

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q15797	Mothers against decapentaplegic homolog 1	5.94	6
Q174W1	L-xylulose reductase	1.27	4
Q171L0	Histone H2A type 3	1.17	1
Q3B7A7	Phosphoribosylglycinamide formyltransferase phosphoribosylglycinamide synthetase	1.11	2
P53396	ATP-citrate synthase	1.09	2
P11413	Glucose-6-phosphate 1-dehydrogenase	1.00	3
P83731	60S ribosomal protein L24	0.96	2
Q9H9B4	Sideroflexin 1	0.95	3
Q00571	ATP-dependent RNA helicase DDX3X	0.84	27
Q06520	Bile salt sulfotransferase	0.84	4
Q92945	Far upstream element-binding protein 2	0.81	3
O00425	Insulin-like growth factor 2 mRNA-binding protein 3	0.79	3
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.79	4
P31151	Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B alpha isoform	0.76	1
Q688L7	Olfactomedin-like protein 2A	0.74	1
Q8NYU2	UDP-glucosylcoprotein glucosyltransferase 1	0.74	1
Q9H479	Fructosamine-3-kinase	0.73	4
Q8WU79	Stromal membrane-associated protein 2	0.72	1
P82837	Pepdyl-prolyl cis-trans isomerase A	0.71	2
Q43813	Lamin C-like protein 1	0.70	2
Q12846	Syntaxin-4	0.68	1
Q15436	Protein transport protein Sec23A	0.68	4
P30041	Peroxisomal protein 6	0.63	2
Q6PJU7	Epididymis luminal protein 215	0.62	2
P06493	Cyclin-dependent kinase 1	0.61	1
Q9H6R7	WD repeat and coiled-coil-containing protein C2orf44	0.60	2
Q9H3K6	BoA-like protein 2	0.60	1
Q14204	Cytoplasmic dynein 1 heavy chain 1	0.59	1
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.58	3
Q2T818	Protein asteroid homolog 1	0.58	1
Q98466	Neurogenic locus notch homolog protein 4	0.57	1
P63261	Actin cytoplasmic 2	0.56	23
P16403	Histone H12	0.55	1
Q8IWP6	Class IVb beta tubulin	0.54	17
Q14157	Ubiquitin-associated protein 2-like	0.54	2
P03367	Glutamate dehydrogenase 1 mitochondrial	0.52	1
Q5H9R7	Serine/threonine-protein phosphatase 6 regulatory subunit 3	0.52	7
P10809	60 kDa heat shock protein mitochondrial	0.51	10
P50454	Serpin H1	0.51	2
Q9H8H3	Methyltransferase-like protein 7A	0.51	1
P49327	Fatty acid synthase	0.50	8
P23228	Coflin 1	0.50	1
P17844	Probable ATP-dependent RNA helicase DDX5	0.49	5
O15094	Serine/threonine-protein phosphatase 6 regulatory ankyrin repeat subunit A	0.48	1
Q96PK6	RNA-binding protein 14	0.48	5
P30101	Protein disulfide-isomerase A3	0.48	1
Q00341	Villin	0.47	1
P30837	Aldehyde dehydrogenase X mitochondrial	0.47	1
P11586	C-1-tetrahydrofolate synthase cytoplasmic	0.46	1
P21980	Protein-glutamine gamma-glutamyltransferase 2	0.46	1
Q01085	Nucleolin TIAR	0.45	2
P23284	Pepdyl-prolyl cis-trans isomerase B	0.45	1
P45854	Shunt-anchored chain specific acyl-CoA dehydrogenase mitochondrial	0.45	2
P81978	Heterogeneous nuclear ribonucleoprotein K	0.44	4
Q61B91	PKC2 protein	0.44	3
Q13838	Spliceosome RNA helicase DDX39B	0.44	1
P35637	RNA-binding protein FUS	0.44	1
P26641	Elongation factor 1-gamma	0.44	1
P07437	Tubulin beta chain	0.44	21
P36542	ATP synthase subunit gamma mitochondrial	0.43	2
Q13268	Dehydrogenase reductase SDR family member 2 mitochondrial	0.43	9
P00338	L-lactate dehydrogenase A chain	0.43	11
Q04843	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1	0.42	1
Q27251	Fibronectin	0.42	14
Q8BYG8	Kinesin-like protein KIFC3	0.40	8
P52597	Heterogeneous nuclear ribonucleoprotein F	0.40	3
P55072	Transitional endoplasmic reticulum ATPase	0.40	1
Q86TG7	Retrotransposon-derived protein PEG10	0.40	1
P08727	Keratin type I cytoskeletal 19	0.40	12
Q99623	Prohibitin-2	0.39	3
P14518	Pyruvate kinase PKM	0.39	5
Q15365	PolyrC-binding protein 1	0.39	4
P46776	60S ribosomal protein L27a	0.38	2
P60842	Eukaryotic initiation factor 4A-1	0.38	4
P26279	Fibronogen gamma chain	0.38	10
Q13359	Tubulin beta 3 chain	0.37	7
P02675	Fibrinogen beta chain	0.37	6
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.37	5
Q04837	Single-stranded DNA-binding protein mitochondrial	0.37	3
P27708	CAD protein	0.37	1
Q9HCC0	Methylcrotonoyl-CoA carboxylase beta chain mitochondrial	0.37	1
P02671	Ribonin alpha chain	0.36	12
Q06830	Peroxisomal protein 1	0.36	3
P05783	Keratin type I cytoskeletal 18	0.36	24
P18085	ADP-ribosylation factor 4	0.35	1
P11021	78 kDa glucose-regulated protein	0.35	20
Q32841	Probable ATP-dependent RNA helicase DDX17	0.32	2
P88363	Tubulin alpha-1B chain	0.32	11
P62829	60S ribosomal protein L23	0.32	1
Q9N2I8	Insulin-like growth factor 2 mRNA-binding protein 1	0.32	2
H7B758	Protein-L-isopartate O-methyltransferase	0.32	11
P45890	Voltage-dependent anion-selective channel protein 2	0.31	1
Q98Y11	Dnal homolog subfamily A member 3 mitochondrial	0.31	2
Q43852	Calumenin	0.31	1
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.31	3
Q08211	ATP-dependent RNA helicase A	0.31	1
P38646	Stress-70 protein mitochondrial	0.31	2
P06576	ATP synthase subunit beta mitochondrial	0.30	4
Q93382	Galectin-1	0.28	2
P05787	Keratin type II cytoskeletal 8	0.28	22
P62987	Ubiquitin-60S ribosomal protein L40	0.28	1
P02768	Serum albumin	0.28	12
P01024	Complement C3	0.27	1
P13639	Elongation factor 2	0.25	1
P49411	Elongation factor Tu mitochondrial	0.25	3
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.23	9
P01617	Ig kappa chain V-II region TEV	0.21	2
P08238	Heat shock protein HSP 90-beta	0.20	4
P11142	Heat shock cognate 71 kDa protein	0.20	4
Q15084	Protein disulfide-isomerase A6	0.19	1
Q4W4Y1	Dopamine receptor interacting protein 4	0.18	1
P78527	DNA-dependent protein kinase catalytic subunit	0.17	1
Q60701	UDP-glucose 6-dehydrogenase	0.16	2
P68104	Elongation factor 1-alpha 1	0.15	3
P05141	ADP-ATP translocase 2	0.15	1
P52732	Kinesin-like protein KIF11	0.15	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.14	4
P04264	Keratin type II cytoskeletal 1	0.14	13
P08107	Heat shock 70 kDa protein 1A1B	0.14	2
P02533	Keratin type I cytoskeletal 14	0.13	3
P25705	ATP synthase subunit alpha mitochondrial	0.13	1
Q5BKZ1	DBIRD complex subunit ZNF326	0.12	1
P13845	Keratin type I cytoskeletal 10	0.12	9
P07900	Heat shock protein HSP 90-alpha	0.11	1
P31943	Heterogeneous nuclear ribonucleoprotein H	0.11	1
P09327	Villin-1	0.11	1
P35579	Myosin-9	0.11	5
P51659	Peroxisomal multifunctional enzyme type 2	0.08	5
P02769	Serum albumin	0.08	24
P15924	Desmoplakin	0.08	2
P52272	Heterogeneous nuclear ribonucleoprotein M	0.08	1
P35580	Myosin-10	0.08	5
P35908	Keratin type II cytoskeletal 2 epidermal	0.08	4
Q15149	Plectin	0.05	1
Q149N5	DOCK4 protein	0.05	1
Q00610	Clathrin heavy chain 1	0.05	2
P35527	Keratin type I cytoskeletal 9	0.04	2