

Accession Number	Description	Fold Enrichment	UniquePeptideCount
P52746	Zinc finger protein 142	3.57	2
O95390	Growth differentiation factor 11	1.48	2
P35232	Prohibitin	0.88	2
Q13838	Spliceosome RNA helicase DDX39B	0.86	2
Q16659	Mitogen-activated protein kinase 6	0.84	1
P01614	Ig kappa chain V-II region Cum	0.77	2
Q6S8J3	POTE ankyrin domain family member E	0.77	3
Q14157	Ubiquitin-associated protein 2-like	0.73	2
A6NLN1	Polypyrimidine tract binding protein 1 isoform CRA_b	0.70	1
P21980	Protein-glutamine gamma-glutamyltransferase 2	0.65	1
Q06830	Peroxisome oxidin-1	0.63	4
P45880	Voltage-dependent anion-selective channel protein 2	0.62	2
P62937	Peptidyl-prolyl cis-trans isomerase A	0.61	1
P31943	Heterogeneous nuclear ribonucleoprotein H	0.59	7
P36542	ATP synthase subunit gamma mitochondrial	0.58	2
O43795	Unconventional myosin-Ib	0.57	2
P08107	Heat shock 70 kDa protein 1A1B	0.55	7
P04843	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1	0.53	1
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.52	3
Q13268	Dehydrogenase reductase SDR family member 2 mitochondrial	0.51	7
P51659	Peroxisomal multifunctional enzyme type 2	0.50	23
Q9ULV0	Unconventional myosin-Vb	0.46	16
Q6P2N6	Protein ADORA3 isoform 3	0.46	1
P07237	Protein disulfide-isomerase	0.45	2
P50454	Serpin H1	0.44	1
P22061	Protein-L-isoaspartate D-aspartate O-methyltransferase	0.44	10
P05388	60S acidic ribosomal protein P0	0.44	1
Q15365	PolyrC-binding protein 1	0.44	3
Q92841	Probable ATP-dependent RNA helicase DDX17	0.43	2
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.42	2
P52597	Heterogeneous nuclear ribonucleoprotein F	0.41	2
P60709	Actin cytoplasmic 1	0.41	13
P62829	60S ribosomal protein L23	0.40	1
P07437	Tubulin beta chain	0.39	12
P15880	40S ribosomal protein S2	0.39	1
Q16822	Phosphoenolpyruvate carboxykinase [GTP] mitochondrial	0.38	2
P61978	Heterogeneous nuclear ribonucleoprotein K	0.37	2
H6VRG3	Keratin 1	0.36	23
P68371	Tubulin beta-4B chain	0.36	10
D3DPG0	Titin isoform CRA_a	0.36	2
Q149N6	Dedicator of cytokinesis 4	0.35	10
P12004	Proliferating cell nuclear antigen	0.35	1
P17844	Probable ATP-dependent RNA helicase DDX5	0.35	2
P02769	Serum albumin	0.34	66
P06576	ATP synthase subunit beta mitochondrial	0.34	3
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.34	2
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.32	2
P68363	Tubulin alpha-1B chain	0.31	7
P60842	Eukaryotic initiation factor 4A-I	0.31	2
P02533	Keratin type I cytoskeletal 14	0.29	5
P01700	Ig lambda chain V-I region HA	0.28	2
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.28	7
P35908	Keratin type II cytoskeletal 2 epidermal	0.27	14
P05783	Keratin type I cytoskeletal 18	0.26	11
Q99623	Prohibitin-2	0.26	1
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.26	2
P02671	Fibrinogen alpha chain	0.25	5
P02679	Fibrinogen gamma chain	0.25	4
O94776	Metastasis-associated protein MTA2	0.25	1
Q15084	Protein disulfide-isomerase A6	0.24	1
P49327	Fatty acid synthase	0.24	2
O60701	UDP-glucose 6-dehydrogenase	0.21	2
P10809	60 kDa heat shock protein mitochondrial	0.20	2
P05141	ADP/ATP translocase 2	0.18	1
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.18	2
P11142	Heat shock cognate 71 kDa protein	0.17	2
P49411	Elongation factor Tu mitochondrial	0.16	1
P68104	Elongation factor 1-alpha 1	0.16	2
P13645	Keratin type I cytoskeletal 10	0.15	8
P35527	Keratin type I cytoskeletal 9	0.15	7
P13647	Keratin type II cytoskeletal 5	0.14	2
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.13	1
P11021	78 kDa glucose-regulated protein	0.12	4
P19474	E3 ubiquitin-protein ligase TRIM21	0.12	2
P08779	Keratin type I cytoskeletal 16	0.12	1
Q15058	Kinesin-like protein KIF14	0.08	3
P07814	Bifunctional glutamate-proline--tRNA ligase	0.08	1
P05787	Keratin type II cytoskeletal 8	0.07	3
P02768	Serum albumin	0.06	1
P43243	Matrin-3	0.06	1
Q00610	Clathrin heavy chain 1	0.04	1