

Accession Number	Description	Fold Enrichment	UniquePeptideCount
A7MD48	Serinearginine repetitive matrix protein 4	2.38	1
O60667	Fas apoptotic inhibitory molecule 3	2.38	1
O96009	Napsin-A	2.38	1
P10124	Serglycin	2.38	1
Q567U6	Coiled-coil domain-containing protein 93	2.38	1
Q7Z398	Zinc finger protein 550	2.38	1
Q81YT1	Protein FAM71A	2.38	1
Q8NEF3	Coiled-coil domain-containing protein 112	2.38	1
Q8NG97	Olfactory receptor 22I	2.38	1
Q8NGZ3	Olfactory receptor 13G1	2.38	1
Q9BX95	Sphingosine-1-phosphate phosphatase 1	2.38	1
Q9UNP4	Lactosylceramide alpha-23-sialyltransferase	2.38	1
Q7Z7G8	Vacuolar protein sorting-associated protein 13B	2.04	1
Q8N755	PQ-loop repeat-containing protein 3	2.00	1
Q8WWW9	Selenoprotein M	2.00	1
P51798	HCl- exchange transporter 7	1.91	1
P13726	Tissue factor	1.83	1
P55774	C-C motif chemokine 18	1.71	2
Q6NZ52	Ribosomal protein L27a	1.68	1
O95211	Pituitary tumor transforming gene protein	1.61	1
Q92620	Pre-mRNA-splicing factor ATP-dependent RNA helicase PRP16	1.51	1
Q92935	Exostosin-like 1	1.40	1
Q9H5V8	CUB domain-containing protein 1	1.36	1
Q13936	Voltage-dependent L-type calcium channel subunit alpha-1C	1.34	1
P42356	Phosphatidylinositol 4-kinase alpha	1.28	1
Q58DX5	Inactive N-acetylated-alpha-linked acidic dipeptidase-like protein 2	1.15	1
P80748	Ig lambda chain V-III region LOI	1.15	4
P14091	Cathepsin E	1.09	1
P01769	Ig heavy chain V-III region GA	1.09	1
Q51ZF3	Ankyrin repeat domain-containing protein 45	1.02	1
A6NMU1	Putative olfactory receptor 52A4	1.01	1
Q8NEJ9	Neuroguidin	0.97	1
Q81EX9	Importin-4	0.96	1
Q99623	Prohibitin-2	0.93	5
Q5VTE0	Putative elongation factor 1-alpha-like 3	0.89	8
P02675	Fibrinogen beta chain	0.84	9
P02679	Fibrinogen gamma chain	0.77	13
P50225	Sulfotransferase 1A1	0.74	1
Q96A28	SLAM family member 9	0.68	1
B9A064	Immunoglobulin lambda-like polypeptide 5	0.67	2
P62266	40S ribosomal protein S23	0.66	1
Q14973	Sodiumbile acid cotransporter	0.66	1
Q9ULV0	Unconventional myosin-Vb	0.62	20
Q92734	Protein TFG	0.60	1
P52597	Heterogeneous nuclear ribonucleoprotein F	0.60	3
P68363	Tubulin alpha-1B chain	0.60	13
Q12771	P37 AUF1	0.59	1
P26641	Elongation factor 1-gamma	0.57	1
B2RWN5	HEAT repeat containing 1	0.54	1
Q9Y277	Voltage-dependent anion-selective channel protein 3	0.53	1
Q86TG7	Retrotransposon-derived protein PEG10	0.52	1
Q04828	Aldo-keto reductase family 1 member C1	0.52	1
P35030	Trypsin-3	0.50	1
Q13201	Multimerin-1	0.48	1
P06576	ATP synthase subunit beta mitochondrial	0.45	4
A2NUT2	Lambda-chain AA -20 to 215	0.45	5
P62829	60S ribosomal protein L23	0.41	1
P19338	Nucleolin	0.41	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.40	8
P00338	L-lactate dehydrogenase A chain	0.39	6
P68371	Tubulin beta-4B chain	0.39	10
Q06830	Peroxioredoxin-1	0.39	2
P31943	Heterogeneous nuclear ribonucleoprotein H	0.39	4
Q81ZT6	Abnormal spindle-like microcephaly-associated protein	0.38	1
P62987	Ubiquitin-60S ribosomal protein L40	0.36	1
P31483	Nucleolysin TIA-1 isoform p40	0.36	1
P07437	Tubulin beta chain	0.35	10
P02768	Serum albumin	0.34	9
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.33	4
Q86Y82	Syntaxin-12	0.33	1
P02671	Fibrinogen alpha chain	0.31	6
Q149N6	Dedicator of cytokinesis 4	0.31	8
P05141	ADP/ATP translocase 2	0.30	2
P01617	Ig kappa chain V-II region TEW	0.30	2
P01700	Ig lambda chain V-I region HA	0.30	2
P10809	60 kDa heat shock protein mitochondrial	0.29	3
P38646	Stress-70 protein mitochondrial	0.28	1
Q9BVG8	Kinesin-like protein KIFC3	0.27	3
P49411	Elongation factor Tu mitochondrial	0.27	2
P25705	ATP synthase subunit alpha mitochondrial	0.27	2
P04075	Fructose-bisphosphate aldolase A	0.26	1
P02769	Serum albumin	0.26	46
P05783	Keratin type I cytoskeletal 18	0.25	10
P54136	Arginine--tRNA ligase cytoplasmic	0.25	1
Q16822	Phosphoenolpyruvate carboxykinase [GTP] mitochondrial	0.25	1
Q4W4Y1	Dopamine receptor interacting protein 4	0.24	1
P05787	Keratin type II cytoskeletal 8	0.24	11
P07900	Heat shock protein HSP 90-alpha	0.24	2
Q15365	PolyrC-binding protein 1	0.21	1
P60842	Eukaryotic initiation factor 4A-I	0.21	1
P06748	Nucleophosmin	0.20	2
P08107	Heat shock 70 kDa protein 1A1B	0.20	2
P14625	Endoplasmic	0.19	1
P11021	78 kDa glucose-regulated protein	0.19	6
P14618	Pyruvate kinase PKM	0.18	1
P11142	Heat shock cognate 71 kDa protein	0.18	2
H6VRG3	Keratin 1	0.18	10
P22061	Protein-L-isoaspartateD-aspartate O-methyltransferase	0.16	3
P02533	Keratin type I cytoskeletal 14	0.15	2
O60701	UDP-glucose 6-dehydrogenase	0.14	1
O00571	ATP-dependent RNA helicase DDX3X	0.14	2
P51659	Peroxisomal multifunctional enzyme type 2	0.13	5
Q02538	Keratin type II cytoskeletal 6A	0.12	1
P06733	Alpha-enolase	0.12	1
P35527	Keratin type I cytoskeletal 9	0.12	5
P08238	Heat shock protein HSP 90-beta	0.11	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.09	4
P08727	Keratin type I cytoskeletal 19	0.09	1
P02751	Fibronectin	0.08	1
P15924	Desmoplakin	0.07	1
P63261	Actin cytoplasmic 2	0.07	1
Q18058	Kinesin-like protein KIF14	0.07	2
P13645	Keratin type I cytoskeletal 10	0.03	1