

Accession Number	Description	Fold Enrichment	UniquePeptideCount
O15156	Zinc finger and BTB domain-containing protein 7B	5.93	6
P30041	Peroxisomal protein PEX6	1.06	4
P51170	Amiloride-sensitive sodium channel subunit gamma	0.98	1
Q13509	Tubulin beta-3 chain	0.98	20
Q9H583	HEAT repeat-containing protein 1	0.96	1
O43813	LanC-like protein 1	0.92	3
O95299	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 10 mitochondrial	0.91	1
Q13885	Tubulin beta-2A chain	0.90	20
Q9H3K6	BoA-like protein 2	0.87	2
O15523	ATP-dependent RNA helicase DDX3Y	0.82	11
P55795	Heterogeneous nuclear ribonucleoprotein H2	0.81	5
P33993	DNA replication licensing factor MCM7	0.73	2
Q8NBS9	Thioredoxin domain-containing protein 5	0.69	1
Q49AG4	DIS3 protein	0.67	1
Q9Y5V3	Melanoma-associated antigen D1	0.64	2
Q92841	Probable ATP-dependent RNA helicase DDX17	0.62	5
P61978	Heterogeneous nuclear ribonucleoprotein K	0.61	6
P51659	Peroxisomal multifunctional enzyme type 2	0.61	44
P62829	60S ribosomal protein L23	0.60	3
Q9Y277	Voltage-dependent anion-selective channel protein 3	0.59	2
P68371	Tubulin beta-4B chain	0.59	26
O95433	Activator of 90 kDa heat shock protein ATPase homolog 1	0.58	1
G3V1B5	Down syndrome cell adhesion molecule like 1 isoform CRA_a	0.57	1
A6NLN1	Polypyrimidine tract binding protein 1 isoform CRA_b	0.56	1
O43795	Unconventional myosin-Ib	0.55	3
Q12904	Aminoacyl tRNA synthase complex-interacting multifunctional protein 1	0.55	1
Q6IB91	PCK2 protein	0.55	4
O00148	ATP-dependent RNA helicase DDX39A	0.53	1
Q9UDY2	Tight junction protein ZO-2	0.51	2
P11413	Glucose-6-phosphate 1-dehydrogenase	0.51	1
P52597	Heterogeneous nuclear ribonucleoprotein F	0.50	4
P07437	Tubulin beta chain	0.49	24
P17844	Probable ATP-dependent RNA helicase DDX5	0.49	5
P62937	Peptidyl-prolyl cis-trans isomerase A	0.49	1
Q06210	Glutamine-fructose-6-phosphate aminotransferase [isomerizing] 1	0.48	2
P68363	Tubulin alpha-1B chain	0.47	17
Q15365	PolyC-binding protein 1	0.46	5
Q15436	Protein transport protein Sec23A	0.46	2
O00571	ATP-dependent RNA helicase DDX3X	0.45	14
Q9UBB4	Ataxin-10	0.45	1
P14868	Aspartate-tRNA ligase cytoplasmic	0.45	2
Q3LXA3	Bifunctional ATP-dependent dihydroxyacetone kinase/FAD-AMP lyase cyclizing	0.45	2
P35232	Prohibitin	0.45	1
Q06830	Peroxisomal protein PEX1	0.44	4
P02751	Fibronectin	0.43	11
P49411	Elongation factor Tu mitochondrial	0.43	6
P11182	Lipoamide acyltransferase component of branched-chain alpha-keto acid dehydrogenase	0.42	2
P25705	ATP synthase subunit alpha mitochondrial	0.42	6
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.41	3
O95793	Double-stranded RNA-binding protein Staufen homolog 1	0.41	2
O60701	UDP-glucose 6-dehydrogenase	0.40	7
P31943	Heterogeneous nuclear ribonucleoprotein H	0.40	7
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.38	8
P41252	Isoleucine-tRNA ligase cytoplasmic	0.38	1
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.37	3
P13639	Elongation factor 2	0.37	2
B7ZLE5	FN1 protein	0.37	9
Q15084	Protein disulfide-isomerase A6	0.36	3
Q06520	Bile salt sulfotransferase	0.36	1
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.34	1
Q5H9R7	Serine/threonine-protein phosphatase 6 regulatory subunit 3	0.33	4
P48643	T-complex protein 1 subunit epsilon	0.32	1
P45880	Voltage-dependent anion-selective channel protein 2	0.31	1
Q9H479	Fructosamine-3-kinase	0.31	1
P45954	Shortbranched chain specific acyl-CoA dehydrogenase mitochondrial	0.31	1
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.31	1
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.30	12
Q99623	Prohibitin-2	0.30	2
P36542	ATP synthase subunit gamma mitochondrial	0.30	1
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.29	10
Q9ULV0	Unconventional myosin-Vb	0.29	15
P62826	GTP-binding nuclear protein Ran	0.28	1
P49327	Fatty acid synthase	0.28	4
Q04837	Single-stranded DNA-binding protein mitochondrial	0.28	2
P62987	Ubiquitin-60S ribosomal protein L40	0.28	1
P31483	Nucleolin TIA-1 isoform p40	0.27	1
Q149N6	Dedicator of cytokinesis 4	0.25	11
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.25	2
Q96PK6	RNA-binding protein 14	0.25	2
P08727	Keratin type I cytoskeletal 19	0.25	7
A2NUT2	Lambda-chain AA -20 to 215	0.24	4
P06576	ATP synthase subunit beta mitochondrial	0.24	3
P60842	Eukaryotic initiation factor 4A-1	0.23	2
Q5BKZ1	DBIRD complex subunit ZNF326	0.23	3
P19474	E3 ubiquitin-protein ligase TRIM21	0.22	7
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.22	4
P05141	ADP/ATP translocase 2	0.21	2
P08238	Heat shock protein HSP 90-beta	0.20	4
P05783	Keratin type I cytoskeletal 18	0.20	13
P68104	Elongation factor 1-alpha 1	0.19	4
D3DPG0	Titin isoform CRA_a	0.18	1
P60709	Actin cytoplasmic 1	0.17	8
P78527	DNA-dependent protein kinase catalytic subunit	0.17	1
P02675	Fibrinogen beta chain	0.17	2
P11142	Heat shock cognate 71 kDa protein	0.17	3
P02769	Serum albumin	0.15	44
P05787	Keratin type II cytoskeletal 8	0.15	11
P52732	Kinesin-like protein KIF11	0.15	1
Q15058	Kinesin-like protein KIF14	0.15	9
P08107	Heat shock 70 kDa protein 1A1B	0.15	2
P14625	Endoplasmic reticulum chaperone protein	0.15	1
P13645	Keratin type I cytoskeletal 10	0.14	11
P11021	78 kDa glucose-regulated protein	0.12	6
P02671	Fibrinogen alpha chain	0.12	3
P00338	L-lactate dehydrogenase A chain	0.11	2
P02768	Serum albumin	0.11	4
P02679	Fibrinogen gamma chain	0.11	2
P10809	60 kDa heat shock protein mitochondrial	0.10	1
P06748	Nucleophosmin	0.10	1
H6VRG3	Keratin 1	0.09	8
Q00610	Clathrin heavy chain 1	0.09	5
P15924	Desmoplakin	0.08	2
P07814	Bifunctional glutamateproline-tRNA ligase	0.07	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.04	2