 3.5.1.-) (SIR2-like	1	1.065	1.065	1.065
P28331	NADH-ubiquinone oxidoreductase 75 kDa subunit, mitochon	1	2.587	2.587	2.587
O43676	NADH-ubiquinone oxidoreductase B12 subunit (EC 1.6.5.3) (1	0.259	0.259	0.259
P03905	NADH-ubiquinone oxidoreductase chain 4 (EC 1.6.5.3) (NAD	1	0.469	0.469	0.469
P16435	NADPH--cytochrome P450 reductase (EC 1.6.2.4) (CPR) (P4	11	0.114	2.038	0.625
Q13765	Nascent polypeptide-associated complex alpha subunit (NAC	1	0.976	0.976	0.976
Q9HCC7	NEDD4-like ubiquitin ligase 1	1	0.78	0.78	0.78
Q14697	Neutral alpha-glucosidase AB precursor (EC 3.2.1.84) (Glucos	1	1.208	1.208	1.208
Q15758	Neutral amino acid transporter B(0) (ATB(0)) (Sodium-depen	5	0.45	1.484	0.863
Q9H2Q2	NGT	1	0.569	0.569	0.569
Q5VZP5	Novel protein	1	0.675	0.675	0.675
Q5JP08	Novel protein	1	0.261	0.261	0.261
Q9NR30	Nucleolar RNA helicase 2 (EC 3.6.1.-) (Nucleolar RNA helica	1	0.779	0.779	0.779
Q8NI70	OK/SW-CL.16	1	0.371	0.371	0.371
Q5JUJ7	OTTHUMP00000017314 (Fragment)	1	0.932	0.932	0.932
Q4VXA4	OTTHUMP00000021035	1	0.435	0.435	0.435

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
P23284	Peptidyl-prolyl cis-trans isomerase B precursor (EC 5.2.1.8) (1	0.753	0.753	0.753
O95613	Pericentrin (Pericentrin B) (Kendrin)	1	0.997	0.997	0.997
Q00325	Phosphate carrier protein, mitochondrial precursor (PTP)	3	0.498	0.717	0.616
O95674	Phosphatidate cytidyltransferase 2 (EC 2.7.7.41) (CDP-digl	4	0.528	3.035	1.277
Q9BVG9	Phosphatidylserine synthase 2 (EC 2.7.8.-) (PtdSer synthase	3	0.456	0.982	0.696
Q16526	Photolyase (Cryptochrome 1)	2	0.233	0.306	0.267
Q9H307	Pinin (140 kDa nuclear and cell adhesion-related phosphopr	2	0.757	1.307	0.995
Q9HB66	PNAS-117	2	0.338	0.524	0.421
Q6NZI2	Polymerase I and transcript release factor (PTRF protein)	1	0.55	0.55	0.55
Q9NS91	Postreplication repair protein RAD18 (hRAD18) (hHR18) (RI	1	0.178	0.178	0.178
Q96CF8	PRKR interacting protein 1 (IL11 inducible)	1	1.006	1.006	1.006
Q9HD20	Probable cation-transporting ATPase 13A1 (EC 3.6.3.-)	3	0.549	0.774	0.646
O94919	Probable endonuclease KIAA0830 precursor (EC 3.1.30.-)	1	0.916	0.916	0.916
O96008	Probable mitochondrial import receptor subunit TOM40 homo	2	0.264	0.62	0.405
Q8WUM4	Programmed cell death 6-interacting protein (PDCD6-interact	1	0.558	0.558	0.558
P35232	Prohibitin	9	0.252	1.36	0.431
Q99623	Prohibitin-2 (B-cell receptor-associated protein BAP37) (Repr	25	0.466	2.843	0.858
Q9HCU5	Prolactin regulatory element-binding protein (Mammalian gua	1	0.917	0.917	0.917
O14684	Prostaglandin E synthase (EC 5.3.99.3) (Microsomal glutathi	1	0.201	0.201	0.201
Q6P0N0	Protein C14orf106 (P243)	1	0.667	0.667	0.667
Q8IVT2	Protein C19orf21	1	0.717	0.717	0.717
Q96CS3	Protein expressed in T-cells and eosinophils in atopic dermat	1	0.229	0.229	0.229
Q14165	Protein KIAA0152 precursor	1	0.835	0.835	0.835
Q9UPV7	Protein KIAA1045	1	0.118	0.118	0.118
P60468	Protein transport protein Sec61 beta subunit	5	0.257	0.461	0.353
Q96QD8	Putative 40-9-1 protein	1	0.734	0.734	0.734
Q8IXJ9	Putative Polycomb group protein ASXL1 (Additional sex com	1	0.202	0.202	0.202
Q9Y4G8	Rap guanine nucleotide exchange factor 2 (PDZ domain cont	1	0.333	0.333	0.333
P46940	Ras GTPase-activating-like protein IQGAP1 (p195)	1	0.435	0.435	0.435
P61026	Ras-related protein Rab-10	1	0.613	0.613	0.613
P61106	Ras-related protein Rab-14	1	0.454	0.454	0.454

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
P61019	Ras-related protein Rab-2A	3	0.677	1.238	0.833
Q9BZG1	Ras-related protein Rab-34 (Rab-39) (Ras-related protein Ra	1	0.527	0.527	0.527
P51148	Ras-related protein Rab-5C (RAB5L) (L1880)	2	1.167	3.965	2.151
P51149	Ras-related protein Rab-7	10	0.401	3.944	1.131
P51151	Ras-related protein Rab-9A (Rab-9)	1	0.237	0.237	0.237
Q8NFJ5	Retinoic acid-induced protein 3 (G-protein coupled receptor f	5	0.597	1.111	0.912
Q15287	RNA-binding protein with serine-rich domain 1 (SR-related pr	1	0.321	0.321	0.321
Q49A11	RPS27L protein	1	0.42	0.42	0.42
Q15413	Ryanodine receptor 3 (Brain-type ryanodine receptor) (RyR3)	1	1.769	1.769	1.769
P16615	Sarcoplasmic/endoplasmic reticulum calcium ATPase 2 (EC	5	0.422	1.115	0.588
Q8WTV0	Scavenger receptor class B member 1 (SRB1) (SR-BI) (CD3	1	0.619	0.619	0.619
O75880	SCO1 protein homolog, mitochondrial precursor	1	0.882	0.882	0.882
Q4VX17	Secretory carrier membrane protein 3	1	0.406	0.406	0.406
Q13501	Sequestosome-1 (Phosphotyrosine-independent ligand for th	1	0.928	0.928	0.928
Q52WX2	SH3-binding kinase	1	0.441	0.441	0.441
Q9H9B4	Sideroflexin-1 (Tricarboxylate carrier protein) (TCC)	3	0.394	1.277	0.787
Q15005	Signal peptidase complex subunit 2 (EC 3.4.-.-) (Microsomal	4	0.096	0.662	0.361
P61009	Signal peptidase complex subunit 3 (EC 3.4.-.-) (Microsomal	1	1.428	1.428	1.428
Q9Y5M8	Signal recognition particle receptor beta subunit (SR-beta) (P	2	0.239	0.61	0.382
Q8N5K1	Similar to mouse 1500009M05Rik protein	1	0.5	0.5	0.5
P05023	Sodium/potassium-transporting ATPase alpha-1 chain precur	2	0.285	0.408	0.341
P11166	Solute carrier family 2, facilitated glucose transporter membe	2	0.338	0.699	0.486
Q6NUK1	Solute carrier family 25 member 24, isoform 1	1	0.07	0.07	0.07
Q9Y512	Sorting and assembly machinery component 50 homolog	7	0.313	1.234	0.571
Q8TCJ2	Source of immunodominant MHC-associated peptides	2	0.949	3.79	1.896
Q13813	Spectrin alpha chain, brain (Spectrin, non-erythroid alpha ch	14	0.068	1.745	0.552
Q01082	Spectrin beta chain, brain 1 (Spectrin, non-erythroid beta cha	2	0.664	1.079	0.846
O75477	SPFH domain protein 1 precursor (Protein KE04)	1	2.052	2.052	2.052
O94905	SPFH domain protein 2 precursor	1	0.502	0.502	0.502
Q9UHA8	Splicing coactivator subunit SRm300	3	0.368	0.823	0.596
Q53GQ0	Steroid dehydrogenase homolog (Fragment)	1	0.361	0.361	0.361

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
O95425	Supervillin (Archvillin) (p205/p250)	1	0.6	0.6	0.6
Q9NZ01	Synaptic glycoprotein SC2	1	0.244	0.244	0.244
Q15006	Tetratricopeptide repeat protein KIAA0103	2	0.876	1.144	1.001
Q9H0W7	THAP domain-containing protein 2	2	0.456	0.516	0.485
Q9H3N1	Thioredoxin domain-containing protein 1 precursor (Transme	1	0.342	0.342	0.342
Q6F5E7	Thioredoxin reductase 2 intronic transcript 1	1	0.447	0.447	0.447
Q15643	Thyroid receptor-interacting protein 11 (TRIP-11) (Golgi-asso	1	0.057	0.057	0.057
O00463	TNF receptor-associated factor 5 (RING finger protein 84)	1	1.03	1.03	1.03
Q9UGU0	Transcription factor 20 (Stromelysin 1 PDGF-responsive ele	1	2.5	2.5	2.5
P02786	Transferrin receptor protein 1 (TfR1) (TR) (TfR) (Tfr) (CD71	6	0.135	0.888	0.448
Q9Y4A5	Transformation/transcription domain-associated protein (350/	1	1.161	1.161	1.161
Q9UGP8	Translocation protein SEC63 homolog	1	0.908	0.908	0.908
P43307	Translocon-associated protein alpha subunit precursor (TRA	1	0.64	0.64	0.64
P51571	Translocon-associated protein delta subunit precursor (TRAP	1	0.267	0.267	0.267
Q9UNL2	Translocon-associated protein gamma subunit (TRAP-gamm	2	0.26	0.484	0.355
Q99805	Transmembrane 9 superfamily protein member 2 precursor (1	0.747	0.747	0.747
P84157	Transmembrane anchor protein 1	2	0.908	1.141	1.018
P49755	Transmembrane emp24 domain containing protein 10 precur	1	0.527	0.527	0.527
Q7Z7H5	Transmembrane emp24 domain containing protein 4 precurs	2	1.171	1.332	1.249
Q9Y3A6	Transmembrane emp24 domain containing protein 5 precurs	1	1.324	1.324	1.324
Q9BVK6	Transmembrane emp24 domain containing protein 9 precurs	1	1.372	1.372	1.372
P17152	Transmembrane protein 11 (Protein PM1)	1	0.311	0.311	0.311
P57088	Transmembrane protein 33 (DB83 protein)	4	0.395	0.806	0.53
Q96GC9	Transmembrane protein 49 (TDC1)	1	0.467	0.467	0.467
Q9HC07	Transmembrane protein PT27 (TPA regulated locus)	1	0.29	0.29	0.29
P53007	Tricarboxylate transport protein, mitochondrial precursor (Citr	2	0.179	0.513	0.303
P23510	Tumor necrosis factor ligand superfamily member 4 (OX40 li	1	1.029	1.029	1.029
Q561W0	TXNDC14 protein	1	0.979	0.979	0.979
Q6RW13	Type-1 angiotensin II receptor-associated protein (AT1 recep	1	1.019	1.019	1.019
Q9UDW1	Ubiquinol-cytochrome c reductase complex 7.2 kDa protein (1	3.456	3.456	3.456
P22695	Ubiquinol-cytochrome-c reductase complex core protein 2, m	4	0.469	0.961	0.689

Acc No	Protein name	TPC	H/L(low)	H/L(high)	H/L(avg)
P31930	Ubiquinol-cytochrome-c reductase complex core protein I, mi	3	0.242	0.97	0.582
Q9Y3E0	UPF0198 protein CGI-141	1	0.939	0.939	0.939
P61421	Vacuolar ATP synthase subunit d (EC 3.6.3.14) (V-ATPase d	1	0.348	0.348	0.348
O75396	Vesicle trafficking protein SEC22b (SEC22 vesicle trafficking	4	0.323	0.709	0.54
Q9P0L0	Vesicle-associated membrane protein-associated protein A (2	0.303	0.823	0.499
P21796	Voltage-dependent anion-selective channel protein 1 (VDAC-	14	0.26	3.418	0.933
Q9Y277	Voltage-dependent anion-selective channel protein 3 (VDAC-	7	0.141	2.046	0.369
Q96DN5	WD-repeat protein 67	1	0.073	0.073	0.073
Q96ME7	Zinc finger protein 512	1	0.44	0.44	0.44