Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q7Z589	Protein EMSY	4.27	3
Q96QT6	PHD finger protein 12	3.18	2
B2R649	3-hydroxy-3-methylglutaryl coenzyme A reductase	2.09	1
B2ZZ86	Collagen type V alpha 1	2.09	1
O43916	Carbohydrate sulfotransferase 1	2.09	1
P00156	Cytochrome b	2.09	1
P20142	Gastricsin	2.09	1
P25025	C-X-C chemokine receptor type 2	2.09	1
P53992	Protein transport protein Sec24C	2.09	1
P62191	26S protease regulatory subunit 4	2.09	1
Q02224	Centromere-associated protein E	2.09	1
Q6PP77	XK-related protein 2	2.09	1
Q7Z3Z0	Keratin type I cytoskeletal 25	2.09	1
Q8IUR5	Transmembrane and TPR repeat-containing protein 1	2.09	1
Q8NFW8	N-acylneuraminate cytidylyltransferase	2.09	1
Q8TDS4	Hydroxycarboxylic acid receptor 2	2.09	1
Q8TF45	Zinc finger protein 418	2.09	1
Q8WUU5	**GATA zinc finger domain-containing protein 1**	**2.09**	**1**
Q92543	Sorting nexin-19	2.09	1
Q92545	Transmembrane protein 131	2.09	1
Q96FB5	Protein RRNAD1	2.09	1
Q96PU4	E3 ubiquitin-protein ligase UHRF2	2.09	1
Q9BVX2	Transmembrane protein 106C	2.09	1
Q9H160	Inhibitor of growth protein 2	2.09	1
Q9H5Q4	Dimethyladenosine transferase 2 mitochondrial	2.09	1
Q9NVH0	Exonuclease 3-5 domain-containing protein 2	2.09	1
Q9NY46	Sodium channel protein type 3 subunit alpha	2.09	1
Q9UPW8	Protein unc-13 homolog A	2.09	1
O75182	Paired amphipathic helix protein Sin3b	1.92	1
Q8NBN7	Retinol dehydrogenase 13	1.81	1
Q8ND30	Liprin-beta-2	1.76	1
P48047	ATP synthase subunit O mitochondrial	1.70	1
Q9BS86	Zona pellucida-binding protein 1	1.67	1
Q16280	Cyclic nucleotide-gated olfactory channel	1.64	1
A6N6J7	JARID1C protein	1.61	1
Q16659	Mitogen-activated protein kinase 6	1.60	1
Q13308	Inactive tyrosine-protein kinase 7	1.59	1
P09619	Platelet-derived growth factor receptor beta	1.58	1
O75354	Ectonucleoside triphosphate diphosphohydrolase 6	1.57	1
A6NKF7	Transmembrane protein 88B	1.55	1
Q14973	Sodiumbile acid cotransporter	1.51	1
Q96T52	Mitochondrial inner membrane protease subunit 2	1.51	1
Q9NXF1	Testis-expressed sequence 10 protein	1.35	1
O43592	Exportin-T	1.14	1
P32322	Pyrroline-5-carboxylate reductase 1 mitochondrial	1.10	1
Q8TD26	Chromodomain-helicase-DNA-binding protein 6	1.00	1
P17987	T-complex protein 1 subunit alpha	0.93	1
Q13155	Aminoacyl tRNA synthase complex-interacting multifunctional protein 2	0.82	1
P54136	Arginine--tRNA ligase cytoplasmic	0.75	2
P06748	Nucleophosmin	0.73	3
Q8WY98	Transmembrane protein 234	0.69	1
O15523	ATP-dependent RNA helicase DDX3Y	0.64	1
Q86Y82	Syntaxin-12	0.64	1
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.63	2
Q149N5	DOCK4 protein	0.62	6
Q92841	Probable ATP-dependent RNA helicase DDX17	0.61	1
Q15365	Poly(rC)-binding protein 1	0.56	1
P49327	Fatty acid synthase	0.55	1
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.51	1
A1A508	PRSS3 protein	0.49	1
P06576	ATP synthase subunit beta mitochondrial	0.49	1
A2NUT2	Lambda-chain (AA -20 to 215)	0.47	1
D3DPG0	Titin isoform CRA_a	0.45	1
P11142	Heat shock cognate 71 kDa protein	0.39	1
P07814	Bifunctional glutamateproline--tRNA ligase	0.39	2
P25705	ATP synthase subunit alpha mitochondrial	0.38	1
P60709	Actin cytoplasmic 1	0.37	2
P31943	Heterogeneous nuclear ribonucleoprotein H	0.35	1
P68363	Tubulin alpha-1B chain	0.35	2
O60701	UDP-glucose 6-dehydrogenase	0.35	1
P13645	Keratin type I cytoskeletal 10	0.34	5
P68104	Elongation factor 1-alpha 1	0.32	1
P08727	Keratin type I cytoskeletal 19	0.32	1
P68371	Tubulin beta-4B chain	0.29	2
P02769	Serum albumin	0.29	15
Q9ULV0	Unconventional myosin-Vb	0.28	2
P04264	Keratin type II cytoskeletal 1	0.28	5
P08238	Heat shock protein HSP 90-beta	0.27	1
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.26	2
P35527	Keratin type I cytoskeletal 9	0.24	3
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.23	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.22	3
P05783	Keratin type I cytoskeletal 18	0.17	1
P07437	Tubulin beta chain	0.16	1
P05787	Keratin type II cytoskeletal 8	0.14	1
P11021	78 kDa glucose-regulated protein	0.14	1
P43243	Matrin-3	0.14	1
P51659	Peroxisomal multifunctional enzyme type 2	0.12	1