

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q14188	Transcription factor Dp-2	9.09	12
A1L0T0	Acetolactate synthase-like protein	1.92	2
Q9H583	HEAT repeat-containing protein 1	1.43	2
Q9N2G9	SIN3/CNF-related matrix-associated actin-dependent regulator of chromatin subfamily A-like protein 1	1.10	2
Q9UBX3	Mitochondrial dicarboxylate carrier	1.09	2
P31930	Cytochrome b-c1 complex subunit 1 mitochondrial	1.06	1
D3DR22	Hydroxysteroid (17-beta) dehydrogenase 12 isoform CRA_a	1.05	4
Q9Y277	Voltage-dependent anion-selective channel protein 3	1.04	3
Q43795	Unconventional myosin-Ib	1.03	7
Q9UJS0	Calcium-binding mitochondrial carrier protein Aralar2	0.97	3
Q6P1L8	39S ribosomal protein L14 mitochondrial	0.87	2
Q9HB29	RNA helicase	0.86	1
Q81D30	Alanine aminotransferase 2	0.84	1
Q9ULV0	Unconventional myosin-Vb	0.84	41
Q16254	Transcription factor E2F4	0.83	2
P62829	60S ribosomal protein L23	0.83	4
P27694	Replication protein A 70 kDa DNA-binding subunit	0.80	4
Q9HCC0	Methylcrotonoyl-CoA carboxylase beta chain mitochondrial	0.79	4
P50991	T-complex protein 1 subunit delta	0.76	3
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.76	3
P52597	Heterogeneous nuclear ribonucleoprotein F	0.75	6
Q6Y1N16	Hydroxysteroid dehydrogenase-like protein 2	0.75	1
Q9H9B4	Sideroflexin-1	0.74	2
Q95433	Activator of 90 kDa heat shock protein ATPase homolog 1	0.73	1
P31943	Heterogeneous nuclear ribonucleoprotein H	0.71	15
Q94905	Erlin-2	0.70	2
Q13509	Tubulin beta-3 chain	0.66	14
Q9H3K6	BolA-like protein 2	0.65	1
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.64	6
Q9NOX4	Unconventional myosin-Vc	0.64	3
P45954	Shortbranched chain specific acyl-CoA dehydrogenase mitochondrial	0.64	3
Q15436	Protein transport protein Sec23A	0.63	3
Q14157	Ubiquitin-associated protein 2-like	0.62	2
P06576	ATP synthase subunit beta mitochondrial	0.61	9
Q5H9R7	Serine/threonine-protein phosphatase 6 regulatory subunit 3	0.60	4
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.59	14
Q99623	Prohibitin-2	0.59	4
Q13907	Isopentenyl-diphosphate Delta-isomerase 1	0.57	1
P51571	Translocase-associated protein subunit delta	0.56	2
P07437	Tubulin beta chain	0.55	26
P11182	Lipoamide acyltransferase component of branched-chain alpha-keto acid dehydrogenase complex mitochondrial	0.55	1
Q60701	UDP-glucose 6-dehydrogenase	0.54	11
P60842	Eukaryotic initiation factor 4A-I	0.54	7
P68363	Tubulin alpha-1B chain	0.54	22
Q15523	ATP-dependent RNA helicase DDX3Y	0.53	6
P01859	Ig gamma-2 chain C region	0.52	2
Q9H8H3	Methyltransferase-like protein 7A	0.51	1
P54136	Arginine--tRNA ligase cytoplasmic	0.50	5
Q04837	Single-stranded DNA-binding protein mitochondrial	0.50	5
P49411	Elongation factor Tu mitochondrial	0.50	8
P32322	Pyroline-5-carboxylate reductase 1 mitochondrial	0.49	2
P33993	DNA replication licensing factor MCM7	0.49	1
Q06520	Bile salt sulfotransferase	0.49	2
P68371	Tubulin beta-4B chain	0.49	22
P62937	Peptidyl-prolyl cis-trans isomerase A	0.48	1
Q43813	LanC-like protein 1	0.47	1
P02675	Fibrinogen beta chain	0.47	4
Q43175	D-3-phosphoglycerate dehydrogenase	0.47	1
P48643	T-complex protein 1 subunit epsilon	0.45	2
Q02671	Fibrinogen alpha chain	0.44	14
Q9Y5V3	Melanoma-associated antigen D1	0.43	1
P26641	Elongation factor 1-gamma	0.43	1
Q08211	ATP-dependent RNA helicase A	0.43	2
P14866	Aspartate--tRNA ligase cytoplasmic	0.43	2
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.42	2
P51659	Peroxisomal multifunctional enzyme type 2	0.41	33
P36542	ATP synthase subunit gamma mitochondrial	0.41	2
P62826	GTP-binding nuclear protein Ran	0.40	2
Q9N2I8	Insulin-like growth factor 2 mRNA-binding protein 1	0.39	3
P25705	ATP synthase subunit alpha mitochondrial	0.39	6
P05141	ADP/ATP translocase 2	0.38	5
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.38	4
P41252	Isoleucine--tRNA ligase cytoplasmic	0.38	1
P68104	Elongation factor 1-alpha 1	0.37	10
P49368	T-complex protein 1 subunit gamma	0.36	1
A7B136	p180ribosome receptor	0.36	1
P18621	60S ribosomal protein L17	0.36	2
Q60547	GDP-mannose 46 dehydratase	0.36	1
P78527	DNA-dependent protein kinase catalytic subunit	0.35	3
A2NUT2	Lambda-chain (AA-20 to 215)	0.34	6
P17987	T-complex protein 1 subunit alpha	0.34	1
P49327	Fatty acid synthase	0.34	4
Q15084	Protein disulfide-isomerase A6	0.34	3
Q00571	ATP-dependent RNA helicase DDX3X	0.33	5
P02679	Fibrinogen gamma chain	0.31	6
P10809	60 kDa heat shock protein mitochondrial	0.30	6
P12004	Proliferating cell nuclear antigen	0.30	1
Q15365	Poly(rC)-binding protein 1	0.30	3
P11142	Heat shock cognate 71 kDa protein	0.30	7
P17944	Probable ATP-dependent RNA helicase DDX5	0.29	1
P02768	Serum albumin	0.29	14
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.28	6
Q9UBM7	7-dehydrocholesterol reductase	0.26	1
Q15058	Kinesin-like protein KIF14	0.26	19
Q06830	Peroxiredoxin-1	0.26	2
Q96PK6	RNA-binding protein 14	0.24	1
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.23	10
Q6IB91	PCK2 protein	0.23	1
Q149N6	Dedicator of cytokinesis 4	0.21	10
P61878	Heterogeneous nuclear ribonucleoprotein K	0.21	1
P38846	Stress-70 protein mitochondrial	0.21	1
Q9Y4L1	Hypoxia up-regulated protein 1	0.21	1
P01700	Ig lambda chain V-I region HA	0.20	2
P05787	Keratin type II cytoskeletal 8	0.20	17
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.19	7
P15924	Desmoplakin	0.19	7
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.18	2
P58107	Epiplakin	0.17	2
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.16	1
P08238	Heat shock protein HSP 90-beta	0.16	3
Q9H0D6	5-3 exoribonuclease 2	0.15	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.13	4
P05783	Keratin type I cytoskeletal 18	0.13	9
P02769	Serum albumin	0.13	42
P60709	Actin cytoplasmic 1	0.13	6
P11021	78 kDa glucose-regulated protein	0.12	7
P07900	Heat shock protein HSP 90-alpha	0.12	1
Q58K21	DBIRD complex subunit ZNF326	0.12	1
HEVVRG3	Keratin 1	0.11	11
P00338	L-lactate dehydrogenase A chain	0.11	2
P08107	Heat shock 70 kDa protein 1A1B	0.10	1
P06748	Nucleophosmin	0.10	1
P08727	Keratin type I cytoskeletal 19	0.09	2
P35908	Keratin type II cytoskeletal 2 epidermal	0.06	4
P35527	Keratin type I cytoskeletal 9	0.04	2
P13645	Keratin type I cytoskeletal 10	0.02	1