

Accession Number	Description	Fold Enrichment	UniquePeptideCount
<b>Q9UBC0</b>	<b>Hepatocyte nuclear factor 6</b>	<b>5.46</b>	<b>6</b>
P15822	Zinc finger protein 40	1.39	1
P62917	60S ribosomal protein L8	1.06	3
P23528	Cofilin-1	1.06	2
Q6IPH7	RPL14 protein	1.04	3
Q99623	Prohibitin-2	0.99	5
P30837	Aldehyde dehydrogenase X mitochondrial	0.97	2
P62266	40S ribosomal protein S23	0.96	2
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.93	4
P62701	40S ribosomal protein S4 X isoform	0.92	5
P62241	40S ribosomal protein S8	0.92	1
P60866	40S ribosomal protein S20	0.90	1
P06576	ATP synthase subunit beta mitochondrial	0.89	9
Q13838	Spliceosome RNA helicase DDX39B	0.88	2
P46782	40S ribosomal protein S5	0.82	1
Q06830	Peroxisome oxidin-1	0.82	5
Q15393	Splicing factor 3B subunit 3	0.82	1
P26641	Elongation factor 1-gamma	0.81	2
P61353	60S ribosomal protein L27	0.79	2
P45880	Voltage-dependent anion-selective channel protein 2	0.79	3
P32969	60S ribosomal protein L9	0.79	1
P30048	Thioredoxin-dependent peroxide reductase mitochondrial	0.78	1
P46779	60S ribosomal protein L28	0.78	1
Q13907	Isopentenyl-diphosphate Delta-isomerase 1	0.77	1
P04843	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1	0.75	1
P05141	ADP-ATP translocase 2	0.75	7
Q6IB91	PCK2 protein	0.74	4
P61626	Lysozyme C	0.73	1
P26373	60S ribosomal protein L13	0.73	1
P49368	T-complex protein 1 subunit gamma	0.69	2
Q06520	Bile salt sulfotransferase	0.68	2
P49327	Fatty acid synthase	0.67	6
Q06210	Glutamine--fructose-6-phosphate aminotransferase [isomerizing] 1	0.67	1
Q15436	Protein transport protein Sec23A	0.66	1
P13639	Elongation factor 2	0.66	3
P62249	40S ribosomal protein S16	0.65	4
P18621	60S ribosomal protein L17	0.65	3
A2NUT2	Lambda-chain (AA -20 to 215)	0.64	8
P68371	Tubulin beta-4B chain	0.64	19
P62854	40S ribosomal protein S26	0.63	1
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.63	10
O43175	D-3-phosphoglycerate dehydrogenase	0.62	1
P30041	Peroxisome oxidin-6	0.62	1
P78527	DNA-dependent protein kinase catalytic subunit	0.62	3
P51571	Translocon-associated protein subunit delta	0.61	1
P02679	Fibrinogen gamma chain	0.59	8
P45954	Shortbranched chain specific acyl-CoA dehydrogenase mitochondrial	0.57	1
O43795	Unconventional myosin-Ib	0.57	1
P32322	Pyroline-5-carboxylate reductase 1 mitochondrial	0.56	1
P07437	Tubulin beta chain	0.55	17
Q9NZI8	Insulin-like growth factor 2 mRNA-binding protein 1	0.55	3
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.54	4
Q16777	Histone H2A type 2-C	0.53	1
P62829	60S ribosomal protein L23	0.51	1
P46776	60S ribosomal protein L27a	0.50	2
P27635	60S ribosomal protein L10	0.50	2
P18085	ADP-ribosylation factor 4	0.50	1
P62081	40S ribosomal protein S7	0.48	1
P62987	Ubiquitin-60S ribosomal protein L40	0.48	1
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.48	7
P02675	Fibrinogen beta chain	0.48	2
O60701	UDP-glucose 6-dehydrogenase	0.48	6
Q08211	ATP-dependent RNA helicase A	0.48	1
P22626	Heterogeneous nuclear ribonucleoproteins A2B1	0.47	1
P23396	40S ribosomal protein S3	0.47	2
P68363	Tubulin alpha-1B chain	0.46	12
P14868	Aspartate--tRNA ligase cytoplasmic	0.46	1
P61978	Heterogeneous nuclear ribonucleoprotein K	0.46	2
P08238	Heat shock protein HSP 90-beta	0.44	7
Q15084	Protein disulfide-isomerase A6	0.43	2
P17844	Probable ATP-dependent RNA helicase DDX5	0.42	1
P60709	Actin cytoplasmic 1	0.42	15
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.40	1
P62805	Histone H4	0.38	1
P22061	Protein-L-isoaspartate(D-aspartate) O-methyltransferase	0.37	9
P14618	Pyruvate kinase PKM	0.36	3
P68104	Elongation factor 1-alpha 1	0.36	6
P49411	Elongation factor Tu mitochondrial	0.34	3
P25705	ATP synthase subunit alpha mitochondrial	0.33	3
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.31	1
P38646	Stress-70 protein mitochondrial	0.29	1
P01700	Ig lambda chain V-I region HA	0.28	2
P05783	Keratin type I cytoskeletal 18	0.27	7
P60842	Eukaryotic initiation factor 4A-1	0.27	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.27	6
P54136	Arginine--tRNA ligase cytoplasmic	0.27	1
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.26	1
P10809	60 kDa heat shock protein mitochondrial	0.26	3
O75533	Splicing factor 3B subunit 1	0.26	1
P02769	Serum albumin	0.24	53
P02671	Fibrinogen alpha chain	0.23	4
Q149N5	DOCK4 protein	0.23	6
H6VRG3	Keratin 1	0.23	16
Q15365	Poly(rC)-binding protein 1	0.23	1
P51659	Peroxisomal multifunctional enzyme type 2	0.23	10
Q02543	60S ribosomal protein L18a	0.22	1
P31943	Heterogeneous nuclear ribonucleoprotein H	0.22	2
P02768	Serum albumin	0.21	6
P08107	Heat shock 70 kDa protein 1A1B	0.20	2
P05787	Keratin type II cytoskeletal 8	0.20	8
O00571	ATP-dependent RNA helicase DDX3X	0.18	1
P07814	Bifunctional glutamateproline--tRNA ligase	0.17	3
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.16	4
P06748	Nucleophosmin	0.14	1
P06733	Alpha-enolase	0.13	1
P11142	Heat shock cognate 71 kDa protein	0.12	1
P08727	Keratin type I cytoskeletal 19	0.12	1
Q15058	Kinesin-like protein KIF14	0.10	4
P13645	Keratin type I cytoskeletal 10	0.10	5
P35908	Keratin type II cytoskeletal 2 epidermal	0.08	4
Q9ULV0	Unconventional myosin-Vb	0.08	1
P11021	78 kDa glucose-regulated protein	0.07	2
P35527	Keratin type I cytoskeletal 9	0.04	1