

Accession Number	Description	Fold Enrichment	UniquePeptideCount
<b>Q9UNY5</b>	<b>Zinc finger protein 232</b>	<b>8.47</b>	<b>5</b>
Q6ZQQ6	WD repeat-containing protein 87	1.43	2
P29322	Ephrin type-A receptor 8	1.17	1
P31483	Nucleolysin TIA-1 isoform p40	1.16	4
Q14847	LIM and SH3 domain protein 1	1.13	6
P07306	Asialoglycoprotein receptor 1	0.97	2
Q9H0A3	Transmembrane protein 191A	0.96	1
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.88	4
Q06830	Peroxioredoxin-1	0.87	5
Q9BVG8	Kinesin-like protein KIFC3	0.85	10
P52597	Heterogeneous nuclear ribonucleoprotein F	0.81	4
P19474	E3 ubiquitin-protein ligase TRIM21	0.73	14
P02675	Fibrinogen beta chain	0.71	7
Q9UBS4	DnaJ homolog subfamily B member 11	0.70	1
Q6P996	Pyridoxal-dependent decarboxylase domain-containing protein 1	0.68	3
Q9C0G6	Dynein heavy chain 6 axonemal	0.68	1
P68371	Tubulin beta-4B chain	0.66	16
Q15436	Protein transport protein Sec23A	0.63	2
P35637	RNA-binding protein FUS	0.60	1
P07437	Tubulin beta chain	0.59	16
Q8WWZ4	ATP-binding cassette sub-family A member 10	0.57	1
P05141	ADP/ATP translocase 2	0.56	4
P13639	Elongation factor 2	0.55	2
P02671	Fibrinogen alpha chain	0.53	10
P35030	Trypsin-3	0.53	1
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.51	6
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.50	10
Q13201	Multimerin-1	0.50	1
Q12931	Heat shock protein 75 kDa mitochondrial	0.50	2
P06576	ATP synthase subunit beta mitochondrial	0.49	4
Q07954	Prolow-density lipoprotein receptor-related protein 1	0.47	1
P38646	Stress-70 protein mitochondrial	0.46	2
P48643	T-complex protein 1 subunit epsilon	0.44	1
P09382	Galectin-1	0.43	2
Q9H479	Fructosamine-3-kinase	0.42	1
Q16822	Phosphoenolpyruvate carboxykinase [GTP] mitochondrial	0.42	2
P14618	Pyruvate kinase PKM	0.42	3
P63261	Actin cytoplasmic 2	0.40	9
P08238	Heat shock protein HSP 90-beta	0.40	5
Q8IZT6	Abnormal spindle-like microcephaly-associated protein	0.40	1
P49411	Elongation factor Tu mitochondrial	0.40	3
Q15366	PolyrC-binding protein 2	0.39	3
P10809	60 kDa heat shock protein mitochondrial	0.39	4
P49327	Fatty acid synthase	0.36	3
P02679	Fibrinogen gamma chain	0.34	5
P11142	Heat shock cognate 71 kDa protein	0.33	4
Q149N6	Dedicator of cytokinesis 4	0.33	8
A2NUT2	Lambda-chain AA -20 to 215	0.31	3
O00571	ATP-dependent RNA helicase DDX3X	0.31	5
Q9NZI8	Insulin-like growth factor 2 mRNA-binding protein 1	0.30	1
P02768	Serum albumin	0.29	7
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.29	2
P62249	40S ribosomal protein S16	0.29	1
P06733	Alpha-enolase	0.28	3
Q99623	Prohibitin-2	0.28	1
P54136	Arginine--tRNA ligase cytoplasmic	0.27	1
Q04837	Single-stranded DNA-binding protein mitochondrial	0.27	1
P61978	Heterogeneous nuclear ribonucleoprotein K	0.26	1
P08727	Keratin type I cytoskeletal 19	0.25	4
D3DPG0	Titin isoform CRA a	0.25	1
P17844	Probable ATP-dependent RNA helicase DDX5	0.24	1
O60701	UDP-glucose 6-dehydrogenase	0.24	2
P31943	Heterogeneous nuclear ribonucleoprotein H	0.24	2
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.22	1
Q15365	PolyrC-binding protein 1	0.22	1
P60842	Eukaryotic initiation factor 4A-I	0.22	1
P02769	Serum albumin	0.21	35
P01700	Ig lambda chain V-I region HA	0.20	1
Q5H9R7	Serine/threonine-protein phosphatase 6 regulatory subunit 3	0.19	1
P25705	ATP synthase subunit alpha mitochondrial	0.18	1
P68104	Elongation factor 1-alpha 1	0.18	2
P05783	Keratin type I cytoskeletal 18	0.17	6
P11021	78 kDa glucose-regulated protein	0.17	5
P05787	Keratin type II cytoskeletal 8	0.17	7
P51659	Peroxisomal multifunctional enzyme type 2	0.14	5
P08107	Heat shock 70 kDa protein 1A1B	0.13	1
P68363	Tubulin alpha-1B chain	0.13	2
Q00610	Clathrin heavy chain 1	0.12	4
Q86YZ3	Hornerin	0.10	1
P13645	Keratin type I cytoskeletal 10	0.10	4
P35527	Keratin type I cytoskeletal 9	0.08	3
H6VRG3	Keratin 1	0.08	4
P35908	Keratin type II cytoskeletal 2 epidermal	0.08	3