

Accession Number	Description	Fold Enrichment	Unique Peptide Count
Q09472	Histone acetyltransferase p300	10.5	6
A4QPB0	IQ motif containing GTPase activating protein 1	7.3	4
Q9NYL9	Tropomodulin-3	5.8	3
P12814	Alpha-actinin-1	4.5	7
O60271	C-Jun-amino-terminal kinase-interacting protein 4	4.2	2
P06748	Nucleophosmin	4.2	2
P14649	Myosin light chain 6B	4.2	2
P21333	Filamin-A	4.2	2
P23528	Cofilin-1	4.2	2
P62987	Ubiquitin-60S ribosomal protein L40	4.2	2
Q96ST8	Centrosomal protein of 89 kDa	4.2	2
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like pro	4.2	2
Q9UM54	Unconventional myosin-VI	4.2	2
P35580	Myosin-10	3.5	9
P06396	Gelsolin	3.0	5
Q9Y4I1	Unconventional myosin-Va	2.7	2
O00571	ATP-dependent RNA helicase DDX3X	2.6	1
P62736	Actin aortic smooth muscle	2.6	1
Q05639	Elongation factor 1-alpha 2	2.6	1
Q14240	Eukaryotic initiation factor 4A-II	2.6	1
Q15366	Poly(rC)-binding protein 2	2.6	1
Q8NDV7	Trinucleotide repeat-containing gene 6A protein	2.6	1
Q92804	TATA-binding protein-associated factor 2N	2.6	1
S6C4S2	IgG L chain	2.6	1
Q7Z406	Myosin-14	2.3	21
O43707	Alpha-actinin-4	2.2	12
P05787	Keratin type II cytoskeletal 8	2.1	30
P60709	Actin cytoplasmic 1	1.9	29
P68371	Tubulin beta-4B chain	1.9	3
P05783	Keratin type I cytoskeletal 18	1.5	16
P04792	Heat shock protein beta-1	1.5	3
P08727	Keratin type I cytoskeletal 19	1.4	11
Q15365	Poly(rC)-binding protein 1	1.4	2
P02769	Serum albumin	1.4	22
O60701	UDP-glucose 6-dehydrogenase	1.4	2
P07437	Tubulin beta chain	1.4	16
P68104	Elongation factor 1-alpha 1	1.3	3
P68363	Tubulin alpha-1B chain	1.3	7
P11142	Heat shock cognate 71 kDa protein	1.2	4
P04075	Fructose-bisphosphate aldolase A	1.2	1
Q9BQS8	FYVE and coiled-coil domain-containing protein 1	1.1	4

P00338	L-lactate dehydrogenase A chain	1.1	3
P08238	Heat shock protein HSP 90-beta	1.1	5
P78527	DNA-dependent protein kinase catalytic subunit	0.9	3
P14618	Pyruvate kinase PKM	0.8	7
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.8	2
Q562R1	Beta-actin-like protein 2	0.7	1
P27708	CAD protein	0.7	1
P07900	Heat shock protein HSP 90-alpha	0.7	1
P08107	Heat shock 70 kDa protein 1A/1B	0.7	2
H6VRG3	Keratin 1	0.7	4
P35579	Myosin-9	0.7	11
P38646	Stress-70 protein mitochondrial	0.6	1
P11021	78 kDa glucose-regulated protein	0.6	1
Q9H9B4	Sideroflexin-1	0.6	1
Q00610	Clathrin heavy chain 1	0.6	1
P49327	Fatty acid synthase	0.5	3
P25705	ATP synthase subunit alpha mitochondrial	0.5	4
P49411	Elongation factor Tu mitochondrial	0.5	2
P15924	Desmoplakin	0.4	1
P13645	Keratin type I cytoskeletal 10	0.4	3
Q99623	Prohibitin-2	0.4	1
P31943	Heterogeneous nuclear ribonucleoprotein H	0.4	1
P23396	40S ribosomal protein S3	0.3	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.3	1
P35527	Keratin type I cytoskeletal 9	0.3	1
P06733	Alpha-enolase	0.3	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.2	2