

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q95948	One cut domain family member 2	4.95	5
Q13347	Eukaryotic translation initiation factor 3 subunit I	1.06	1
O75147	Obscurin-like protein 1	1.04	1
Q9Y265	RuvB-like 1	1.00	2
P30041	Peroxiredoxin-6	0.99	4
Q8NBS9	Thioredoxin domain-containing protein 5	0.96	2
Q9Y678	Coatomer subunit gamma-1	0.92	2
P23528	Cofilin-1	0.90	3
Q99623	Prohibitin-2	0.90	7
O15523	ATP-dependent RNA helicase DDX3Y	0.84	12
P55795	Heterogeneous nuclear ribonucleoprotein H2	0.77	5
Q8T1B2	Spermatogenesis-associated protein 20	0.75	4
P63151	Serinethreonine-protein phosphatase 2A 55 kDa regulatory subunit B alpha isoform	0.74	1
Q13509	Tubulin beta-3 chain	0.74	15
P32119	Peroxiredoxin-2	0.73	4
Q8HWP6	Class IVb beta tubulin	0.65	22
Q9BVG8	Kinesin-like protein KIFC3	0.65	14
Q14157	Ubiquitin-associated protein 2-like	0.63	2
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.62	14
Q13838	Spliceosome RNA helicase DDX39B	0.61	2
P35637	RNA-binding protein FUS	0.61	2
Q6P996	Pyridoxal-dependent decarboxylase domain-containing protein 1	0.61	3
P38919	Eukaryotic initiation factor 4A-III	0.60	1
Q06210	Glutamine--fructose-6-phosphate aminotransferase (isomerizing) 1	0.60	3
Q86X52	Chondroitin sulfate synthase 1	0.60	1
Q9UBS4	DnaJ homolog subfamily B member 11	0.58	1
Q9H6R7	WD repeat and coiled-coil-containing protein C2orf44	0.58	2
Q01085	Nucleolysin TIAR	0.58	3
Q06830	Peroxiredoxin-1	0.58	6
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.57	5
Q15436	Protein transport protein Sec23A	0.57	3
Q95433	Activator of 90 kDa heat shock protein ATPase homolog 1	0.56	1
P52597	Heterogeneous nuclear ribonucleoprotein F	0.56	5
P51571	Translocon-associated protein subunit delta	0.53	2
Q96PK6	RNA-binding protein 14	0.51	5
Q9H8H3	Methyltransferase-like protein 7A	0.49	1
P02671	Fibrinogen alpha chain	0.49	15
Q9Y411	Unconventional myosin-Va	0.49	2
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.48	2
Q8ULV0	Unconventional myosin-Vb	0.48	22
O00743	Serinethreonine-protein phosphatase 6 catalytic subunit	0.47	1
Q00341	Vigilin	0.47	1
P15924	Desmoplakin	0.47	19
P61204	ADP-ribosylation factor 3	0.47	1
Q13162	Peroxiredoxin-4	0.47	1
P02679	Fibrinogen gamma chain	0.46	9
P07437	Tubulin beta chain	0.45	20
P23284	Peptidyl-prolyl cis-trans isomerase B	0.44	1
B2RWN5	HEAT repeat containing 1	0.43	1
Q9H9R7	Serinethreonine-protein phosphatase 6 regulatory subunit 3	0.43	6
P12004	Proliferating cell nuclear antigen	0.43	2
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.42	2
P02675	Fibrinogen beta chain	0.42	4
Q15084	Protein disulfide-isomerase A6	0.42	4
P36542	ATP synthase subunit gamma mitochondrial	0.41	2
Q86TG7	Retrotransposon-derived protein PEG10	0.41	1
P09382	Galectin-1	0.41	2
P04792	Heat shock protein beta-1	0.40	1
O00425	Insulin-like growth factor 2 mRNA-binding protein 3	0.40	1
P55072	Transitional endoplasmic reticulum ATPase	0.40	1
Q00571	ATP-dependent RNA helicase DDX3X	0.40	13
Q06YQ0	Rat GTPase-activating protein subunit alpha-1	0.40	1
P17844	Probable ATP-dependent RNA helicase DDX5	0.39	4
P63261	Actin cytoplasmic 2	0.39	16
P51659	Peroxisomal multifunctional enzyme type 2	0.38	29
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.37	16
Q8WUM4	Programmed cell death 6-interacting protein	0.36	1
P61978	Heterogeneous nuclear ribonucleoprotein K	0.34	3
P19474	E3 ubiquitin-protein ligase TRIM21	0.34	12
O75379	Vesicle-associated membrane protein 4	0.34	2
P49327	Fatty acid synthase	0.34	4
P68363	Tubulin alpha-1B chain	0.32	12
P05787	Keratin type II cytoskeletal 8	0.32	27
Q15027	Protein transport protein Sec16A	0.31	1
Q3LXA3	Bifunctional ATP-dependent dihydroxyacetone kinaseFAD-AMP lyase cyclizing	0.31	1
P31483	Nucleolysin TIA-1 isoform p40	0.30	1
P49411	Elongation factor Tu mitochondrial	0.29	4
P60842	Eukaryotic initiation factor 4A-I	0.29	3
P25705	ATP synthase subunit alpha mitochondrial	0.29	4
Q95793	Double-stranded RNA-binding protein Staufen homolog 1	0.27	1
P05141	ADP-ATP translocase 2	0.26	3
P13639	Elongation factor 2	0.24	1
P31943	Heterogeneous nuclear ribonucleoprotein H	0.24	4
P78527	DNA-dependent protein kinase catalytic subunit	0.24	2
P08721	Keratin type I cytoskeletal 19	0.23	7
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.23	2
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.22	8
P02768	Serum albumin	0.22	10
Q92841	Probable ATP-dependent RNA helicase DDX17	0.21	1
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.21	4
P11021	78 kDa glucose-regulated protein	0.20	12
P14625	Endoplasmic	0.20	2
P01700	Ig lambda chain V-I region HA	0.20	2
Q14847	LIM and SH3 domain protein 1	0.20	1
P04075	Fructose-bisphosphate aldolase A	0.19	1
P14618	Pyruvate kinase PKM	0.19	2
P05783	Keratin type I cytoskeletal 18	0.19	13
P54136	Arginine--tRNA ligase cytoplasmic	0.19	1
D3DPG0	Titin isoform CRA_a	0.19	1
P10809	60 kDa heat shock protein mitochondrial	0.18	3
P08107	Heat shock 70 kDa protein 1A1B	0.18	3
Q15366	PolyC-binding protein 2	0.18	2
P02769	Serum albumin	0.18	57
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.17	1
Q149N6	Dedicator of cytokinesis 4	0.16	7
Q15365	PolyC-binding protein 1	0.16	1
A2NUT2	Lambda-chain AA_20 to 215	0.16	2
P08238	Heat shock protein HSP 90-beta	0.16	3
O15090	Zinc finger protein 536	0.15	1
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.14	2
P11142	Heat shock cognate 71 kDa protein	0.12	2
Q5BKZ1	DBIRD complex subunit ZNF326	0.11	1
H6VRG3	Keratin 1	0.11	11
P00338	L-lactate dehydrogenase A chain	0.11	2
O60701	UDP-glucose 6-dehydrogenase	0.10	1
P68104	Elongation factor 1-alpha 1	0.08	1
P02751	Fibronectin	0.08	1
Q15149	Plectin	0.07	2
P35908	Keratin type II cytoskeletal 2 epidermal	0.07	5
P13645	Keratin type I cytoskeletal 10	0.06	4
Q00610	Clathrin heavy chain 1	0.04	2
P35527	Keratin type I cytoskeletal 9	0.04	2