

Accession Number	Description	Fold Enrichment	UniquePeptideCount
<b>P57071</b>	<b>PR domain zinc finger protein 15</b>	<b>2.57</b>	<b>2</b>
P62917	60S ribosomal protein L8	1.49	3
P62805	Histone H4	1.24	4
Q13838	Spliceosome RNA helicase DDX39B	1.13	2
O00410	Importin-5	1.13	1
P18621	60S ribosomal protein L17	1.07	4
Q00325	Phosphate carrier protein mitochondrial	1.00	1
P46782	40S ribosomal protein S5	0.99	1
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.97	5
P06576	ATP synthase subunit beta mitochondrial	0.97	7
Q6IPH7	RPL14 protein	0.97	2
P61254	60S ribosomal protein L26	0.97	1
P18085	ADP-ribosylation factor 4	0.94	2
P62847	40S ribosomal protein S24	0.92	1
P04843	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1	0.89	1
P49368	T-complex protein 1 subunit gamma	0.86	2
P07437	Tubulin beta chain	0.84	19
Q08211	ATP-dependent RNA helicase A	0.84	2
Q06814	Histone H2B type 1-K	0.83	1
P23528	Cofilin-1	0.82	1
P48643	T-complex protein 1 subunit epsilon	0.75	2
P62854	40S ribosomal protein S26	0.74	1
P45880	Voltage-dependent anion-selective channel protein 2	0.74	2
P15880	40S ribosomal protein S2	0.73	2
P30041	Peroxisomal protein S2	0.72	1
Q15084	Protein disulfide-isomerase A6	0.70	3
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.70	4
P62826	GTP-binding nuclear protein Ran	0.68	2
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	0.68	1
P62249	40S ribosomal protein S16	0.67	3
P68371	Tubulin beta-4B chain	0.65	14
P61353	60S ribosomal protein L27	0.65	1
Q99623	Prohibitin-2	0.64	2
P62263	40S ribosomal protein S14	0.64	2
P49327	Fatty acid synthase	0.63	4
P46776	60S ribosomal protein L27a	0.62	2
P62829	60S ribosomal protein L23	0.60	1
P62701	40S ribosomal protein S4 X isoform	0.59	2
P18124	60S ribosomal protein L7	0.58	2
P08727	Keratin type I cytoskeletal 19	0.58	7
P01859	Ig gamma-2 chain C region	0.58	1
P62081	40S ribosomal protein S7	0.56	1
P14868	Aspartate--tRNA ligase cytoplasmic	0.54	1
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.54	6
P31943	Heterogeneous nuclear ribonucleoprotein H	0.54	5
P0CG38	POTE ankyrin domain family member I	0.53	2
Q9BVG8	Kinesin-like protein KIFC3	0.51	2
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.47	1
P68363	Tubulin alpha-1B chain	0.43	8
P13639	Elongation factor 2	0.41	1
P27635	60S ribosomal protein L10	0.41	1
P05783	Keratin type I cytoskeletal 18	0.41	8
P68104	Elongation factor 1-alpha 1	0.40	5
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.40	4
Q149N5	DOCK4 protein	0.39	8
P23396	40S ribosomal protein S3	0.38	1
Q02543	60S ribosomal protein L18a	0.38	2
A2NUT2	Lambda-chain (AA -20 to 215)	0.38	3
P61978	Heterogeneous nuclear ribonucleoprotein K	0.38	1
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.36	1
P05141	ADP/ATP translocase 2	0.36	2
P60709	Actin cytoplasmic 1	0.35	9
P01700	Ig lambda chain V-I region HA	0.35	2
Q14847	LIM and SH3 domain protein 1	0.33	1
P25705	ATP synthase subunit alpha mitochondrial	0.32	2
P06733	Alpha-enolase	0.31	3
P54136	Arginine--tRNA ligase cytoplasmic	0.31	1
P08238	Heat shock protein HSP 90-beta	0.29	3
P02769	Serum albumin	0.27	42
Q06701	UDP-glucose 6-dehydrogenase	0.27	2
Q15365	Poly(rC)-binding protein 1	0.27	1
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.26	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.25	4
P22061	Protein-L-isoaspartate(D-aspartate) O-methyltransferase	0.24	4
P02533	Keratin type I cytoskeletal 14	0.24	3
Q9BUZ4	TNF receptor-associated factor 4	0.21	1
H6VRG3	Keratin 1	0.20	10
P02768	Serum albumin	0.19	4
P51659	Peroxisomal multifunctional enzyme type 2	0.19	6
P02679	Fibrinogen gamma chain	0.17	1
P08107	Heat shock 70 kDa protein 1A1B	0.16	1
P06748	Nucleophosmin	0.16	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.15	6
P11142	Heat shock cognate 71 kDa protein	0.14	1
P35527	Keratin type I cytoskeletal 9	0.14	5
P00338	L-lactate dehydrogenase A chain	0.13	1
P02671	Fibrinogen alpha chain	0.12	1
P05787	Keratin type II cytoskeletal 8	0.12	3
P35579	Myosin-9	0.11	1
Q15058	Kinesin-like protein KIF14	0.11	3
P11021	78 kDa glucose-regulated protein	0.09	2
P13645	Keratin type I cytoskeletal 10	0.08	3
Q00610	Clathrin heavy chain 1	0.05	1