

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q9H5V7	Zinc finger protein Pegasus	2.89	2
P01709	Ig lambda chain V-II region MGC	0.99	2
Q16647	Prostacyclin synthase	0.93	1
P05783	Keratin type I cytoskeletal 18	0.84	10
Q8NBT3	Transmembrane protein 145	0.83	1
P04264	Keratin type II cytoskeletal 1	0.81	19
Q6NUR1	Non-SMC condensin I complex subunit G	0.71	1
P60709	Actin cytoplasmic 1	0.66	5
P05787	Keratin type II cytoskeletal 8	0.65	9
Q86YZ3	Hornerin	0.63	5
Q149N6	Dedicator of cytokinesis 4	0.60	7
P01617	Ig kappa chain V-II region TEW	0.59	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.53	10
P07437	Tubulin beta chain	0.51	6
P49327	Fatty acid synthase	0.51	1
P13645	Keratin type I cytoskeletal 10	0.50	9
Q02878	60S ribosomal protein L6	0.50	1
P68104	Elongation factor 1-alpha 1	0.47	2
A2NUT2	Lambda-chain (AA -20 to 215)	0.44	1
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.44	2
P49411	Elongation factor Tu mitochondrial	0.43	1
P13647	Keratin type II cytoskeletal 5	0.43	2
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.42	3
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.39	1
P08238	Heat shock protein HSP 90-beta	0.38	2
P02769	Serum albumin	0.36	22
P25705	ATP synthase subunit alpha mitochondrial	0.36	1
P02533	Keratin type I cytoskeletal 14	0.36	2
P14618	Pyruvate kinase PKM	0.35	1
P35527	Keratin type I cytoskeletal 9	0.33	5
Q60701	UDP-glucose 6-dehydrogenase	0.32	1
P68363	Tubulin alpha-1B chain	0.32	2
P08727	Keratin type I cytoskeletal 19	0.29	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.29	2
P11021	78 kDa glucose-regulated protein	0.26	3
P00338	L-lactate dehydrogenase A chain	0.25	1
P51659	Peroxisomal multifunctional enzyme type 2	0.22	3
P68371	Tubulin beta-4B chain	0.18	1
Q9ULV0	Unconventional myosin-Vb	0.17	1
Q00610	Clathrin heavy chain 1	0.17	2