

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q6ZW49	PAX-interacting protein 1	11.78	22
P68104	Elongation factor 1-alpha 1	1.71	5
Q8TDN2	Potassium voltage-gated channel subfamily V member 2	1.19	1
Q9ULQ0	Striatin-interacting protein 2	0.84	1
P02538	Keratin type II cytoskeletal 6A	0.75	8
Q8N110	Dedicator of cytokinesis protein 4	0.68	14
Q86YZ3	Hornerin	0.64	9
P68371	Tubulin beta-4B chain	0.60	12
Q9Y3T6	R3H and coiled-coil domain-containing protein 1	0.57	1
P13647	Keratin type II cytoskeletal 5	0.57	8
P02533	Keratin type I cytoskeletal 14	0.55	8
P62081	40S ribosomal protein S7	0.54	1
P38646	Stress-70 protein mitochondrial	0.50	2
Q14847	LIM and SH3 domain protein 1	0.49	2
P08779	Keratin type I cytoskeletal 16	0.47	5
P04264	Keratin type II cytoskeletal 1	0.46	22
P22061	Protein-L-isoaspartateD-aspartate O-methyltransferase	0.46	8
Q7Z5J8	Ankyrin and armadillo repeat-containing protein	0.45	1
Q99808	Equilibrative nucleoside transporter 1	0.44	1
Q8IZT6	Abnormal spindle-like microcephaly-associated protein	0.42	1
P68363	Tubulin alpha-1B chain	0.40	7
P00338	L-lactate dehydrogenase A chain	0.39	5
P07437	Tubulin beta chain	0.39	8
Q99623	Prohibitin-2	0.37	1
P49327	Fatty acid synthase	0.36	2
P18124	60S ribosomal protein L7	0.35	1
Q6NUR1	Non-SMC condensin I complex subunit G	0.33	1
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.33	1
P14618	Pyruvate kinase PKM	0.33	2
P63261	Actin cytoplasmic 2	0.32	6
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.31	3
P49411	Elongation factor Tu mitochondrial	0.31	2
P19474	E3 ubiquitin-protein ligase TRIM21	0.31	5
Q15366	PolrC-binding protein 2	0.31	2
P25705	ATP synthase subunit alpha mitochondrial	0.30	2
D3DPG0	Titin isoform CRA_a	0.28	1
Q16822	Phosphoenolpyruvate carboxykinase [GTP] mitochondrial	0.28	1
Q15084	Protein disulfide-isomerase A6	0.27	1
P61978	Heterogeneous nuclear ribonucleoprotein K	0.27	1
P13645	Keratin type I cytoskeletal 10	0.26	11
P05787	Keratin type II cytoskeletal 8	0.26	10
O60701	UDP-glucose 6-dehydrogenase	0.26	2
P11021	78 kDa glucose-regulated protein	0.25	7
P02679	Fibrinogen gamma chain	0.25	2
Q15365	PolrC-binding protein 1	0.24	1
P08107	Heat shock 70 kDa protein 1A1B	0.24	2
P02671	Fibrinogen alpha chain	0.23	3
P02769	Serum albumin	0.23	34
P35527	Keratin type I cytoskeletal 9	0.22	8
P05783	Keratin type I cytoskeletal 18	0.22	7
P14625	Endoplasmin	0.21	1
P11142	Heat shock cognate 71 kDa protein	0.21	2
P08238	Heat shock protein HSP 90-beta	0.20	2
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.17	3
P31943	Heterogeneous nuclear ribonucleoprotein H	0.16	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.15	6
P02768	Serum albumin	0.15	3
P51659	Peroxisomal multifunctional enzyme type 2	0.13	4
P08727	Keratin type I cytoskeletal 19	0.10	1