

Accession Number	Description	Fold Enrichment	UniquePeptideCount
P58317	Zinc finger protein 121	6.66	6
Q6IAJ9	PIPOX protein	3.83	3
B9EG68	PEG3 protein	2.10	11
P00367	Glutamate dehydrogenase 1 mitochondrial	1.18	2
Q96PR1	Potassium voltage-gated channel subfamily C member 2	1.14	2
P11413	Glucose-6-phosphate 1-dehydrogenase	1.11	3
P16284	Platelet endothelial cell adhesion molecule	1.07	2
Q15223	Nectin-1	1.02	1
Q00325	Phosphate carrier protein mitochondrial	1.02	2
Q96PK6	RNA-binding protein 14	1.00	9
Q06520	Bile salt sulfotransferase	0.91	4
P14373	Zinc finger protein RFP	0.88	3
P10620	Microsomal glutathione S-transferase 1	0.85	1
P32119	Peroxiredoxin-2	0.84	4
P63151	Serine/threonine-protein phosphatase 2A 55 kDa regulatory subunit B alpha isoform	0.80	1
Q99623	Prohibitin-2	0.79	5
P23528	Cofilin-1	0.77	2
P30837	Aldehyde dehydrogenase X mitochondrial	0.73	2
Q6P996	Pyridoxal-dependent decarboxylase domain-containing protein 1	0.69	3
P49327	Fatty acid synthase	0.69	8
P35637	RNA-binding protein FUS	0.68	2
Q06210	Glutamine-fructose-6-phosphate aminotransferase [isomerizing] 1	0.68	3
P08727	Keratin type I cytoskeletal 19	0.67	19
P63261	Actin cytoplasmic 2	0.67	24
Q06830	Peroxiredoxin-1	0.67	6
P04643	Dolichyl-diphosphooligosaccharide-protein glycosyltransferase subunit 1	0.66	2
Q01085	Nucleolin TIAR	0.66	3
P52597	Heterogeneous nuclear ribonucleoprotein F	0.65	5
Q9H3K6	BoA-like protein 2	0.63	1
Q6IB91	PCK2 protein	0.59	4
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.58	11
Q95793	Double-stranded RNA-binding protein Staufen homolog 1	0.58	3
P02675	Fibrinogen beta chain	0.58	5
Q43242	26S proteasome non-ATPase regulatory subunit 3	0.57	1
P16403	Histone H12	0.57	1
Q81937	Probable ATP-dependent RNA helicase DHX37	0.57	1
Q92841	Probable ATP-dependent RNA helicase DDX17	0.56	4
Q00571	ATP-dependent RNA helicase DDX3X	0.54	15
Q6NVV3	Magnesium transporter NIPA3	0.52	1
P62937	Peptidyl-prolyl cis-trans isomerase A	0.51	1
Q00341	Vigilin	0.51	1
P32322	Pyroline-5-carboxylate reductase 1 mitochondrial	0.51	2
P11586	C-1-tetrahydrofolate synthase cytoplasmic	0.48	1
Q9H479	Fructosamine-3-kinase	0.48	2
Q14157	Ubiquitin-associated protein 2-like	0.47	1
Q13838	Spliceosome RNA helicase DDX39B	0.46	1
Q9BVG8	Kinesin-like protein KIFC3	0.46	8
P26641	Elongation factor 1-gamma	0.46	1
P30041	Peroxiredoxin-6	0.45	1
P27708	CAD protein	0.45	1
Q9NZI8	Insulin-like growth factor 2 mRNA-binding protein 1	0.45	3
Q9H6R7	WD repeat and coiled-coil-containing protein C2orf44	0.43	1
P02679	Fibrinogen gamma chain	0.43	7
P10809	60 kDa heat shock protein mitochondrial	0.41	7
P11021	78 kDa glucose-regulated protein	0.41	21
P07437	Tubulin beta chain	0.40	15
P49411	Elongation factor Tu mitochondrial	0.40	5
P51571	Translocon-associated protein subunit delta	0.40	1
P13639	Elongation factor 2	0.39	2
P06576	ATP synthase subunit beta mitochondrial	0.39	5
Q15084	Protein disulfide-isomerase A6	0.39	3
P31943	Heterogeneous nuclear ribonucleoprotein H	0.39	6
Q9HCC0	Methylcrotonyl-CoA carboxylase beta chain mitochondrial	0.38	1
Q8IWP6	Class IVb beta tubulin	0.37	10
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.36	1
P68363	Tubulin alpha-1B chain	0.35	11
Q8TB22	Spermatogenesis-associated protein 20	0.34	1
Q15366	PolyrC-binding protein 2	0.33	4
Q15436	Protein transport protein Sec23A	0.33	1
P62929	60S ribosomal protein L23	0.33	1
P45880	Voltage-dependent anion-selective channel protein 2	0.33	1
P60842	Eukaryotic initiation factor 4A-1	0.33	3
P05787	Keratin type II cytoskeletal 8	0.32	23
P14868	Aspartate-tRNA ligase cytoplasmic	0.32	1
P12004	Proliferating cell nuclear antigen	0.32	1
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.32	1
P00338	L-lactate dehydrogenase A chain	0.31	7
P11142	Heat shock cognate 71 kDa protein	0.31	6
P11182	Lipoamide acyltransferase component of branched-chain alpha-keto acid dehydrogenase	0.30	1
P05763	Keratin type I cytoskeletal 18	0.30	18
D3DFG0	Titin isoform CRA a	0.30	2
Q4W4Y1	Dopamine receptor interacting protein 4	0.30	2
P68104	Elongation factor 1-alpha 1	0.29	6
Q5H9R7	Serine/threonine-protein phosphatase 6 regulatory subunit 3	0.28	3
P14618	Pyruvate kinase PKM	0.28	3
P02768	Serum albumin	0.28	11
Q06701	UDP-glucose 6-dehydrogenase	0.28	4
P17844	Probable ATP-dependent RNA helicase DDX5	0.27	2
P01617	Ig kappa chain V-II region TEW	0.27	2
Q15365	PolyrC-binding protein 1	0.26	2
P02671	Fibrinogen alpha chain	0.26	6
Q12931	Heat shock protein 75 kDa mitochondrial	0.25	1
P52272	Heterogeneous nuclear ribonucleoprotein M	0.24	5
P14625	Endoplasmic	0.23	2
P08238	Heat shock protein HSP 90-beta	0.22	4
P38646	Stress-70 protein mitochondrial	0.22	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.22	6
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.20	6
P54136	Arginine-tRNA ligase cytoplasmic	0.20	1
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.20	1
P25705	ATP synthase subunit alpha mitochondrial	0.20	2
P61978	Heterogeneous nuclear ribonucleoprotein K	0.20	1
P78527	DNA-dependent protein kinase catalytic subunit	0.18	1
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.17	1
Q15149	Plectin	0.17	5
P02751	Fibronectin	0.16	3
Q149N5	DOCK4 protein	0.16	5
P02769	Serum albumin	0.15	41
P05141	ADPATP translocase 2	0.15	1
A2NUT2	Lambda-chain AA -20 to 215	0.12	1
P58107	Epipalatin	0.12	1
Q13509	Tubulin beta-3 chain	0.11	1
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.10	3
P07814	Bifunctional glutamate-proline-tRNA ligase	0.10	2
P35580	Myosin-10	0.09	7
P51654	Glypican-3	0.09	1
P15924	Desmoplakin	0.08	2
P35579	Myosin-9	0.06	2
Q5T9A4	ATPase family AAA domain-containing protein 3B	0.06	1
P04264	Keratin type II cytoskeletal 1	0.06	4
Q00610	Clathrin heavy chain 1	0.05	2
P35908	Keratin type II cytoskeletal 2 epidermal	0.03	1
P13645	Keratin type I cytoskeletal 10	0.03	1