

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q03184	Histone-lysine N-methyltransferase 2A	3.01	16
Q08210	Glycyl-tRNA synthetase [isomerizing] 1	1.25	9
Q8NBS9	Thioredoxin domain-containing protein 5	1.16	3
Q75874	Isochorate dehydrogenase [NAD(P)] cytoplasmic	1.04	1
P11586	C-1-tetrahydrofolate synthase cytoplasmic	1.00	3
Q8Y116	Hydroxyteroid dehydrogenase-like protein 2	0.97	2
Q95433	Activator of 90 kDa heat shock protein ATPase homolog 1	0.92	2
Q15523	ATP-dependent RNA helicase DDX3Y	0.92	14
Q92900	Regulator of nonsense transcripts 1	0.91	1
Q81030	Alanine aminotransferase 2	0.79	1
P51571	Translocon-associated protein subunit delta	0.77	4
B0Y1W6	Archain 1 isoform CRA_a	0.77	1
Q43813	LanC-like protein 1	0.76	3
Q43795	Unconventional myosin-Ib	0.74	6
P13639	Elongation factor 2	0.74	7
P00558	Phosphoglycerate kinase 1	0.74	2
P52597	Heterogeneous nuclear ribonucleoprotein F	0.73	7
Q13885	Tubulin beta-2A chain	0.71	21
Q49327	Fatty acid synthase	0.71	12
Q15084	Serine/threonine-protein phosphatase 6 regulatory ankyrin repeat subunit A	0.70	1
P53396	ATP-citrate synthase	0.70	1
P47897	Glutamine-tRNA ligase	0.70	4
P35232	Prohibitin	0.70	3
Q99623	Prohibitin-2	0.69	6
Q9UBX3	Mitochondrial dicarboxylate carrier	0.69	1
ANLN1	Polypyrimidine tract binding protein 1 isoform CRA_b	0.69	2
P26641	Elongation factor 1 gamma	0.69	3
Q9NZC9	SWI5NF-related matrix-associated actin-dependent regulator of chromatin subfamily A-like protein 1	0.67	1
Q9ULV0	Unconventional myosin-Vb	0.65	38
Q9H9H3	Methyltransferase-like protein 7A	0.65	2
Q94905	Erlin-2	0.63	2
P23284	Peptidyl-prolyl cis-trans isomerase B	0.61	1
Q9H3K6	BoA-like protein 2	0.60	1
Q00571	ATP-dependent RNA helicase DDX3X	0.59	12
D3D922	Hydroxyteroid (17-beta) dehydrogenase 12 isoform CRA_a	0.58	2
P07437	Tubulin beta chain	0.57	33
Q9NQX4	Unconventional myosin-Vc	0.57	3
Q02671	Fibrinogen alpha chain	0.57	22
R45564	Shunt branched chain specific acyl-CoA dehydrogenase mitochondrial	0.57	3
Q92841	Probable ATP-dependent RNA helicase DDX17	0.56	5
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.56	6
Q9Y411	Unconventional myosin-Va	0.55	3
Q30101	Protein disulfide-isomerase A3	0.55	1
Q9Y5V3	Melanoma-associated antigen D1	0.54	2
Q95793	Double-stranded RNA-binding protein Staufin homolog 1	0.54	4
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.53	2
P68371	Tubulin beta-4B chain	0.53	29
P68363	Tubulin alpha-1B chain	0.53	26
Q02227	T-complex protein 1 subunit zeta	0.52	1
Q8IB91	PKC2 protein	0.52	5
Q12904	Aminase/17RNA synthase complex-interacting multifunctional protein 1	0.50	1
Q3LXA3	Bifunctional ATP-dependent dihydroxyacetone kinase/FAD-AMP lyase (cycling)	0.50	3
Q43251	RNA binding protein fox-1 homolog 2	0.50	1
Q9H984	Sideroflexin-1	0.49	1
P02751	Fibronectin	0.49	17
Q08211	ATP-dependent RNA helicase A	0.49	3
Q06830	Peroxisome-1	0.48	6
Q48428	Aldo-keto reductase family 1 member C1	0.48	1
P18085	ADP-ribosylation factor 4	0.47	2
P53618	Coatomer subunit beta	0.47	1
P01859	Ig gamma-2 chain C region	0.47	2
P17844	Probable ATP-dependent RNA helicase DDX5	0.46	3
P31943	Heterogeneous nuclear ribonucleoprotein H	0.46	11
Q05547	GDP-mannose 4-epimerase	0.45	2
P62937	Peptidyl-prolyl cis-trans isomerase A	0.44	1
Q06520	Bile salt sulfotransferase	0.44	2
Q15436	Protein transport protein Sec23A	0.44	2
Q43175	D-3-phosphoglycerate dehydrogenase	0.44	1
Q9H9R7	Serine/threonine-protein phosphatase 6 regulatory subunit 3	0.43	3
P30041	Peroxisome-6	0.41	1
P35637	RNA-binding protein FUS	0.41	1
Q43143	Protein pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15	0.41	1
P48643	T-complex protein 1 subunit epsilon	0.40	2
P08236	Heat shock protein HSP 90-beta	0.40	12
P50891	T-complex protein 1 subunit delta	0.39	1
Q92945	Fc gammaRIIb binding protein 2	0.39	1
P45880	Voltage-dependent anion-selective channel protein 2	0.39	2
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.39	7
P14868	Asparagine-tRNA ligase cytoplasmic	0.38	2
Q12931	Heat shock protein 75 kDa mitochondrial	0.38	2
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.38	2
Q15365	Poly(C)-binding protein 1	0.37	5
P36542	ATP synthase subunit gamma mitochondrial	0.37	2
Q04837	Single-stranded DNA-binding protein mitochondrial	0.37	3
P07237	Protein disulfide-isomerase	0.36	3
Q15084	Protein disulfide-isomerase A6	0.36	4
P62829	60S ribosomal protein L23	0.35	1
P2679	Fibrinogen gamma chain	0.34	9
P49368	T-complex protein 1 subunit gamma	0.34	1
P27694	Replication protein A 70 kDa DNA-binding subunit	0.34	1
P62967	Ubiquitin-60S ribosomal protein L40	0.34	1
Q9UBM7	7-dehydrocholesterol reductase	0.33	2
P32322	Pyrroline-5-carboxylate reductase 1 mitochondrial	0.33	1
P18621	60S ribosomal protein L17	0.32	2
Q13268	Dehydrogenase/reductase SDR family member 2 mitochondrial	0.31	8
P07400	Heat shock protein HSP 90-alpha	0.31	6
P60842	Eukaryotic initiation factor 4A-1	0.30	4
A2NUT2	Lambda-chain (AA-20 to 215)	0.30	6
P10809	60 kDa heat shock protein mitochondrial	0.29	7
P19338	Nucleolin	0.29	1
Q9H0D6	5-3 exonuclease 2	0.29	4
P12004	Proliferating cell nuclear antigen	0.28	1
HFY758	Protein-L-isopartate O-methyltransferase	0.28	13
P14625	Endoleasmin	0.28	4
P19474	E3 ubiquitin-protein ligase TRIM21	0.28	12
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.27	3
P02675	Fibrinogen beta chain	0.27	2
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.27	7
P51978	Heterogeneous nuclear ribonucleoprotein K	0.27	2
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.26	14
P68104	Elongation factor 1-alpha 1	0.26	8
Q14847	LIM and SH3 domain protein 1	0.26	2
Q15058	Kinesin-like protein Kif-14	0.25	22
P78527	DNA-dependent protein kinase catalytic subunit	0.25	2
Q60701	UDP-glucose 6-dehydrogenase	0.24	5
P05141	ADP/ATP translocase 2	0.23	3
P05576	ATP synthase subunit beta mitochondrial	0.23	3
Q96PK6	RNA-binding protein 14	0.22	1
P51659	Peroxisomal multifunctional enzyme type 2	0.22	20
P49411	Elongation factor Tu mitochondrial	0.21	3
P04408	Glyceraldehyde-3-phosphate dehydrogenase	0.20	4
P25705	ATP synthase subunit alpha mitochondrial	0.20	3
P38646	Stress-70 protein mitochondrial	0.19	1
P08107	Heat shock 70 kDa protein 1A1B	0.19	4
P54138	Arginine-tRNA ligase cytoplasmic	0.18	1
D3DPG0	Titin isoform CRA_a	0.18	1
P60709	Actin cytoplasmic 1	0.18	11
P02768	Serum albumin	0.16	9
P05187	Keratin type II cytoskeletal 8	0.16	16
P11021	78 kDa glucose-regulated protein	0.16	11
Q9BKZ1	DBIRD complex subunit ZNF326	0.15	2
P15924	Desmoplakin	0.14	6
Q05183	Keratin type I cytoskeletal 1B	0.13	11
P14618	Pyruvate kinase PKM	0.13	1
P02769	Serum albumin	0.13	50
Q06748	Nucleophosmin	0.12	2
Q149N8	Dedicator of cytokinesis 4	0.12	6
H6VRC3	Keratin 1	0.11	14
P11142	Heat shock cognate 71 kDa protein	0.11	2
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.10	1
P52272	Heterogeneous nuclear ribonucleoprotein M	0.07	1
P00338	L-lactate dehydrogenase A chain	0.07	1
P08727	Keratin type I cytoskeletal 19	0.06	1
P07814	Bifunctional glutamateproline-tRNA ligase	0.06	1
P13645	Keratin type I cytoskeletal 10	0.06	6
P35527	Keratin type I cytoskeletal 9	0.05	4
Q00610	Clathrin heavy chain 1	0.05	3
P35608	Keratin type II cytoskeletal 2 epidermal	0.02	1