

Accession Number	Description	Fold Enrichment	UniquePeptideCount
I1VZV6	Hemoglobin alpha 1	5.62	5
O60547	GDP-mannose 46 dehydratase	3.98	4
P15923	Transcription factor E2-alpha	2.99	3
Q14978	Nucleolar and coiled-body phosphoprotein 1	2.51	2
Q9H5Z1	Probable ATP-dependent RNA helicase DHX35	2.36	2
A4D159	Serine/threonine kinase 31	1.92	1
P27797	Calreticulin	1.92	1
P36268	Inactive gamma-glutamyltranspeptidase 2	1.92	1
Q3T906	N-acetylglucosamine-1-phosphotransferase subunits alpha/beta	1.92	1
Q495A1	T-cell immunoreceptor with Ig and ITIM domains	1.92	1
Q5JV73	FERM and PDZ domain-containing protein 3	1.92	1
Q9BTF0	THUMP domain-containing protein 2	1.92	1
Q9HAV5	Tumor necrosis factor receptor superfamily member 27	1.92	1
Q9P2H3	Intraflagellar transport protein 80 homolog	1.92	1
P06310	Ig kappa chain V-II region RPM1 6410	1.87	2
Q99729	Heterogeneous nuclear ribonucleoprotein A/B	1.61	1
P50991	T-complex protein 1 subunit delta	1.60	1
Q7RTU4	Class A basic helix-loop-helix protein 9	1.59	1
Q9H469	F-box/LRR-repeat protein 15	1.57	2
Q9NVP1	ATP-dependent RNA helicase DDX18	1.48	1
Q9GZU2	Paternally-expressed gene 3 protein	1.41	1
Q7L4E1	Protein FAM73B	1.38	1
Q13459	Unconventional myosin-IXb	1.35	1
A6N6J7	JARID1C protein	1.28	1
Q16777	Histone H2A type 2-C	1.26	2
Q14103	Heterogeneous nuclear ribonucleoprotein D0	1.08	3
O43242	26S proteasome non-ATPase regulatory subunit 3	0.97	1
Q9JEU0	Vesicle transport through interaction with t-SNAREs homolog 1B	0.96	1
P30041	Peroxiredoxin-6	0.92	1
Q8N2S1	Latent-transforming growth factor beta-binding protein 4	0.92	1
O75096	Low-density lipoprotein receptor-related protein 4	0.88	1
Q16647	Prostacyclin synthase	0.82	1
Q03164	Histone-lysine N-methyltransferase 2A	0.81	1
P62913	60S ribosomal protein L11	0.77	2
P36578	60S ribosomal protein L4	0.72	1
Q13201	Multimerin-1	0.67	1
P46776	60S ribosomal protein L27a	0.66	2
P01709	Ig lambda chain V-II region MGC	0.65	4
P49327	Fatty acid synthase	0.64	3
O60337	E3 ubiquitin-protein ligase MARCH6	0.64	1
Q06830	Peroxiredoxin-1	0.62	2
P02656	Apolipoprotein C-III	0.62	3
P06576	ATP synthase subunit beta mitochondrial	0.61	3
P02768	Serum albumin	0.58	13
P08727	Keratin type I cytoskeletal 19	0.54	6
P04075	Fructose-bisphosphate aldolase A	0.52	2
P62826	GTP-binding nuclear protein Ran	0.52	1
P13639	Elongation factor 2	0.50	1
O75533	Splicing factor 3B subunit 1	0.50	2
P14618	Pyruvate kinase PKM	0.49	3
P02769	Serum albumin	0.46	68
Q13268	Dehydrogenase/reductase SDR family member 2 mitochondrial	0.43	4
P07900	Heat shock protein HSP 90-alpha	0.43	3
P62249	40S ribosomal protein S16	0.42	1
A2NUT2	Lambda-chain (AA-20 to 215)	0.41	3
P60709	Actin cytoplasmic 1	0.40	8
P02671	Fibrinogen alpha chain	0.39	5
P19474	E3 ubiquitin-protein ligase TRIM21	0.38	6
P68371	Tubulin beta-4B chain	0.38	7
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.32	6
O60701	UDP-glucose 6-dehydrogenase	0.32	2
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.29	1
A1A508	PRSS3 protein	0.28	1
P08107	Heat shock 70 kDa protein 1A/1B	0.27	2
P07437	Tubulin beta chain	0.27	5
P02538	Keratin type II cytoskeletal 6A	0.26	2
P06733	Alpha-enolase	0.26	2
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.25	1
P08238	Heat shock protein HSP 90-beta	0.24	2
P05787	Keratin type II cytoskeletal 8	0.22	6
P35908	Keratin type II cytoskeletal 2 epidermal	0.21	8
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.21	3
P15924	Desmoplakin	0.20	2
H6VRG3	Keratin 1	0.20	9
Q9ULV0	Unconventional myosin-Vb	0.20	3
P31943	Heterogeneous nuclear ribonucleoprotein H	0.20	1
P02533	Keratin type I cytoskeletal 14	0.19	2
Q149N5	DOCK4 protein	0.19	3
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.19	1
P13645	Keratin type I cytoskeletal 10	0.16	6
P05783	Keratin type I cytoskeletal 18	0.15	2
Q00610	Clathrin heavy chain 1	0.15	4
Q15058	Kinesin-like protein KIF14	0.15	4
P11021	78 kDa glucose-regulated protein	0.13	3
P51659	Peroxisomal multifunctional enzyme type 2	0.12	3
P68363	Tubulin alpha-1B chain	0.11	1
P35527	Keratin type I cytoskeletal 9	0.08	2
P35580	Myosin-10	0.04	1