

Protein Name	Accession Number	Protein PI	Peptide Count	Total Ion Score
10 kDa heat shock protein, mitochondrial	P61604	8,89	2	89,47
14-3-3 protein zeta/delta	P29361	4,77	2	71,89
28 kDa heat- and acid-stable phosphoprotein	Q13442	8,84	4	150,30
28S ribosomal protein S15, mitochondrial	Q9DC71	10,13	3	74,61
40S ribosomal protein S10	P46783	10,15	5	178,96
40S ribosomal protein S12	P25398	6,81	3	133,25
40S ribosomal protein S18	Q5TJE9	10,99	5	107,09
40S ribosomal protein S19 (Fragment)	Q29308	10,2	8	286,11
40S ribosomal protein S19	P39019	10,31	8	322,69
40S ribosomal protein S21	P63220	8,68	3	102,63
40S ribosomal protein S29	P62273	10,17	5	165,20
40S ribosomal protein S30	P62860	12,15	3	116,07
40S ribosomal protein S5	P46782	9,73	3	137,66
40S ribosomal protein SA	Q2L9X0	4,8	4	131,97
60 kDa heat shock protein, mitochondrial	P10809	5,7	4	170,52
60S acidic ribosomal protein P2	P05387	4,42	3	107,15
60S ribosomal protein L12	Q6QMZ7	9,48	1	64,42
60S ribosomal protein L13	Q56JZ1	11,59	4	75,84
60S ribosomal protein L24	Q66WF5	11,26	4	86,30
60S ribosomal protein L26	P61254	10,55	4	120,74
60S ribosomal protein L26-like 1	Q9UNX3	10,55	3	72,65
60S ribosomal protein L27	P61357	10,56	3	73,50
60S ribosomal protein L31	P62899	10,54	2	51,34
60S ribosomal protein L36	Q9Y3U8	11,59	1	58,63
60S ribosomal protein L6	Q2YGT9	10,73	3	132,37
78 kDa glucose-regulated protein	Q3S4T7	5,07	17	671,28
Abnormal spindle-like microcephaly-associated protein homolog (Fragment)	P62297	11,06	20	158,21
Actin, alpha cardiac muscle 1	P68032	5,23	12	796,27
Actin, aortic smooth muscle	P62736	5,23	12	789,44
Actin, cytoplasmic 1	O18840	5,29	13	888,06

Actin-related protein 2/3 complex subunit 4	P59998	8,53	1	44,36
Activated RNA polymerase II transcriptional coactivator p15	Q4R947	9,6	2	60,50
Adenylyl cyclase-associated protein 1	Q3SYV4	7,16	4	189,63
Alpha-2-macroglobulin receptor-associated protein	Q99068	6,85	1	61,80
Alpha-actinin-1	Q3B7N2	5,25	17	655,82
Alpha-actinin-2	Q9JI91	5,36	8	172,74
Alpha-actinin-4	O43707	5,27	9	301,01
Alpha-crystallin B chain	Q7M2W6	6,76	7	287,16
Alpha-enolase	Q9XSJ4	6,37	7	200,23
Ankyrin repeat domain-containing protein 35	Q8N283	5,76	4	95,36
Annexin A5	Q5RIW0	4,94	6	441,80
Arf-GAP with SH3 domain, ANK repeat and PH domain-containing protein 2	Q7SIG6	6,22	3	56,79
Astrocytic phosphoprotein PEA-15	Q15121	4,93	4	106,50
ATP synthase protein 8	Q35914	9,16	1	44,56
ATP synthase subunit alpha, mitochondrial	Q03265	9,22	5	91,80
ATP synthase subunit beta, mitochondrial	P00829	5,15	1	51,56
ATP synthase subunit e, mitochondrial	Q9MYT8	9,3	3	107,51
ATP synthase-coupling factor 6, mitochondrial	P13618	5,41	2	123,62
ATPase inhibitor, mitochondrial	Q29307	9,34	3	187,43
Atrial natriuretic peptide-converting enzyme	Q80YN4	5,18	2	60,03
Baculoviral IAP repeat-containing protein 5	Q9JHY7	5,85	2	52,79
Barrier-to-autointegration factor	O75531	5,81	3	212,75
Beta-actin-like protein 2	Q8BFZ3	5,3	7	334,12
Beta-glucuronidase	O97524	6,13	5	103,03
Beta-lactoglobulin	P02756	5,5	1	57,92
Bromodomain testis-specific protein	Q4R8Y1	9,23	6	117,59
C2 domain-containing protein 3	Q4AC94	6,65	6	88,76
Calcium-regulated heat stable protein 1	Q9CR86	8,41	1	87,90
Caldesmon	Q05682	5,63	27	1024,06
Calmodulin	P62158	4,09	2	151,18
Calpastatin	P12675	5,21	6	180,61
Calponin-1	Q08092	8,91	12	427,15

Calponin-2 (Fragment)	Q08094	8,58	7	149,00
Calponin-2	Q08093	7,53	8	167,80
Cathepsin D	Q4LAL9	7,97	4	89,73
Cationic trypsin	P00760	8,4	4	281,65
Cellular nucleic acid-binding protein	P62633	8	5	218,29
Centromere protein F	P49454	5,03	15	168,42
Chloride intracellular channel protein 4	Q9QYB1	5,44	3	76,64
Chloride intracellular channel protein 6	Q811Q2	4,29	2	55,57
Cingulin	B0KWC9	5,46	7	128,90
Clathrin light chain A	O08585	4,45	2	60,82
Clathrin light chain B	P09497	4,57	3	109,92
Cofilin-1	Q4R5C0	8,16	5	133,69
Coiled-coil domain-containing protein 88B	A6NC98	5,09	5	80,69
Collagen alpha-1(I) chain	Q9XSJ7	5,67	20	805,18
Collagen alpha-1(III) chain	P08121	6,11	7	148,66
Collagen alpha-2(I) chain	O46392	9,18	8	205,45
Collagen alpha-5(VI) chain	A8TX70	6,5	7	99,08
Core histone macro-H2A.1	Q9QZQ8	9,81	4	101,69
Cysteine and glycine-rich protein 1	Q3MHY1	8,91	5	250,93
Cysteine and glycine-rich protein 2	Q16527	8,95	2	56,94
Cytochrome b-c1 complex subunit 6, mitochondrial	Q5M9I5	4,9	1	75,78
Cytochrome c oxidase subunit 4 isoform 1, mitochondrial (Fragment)	Q95283	9,37	2	98,80
Cytochrome c oxidase subunit 4 isoform 1, mitochondrial	P19783	9,25	1	45,48
Cytochrome c oxidase subunit 5A, mitochondrial	B0VYY4	6,42	2	66,43
Cytochrome c oxidase subunit 6B1	P00429	8,76	3	123,65
Dermcidin	P81605	6,08	3	195,61
Desmin	P02540	5,21	7	208,70
Desmoglein-1	Q02413	4,9	3	63,65
Desmoplakin	P15924	6,44	20	309,97
Dihydrolipoyl dehydrogenase, mitochondrial (Fragment)	Q811C4	6,94	4	83,49
Dihydropyrimidinase-related protein 2	P47942	5,95	5	106,45
Dimethyladenosine transferase 1, mitochondrial	Q811P6	9,57	4	99,57

DNA-binding protein A	P16989	9,77	12	450,60
DNA-dependent protein kinase catalytic subunit	P78527	6,75	12	137,52
DnaJ homolog subfamily C member 8	Q6NZB0	9,04	1	42,63
E3 ubiquitin-protein ligase SHPRH	Q149N8	7,3	5	91,71
EF-hand calcium-binding domain-containing protein 3	Q8N7B9	9,31	2	58,78
Elongation factor 1-alpha 1	P62629	9,1	2	48,85
Elongation factor 1-alpha 2	Q05639	9,11	3	83,65
Elongation factor 1-beta	P29412	4,57	2	50,91
Elongation factor 2	P09445	6,22	6	143,08
Endoplasmic reticulum resident protein 29	P57759	5,9	2	65,90
Enhancer of rudimentary homolog	P84090	5,63	4	109,40
Eukaryotic translation initiation factor 1	P41567	6,89	3	61,34
Eukaryotic translation initiation factor 3 subunit J	O75822	4,72	1	41,61
Eukaryotic translation initiation factor 4B	Q8BGD9	5,47	6	135,11
Eukaryotic translation initiation factor 4E-binding protein 1	Q13541	5,32	2	65,56
Eukaryotic translation initiation factor 5A-1	P63241	5,08	3	150,38
Ezrin	P15311	5,94	7	141,33
FH1/FH2 domain-containing protein 1	Q6P9Q4	5,93	1	57,57
Fibronectin	P07589	5,33	36	1450,92
Filaggrin-2	Q5D862	8,45	7	117,60
Filamin-A	P21333	5,7	58	2520,15
Filamin-B	O75369	5,47	14	302,52
Filamin-binding LIM protein 1	Q5REN1	5,9	5	187,44
Filamin-C	Q14315	5,65	15	475,81
Four and a half LIM domains protein 1	Q9WUH4	8,76	3	64,07
Galectin-1	Q49I35	4,99	7	396,14
Gamma-enolase	P17183	4,99	5	159,50
Glia maturation factor beta	P60983	5,19	1	46,72
Glial fibrillary acidic protein	Q5RA72	5,42	5	176,08
Glyceraldehyde-3-phosphate dehydrogenase	A3FKF7	8,2	5	123,16
H/ACA ribonucleoprotein complex subunit 2	Q5E950	8,43	2	61,07
Heat shock 70 kDa protein 1A/1B	Q07439	5,61	4	101,67
Heat shock 70 kDa protein 1-like	P55063	5,91	4	116,51

Heat shock 70 kDa protein 6	Q04967	5,77	4	111,17
Heat shock cognate 71 kDa protein	P19378	5,24	11	285,93
Heat shock protein beta-1	Q5S1U1	6,23	12	695,55
Heat shock protein beta-6	O14558	5,95	3	108,47
Heat shock-related 70 kDa protein 2	P34933	5,44	6	131,23
Hemoglobin subunit alpha	P01965	8,76	1	41,61
Heterogeneous nuclear ribonucleoprotein A/B	Q99729	8,22	4	216,13
Heterogeneous nuclear ribonucleoprotein A1	P04256	9,2	9	452,20
Heterogeneous nuclear ribonucleoprotein A3	Q8BG05	9,1	5	171,78
Heterogeneous nuclear ribonucleoprotein C	Q5RA82	4,95	3	91,87
Heterogeneous nuclear ribonucleoprotein G	A5A6M3	10,06	16	405,88
Heterogeneous nuclear ribonucleoprotein K	P61978	5,39	5	147,81
Heterogeneous nuclear ribonucleoproteins A2/B1	P22626	8,97	8	365,95
High mobility group protein B1	B0CM99	5,62	8	381,36
High mobility group protein B2	P26583	7,62	6	186,38
High mobility group protein HMG-I/HMG-Y	Q9QXP3	10,32	2	66,89
Histone H1.2	P15864	11	8	324,72
Histone H1.3	P43277	11,03	8	336,29
Histone H1.4	P43274	11,1	5	225,41
Histone H1.5	P16401	10,91	5	102,16
Histone H1t	P06348	11,58	5	230,38
Histone H2A type 1-A	Q96QV6	10,86	6	267,13
Histone H2A type 1-H	Q96KK5	10,88	7	396,53
Histone H2A type 4	Q00728	11,02	6	372,72
Histone H2A.V	Q71UI9	10,58	4	246,04
Histone H2A.x	P16104	10,74	5	244,33
Histone H2B type 1-A	P70696	10,29	5	222,44
Histone H2B type 1-J	P06899	10,31	6	390,61
Histone H2B type 1-L	Q99880	10,31	7	424,47
Histone H2B type 1-N	Q32L48	10,31	6	351,51
Histone H2B type 3-B	Q8N257	10,31	6	353,53
Histone H3.1	P68431	11,13	10	370,62
Histone H3.3	Q5E9F8	11,27	10	344,63

Histone H3.3C	Q3SZB8	10,9	5	168,32
Histone H4	P62805	11,36	9	503,84
Hornerin	Q86YZ3	10,05	15	418,90
Hyaluronan mediated motility receptor	O75330	5,68	5	93,61
Immunity-related GTPase family M protein 1	Q60766	8,56	2	55,17
Immunity-related GTPase family M protein	Q6AYC2	8,39	1	51,35
Integrin beta-1	Q9GLP0	5,32	7	294,46
Interferon-induced protein with tetratricopeptide repeats 2	Q60462	8,73	2	53,05
Interleukin-10	P18893	8,44	1	41,80
Interleukin-2	P36835	6,73	1	42,11
Jerky protein homolog-like	Q9Y4A0	8,25	1	42,76
Keratin, type I cytoskeletal 10	P13645	5,13	25	1238,78
Keratin, type I cytoskeletal 12 (Fragment)	Q28706	4,63	5	206,33
Keratin, type I cytoskeletal 13	P08730	4,79	6	248,86
Keratin, type I cytoskeletal 14	P02533	5,09	24	954,47
Keratin, type I cytoskeletal 16	P08779	4,99	23	940,42
Keratin, type I cytoskeletal 17	Q04695	4,97	17	608,01
Keratin, type I cytoskeletal 18	P05784	5,22	5	142,18
Keratin, type I cytoskeletal 19	Q63279	5,21	12	373,88
Keratin, type I cytoskeletal 20	P35900	5,52	4	116,82
Keratin, type I cytoskeletal 25	Q0P5J4	4,97	6	163,23
Keratin, type I cytoskeletal 27	Q6R649	4,97	4	74,57
Keratin, type I cytoskeletal 28	Q7Z3Y7	5,33	4	101,12
Keratin, type I cytoskeletal 42	Q6IFX2	5,09	16	472,27
Keratin, type I cytoskeletal 9	P35527	5,14	31	1787,90
Keratin, type II cytoskeletal 1	P04264	8,15	36	2881,99
Keratin, type II cytoskeletal 1b	Q6IFZ6	7,74	10	472,26
Keratin, type II cytoskeletal 3	Q29426	7,6	13	408,11
Keratin, type II cytoskeletal 4	P19013	6,25	7	271,87
Keratin, type II cytoskeletal 5	P13647	7,59	24	1148,82
Keratin, type II cytoskeletal 6A	P02538	8,09	26	1298,21
Keratin, type II cytoskeletal 6B	P04259	8,09	23	1239,98
Keratin, type II cytoskeletal 7	Q29S21	5,79	11	436,14

Keratin, type II cytoskeletal 71	Q9R0H5	6,6	5	223,84
Keratin, type II cytoskeletal 72	Q6IG04	7,05	4	142,51
Keratin, type II cytoskeletal 73	Q6NXH9	8,36	10	440,37
Keratin, type II cytoskeletal 74	Q7RTS7	7,59	8	244,62
Keratin, type II cytoskeletal 75	Q08D91	7,56	14	522,10
Keratin, type II cytoskeletal 79	Q148H7	7,1	12	482,86
Keratin, type II cytoskeletal 8	Q10758	5,83	13	463,18
Keratin, type II cytoskeletal 80	A0JND2	5,23	2	112,27
Keratin-like protein KRT222	Q8CCX5	5,6	2	57,41
Keratinocyte proline-rich protein	Q5T749	8,72	8	220,52
Kinesin-like protein KIF2B	A6H750	9,3	4	86,88
Kit ligand	P79169	5,88	1	41,41
Lamina-associated polypeptide 2, isoform alpha	P42166	7,56	4	104,53
Leucine-rich repeat-containing protein 59	Q5RJR8	9,57	4	112,02
LIM and SH3 domain protein 1	Q99MZ8	6,61	13	647,35
Liprin-alpha-2	Q8BSS9	5,79	5	76,78
Lysosomal alpha-mannosidase	Q29451	9,2	6	136,73
Macrophage migration inhibitory factor	Q4R549	7,74	1	43,27
MAM domain-containing glycosylphosphatidylinositol anchor protein 2	P60755	6,65	3	82,31
Membrane-associated progesterone receptor component 1	O00264	4,56	2	79,79
Membrane-associated progesterone receptor component 2	O15173	4,76	2	64,23
Microtubule-associated protein 4	P27546	4,9	4	87,23
Microtubule-associated protein RP/EB family member 1	Q3ZBD9	5,2	3	62,75
Mitochondrial import inner membrane translocase subunit Tim13	P62076	8,42	2	67,29
Mitochondrial import inner membrane translocase subunit Tim8 A	O60220	5,09	2	98,60
Mitochondrial import inner membrane translocase subunit Tim9	Q9Y5J7	6,71	3	146,69
Moesin	P26042	6,3	11	234,18
Myeloperoxidase	P05164	9,19	3	51,88
Myosin heavy chain, embryonic smooth muscle isoform (Fragment)	Q99105	5,27	5	172,93
Myosin light chain 1/3, skeletal muscle isoform	P02602	4,97	2	66,70
Myosin light chain 3	P16409	5,03	5	177,94

Myosin light polypeptide 6	P60660	4,56	6	338,89
Myosin regulatory light chain 12A	Q5RC34	4,67	5	214,14
Myosin regulatory light polypeptide 9	Q9CQ19	4,8	3	91,01
Myosin-10	Q27991	5,44	15	535,44
Myosin-11	P35748	5,39	10	218,89
Myosin-14	Q6URW6	5,49	8	178,64
Myosin-9	Q258K2	5,5	36	1419,56
Myotrophin	Q863Z4	5,27	3	135,69
N6-adenosine-methyltransferase 70 kDa subunit	Q8C3P7	6,05	2	59,61
N-acetylglucosamine-6-sulfatase	P50426	8,54	3	175,62
NACHT, LRR and PYD domains-containing protein 3	Q96P20	6,22	3	57,14
NADH dehydrogenase [ubiquinone] iron-sulfur protein 8, mitochondrial	Q60HE3	6,45	2	57,49
Nesprin-1	Q6ZWR6	5,43	15	165,76
Neuroblast differentiation-associated protein AHNAK	Q09666	5,8	37	1035,38
Nexilin	Q0ZGT2	5,31	3	71,32
Non-muscle caldesmon	Q62736	6,34	20	781,92
Nuclear ubiquitous casein and cyclin-dependent kinases substrate	Q9H1E3	5	4	174,95
Nuclease-sensitive element-binding protein 1	P67809	9,87	18	754,39
Nucleobindin-1	Q63083	5,04	4	122,12
Nucleolin	P08199	4,7	5	149,63
Nucleophosmin	Q3T160	4,62	5	106,06
P protein	Q04671	6,84	1	64,14
Palladin	Q8WX93	6,67	5	90,24
PDZ and LIM domain protein 1	Q5E9E1	6,76	6	313,19
PDZ and LIM domain protein 7	Q3TJD7	8,82	4	79,71
Peptidyl-prolyl cis-trans isomerase A	P62936	8,34	6	286,32
Peptidylprolyl cis-trans isomerase A-like 4A/B/C	Q9Y536	9,32	2	73,63
Peptidyl-prolyl cis-trans isomerase FKBP10	Q2HJ89	5,56	5	178,80
Peptidyl-prolyl cis-trans isomerase FKBP7	Q5RET2	6,11	3	70,91
Pericentrin	P48725	5,27	8	119,91
Peroxiredoxin-1	Q5E947	8,59	4	198,23
Peroxiredoxin-4	Q13162	5,86	1	76,84

Phosphatidylinositol 3-kinase regulatory subunit alpha	Q63787	5,9	3	88,79
Phosphoglycerate mutase 1	P18669	6,67	10	480,54
Plasminogen activator inhibitor 1 RNA-binding protein	Q8NC51	8,66	9	363,60
Polyadenylate-binding protein 1	P11940	9,52	4	133,68
Polyadenylate-binding protein 3	Q9H361	9,68	6	142,05
POTE ankyrin domain family member E	Q6S8J3	5,83	8	492,81
POTE ankyrin domain family member I	P0CG38	5,83	7	393,88
Prefoldin subunit 2	Q9UHV9	6,2	2	87,89
Prelamin-A/C	Q3ZD69	6,73	9	307,39
Proactivator polypeptide	P07602	5,06	6	190,90
Profilin-1	P62962	8,46	3	153,00
Programmed cell death protein 5	P56812	5,55	4	160,52
Proteasome activator complex subunit 1	Q06323	5,78	4	178,09
Proteasome activator complex subunit 2	Q5E9G3	5,31	1	45,47
Proteasome subunit beta type-1	Q2TBX6	8,26	3	77,00
Protein canopy homolog 2	Q9QXT0	4,95	3	142,54
Protein canopy homolog 4	Q3SWX1	4,71	1	66,17
Protein disulfide-isomerase A3	P30101	5,98	16	757,49
Protein disulfide-isomerase A6	P38660	5,04	2	68,78
Protein disulfide-isomerase	P05307	4,8	8	320,67
Protein enabled homolog	Q03173	7,62	5	163,76
Protein piccolo	Q9Y6V0	6,12	9	124,09
Protein S100-A11	P31950	6,07	2	86,63
Protein S100-A2	P10462	4,68	2	74,26
Protein transport protein Sec61 subunit beta	P60468	11,57	1	62,17
Putative high mobility group protein B1-like 1	B2RPK0	5,92	3	98,52
Putative histone H2B type 2-D	Q6DRA6	10,6	3	114,00
Putative tropomyosin alpha-3 chain-like protein	A6NL28	4,47	2	70,45
PX domain-containing protein kinase-like protein	Q4FZZ1	9,5	5	91,29
Radixin	Q32LP2	5,95	5	133,88
Regulator of nonsense transcripts 3A	Q9H1J1	9,12	2	81,26
Reticulocalbin-3	Q8BH97	4,74	4	203,78
Reticulon-4	Q9NQC3	4,43	3	67,99

Rho GDP-dissociation inhibitor 1	P52565	5,02	2	75,89
Ribosomal protein S6 kinase alpha-6	Q9UK32	5,92	4	51,42
RNA methyltransferase-like protein 1	Q9HC36	8,89	3	66,74
RNA-binding motif protein, X-linked-like-2	Q4R813	10,28	4	100,67
RNA-binding motif protein, X-linked-like-3	Q8N7X1	9,19	8	123,11
RNA-binding protein EWS	Q61545	9,37	2	53,17
Saposin-B-Val	P81405	4,41	3	136,45
Serine/arginine-rich splicing factor 1	Q07955	10,37	10	452,98
Serine/arginine-rich splicing factor 10	Q9R0U0	11,26	2	75,93
Serine/arginine-rich splicing factor 12	Q8WXF0	11,69	2	58,93
Serine/arginine-rich splicing factor 2	Q01130	11,86	3	81,08
Serine/arginine-rich splicing factor 3	P84103	11,64	4	171,94
Serine/arginine-rich splicing factor 7	Q16629	11,83	3	120,21
Serpin H1	Q5RBS3	8,37	3	132,37
Serum albumin	P08835	6,08	13	453,27
SH3 domain-binding glutamic acid-rich-like protein 3	Q9H299	4,82	2	109,52
Small nuclear ribonucleoprotein Sm D3	P62320	10,33	5	118,63
Small ubiquitin-related modifier 2	Q6LDZ8	5,32	1	78,47
Sodium/hydrogen exchanger 4	P26434	9,16	2	51,95
Sorting nexin-18	Q96RF0	5,44	3	80,75
Sorting nexin-25	Q3ZT31	5,83	2	49,37
Splicing factor 1	Q64213	8,98	1	40,00
Src substrate cortactin	Q60598	5,24	6	211,60
Stathmin	P16949	5,76	3	161,63
Stress-70 protein, mitochondrial	P48721	5,97	8	177,35
Stress-induced-phosphoprotein 1	Q3ZBZ8	6,08	11	242,21
Sulfated glycoprotein 1	P10960	5,13	5	128,60
Syntaxin-7	Q5R602	5,44	2	55,49
Talin-1	Q9Y490	5,77	21	687,40
T-complex protein 1 subunit alpha	P11983	5,82	4	61,00
T-complex protein 1 subunit beta	Q3ZBH0	6,18	6	155,73
Tetratricopeptide repeat protein 29	Q4R6M4	5,74	3	65,57
Thioredoxin	P82460	4,97	9	546,50

THO complex subunit 4	Q86V81	11,15	4	86,58
Thrombospondin-1	P07996	4,71	3	73,40
Thymosin beta-10	P21753	6,18	2	74,74
Thymosin beta-4	Q6S9C5	5,02	6	316,95
Titin	A2ASS6	5,91	68	502,67
Transaldolase	Q2TBL6	7,03	4	135,61
Transcription factor BTF3 homolog 4	Q96K17	5,95	3	64,90
Transcription factor BTF3	Q64152	9,52	2	85,14
Transforming growth factor beta-1-induced transcript 1 protein	O43294	6,62	3	107,50
Transgelin	Q01995	8,87	14	640,74
Transgelin-2	Q5E9F5	8,39	13	699,78
Translocation protein SEC63 homolog	Q5R660	5,25	2	58,88
Trichohyalin	P37709	5,61	4	88,04
Triosephosphate isomerase	Q29371	7,01	10	393,29
Tropomodulin-3	Q9NYL9	5,08	4	97,59
Tropomyosin alpha-1 chain	Q5KR49	4,69	17	1065,18
Tropomyosin alpha-3 chain	P06753	4,68	15	784,61
Tropomyosin alpha-4 chain	P67937	4,67	22	1199,59
Tropomyosin beta chain	Q5KR48	4,66	19	1050,71
Tubulin alpha-1A chain	Q71U36	4,94	2	70,58
Ubiquilin-1	Q5R684	5,02	3	91,86
Ubiquitin-60S ribosomal protein L40	P63050	9,87	2	66,59
Ubiquitin-associated protein 2-like	Q80X50	6,61	1	44,32
UV excision repair protein RAD23 homolog B	P54727	4,79	3	121,47
Vacuolar protein sorting-associated protein 13C	Q709C8	6,38	7	91,68
Vimentin	P31000	5,06	27	1173,15
Vinculin	P26234	5,62	24	744,19
Voltage-dependent N-type calcium channel subunit alpha-1B	Q02294	8,82	8	109,30
Voltage-dependent P/Q-type calcium channel subunit alpha-1A	P97445	9,03	7	123,61
Xylulose kinase	O75191	5,55	3	61,59
Y-box-binding protein 2	Q9Z2C8	10,95	6	287,70
Zinc finger ZZ-type and EF-hand domain-containing protein 1	Q5SSH7	5,74	7	111,43
Zyxin	Q15942	6,22	3	92,51

