

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q9UPP1	Histone lysine demethylase PHF8	10.75	12
Q15436	Protein transport protein Sec23A	1.54	2
Q86VU5	Catechol O-methyltransferase domain-containing protein 1	1.41	1
O60814	Histone H2B type 1-K	1.36	2
K9N0C7	Interleukin 15 receptor alpha isoform IC4	1.01	1
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.96	3
Q08211	ATP-dependent RNA helicase A	0.89	2
P30041	Peroxiredoxin-6	0.85	1
Q16822	Phosphoenolpyruvate carboxykinase [GTP] mitochondrial	0.80	3
D3DPG0	Titin isoform CRA_a	0.75	4
P54886	Delta-1-pyrroline-5-carboxylate synthase	0.72	3
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.71	3
P62826	GTP-binding nuclear protein Ran	0.70	2
P13639	Elongation factor 2	0.69	2
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.64	4
Q4W4Y1	Dopamine receptor interacting protein 4	0.61	2
O75533	Splicing factor 3B subunit 1	0.60	3
P68104	Elongation factor 1-alpha 1	0.59	2
P01709	Ig lambda chain V-II region MGC	0.59	4
P19474	E3 ubiquitin-protein ligase TRIM21	0.58	11
P62249	40S ribosomal protein S16	0.57	2
Q07954	Prolow-density lipoprotein receptor-related protein 1	0.56	1
P00338	L-lactate dehydrogenase A chain	0.56	8
P02671	Fibrinogen alpha chain	0.52	8
Q9NZ18	Insulin-like growth factor 2 mRNA-binding protein 1	0.52	2
P68371	Tubulin beta-4B chain	0.50	11
Q15523	ATP-dependent RNA helicase DDX3Y	0.49	2
P08238	Heat shock protein HSP 90-beta	0.49	6
P04075	Fructose-bisphosphate aldolase A	0.48	2
P60842	Eukaryotic initiation factor 4A-I	0.48	2
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.46	3
P49327	Fatty acid synthase	0.45	2
P14618	Pyruvate kinase PKM	0.45	3
P05787	Keratin type II cytoskeletal 8	0.42	14
P18124	60S ribosomal protein L7	0.40	1
Q06830	Peroxiredoxin-1	0.40	1
P23396	40S ribosomal protein S3	0.40	1
P05783	Keratin type I cytoskeletal 18	0.39	8
P07900	Heat shock protein HSP 90-alpha	0.39	3
A1A508	PRSS3 protein	0.38	2
O60701	UDP-glucose 6-dehydrogenase	0.38	3
A2NUT2	Lambda-chain (AA -20 to 215)	0.37	3
P07437	Tubulin beta chain	0.36	8
P60709	Actin cytoplasmic 1	0.36	8
P08727	Keratin type I cytoskeletal 19	0.35	4
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.33	7
P06733	Alpha-enolase	0.31	3
P02769	Serum albumin	0.30	50
P02768	Serum albumin	0.30	7
P51659	Peroxisomal multifunctional enzyme type 2	0.29	10
Q15058	Kinesin-like protein KIF14	0.28	10
Q15365	Poly(rC)-binding protein 1	0.28	1
P07814	Bifunctional glutamate/proline--tRNA ligase	0.27	4
P31943	Heterogeneous nuclear ribonucleoprotein H	0.27	2
Q9ULV0	Unconventional myosin-Vb	0.27	5
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.26	2
P14625	Endoplasmic	0.25	1
Q86YZ3	Hornerin	0.24	3
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.24	4
P49411	Elongation factor Tu mitochondrial	0.22	1
P25705	ATP synthase subunit alpha mitochondrial	0.22	1
Q00610	Clathrin heavy chain 1	0.21	7
P13645	Keratin type I cytoskeletal 10	0.20	9
P15924	Desmoplakin	0.19	2
P35527	Keratin type I cytoskeletal 9	0.18	7
P02679	Fibrinogen gamma chain	0.18	1
P11021	78 kDa glucose-regulated protein	0.18	5
P10809	60 kDa heat shock protein mitochondrial	0.18	1
Q149N5	DOCK4 protein	0.17	3
Q13268	Dehydrogenase/reductase SDR family member 2 mitochondrial	0.17	1
H6VRG3	Keratin 1	0.14	7
P68363	Tubulin alpha-1B chain	0.10	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.05	1
P35580	Myosin-10	0.04	1