

Accession Number	Description	Fold Enrichment	UniquePeptideCount
<b>P84022</b>	<b>Mothers against decapentaplegic homolog 3</b>	<b>17.64</b>	<b>22</b>
Q49AG4	Df53 protein	1.99	4
Q08210	Glutamine-fructose-6-phosphate aminotransferase [isomerizing] 1	1.05	6
Q96UJ3	Engulfment and cell motility protein 2	1.03	3
Q92900	Regulator of nonsense transcripts 1	0.99	1
Q95299	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 10 mitochondrial	0.87	1
P62937	Peptidyl-prolyl cis-trans isomerase A	0.86	3
P34897	Serine hydroxymethyltransferase mitochondrial	0.83	2
Q61891	PKC $\zeta$ protein	0.78	7
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.75	4
Q60488	Long-chain-fatty-acid-CoA ligase 4	0.75	1
Q43251	RNA binding protein fox-1 homolog 2	0.74	2
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.73	7
Q9UBX3	Mitochondrial dicarboxylate carrier	0.73	1
Q04828	Aldo-keto reductase family 1 member C1	0.72	2
Q06830	Peroxisome protein 1	0.71	8
Q9H8H3	Methyltransferase-like protein 7A	0.69	2
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.68	28
B0Y1W6	Archain 1 isoform CRA_a	0.67	1
D3DR22	Hydroxysteroid 17-beta dehydrogenase 12 isoform CRA_a	0.64	2
P49327	Fatty acid synthase	0.63	9
Q06520	Bile salt sulfotransferase	0.62	3
P11498	Pyruvate carboxylase mitochondrial	0.62	1
Q01955	Collagen alpha-3(V) chain	0.60	1
Q02671	Fibrinogen alpha chain	0.59	19
Q02945	F-ar upstream element-binding protein 2	0.58	2
Q9H3K6	BoA-like protein 2	0.57	1
P02679	Fibrinogen gamma chain	0.57	12
Q15523	ATP-dependent RNA helicase DDX3Y	0.56	8
Q00425	Insulin-like growth factor 2 mRNA-binding protein 3	0.56	2
Q14204	Cytoplasmic dynein 1 heavy chain 1	0.55	1
Q96623	Prohibitin-2	0.55	4
Q95433	Activator of 90 kDa heat shock protein ATPase homolog 1	0.55	1
P45880	Voltage-dependent anion-selective channel protein 2	0.55	3
P50225	Sulfotransferase 1A1	0.54	1
Q12904	Aminoacyl tRNA synthase complex-interacting multifunctional protein 1	0.52	1
P36542	ATP synthase subunit gamma mitochondrial	0.52	3
P41252	Methionine-tRNA ligase cytoplasmic	0.52	2
P51571	Translocin-associated protein subunit delta	0.52	2
P01859	Ig gamma-2 chain C region	0.51	2
P62826	GTP-binding nuclear protein Ran	0.51	3
Q43795	Unconventional myosin-1b	0.51	3
P31943	Heterogeneous nuclear ribonucleoprotein H	0.50	10
Q9HCC0	Methylcrotonyl-CoA carboxylase beta chain mitochondrial	0.49	2
P47897	Glutamine-tRNA ligase	0.49	2
Q94905	Erlin-2	0.48	1
P60842	Eukaryotic initiation factor 4A-1	0.47	6
Q13885	Tubulin beta-2A chain	0.47	11
Q724W1	L-xylulose reductase	0.46	1
P30101	Protein disulfide-isomerase A3	0.46	1
P53618	Coatomer subunit beta	0.44	1
P56192	Methionine-tRNA ligase cytoplasmic	0.43	1
Q15365	PolyrC-binding protein 1	0.43	5
Q43143	Putative pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15	0.42	1
P54136	Arginine-tRNA ligase cytoplasmic	0.42	4
P35637	RNA-binding protein FUS	0.42	1
P26841	Elongation factor 1-gamma	0.41	1
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.41	2
P30041	Peroxisome protein 6	0.41	1
P50991	T-complex protein 1 subunit delta	0.41	1
P51078	Heterogeneous nuclear ribonucleoprotein K	0.41	4
P68363	Tubulin alpha-1B chain	0.40	16
P04843	Dolichyl-diphosphooligosaccharide-protein glycosyltransferase subunit 1	0.40	1
P49411	Elongation factor Tu mitochondrial	0.39	6
P62987	Ubiquitin-60S ribosomal protein L40	0.35	1
P68371	Tubulin beta-4B chain	0.35	15
P45954	Shortbranched chain specific acyl-CoA dehydrogenase mitochondrial	0.34	1
P25705	ATP synthase subunit alpha mitochondrial	0.33	5
Q15084	Protein disulfide-isomerase A6	0.33	3
P19474	E3 ubiquitin-protein ligase TRIM21	0.33	12
Q13268	Dehydrogenase reductase SDR family member 2 mitochondrial	0.32	7
Q13509	Insulin beta-3 chain	0.32	6
P05141	ADP-ATP translocase 2	0.32	4
P32119	Peroxisome protein 2	0.31	1
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.30	3
Q15436	Protein transport protein Sec23A	0.30	1
P62829	60S ribosomal protein L23	0.30	1
P31483	Nucleolin T1A-1 isoform p40	0.30	1
Q9NZ18	Insulin-like growth factor 2 mRNA-binding protein 1	0.29	2
P14868	Aspartate-tRNA ligase cytoplasmic	0.29	1
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.29	3
P38846	Stress-70 protein mitochondrial	0.28	2
Q15386	PolyrC-binding protein 2	0.28	4
P11182	Lipoamide acyltransferase component of branched-chain alpha-keto acid dehydrogenase	0.27	1
P06576	ATP synthase subunit beta mitochondrial	0.27	4
Q149N6	Dedicator of cytokinesis 4	0.27	13
Q04837	Single-stranded DNA-binding protein mitochondrial	0.26	2
P08107	Heat shock 70 kDa protein 1A1B	0.26	5
P02768	Serum albumin	0.25	12
P07437	Tubulin beta chain	0.25	11
A2NUT2	Lambda-chain AA -20 to 215	0.25	4
P18621	60S ribosomal protein L17	0.24	1
Q5H9R7	Serine-threonine-protein phosphatase 6 regulatory subunit 3	0.24	3
P14818	Pyruvate kinase PKM	0.24	3
P13639	Elongation factor 2	0.24	1
Q60701	UDP-glucose 6-dehydrogenase	0.23	4
P78527	DNA-dependent protein kinase catalytic subunit	0.23	2
P01617	Ig kappa chain V-II region TEW	0.23	2
P60709	Actin cytoplasmic 1	0.22	12
Q00571	ATP-dependent RNA helicase DDX3X	0.22	7
P10809	60 kDa heat shock protein mitochondrial	0.22	4
Q9BVG8	Kinesin-like protein KIFC3	0.22	4
P07237	Protein disulfide-isomerase	0.22	1
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.21	9
P02769	Serum albumin	0.21	69
Q92841	Probable ATP-dependent RNA helicase DDX17	0.21	1
P88104	Elongation factor 1-alpha 1	0.21	5
P07814	Bifunctional glutamylproline-tRNA ligase	0.20	6
P01700	Ig lambda chain V-I region HA	0.19	2
P11142	Heat shock cognate 71 kDa protein	0.19	4
P11021	78 kDa glucose-regulated protein	0.18	11
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.18	1
Q96PK6	RNA-binding protein 14	0.18	1
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.18	3
P02675	Fibrinogen beta chain	0.18	1
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.18	2
P17844	Probable ATP-dependent RNA helicase DDX5	0.17	1
Q00639	Heterogeneous nuclear ribonucleoprotein U	0.17	3
HBVRG3	Keratin 1	0.16	17
Q5BKZ1	DBIRD complex subunit ZNF326	0.16	2
Q00610	Clathrin heavy chain 1	0.16	11
P08238	Heat shock protein HSP 90-beta	0.15	3
P35908	Keratin type II cytoskeletal 2 epidermal	0.15	13
P05783	Keratin type I cytoskeletal 18	0.14	10
P14625	Endoplasmic reticulum chaperone	0.14	1
P13647	Keratin type II cytoskeletal 5	0.13	3
P51659	Peroxisomal multifunctional enzyme type 2	0.12	9
P13645	Keratin type I cytoskeletal 10	0.12	11
P35527	Keratin type I cytoskeletal 9	0.12	9
P00338	L-lactate dehydrogenase A chain	0.11	2
Q15058	Kinesin-like protein KIF14	0.09	6
P06748	Nucleophosmin	0.09	1
P06733	Alpha-enolase	0.09	1
P05787	Keratin type II cytoskeletal 8	0.08	6