

Accession Number	Description	Fold Enrichment	UniquePeptideCount
<b>Q15853</b>	<b>Upstream stimulatory factor 2</b>	<b>7.07</b>	<b>6</b>
P68371	Tubulin beta-4B chain	4.03	12
P21333	Filamin-A	2.01	1
H7BY58	Protein-L-isoaspartate O-methyltransferase	1.52	4
P62753	40S ribosomal protein S6	1.47	2
O15523	ATP-dependent RNA helicase DDX3Y	1.29	5
O00299	Chloride intracellular channel protein 1	1.25	2
Q15058	Kinesin-like protein KIF14	1.09	31
P62701	40S ribosomal protein S4 X isoform	1.02	2
P68104	Elongation factor 1-alpha 1	1.02	3
P60709	Actin cytoplasmic 1	1.00	6
P19474	E3 ubiquitin-protein ligase TRIM21	0.89	14
P09382	Galectin-1	0.82	2
Q99623	Prohibitin-2	0.80	2
A2NUT2	Lambda-chain (AA -20 to 215)	0.77	6
D3DPG0	Titin isoform CRA_a	0.71	3
P17987	T-complex protein 1 subunit alpha	0.69	1
P49368	T-complex protein 1 subunit gamma	0.69	1
P07437	Tubulin beta chain	0.68	13
Q9H222	ATP-binding cassette sub-family G member 5	0.66	2
Q07954	Prolow-density lipoprotein receptor-related protein 1	0.62	1
P07998	Ribonuclease pancreatic	0.62	1
H6VRG3	Keratin 1	0.56	6
P08238	Heat shock protein HSP 90-beta	0.50	5
P07900	Heat shock protein HSP 90-alpha	0.46	3
P78527	DNA-dependent protein kinase catalytic subunit	0.46	1
P02769	Serum albumin	0.42	57
P13645	Keratin type I cytoskeletal 10	0.42	16
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.40	7
P08107	Heat shock 70 kDa protein 1A/1B	0.39	3
P49411	Elongation factor Tu mitochondrial	0.37	2
P06733	Alpha-enolase	0.36	3
P02533	Keratin type I cytoskeletal 14	0.34	4
P11142	Heat shock cognate 71 kDa protein	0.34	3
P68363	Tubulin alpha-1B chain	0.34	5
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.33	1
P51659	Peroxisomal multifunctional enzyme type 2	0.32	9
P06748	Nucleophosmin	0.28	2
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.27	1
P14618	Pyruvate kinase PKM	0.26	1
P07814	Bifunctional glutamate/proline--tRNA ligase	0.26	3
P25705	ATP synthase subunit alpha mitochondrial	0.24	1
P00338	L-lactate dehydrogenase A chain	0.22	2
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.20	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.15	5
Q00610	Clathrin heavy chain 1	0.13	3
Q9NVI7	ATPase family AAA domain-containing protein 3A	0.10	1
P35527	Keratin type I cytoskeletal 9	0.08	2