Accession Number	Description	Fold Enrichment	UniquePeptideCount
P52732	Kinesin-like protein KIF11	9.35	10
P54132	Bloom syndrome protein	4.34	4
Q9BUZ4	TNF receptor-associated factor 4	3.48	9
Q53FZ2	Acyl-coenzyme A synthetase ACSM3 mitochondrial	2.26	2
A4D2A4	**Zinc finger protein 3 (A8-51)**	**1.84**	**1**
H2DF05	Interleukin 10 alpha receptor	1.84	1
H7BZ11	Protein RPL36A-HNRNPH2	1.84	1
K7EQ86	KIAA0100 isoform CRA_a	1.84	1
P01620	Ig kappa chain V-III region SIE	1.84	1
P09683	Secretin	1.84	1
P0C7X2	Zinc finger protein 688	1.84	1
P22695	Cytochrome b-c1 complex subunit 2 mitochondrial	1.84	1
P26641	Elongation factor 1-gamma	1.84	1
P40763	Signal transducer and activator of transcription 3	1.84	1
P55884	Eukaryotic translation initiation factor 3 subunit B	1.84	1
P63027	Vesicle-associated membrane protein 2	1.84	1
Q68DA7	Formin-1	1.84	1
Q8IUG5	Unconventional myosin-XVIIIb	1.84	1
Q8NHB8	Olfactory receptor 5K2	1.84	1
Q8TD16	Protein bicaudal D homolog 2	1.84	1
Q8TEV8	Smith-Magenis syndrome chromosomal region candidate gene 5 protein	1.84	1
Q8WU79	Stromal membrane-associated protein 2	1.84	1
Q969S0	UDP-xylose and UDP-N-acetylglucosamine transporter	1.84	1
Q96PK6	RNA-binding protein 14	1.84	1
Q9C0B7	Transport and Golgi organization protein 6 homolog	1.84	1
Q9HC07	Transmembrane protein 165	1.84	1
P35606	Coatomer subunit beta	1.69	1
Q08289	Voltage-dependent L-type calcium channel subunit beta-2	1.69	1
Q13145	BMP and activin membrane-bound inhibitor homolog	1.69	1
Q6VY07	Phosphofurin acidic cluster sorting protein 1	1.61	1
P50454	Serpin H1	1.59	1
Q9Y5Y4	Prostaglandin D2 receptor 2	1.57	1
P11171	Protein 41	1.55	1
P52292	Importin subunit alpha-1	1.55	1
P42574	Caspase-3	1.53	1
Q70CQ2	Ubiquitin carboxyl-terminal hydrolase 34	1.52	1
W8FP62	CBFA2T1 isoform r1t1-11a63	1.47	1
Q14157	Ubiquitin-associated protein 2-like	1.45	1
P21333	Filamin-A	1.44	1
Q7L4E1	Protein FAM73B	1.41	1
P55199	RNA polymerase II elongation factor ELL	1.40	1
Q92546	RAB6A-GEF complex partner protein 2	1.39	1
B7ZKS6	TRPM5 protein	1.37	1
Q6P3W7	SCY1-like protein 2	1.33	1
Q8WXX0	Dynein heavy chain 7 axonemal	1.29	1
Q9BUA6	Myosin regulatory light chain 10	1.29	1
Q14973	Sodiumbile acid cotransporter	1.25	1
Q96T52	Mitochondrial inner membrane protease subunit 2	1.25	1
Q93034	Cullin-5	1.24	1
P62633	Cellular nucleic acid-binding protein	1.14	1
Q9BQE3	Tubulin alpha-1C chain	1.10	1
P04792	Heat shock protein beta-1	1.07	1
P08670	Vimentin	1.06	2
P61201	COP9 signalosome complex subunit 2	0.99	1
Q99466	Neurogenic locus notch homolog protein 4	0.96	1
P06748	Nucleophosmin	0.94	5
P12004	Proliferating cell nuclear antigen	0.92	1
P62081	40S ribosomal protein S7	0.88	1
Q15058	Kinesin-like protein KIF14	0.86	18
Q6GYQ0	Ral GTPase-activating protein subunit alpha-1	0.85	1
Q08211	ATP-dependent RNA helicase A	0.80	1
B9A064	Immunoglobulin lambda-like polypeptide 5	0.79	1
P49368	T-complex protein 1 subunit gamma	0.77	1
P17844	Probable ATP-dependent RNA helicase DDX5	0.75	1
P01605	Ig kappa chain V-I region Lay	0.67	1
P08107	Heat shock 70 kDa protein 1A1B	0.66	3
P54136	Arginine--tRNA ligase cytoplasmic	0.63	2
P01709	Ig lambda chain V-II region MGC	0.62	1
P01818	Ig heavy chain V-II region HE	0.61	1
O15523	ATP-dependent RNA helicase DDX3Y	0.57	1
P08238	Heat shock protein HSP 90-beta	0.56	4
P68371	Tubulin beta-4B chain	0.48	5
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.45	1
P60842	Eukaryotic initiation factor 4A-I	0.44	1
A1A508	PRSS3 protein	0.43	1
P07814	Bifunctional glutamateproline--tRNA ligase	0.43	3
P06576	ATP synthase subunit beta mitochondrial	0.43	1
A2NUT2	Lambda-chain (AA -20 to 215)	0.42	1
D3DPG0	Titin isoform CRA_a	0.40	1
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.36	1
P07437	Tubulin beta chain	0.33	4
P31943	Heterogeneous nuclear ribonucleoprotein H	0.31	1
P68363	Tubulin alpha-1B chain	0.29	2
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.28	3
P68104	Elongation factor 1-alpha 1	0.28	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.27	2
P02769	Serum albumin	0.27	18
P35527	Keratin type I cytoskeletal 9	0.15	2
P05787	Keratin type II cytoskeletal 8	0.13	1
P04264	Keratin type II cytoskeletal 1	0.11	2
Q00610	Clathrin heavy chain 1	0.11	1