

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q5TKA1	Protein lin-9 homolog	6.36	8
Q09028	Histone-binding protein RBBP4	3.01	3
<b>P10244</b>	<b>Myb-related protein B</b>	<b>1.86</b>	<b>3</b>
Q94905	Erlin-2	1.17	4
P27694	Replication protein A 70 kDa DNA-binding subunit	1.16	6
A1L0T0	Acetolactate synthase-like protein	1.15	1
Q00341	Vigilin	1.01	1
P23528	Cofilin-1	0.98	3
Q8N6S9	Thioredoxin domain-containing protein 5	0.97	2
O15523	ATP-dependent RNA helicase DDX3Y	0.94	12
Q9ULV0	Unconventional myosin-Vb	0.91	44
Q13885	Tubulin beta-2A chain	0.88	22
Q95299	NADH dehydrogenase [ubiquinone] 1 alpha subcomplex subunit 10 mitochondrial	0.87	1
Q13509	Tubulin beta-3 chain	0.87	19
Q8T030	Alanine aminotransferase 2	0.84	1
Q17519	RNAPNP complex-1-interacting phosphatase	0.83	2
Q9BSJ8	Extended synaptotagmin-1	0.80	1
O00571	ATP-dependent RNA helicase DDX3X	0.79	14
Q9HCC0	Methycrotonyl-CoA carboxylase beta chain mitochondrial	0.79	4
Q9Y5V3	Melanoma-associated antigen D1	0.77	3
Q15436	Protein transport protein Sec23A	0.76	4
Q6YN16	Hydroxysteroid dehydrogenase-like protein 2	0.75	1
Q9NY26	Zinc transporter ZIP1	0.69	1
P62829	60S ribosomal protein L23	0.67	3
O43813	LanC-like protein 1	0.66	2
Q9NQX4	Unconventional myosin-Vc	0.64	3
Q06520	Bile salt sulfotransferase	0.63	3
Q6P1L8	36S ribosomal protein L14 mitochondrial	0.62	1
P11586	C-1-tetrahydrofolate synthase cytoplasmic	0.62	1
P35232	Prohibitin	0.61	2
P07437	Tubulin beta chain	0.60	29
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.59	2
P49327	Fatty acid synthase	0.58	8
P51571	Translocon-associated protein subunit delta	0.56	2
P14868	Aspartate--tRNA ligase cytoplasmic	0.55	3
Q43251	RNA binding protein fox-1 homolog 2	0.53	1
Q9H9B4	Sideroflexin-1	0.53	1
Q9ULS0	Calcium-binding mitochondrial carrier protein Aralar2	0.53	1
P36542	ATP synthase subunit gamma mitochondrial	0.53	3
P17844	Probable ATP-dependent RNA helicase DDX5	0.52	3
Q43795	Unconventional myosin-Ib	0.52	3
Q04828	Aldo-keto reductase family 1 member C1	0.51	1
Q9H8H3	Methyltransferase-like protein 7A	0.51	1
P53618	Coatomer subunit beta	0.51	1
Q6P2N6	Protein ADORA3 isoform 3	0.50	2
O60547	GDP-mannose 46 dehydratase	0.50	2
Q95793	Double-stranded RNA-binding protein Staufen homolog 1	0.50	3
Q04837	Single-stranded DNA-binding protein mitochondrial	0.50	5
P47897	Glutamine--tRNA ligase	0.49	2
P68363	Tubulin alpha-1B chain	0.49	20
Q5H9R7	Serinethreonine-protein phosphatase 6 regulatory subunit 3	0.49	3
P68371	Tubulin beta-4B chain	0.48	22
Q9Y411	Unconventional myosin-Va	0.48	2
Q15084	Protein disulfide-isomerase A6	0.48	5
Q06830	Peroxisome-oxalyl-CoA decarboxylase	0.48	2
Q9H479	Fructosamine-3-kinase	0.48	1
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.47	4
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.45	7
P56192	Methionine--tRNA ligase cytoplasmic	0.44	1
P02671	Fibrinogen alpha chain	0.44	14
P02679	Fibrinogen gamma chain	0.44	9
Q43143	Putative pre-mRNA-splicing factor ATP-dependent RNA helicase DHX15	0.44	1
P45880	Voltage-dependent anion-selective channel protein 2	0.43	2
Q12931	Heat shock protein 75 kDa mitochondrial	0.43	3
P26641	Elongation factor 1-gamma	0.43	1
P12004	Proliferating cell nuclear antigen	0.42	2
P50991	T-complex protein 1 subunit delta	0.42	1
Q92945	Far upstream element-binding protein 2	0.42	1
Q61B91	PCK2 protein	0.41	3
P25705	ATP synthase subunit alpha mitochondrial	0.38	6
P05141	ADP/ATP translocase 2	0.38	5
P41252	Isoleucine--tRNA ligase cytoplasmic	0.37	1
P06576	ATP synthase subunit beta mitochondrial	0.37	5
P01859	Ig gamma-2 chain C region	0.37	1
Q9UBM7	7-dehydrocholesterol reductase	0.37	2
P68104	Elongation factor 1-alpha 1	0.37	10
P62987	Ubiquitin-60S ribosomal protein L40	0.36	1
P52997	Heterogeneous nuclear ribonucleoprotein F	0.35	2
P178527	DNA-dependent protein kinase catalytic subunit	0.35	3
P49411	Elongation factor Tu mitochondrial	0.34	5
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.32	7
P31943	Heterogeneous nuclear ribonucleoprotein H	0.32	6
Q3LXA3	Bifunctional ATP-dependent dihydroxyacetone kinase/FAD-AMP lyase (cycling)	0.31	1
Q08211	ATP-dependent RNA helicase A	0.30	1
P02675	Fibrinogen beta chain	0.30	2
Q9BLU2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.29	13
P62826	GTP-binding nuclear protein Ran	0.29	1
P60842	Eukaryotic initiation factor 4A-1	0.28	3
Q99623	Prohibitin-2	0.27	1
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.26	10
P08236	Heat shock protein HSP 90-beta	0.26	6
P51659	Peroxisomal multifunctional enzyme type 2	0.26	20
A2NLT2	Lambda-chain (AA_20 to 215)	0.25	4
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.24	5
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.24	2
P01617	Ig kappa chain V-II region TEW	0.24	2
Q96PK6	RNA-binding protein 14	0.24	1
Q60701	UDP-glucose 6-dehydrogenase	0.24	4
Q9H222	ATP-binding cassette sub-family G member 5	0.23	1
Q9NZ18	Insulin-like growth factor 2 mRNA-binding protein 1	0.21	1
P61978	Heterogeneous nuclear ribonucleoprotein K	0.21	1
Q149N6	Dedicator of cytokinesis 4	0.21	10
P15924	Desmoplakin	0.21	8
Q14847	LIM and SH3 domain protein 1	0.20	1
P01700	Ig lambda chain V-I region HA	0.20	2
Q15058	Kinesin-like protein KIF14	0.20	14
P54136	Arginine--tRNA ligase cytoplasmic	0.19	1
P02769	Serum albumin	0.19	64
D3DPC0	Titin isoform CRA_a	0.19	1
P11142	Heat shock cognate 71 kDa protein	0.19	4
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.19	1
P10809	60 kDa heat shock protein mitochondrial	0.18	3
Q15366	Poly(RC)-binding protein 2	0.18	2
P08107	Heat shock 70 kDa protein 1A1B	0.18	3
Q5BKZ1	DBIRD complex subunit ZNF326	0.16	2
Q15365	Poly(RC)-binding protein 1	0.16	1
P05787	Keratin type II cytoskeletal 8	0.15	13
P11021	78 kDa glucose-regulated protein	0.15	9
P02768	Serum albumin	0.14	6
P14618	Pyruvate kinase PKM	0.14	1
P06748	Nucleophosmin	0.14	2
P19474	E3 ubiquitin-protein ligase TRIM21	0.13	4
P05783	Keratin type I cytoskeletal 16	0.13	9
P13647	Keratin type II cytoskeletal 5	0.13	3
P08727	Keratin type I cytoskeletal 19	0.12	3
P00338	L-lactate dehydrogenase A chain	0.11	2
P02751	Fibronectin	0.08	1
P52272	Heterogeneous nuclear ribonucleoprotein M	0.08	1
P60709	Actin cytoplasmic 1	0.07	3
P13645	Keratin type I cytoskeletal 10	0.05	4
HEVRC3	Keratin 1	0.05	4
P35908	Keratin type II cytoskeletal 2 epidermal	0.04	2