

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q6ZMG9	Ceramide synthase 6	4.00	5
P13646	Keratin type I cytoskeletal 13	1.69	3
P68371	Tubulin beta-4B chain	0.99	14
P07437	Tubulin beta chain	0.94	14
P68104	Elongation factor 1-alpha 1	0.90	8
P78527	DNA-dependent protein kinase catalytic subunit	0.80	2
Q149N5	DOCK4 protein	0.71	10
P68363	Tubulin alpha-1B chain	0.71	9
P05787	Keratin type II cytoskeletal 8	0.67	14
P14618	Pyruvate kinase PKM	0.64	3
Q9UKP6	Urotensin-2 receptor	0.61	1
P05783	Keratin type I cytoskeletal 18	0.60	8
P04264	Keratin type II cytoskeletal 1	0.51	17
P06576	ATP synthase subunit beta mitochondrial	0.48	2
P35527	Keratin type I cytoskeletal 9	0.46	12
P04259	Keratin type II cytoskeletal 6B	0.46	3
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.42	5
P08727	Keratin type I cytoskeletal 19	0.41	3
P08238	Heat shock protein HSP 90-beta	0.40	3
O60701	UDP-glucose 6-dehydrogenase	0.36	2
P02769	Serum albumin	0.35	36
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.32	2
A1A508	PRSS3 protein	0.32	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.29	8
P19474	E3 ubiquitin-protein ligase TRIM21	0.28	3
P13645	Keratin type I cytoskeletal 10	0.25	7
P02533	Keratin type I cytoskeletal 14	0.24	2
P07814	Bifunctional glutamateproline--tRNA ligase	0.22	2
Q9ULV0	Unconventional myosin-Vb	0.18	2
P60709	Actin cytoplasmic 1	0.14	2
Q15058	Kinesin-like protein KIF14	0.07	1