

Accession Number	Description	Fold Enrichment	UniquePeptideCount
<b>P08048</b>	<b>Zinc finger Y-chromosomal protein</b>	<b>2.40</b>	<b>2</b>
C9J116	Protein ATP5J2-PTC1	1.70	1
P63173	60S ribosomal protein L38	1.70	1
Q6SZW1	Sterile alpha and TIR motif-containing protein 1	1.70	1
Q7KZ99	Cytochrome c oxidase assembly protein COX15 homolog	1.70	1
Q9BVP2	Guanine nucleotide-binding protein-like 3	1.70	1
Q9H5Y7	SLIT and NTRK-like protein 6	1.70	1
Q9NR30	Nucleolar RNA helicase 2	1.70	1
Q9BQE3	Tubulin alpha-1c chain	1.70	17
P16403	Histone H12	1.67	3
P45954	Shortbranched chain specific acyl-CoA dehydrogenase mitochondrial	1.47	4
P61353	60S ribosomal protein L27	1.38	4
Q7Z460	CLIP-associating protein 1	1.35	1
Q00325	Phosphate carrier protein mitochondrial	1.26	2
P46782	40S ribosomal protein S5	1.24	2
Q60884	DnaJ homolog subfamily A member 2	1.24	2
D3DR22	Hydroxysteroid (17-beta) dehydrogenase 12 isoform CRA_a	1.17	1
P18085	ADP-ribosylation factor 4	0.95	3
Q9Y678	Coatomer subunit gamma-1	0.94	1
P62854	40S ribosomal protein S6	0.91	2
P46781	40S ribosomal protein S9	0.88	2
P80748	Ig lambda chain V-III region LOI	0.88	4
P62081	40S ribosomal protein S7	0.86	3
P62987	Ubiquitin-60S ribosomal protein L40	0.86	3
P61247	40S ribosomal protein S3a	0.85	2
Q13838	Spliceosome RNA helicase DDX39B	0.84	2
P27708	CAD protein	0.84	1
Q06418	Tyrosine-protein kinase receptor TYRO3	0.82	1
Q9HCC0	Methylcrotonoyl-CoA carboxylase beta chain mitochondrial	0.82	1
Q99623	Prohibitin-2	0.80	4
P26441	Elongation factor 1-gamma	0.78	2
P61254	60S ribosomal protein L26	0.78	1
Q14157	Ubiquitin-associated protein 2-like	0.77	1
P32969	60S ribosomal protein L9	0.77	1
P18621	60S ribosomal protein L17	0.76	4
O75477	Erlin-1	0.75	1
Q16777	Histone H2A type 2-C	0.73	2
P48327	Fatty acid synthase	0.73	7
P61626	Lysozyme C	0.71	1
P26373	60S ribosomal protein L13	0.71	1
P61978	Heterogeneous nuclear ribonucleoprotein K	0.70	4
P62829	60S ribosomal protein L23	0.70	2
P40227	T-complex protein 1 subunit zeta	0.70	1
Q60814	Histone H2B type 1-K	0.70	1
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.68	5
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.68	4
P49368	T-complex protein 1 subunit gamma	0.67	2
Q96428	SLAM family member 9	0.66	1
Q06210	6- fructose-6-phosphate aminotransferase [isomerizing] 1	0.65	1
Q02543	60S ribosomal protein L18a	0.65	6
P14868	Aspartate--tRNA ligase cytoplasmic	0.64	2
Q14638	Ectonucleotide pyrophosphatasephosphodiesterase family member 3	0.64	1
P62249	40S ribosomal protein S16	0.62	4
P93041	Peroxisedoxin-6	0.60	1
P78527	DNA-dependent protein kinase catalytic subunit	0.59	3
P51571	Translocon-associated protein subunit delta	0.59	1
P22895	Cytochrome b-c1 complex subunit 2 mitochondrial	0.57	1
P68363	Tubulin alpha-1B chain	0.57	16
Q43795	Unconventional myosin-Ib	0.56	1
P36542	ATP synthase subunit gamma mitochondrial	0.55	2
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.55	2
P50991	T-complex protein 1 subunit delta	0.54	1
Q15084	Protein disulfide-isomerase A6	0.53	3
P62805	Histone H4	0.52	2
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.49	8
P41252	Isoleucine--tRNA ligase cytoplasmic	0.49	1
P68371	Tubulin beta-4B chain	0.49	15
P46776	60S ribosomal protein L27a	0.49	2
P01859	Ig gamma-2 chain C region	0.48	1
P47897	Glutamine--tRNA ligase	0.47	1
P07437	Tubulin beta chain	0.47	15
Q06211	ATP-dependent RNA helicase A	0.46	1
P62701	40S ribosomal protein S4 X isoform	0.46	2
Q06520	Bile salt sulfotransferase	0.46	1
Q61891	PCK2 protein	0.45	2
P02679	Fibrinogen gamma chain	0.44	6
P10809	60 kDa heat shock protein mitochondrial	0.42	6
A2N112	Lambda-chain (AA-20 to 215)	0.41	5
Q15365	Poly(rC)-binding protein 1	0.41	3
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.40	6
Q9BVG8	Kinesin-like protein KIFC3	0.40	2
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.40	1
P62513	60S ribosomal protein L11	0.39	1
Q60701	UDP-glucose 6-dehydrogenase	0.39	5
Q14847	LIM and SH3 domain protein 1	0.39	2
P62826	GTP-binding nuclear protein Ran	0.38	1
Q81216	Abnormal spindle-like microcephaly-associated protein	0.38	1
P54136	Arginine--tRNA ligase cytoplasmic	0.37	2
P08238	Heat shock protein HSP 90-beta	0.36	6
P02769	Serum albumin	0.36	84
O75533	Splicing factor 3B subunit 1	0.36	2
P06576	ATP synthase subunit beta mitochondrial	0.36	3
P01700	Ig lambda chain V-I region HA	0.35	3
P13639	Elongation factor 2	0.35	1
P02656	Aspirinogen C-III	0.34	2
P01605	Ig kappa chain V-I region Lay	0.33	1
Q86Y82	Syntaxin-12	0.33	1
P31943	Heterogeneous nuclear ribonucleoprotein H	0.33	4
P19474	E3 ubiquitin-protein ligase TRIM21	0.32	8
P18124	60S ribosomal protein L7	0.32	1
P23396	40S ribosomal protein S3	0.32	1
P25705	ATP synthase subunit alpha mitochondrial	0.32	3
P60709	Actin cytoplasmic 1	0.30	11
Q06830	Peroxisedoxin-1	0.30	1
Q9N218	Insulin-like growth factor 2 mRNA-binding protein 1	0.29	1
P05141	ADP/ATP translocase 2	0.28	2
P04075	Fructose-bisphosphate aldolase A	0.27	1
P14618	Pyruvate kinase PKM	0.26	2
P60842	Eukaryotic initiation factor 4A-I	0.26	1
P49411	Elongation factor Tu mitochondrial	0.25	2
P68104	Elongation factor 1-alpha 1	0.25	4
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.23	1
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.23	1
P08727	Keratin type I cytoskeletal 19	0.22	3
P02768	Serum albumin	0.20	6
P51659	Peroxisomal multifunctional enzyme type 2	0.20	9
P00338	L-lactate dehydrogenase A chain	0.19	3
P02671	Fibrinogen alpha chain	0.18	3
Q9BUZ4	TNF receptor-associated factor 4	0.18	1
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.16	4
P08107	Heat shock 70 kDa protein 1A1B	0.14	1
P05783	Keratin type I cytoskeletal 18	0.14	3
Q149N5	DCK4 protein	0.13	3
P06733	Alpha-enolase	0.13	1
P05787	Keratin type II cytoskeletal 8	0.11	4
Q9ULV0	Unconventional myosin-Vb	0.11	2
H6VRG3	Keratin 1	0.11	7
P11021	78 kDa glucose-regulated protein	0.09	3
Q07814	Bifunctional glutamaterproline--tRNA ligase	0.09	1
P22061	Protein-L-isaspartate(D-aspartate) O-methyltransferase	0.08	1
P13645	Keratin type I cytoskeletal 10	0.06	3
Q00610	Clastrin heavy chain 1	0.06	2
Q15058	Kinesin-like protein KIF14	0.06	2
P35527	Keratin type I cytoskeletal 9	0.06	2