

Accession Number	Description	Fold Enrichment	UniquePeptideCount
O43296	Zinc finger protein 264	8.43	6
L8B4Z6	Ubiquitin C	1.42	4
P29558	RNA-binding motif single-stranded-interacting protein 1	1.23	2
P43307	Translocon-associated protein subunit alpha	1.09	1
P61978	Heterogeneous nuclear ribonucleoprotein K	1.01	7
F8J2B5	LIM domain only 7 protein	1.00	2
P62937	Peptidyl-prolyl cis-trans isomerase A	0.93	2
O00425	Insulin-like growth factor 2 mRNA-binding protein 3	0.79	2
Q6S8J3	POTE ankyrin domain family member E	0.74	3
P01614	Ig kappa chain V-II region Cum	0.74	2
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.71	4
Q14157	Ubiquitin-associated protein 2-like	0.71	2
Q9HCC0	Methylcrotonoyl-CoA carboxylase beta chain mitochondrial	0.70	2
P68371	Tubulin beta-4B chain	0.68	20
O43251	RNA binding protein fox-1 homolog 2	0.67	1
Q96RD3	Olfactory receptor 52E6	0.63	1
Q06210	Glutamine--fructose-6-phosphate aminotransferase [isomerizing] 1	0.63	2
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.62	9
P68363	Tubulin alpha-1B chain	0.61	15
Q06830	Peroxiredoxin-1	0.61	4
P62829	60S ribosomal protein L23	0.60	2
P45880	Voltage-dependent anion-selective channel protein 2	0.60	2
P14868	Aspartate--tRNA ligase cytoplasmic	0.59	2
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.58	2
Q15365	PolyrC-binding protein 1	0.53	4
P31943	Heterogeneous nuclear ribonucleoprotein H	0.50	6
Q16822	Phosphoenolpyruvate carboxykinase [GTP] mitochondrial	0.50	3
Q15084	Protein disulfide-isomerase A6	0.49	3
P68104	Elongation factor 1-alpha 1	0.48	8
P60709	Actin cytoplasmic 1	0.48	16
P07437	Tubulin beta chain	0.44	14
P52597	Heterogeneous nuclear ribonucleoprotein F	0.40	2
Q149N6	Dedicator of cytokinesis 4	0.40	12
Q99623	Prohibitin-2	0.39	2
P49327	Fatty acid synthase	0.39	4
P02675	Fibrinogen beta chain	0.39	4
Q01085	Nucleolysin TIAR	0.38	1
Q04837	Single-stranded DNA-binding protein mitochondrial	0.37	2
P14618	Pyruvate kinase PKM	0.36	3
O60701	UDP-glucose 6-dehydrogenase	0.35	4
P08238	Heat shock protein HSP 90-beta	0.34	5
P49411	Elongation factor Tu mitochondrial	0.34	3
P25705	ATP synthase subunit alpha mitochondrial	0.33	3
P02671	Fibrinogen alpha chain	0.33	7
P46776	60S ribosomal protein L27a	0.32	1
P06748	Nucleophosmin	0.32	4
P02769	Serum albumin	0.32	63
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.31	2
P13639	Elongation factor 2	0.31	1
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.30	4
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.30	8
O75379	Vesicle-associated membrane protein 4	0.30	1
P02679	Fibrinogen gamma chain	0.29	5
Q9ULV0	Unconventional myosin-Vb	0.29	10
P07237	Protein disulfide-isomerase	0.28	1
P01700	Ig lambda chain V-I region HA	0.27	2
P10809	60 kDa heat shock protein mitochondrial	0.26	3
P08107	Heat shock 70 kDa protein 1A1B	0.26	3
P38646	Stress-70 protein mitochondrial	0.26	1
Q9BVG8	Kinesin-like protein KIFC3	0.25	3
P06576	ATP synthase subunit beta mitochondrial	0.24	2
P51659	Peroxisomal multifunctional enzyme type 2	0.24	11
D3DPG0	Titin isoform CRA a	0.23	1
P17844	Probable ATP-dependent RNA helicase DDX5	0.22	1
Q96PK6	RNA-binding protein 14	0.21	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.20	4
P60842	Eukaryotic initiation factor 4A-I	0.20	1
P22061	Protein-L-isoaspartateD-aspartate O-methyltransferase	0.19	4
P08779	Keratin type I cytoskeletal 16	0.18	2
Q15058	Kinesin-like protein KIF14	0.17	7
P11142	Heat shock cognate 71 kDa protein	0.17	2
P05787	Keratin type II cytoskeletal 8	0.16	8
P00338	L-lactate dehydrogenase A chain	0.15	2
P13647	Keratin type II cytoskeletal 5	0.14	2
P02533	Keratin type I cytoskeletal 14	0.13	2
H6VRG3	Keratin 1	0.13	8
O00571	ATP-dependent RNA helicase DDX3X	0.13	2
P05783	Keratin type I cytoskeletal 18	0.12	5
P02768	Serum albumin	0.12	3
P07814	Bifunctional glutamateproline--tRNA ligase	0.08	1
P11021	78 kDa glucose-regulated protein	0.07	2
P13645	Keratin type I cytoskeletal 10	0.06	3
P43243	Matrin-3	0.06	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.05	2
P35580	Myosin-10	0.03	1