

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q9UBY0	Sodium/hydrogen exchanger 2	3.33	2
Q9Y2P7	Zinc finger protein 256	3.33	2
P01615	Ig kappa chain V-II region FR	3.05	2
Q3T8J9	GON-4-like protein	2.17	1
Q9NQZ2	Something about silencing protein 10	1.86	2
Q6P996	Pyridoxal-dependent decarboxylase domain-containing protein 1	1.28	7
P35368	Alpha-1B adrenergic receptor	1.02	1
Q96TA2	ATP-dependent zinc metalloprotease YME1L1	0.98	1
Q9UBS4	DnaJ homolog subfamily B member 11	0.94	2
P05023	Sodium/potassium-transporting ATPase subunit alpha-1	0.92	1
Q9LULV0	Unconventional myosin-Vb	0.90	35
P12004	Proliferating cell nuclear antigen	0.86	4
P07306	Asialoglycoprotein receptor 1	0.78	2
P43378	Tyrosine-protein phosphatase non-receptor type 9	0.77	1
P27105	Erythrocyte band 7 integral membrane protein	0.72	1
Q969W1	Probable palmitoyltransferase ZDHHC16	0.72	1
Q5RKT7	Ribosomal protein S27a	0.72	2
P35030	Trypsin-3	0.71	2
Q685J3	Mucin-17	0.70	1
P09758	Tumor-associated calcium signal transducer 2	0.68	1
P68363	Tubulin alpha-1B chain	0.62	16
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.61	2
P68371	Tubulin beta-4B chain	0.61	19
P07437	Tubulin beta chain	0.58	20
P62829	60S ribosomal protein L23	0.58	2
O15440	Multidrug resistance-associated protein 5	0.58	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.57	14
Q08211	ATP-dependent RNA helicase A	0.57	2
P14868	Aspartate--tRNA ligase cytoplasmic	0.56	2
P06576	ATP synthase subunit beta mitochondrial	0.56	6
P36542	ATP synthase subunit gamma mitochondrial	0.54	2
Q00325	Phosphate carrier protein mitochondrial	0.54	1
P14618	Pyruvate kinase PKM	0.52	5
P31483	Nucleolin T1A-1 isoform p40	0.50	2
P60842	Eukaryotic initiation factor 4A-1	0.50	4
O43175	D-3-phosphoglycerate dehydrogenase	0.49	1
P25705	ATP synthase subunit alpha mitochondrial	0.48	5
Q16822	Phosphoenolpyruvate carboxykinase [GTP] mitochondrial	0.48	3
Q86TG7	Retrotransposon-derived protein PEG10	0.47	1
P02679	Fibrinogen gamma chain	0.47	9
Q4W4Y1	Dopamine receptor interacting protein 4	0.45	3
Q14157	Ubiquitin-associated protein 2-like	0.44	1
P02671	Fibrinogen alpha chain	0.42	10
Q12931	Heat shock protein 75 kDa mitochondrial	0.42	2
Q07954	Prolow-density lipoprotein receptor-related protein 1	0.41	1
Q99623	Prohibitin-2	0.38	2
Q3LXA3	Bifunctional ATP-dependent dihydroxyacetone kinase/FAD-AMP lyase cyclizing	0.37	1
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.36	1
P09382	Galectin-1	0.36	2
P01700	Ig lambda chain V-I region HA	0.35	3
Q04937	Single-stranded DNA-binding protein mitochondrial	0.35	2
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.35	5
P02768	Serum albumin	0.35	11
Q15084	Protein disulfide-isomerase A6	0.34	2
Q14847	LIM and SH3 domain protein 1	0.34	2
Q15436	Protein transport protein Sec23A	0.34	1
P05783	Keratin type I cytoskeletal 18	0.34	16
O00571	ATP-dependent RNA helicase DDX3X	0.33	7
Q8TB22	Spermatogenesis-associated protein 20	0.33	1
P08238	Heat shock protein HSP 90-beta	0.32	5
P17844	Probable ATP-dependent RNA helicase DDX5	0.32	2
Q149N6	Dedicator of cytokinesis 4	0.32	10
P27635	60S ribosomal protein L10	0.31	1
P68104	Elongation factor 1-alpha 1	0.30	5
Q86Y82	Syntaxin-12	0.30	1
Q15365	PolyrC-binding protein 1	0.30	2
P08727	Keratin type I cytoskeletal 19	0.29	6
P05141	ADP/ATP translocase 2	0.27	2
P31943	Heterogeneous nuclear ribonucleoprotein H	0.27	3
Q9NZ18	Insulin-like growth factor 2 mRNA-binding protein 1	0.26	1
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.26	1
P63261	Actin cytoplasmic 2	0.26	7
P38646	Stress-70 protein mitochondrial	0.25	1
P49411	Elongation factor Tu mitochondrial	0.24	2
Q9BVG8	Kinesin-like protein KIFC3	0.24	3
P08107	Heat shock 70 kDa protein 1A1B	0.24	3
P54136	Arginine--tRNA ligase cytoplasmic	0.23	1
P05787	Keratin type II cytoskeletal 8	0.22	12
P78527	DNA-dependent protein kinase catalytic subunit	0.21	1
A2NUT2	Lambda-chain AA -20 to 215	0.19	2
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.19	2
O15090	Zinc finger protein 536	0.19	1
P02769	Serum albumin	0.19	39
P10809	60 kDa heat shock protein mitochondrial	0.19	2
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.18	4
P51659	Peroxisomal multifunctional enzyme type 2	0.17	8
Q5H9R7	Serinethreonine-protein phosphatase 6 regulatory subunit 3	0.17	1
P11142	Heat shock cognate 71 kDa protein	0.16	2
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.16	1
Q15366	PolyrC-binding protein 2	0.15	1
Q9BUZ4	TNF receptor-associated factor 4	0.15	1
P49327	Fatty acid synthase	0.14	1
P11021	78 kDa glucose-regulated protein	0.14	5
P58107	Epiplakin	0.14	1
P02675	Fibrinogen beta chain	0.13	1
P02533	Keratin type I cytoskeletal 14	0.13	2
O60701	UDP-glucose 6-dehydrogenase	0.13	1
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.12	3
Q15058	Kinesin-like protein KIF14	0.12	5
P35527	Keratin type I cytoskeletal 9	0.11	5
P15924	Desmoplakin	0.06	1