

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q03151	Growth factor independent 1 isoform CRA_a	3.77	4
Q03151	PI2 and LIM domain protein 1	3.08	2
Q9NYU2	UDP-glucosyltransferase 1	1.44	3
Q02318	Sterol 26-hydroxylase mitochondrial	1.39	1
Q9Y878	Coatomer subunit gamma-1	1.16	3
Q8WVM8	Sec1 family domain-containing protein 1	1.15	2
P62330	ADP-ribosylation factor 6	1.10	1
B2RN96	Zinc finger RNA binding protein	1.05	2
Q14522	Receptor-type tyrosine-protein phosphatase T	0.96	2
Q8NH61	Olfactory receptor 51F2	0.94	1
Q15800	Methylsterol monooxygenase 1	0.90	1
P11448	Pyruvate carboxylase mitochondrial	0.84	2
P33993	DNA replication licensing factor MCM7	0.83	3
P63261	Actin cytoplasmic 2	0.80	39
ABQKWD	SHINC3	0.73	1
P08676	ATP synthase subunit beta mitochondrial	0.73	14
O00487	26S proteasome non-ATPase regulatory subunit 14	0.73	1
Q06520	Bile salt sulfotransferase	0.72	4
P51812	Ribosomal protein S6 kinase alpha-3	0.71	2
Q43707	Alpha-actinin-4	0.70	20
P21980	Protein-glutamine gamma-glutamyltransferase 2	0.67	2
P52597	Heterogeneous nuclear ribonucleoprotein F	0.66	7
Q13509	Tubulin beta-3 chain	0.65	15
P62266	40S ribosomal protein S23	0.65	2
Q02787	Serotriferin	0.64	1
Q381A7	Phosphoglycine:phosphoserine phosphotransferase	0.64	1
P15927	Replication protein A 32 kDa subunit	0.64	2
Q15084	Serine-threonine-protein phosphatase 6 regulatory ankyrin repeat subunit A	0.62	2
Q13268	Dihydrogenase reductase SCR family member 2 mitochondrial	0.61	16
Q15393	Sclerostin factor 3B subunit 3	0.62	1
Q09623	Prohibitin-2	0.62	5
Q6H8R7	Serine-threonine-protein phosphatase 6 regulatory subunit 3	0.59	10
P11586	O-1-methylhydrofolate synthase cytoplasmic	0.59	2
Q53GQ0	Very-long-chain 3-oxoacyl-CoA reductase	0.59	1
P29401	Transketolase	0.58	2
P53294	Peptidyl-prolyl cis-trans isomerase B	0.58	2
Q78477	Etn-1	0.58	1
P39656	Dolichyl-diphosphooligosaccharide-protein glycosyltransferase 48 kDa subunit	0.57	1
Q8W1P6	Class IVb beta tubulin	0.57	22
B2RN113	Voltage-gated channel shaker-related subfamily member 7	0.56	3
P34897	Serine hydroxymethyltransferase mitochondrial	0.56	1
Q9JBS4	DnaJ homolog subfamily B member 11	0.55	1
P52322	Pyruvate 5-carboxylate reductase 1 mitochondrial	0.54	3
P27694	Replication protein A 70 kDa DNA-binding subunit	0.53	3
Q3LXA3	Bifunctional ATP-dependent dihydroxyacetone kinaseFAD-AMP lyase cyclizing	0.53	3
Q99486	Neurogenic locus notch homolog protein 4	0.52	1
P08299	60S ribosomal protein L23	0.52	3
P12004	Proliferating cell nuclear antigen	0.51	3
Q9NZR2	Low-density lipoprotein receptor-related protein 1B	0.50	2
P08642	Eukaryotic initiation factor 4A-1	0.44	7
Q9LJ50	Calcium-binding mitochondrial carrier protein Aralar2	0.50	1
P15924	Desmoplakin	0.50	23
Q9J527	Fatty acid synthase	0.47	7
AT7B16	o160S ribosome receptor	0.47	2
R07437	Tubulin beta chain	0.47	24
P61158	Actin-related protein 3	0.47	1
Q04837	Single-stranded DNA-binding protein mitochondrial	0.46	5
Q4W4Y1	Dopamine receptor interacting protein 4	0.46	6
P11413	Glucose-6-phosphate 1-dehydrogenase	0.46	1
Q9NVV3	Magnesium transporter NIPA3	0.46	1
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.45	2
P62937	Peptidyl-prolyl cis-trans isomerase A	0.45	1
Q00341	Vigilin	0.45	1
Q43813	LaC-like protein 1	0.44	1
Q9H2U9	Disintegrin and metalloproteinase domain-containing protein 7	0.43	1
P18621	60S ribosomal protein L17	0.43	3
P53618	Coatomer subunit beta	0.43	1
Q07020	60S ribosomal protein L18	0.42	1
Q81B22	Spermatogenesis-associated protein 20	0.42	2
Q01085	Nucleoglycin TIAR	0.41	2
Q9JLV0	Unconventional myosin-Vb	0.41	21
Q15436	Protein transport protein Sec23A	0.40	2
P45880	Voltage-dependent anion-selective channel protein 2	0.40	2
Q13838	Spliceosome RNA helicase DDX39b	0.40	1
P06441	Elongation factor 1-gamma	0.40	1
Q9UKP6	Urolectin-2 receptor	0.40	2
Q43175	D-3-phosphoglycerate dehydrogenase	0.40	1
P14868	Aspartate-RNA lyase cytoplasmic	0.40	2
P50991	I-complex protein 1 subunit delta	0.39	1
Q86TG7	Retrotransposon-derived protein PEG10	0.39	1
Q6PDU7	Epididymis luminal protein 215	0.39	1
Q92945	Far upstream element-binding protein 2	0.39	1
R04843	Dolichyl-diphosphooligosaccharide-protein glycosyltransferase subunit 1	0.39	1
Q09327	Vilrin-1	0.38	8
P11142	Heat shock cognate 71 kDa protein	0.37	10
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.36	3
P25705	ATP synthase subunit alpha mitochondrial	0.36	6
P50454	Serpin H1	0.36	1
P17844	Probable ATP-dependent RNA helicase DDX5	0.36	4
P12814	Alpha-actinin-1	0.35	6
P05783	Keratin type I cytoskeletal 18	0.34	28
P03011	60S ribosomal protein S7	0.34	1
P45954	Shortbranched chain specific acyl-CoA dehydrogenase mitochondrial	0.33	1
P13639	Elongation factor 2	0.32	2
Q96PK6	RNA-binding protein 14	0.32	3
P17997	I-complex protein 1 subunit alpha	0.32	1
R02788	Serum albumin	0.32	17
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.31	5
R02675	Fibrinogen beta chain	0.31	3
R61978	Heterogeneous nuclear ribonucleoprotein K	0.31	3
R08727	Keratin type I cytoskeletal 19	0.31	11
R05767	Keratin type II cytoskeletal 8	0.30	29
Q6IB91	PKC2 protein	0.30	2
P05141	ADP-ATP translocase 2	0.30	4
Q15149	Plectin	0.29	13
P18527	DNA-dependent protein kinase catalytic subunit	0.29	3
Q9NZ18	Insulin-like growth factor 2 mRNA-binding protein 1	0.28	2
R0382	Galactin-1	0.27	1
Q43795	Unconventional myosin-1b	0.26	1
R03833	Tubulin alpha-1B chain	0.26	11
H7B58	Protein-L-isopartate O-methyltransferase	0.26	11
P19474	E3 ubiquitin-protein ligase TRIM21	0.26	10
Q05793	Double-stranded RNA-binding protein Staufenn homolog 1	0.26	1
Q13155	Aminoacyl tRNA synthase complex-interacting multifunctional protein 2	0.25	1
D3DPC0	Titin isoform CRA_a	0.25	2
Q80782	Syntaxin-12	0.23	1
P19443	Heterogeneous nuclear ribonucleoprotein H	0.23	1
P49411	Elongation factor Tu mitochondrial	0.22	3
R00338	L-lactate dehydrogenase A chain	0.21	6
R07237	Protein disulfide-isomerase	0.21	1
R04259	Keratin type II cytoskeletal 6B	0.20	4
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.20	9
P38946	Stress-70 protein mitochondrial	0.19	1
R02751	Fibronectin	0.19	5
P14625	Endoplasmic	0.19	2
P11021	78 kDa glucose-regulated protein	0.18	12
R54136	Arginine-RNA lyase cytoplasmic	0.18	1
R08238	Heat shock protein HSP 90-beta	0.18	4
R02679	Fibrinogen gamma chain	0.17	3
Q06850	Peroxiredoxin-1	0.17	1
P10809	60 kDa heat shock protein mitochondrial	0.17	3
R07900	Heat shock protein HSP 90-alpha	0.15	2
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.15	3
Q15365	Pot1-binding protein 1	0.15	1
R04406	Glyceraldehyde-3-phosphate dehydrogenase	0.15	1
A2NU12	Lambda-chain AA-20 to 215	0.15	2
P58107	Epiplakin	0.15	2
Q80701	UDP-glucose 6-dehydrogenase	0.14	1
R01709	Ig lambda chain V-I region MGC	0.13	1
Q00571	ATP-dependent RNA helicase DDX3X	0.13	4
R81104	Elongation factor 1-alpha	0.13	3
P51659	Peroxisomal multifunctional enzyme type 2	0.13	10
P14618	Pyruvate kinase PKM	0.13	1
R02671	Fibrinogen alpha chain	0.12	3
P59908	Keratin type II cytoskeletal 2 epidermal	0.10	9
P13645	Keratin type I cytoskeletal 10	0.10	9
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.09	1
R03107	Heat shock 70 kDa protein 1A.1B	0.09	2
Q9BVG8	Kinesin-like protein KIFC3	0.09	1
R02533	Keratin type I cytoskeletal 14	0.09	2
R02769	Serum albumin	0.09	31
Q13813	Spectrin alpha chain non-erythrocytic 1	0.08	2
R04264	Keratin type II cytoskeletal 1	0.08	8
Q01062	Spectrin beta chain non-erythrocytic 1	0.07	2
Q14945	D-CK1 protein	0.07	2
P35579	Myosin-9	0.05	2
P35527	Keratin type I cytoskeletal 9	0.03	1