

Accession Number	Description	Fold Enrichment	UniquePeptideCount
G8Y9Q3	Zinc finger protein 37 homolog	3.63	7
P04362	Serine/threonine-protein kinase Cdc42	2.29	2
P04844	Dolichyl-diphosphooligosaccharide–protein glycosyltransferase subunit 2	1.71	4
I1VZ6	Hemoglobin alpha 1	1.28	7
Q9H3K8	AlphaA-like protein 1	1.12	4
Q93432	Mitochondrial 1-phosphate quinolinate transferase alpha	1.11	1
B0YIW8	Anchian 1 isoform CRA_a	1.08	3
P30085	UHM-CMP kinase	1.08	1
P02740	Serine/threonine-protein phosphatase 6 catalytic subunit	1.03	5
P02533	Calcium/calmodulin-dependent protein kinase Calk kinase II delta isoform CRA_e	1.01	4
P30048	Thioredoxin-dependent peroxide reductase mitochondrial	1.00	4
P15927	Replication protein A 32 kDa subunit	0.91	4
Q92523	AlphaB-crystallin-binding protein CRABP2	0.89	16
I1VZ6	Serine/threonine-protein phosphatase 6 regulatory ankyrin repeat subunit A	0.89	4
Q8N848	Serine/threonine-protein phosphatase 6 regulatory ankyrin repeat subunit C	0.88	1
Q72557	Lysyl oxidase	0.85	4
P28558	Ribonuclease D most single-stranded-interacting protein 1	0.85	3
Q14157	Ubiquitin-associated protein 2-like	0.79	4
P52597	Heterogeneous nuclear ribonucleoprotein F	0.75	10
P02533	Dolichyl-diphosphooligosaccharide–RNA-binding protein Staufen homolog 1	0.75	7
Q867PK6	RNA-binding protein 14	0.75	11
Q867W4	Anchiamatinin-2	0.72	2
Q9U943	Ablaxin-10	0.71	3
P07411	AlphaB-crystallin-binding protein	0.69	1
Q9U854	DnaJ homolog subfamily B member 11	0.69	2
Q06210	Glutamine-fructose-6-phosphate aminotransferase (isomerizing) 1	0.68	5
Q93432	Anchian 1 isoform CRA_b	0.67	2
P11586	C-tetrahydrofolate synthase cytoplasmic	0.67	3
Q9H9R7	Serine/threonine-protein phosphatase 6 regulatory subunit 3	0.66	14
P40227	T-complex protein 1 subunit zeta	0.66	2
P02533	Serine/threonine-protein kinase 2A 55 kDa regulatory subunit B alpha isoform	0.65	1
Q9N2Z8	Insulin-like growth factor 2 mRNA-binding protein 1	0.64	8
Q9Y277	Voltage-dependent anion-selective channel protein 3	0.64	2
Q967L1	Unconventional myosin-IV	0.63	1
Q8WV95	Chaperone IV beta homolog	0.61	29
BAE0N9	Glutamate dehydrogenase	0.61	3
P22955	Cytochrome b-c1 complex subunit 2 mitochondrial	0.61	4
Q92527	Widespread membrane spanning protein C2orf44	0.60	3
P84098	60S ribosomal protein L19	0.60	1
P11182	Lipoamide acyltransferase component of branched-chain alpha-keto acid dehydrogenase	0.60	5
ARK122	KIAA0946 isoform CRA_b	0.60	1
P02533	Mitochondrial 1-phosphate quinolinate transferase 7A	0.59	2
Q9Y265	RuvB-like 1	0.59	1
P11413	Glucose-6-phosphate 1-dehydrogenase	0.58	2
P02533	Dolichyl-diphosphooligosaccharide–RNA-binding protein MCM7	0.58	2
P09042	Eukaryotic initiation factor 4A-I	0.57	10
Q75881	25-hydroxycholesterol 7-alpha-hydroxylase	0.57	1
P62537	Peptidyl-prolyl cis-trans isomerase A	0.56	2
P02533	Recombinant 70 kDa RNA-binding subunit	0.56	4
Q9Y678	Cotranslational subunit gamma-1	0.56	1
PS1571	Translocon-associated protein subunit delta	0.56	3
P15005	Cotranslational subunit beta	0.55	2
P12258	Unconventional myosin-IV	0.55	1
Q9H479	Fructosamine-3-kinase	0.54	4
Q567R6	Single-stranded DNA-binding protein	0.54	4
P02533	Heterogeneous nuclear ribonucleoprotein D0	0.53	4
Q9ULV0	Unconventional myosin-Vb	0.53	34
Q86X52	Chaperonin 60 subunit 1	0.52	1
Q43143	Putative pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15	0.51	2
P02533	Recombinant 70 kDa RNA-binding protein K	0.50	7
P06830	Peroxiredoxin-1	0.50	7
Q43175	D-3-phosphoglycerate dehydrogenase	0.50	2
P02533	Unconventional myosin-Vb	0.49	1
P27708	CAD protein	0.49	2
P04843	Dolichyl-diphosphooligosaccharide–protein glycosyltransferase subunit 1	0.49	2
Q87B22	Spermatogenesis-associated protein 20	0.48	3
P02533	Keratin type I cytoskeletal 1	0.48	2
A8NLN1	Pyrimidine ring binding protein 1 isoform CRA_b	0.47	1
PS1812	Recombinant protein SS kinase alpha-3	0.46	1
Q9H611	Isthmin-2	0.46	1
Q9CQG0	Myosin binding protein 1A	0.46	1
Q00571	ATP-dependent RNA helicase DDX3X	0.46	21
P50454	Serpin H1	0.45	2
P02533	Unconventional myosin-Vb	0.44	1
P15924	Desmoplakin	0.43	24
P49368	T-complex protein 1 subunit gamma	0.43	2
Q43795	Unconventional myosin-IVb	0.42	3
P02533	Cotranslational subunit delta	0.42	1
Q98VG3	Kinesin-like protein Kif-C3	0.42	12
Q00341	Villin	0.41	1
P19404	Alpha/beta cyclin factor 3	0.41	1
P72679	Fractin gamma chain	0.41	11
P30837	Aldehyde dehydrogenase X mitochondrial	0.39	1
P02675	Fibrinogen beta chain	0.39	5
P02533	Unconventional myosin-Vb	0.39	4
P23284	Peptidyl-prolyl cis-trans isomerase B	0.38	1
Q15027	Protein transport protein Sec16A	0.38	2
P02533	Peroxiredoxin-2	0.37	2
Q13388	Spliceosomal RNA helicase DDX3B	0.37	1
Q9Y5V3	Melanoma-associated antigen D1	0.37	1
Q13268	Dehydrogenases/reductase SDR family member 2 mitochondrial	0.37	11
P02533	Peptidyl-prolyl cis-trans isomerase A	0.37	1
P50981	T-complex protein 1 subunit delta	0.37	1
Q01085	Nucleolin TAR	0.37	2
P63261	Actin cytoplasmic 2	0.36	21
P02533	Unconventional myosin-Vb	0.36	1
P02533	Unconventional myosin-Vb	0.35	1
P15436	Protein transport protein Sec23A	0.36	2
Q867G7	Retrotransposon-derived protein PEG10	0.36	1
P02545	Far upstream element-binding protein 2	0.36	1
P10143	Heat shock 70 kDa nucleolar protein H	0.35	9
P12931	Heat shock protein 70 kDa mitochondrial	0.35	3
P55072	Translational endoplasmic reticulum ATPase	0.35	1
P02533	AlphaB-crystallin subunit alpha mitochondrial	0.34	7
P09382	Galedin-1	0.34	2
P02768	Serum albumin	0.33	22
P02533	Fibrinogen alpha chain	0.32	13
Q021044	Probable alpha-1B-glycoprotein 6-interacting protein	0.32	1
P49327	Fatty acid synthase	0.31	5
P19474	E3 ubiquitin-protein ligase TRIM21	0.31	15
Q43747	GDP	0.31	1
P45654	Short-branched chain specific acyl-CoA dehydrogenase mitochondrial	0.30	1
P78527	DNA-dependent protein kinase catalytic subunit	0.30	4
P10809	60 kDa heat shock protein mitochondrial	0.30	8
P02533	Peptidyl-prolyl cis-trans isomerase A	0.29	2
Q15365	Polycy-binding protein 1	0.29	4
Q69996	Pyridoxal-dependent decarboxylase domain-containing protein 1	0.28	1
Q43136	AlphaB-crystallin subunit cytoplasmic	0.28	1
P02533	Ubulin alpha-1B-glycoprotein	0.27	14
Q3LX3	Bifunctional ATP-dependent dihydroxyacetone kinase/FAD-AMP lyase cyclizing	0.27	1
P62629	60S ribosomal protein L23	0.27	1
P02533	Unconventional myosin-Vb	0.26	1
P31483	Nucleolin TATA 1 isoform p40	0.26	1
P14868	Aspartate-tRNA ligase cytosolic	0.26	1
P12026	Proliferating cell nuclear antigen	0.26	1
Q10110	Keratin type I cytoskeletal 1	0.25	15
P36542	ATP synthase subunit gamma mitochondrial	0.25	1
Q82T6	Abnormal spindle-like microcephaly-associated protein	0.24	1
P02533	Sh3bp70D	0.24	2
P49411	Fringing factor Tu mitochondrial	0.23	4
Q15366	Polycy-binding protein 2	0.23	4
Q9UBM7	7-dehydrocholesterol reductase	0.23	1
P02533	Heat shock 70 kDa nucleolar protein U	0.22	6
P68104	Elongation factor 1-alpha 1	0.21	7
P05787	Keratin type I cytoskeletal 8	0.21	24
P01707	Heat shock 70 kDa protein 1A/TATA box-binding protein	0.21	5
P75379	Vesicle-associated membrane protein 4	0.21	1
P05783	Keratin type I cytoskeletal 18	0.19	18
P06576	ATP synthase subunit beta mitochondrial	0.18	3
P02533	CD105 heat shock protein 90-beta 1	0.18	1
P02533	Heat shock protein HSP 90-beta 1	0.18	5
P04075	Fructose-bisphosphatase 3	0.17	1
P25711	ADP/ATP translocase 2	0.17	2
P02533	Unconventional myosin-Vb	0.17	1
P02533	Unconventional myosin-Vb	0.16	1
D3DPG0	Titin isoform CRA_a	0.16	1
Q1KMD5	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.16	1
P02533	Phosphoglycerate kinase	0.16	1
Q15364	Protein disulfide-isomerase AB	0.16	1
P14618	l-Pyruvate kinase PKR	0.16	2
P60701	UDP-glucose 4-dehydrogenase	0.16	3
P02533	Heat shock protein HSP 90-beta 1	0.15	4
P17844	Probable ATP-dependent RNA helicase DDX5	0.15	1
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.14	1
P02533	Glyceraldehyde-3-phosphate dehydrogenase	0.14	1
P785Y8	Phosphatidylethanolamine N-methyltransferase	0.13	6
P58107	Eppikain	0.13	2
P02769	Serum albumin	0.13	56
P02533	Keratin type I cytoskeletal 5	0.13	4
P02533	Keratin Type I cytoskeletal 14	0.12	4
P14625	Endoplasm	0.12	1
HBVRG3	Keratin	0.12	16
P02533	Probable multifunctional enzyme type 2	0.10	9
P33845	Keratin type I cytoskeletal 10	0.10	11
A2NU12	Lambda-chain AA-20 to 215	0.10	1
P02751	beta-1,4-galactosidase	0.10	3
P00338	L-lactate dehydrogenase A chain	0.09	2
Q88Y23	Hormerin	0.08	2
P02533	Neurogranin	0.08	1
B6ZGU2	Nuclear receptor subfamily 2 group F member 6	0.08	1
P35908	Keratin type I cytoskeletal 2 epidermal	0.08	6
Q72557	Defensin-like protein 9	0.08	2
Q9N7V7	ATPase family AAA-domain-containing protein 3A	0.04	1
P00610	Claudin heavy chain 1	0.04	2