

Accession Number	Description	Fold Enrichment	UniquePeptideCount
O75955	Flotillin-1	6.70	5
Q01664	Transcription factor AP-4	5.25	9
P55786	Puromycin-sensitive aminopeptidase	3.64	3
B4DJV2	Citrate synthase	3.28	2
O94874	E3 UFM1-protein ligase 1	3.28	2
P11717	Cation-independent mannose-6-phosphate receptor	3.28	2
P78406	mRNA export factor	3.28	2
Q03426	Mevalonate kinase	3.28	2
Q562R1	Beta-actin-like protein 2	3.28	2
Q9UBM7	7-dehydrocholesterol reductase	3.28	2
Q9NRH3	Tubulin gamma-2 chain	2.88	3
O00303	Eukaryotic translation initiation factor 3 subunit F	2.10	6
Q9P2R7	Succinyl-CoA ligase [ADP-forming] subunit beta mitochondrial	2.09	6
G5E9V5	28S ribosomal protein S22 mitochondrial	1.99	2
P00505	Aspartate aminotransferase mitochondrial	1.99	2
O60547	GDP-mannose 46 dehydratase	1.96	4
B5BU52	C-src tyrosine kinase	1.92	2
Q5JWF2	Guanine nucleotide-binding protein G(s) subunit alpha isoforms XLas	1.77	2
P08865	40S ribosomal protein SA	1.75	3
P00558	Phosphoglycerate kinase 1	1.74	11
Q9Y4P3	Transducin beta-like protein 2	1.73	5
Q99447	Ethanolamine-phosphate cytidyltransferase	1.71	2
P07954	Fumarate hydratase mitochondrial	1.70	5
P16219	Short-chain specific acyl-CoA dehydrogenase mitochondrial	1.68	3
P08319	Alcohol dehydrogenase 4	1.66	2
B4E2W0	3-ketoacyl-CoA thiolase	1.64	7
P51570	Galactokinase	1.62	3
Q14103	Heterogeneous nuclear ribonucleoprotein D0	1.59	2
P68104	Elongation factor 1-alpha 1	1.58	10
P08107	Heat shock 70 kDa protein 1A/1B	1.56	2
O95433	Activator of 90 kDa heat shock protein ATPase homolog 1	1.54	4
Q9NUQ8	ATP-binding cassette sub-family F member 3	1.53	2
P35998	26S protease regulatory subunit 7	1.51	4
P30837	Aldehyde dehydrogenase X mitochondrial	1.50	2
P09543	2'3'-cyclic-nucleotide 3'-phosphodiesterase	1.49	2
O00231	26S proteasome non-ATPase regulatory subunit 11	1.46	2
P45954	Short/branched chain specific acyl-CoA dehydrogenase mitochondrial	1.38	4
Q13363	C-terminal-binding protein 1	1.38	2
Q9Y570	Protein phosphatase methylesterase 1	1.38	2
O75874	Isocitrate dehydrogenase [NADP] cytoplasmic	1.36	5
Q9Y305	Acyl-coenzyme A thioesterase 9 mitochondrial	1.31	2
A8K3Z3	26S protease regulatory subunit 8	1.30	3
Q9UJZ1	Stomatin-like protein 2	1.29	2
P27708	CAD protein	1.29	2
P24752	Acetyl-CoA acetyltransferase mitochondrial	1.28	2
O94905	Erlin-2	1.27	4
Q96199	Succinyl-CoA ligase [GDP-forming] subunit beta mitochondrial	1.26	2
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	1.25	5
O60701	UDP-glucose 6-dehydrogenase	1.24	5
B5BTY4	ATP-dependent RNA helicase DDX3X	1.21	7
Q16851	UTP-glucose-1-phosphate uridylyltransferase	1.20	2
Q6YN16	Hydroxysteroid dehydrogenase-like protein 2	1.16	3
P50454	Serpin H1	1.10	2
P49368	T-complex protein 1 subunit gamma	1.08	4
O60884	DnaJ homolog subfamily A member 2	1.03	2
Q9BV38	WD repeat-containing protein 18	0.96	2
Q9Y617	Phosphoserine aminotransferase	0.96	2
P38919	Eukaryotic initiation factor 4A-III	0.94	3
P42704	Leucine-rich PPR motif-containing protein mitochondrial	0.87	2
P02769	Serum albumin	0.86	16
B4E335	Actin cytoplasmic 1 N-terminally processed	0.86	11
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.84	7
Q9UBB4	Ataxin-10	0.83	3
P11586	C-1-tetrahydrofolate synthase cytoplasmic	0.81	2
P48735	Isocitrate dehydrogenase [NADP] mitochondrial	0.80	2
P31689	DnaJ homolog subfamily A member 1	0.80	2
P19474	E3 ubiquitin-protein ligase TRIM21	0.80	5
P68371	Tubulin beta-4B chain	0.80	5
P48643	T-complex protein 1 subunit epsilon	0.76	2
P60842	Eukaryotic initiation factor 4A-I	0.75	4
P49257	Protein ERGIC-53	0.74	2
P29401	Transketolase	0.73	2
P17844	Probable ATP-dependent RNA helicase DDX5	0.73	2
P52272	Heterogeneous nuclear ribonucleoprotein M	0.67	2
P08238	Heat shock protein HSP 90-beta	0.64	8
P25705	ATP synthase subunit alpha mitochondrial	0.60	6
P26641	Elongation factor 1-gamma	0.58	3
P11142	Heat shock cognate 71 kDa protein	0.55	2
P49327	Fatty acid synthase	0.54	3
P49411	Elongation factor Tu mitochondrial	0.53	4
P06733	Alpha-enolase	0.51	6
P04075	Fructose-bisphosphate aldolase A	0.49	4
P14625	Endoplasmic	0.48	2
P10809	60 kDa heat shock protein mitochondrial	0.44	7
O75477	Erlin-1	0.43	2
P23526	Adenosylhomocysteinase	0.43	2
P14618	Pyruvate kinase isozymes M1/M2	0.41	4
P15924	Desmoplakin	0.40	2
P06576	ATP synthase subunit beta mitochondrial	0.38	4
P00367	Glutamate dehydrogenase 1 mitochondrial	0.30	2