

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q9NUA8	Zinc finger and BTB domain-containing protein 40	14.94	17
P05455	Lupus La protein	1.82	1
P78504	Protein jagged-1	1.43	1
P15880	40S ribosomal protein S2	1.38	4
P60709	Actin cytoplasmic 1	1.20	27
Q96S66	Chloride channel CLIC-like protein 1	1.04	2
P08238	Heat shock protein HSP 90-beta	1.04	13
P36578	60S ribosomal protein L4	1.01	2
P19474	E3 ubiquitin-protein ligase TRIM21	1.00	19
O75477	Erlin-1	0.97	1
P17987	T-complex protein 1 subunit alpha	0.92	2
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.89	4
Q4W4Y1	Dopamine receptor interacting protein 4	0.83	3
O60814	Histone H2B type 1-K	0.83	1
P47914	60S ribosomal protein L29	0.82	2
O15523	ATP-dependent RNA helicase DDX3Y	0.81	4
P62805	Histone H4	0.81	2
Q15058	Kinesin-like protein KIF14	0.77	29
P23396	40S ribosomal protein S3	0.76	3
Q12931	Heat shock protein 75 kDa mitochondrial	0.75	3
P62249	40S ribosomal protein S16	0.75	3
A2NUT2	Lambda-chain (AA -20 to 215)	0.73	7
Q16777	Histone H2A type 2-C	0.70	1
P13639	Elongation factor 2	0.69	2
P07900	Heat shock protein HSP 90-alpha	0.67	6
P07437	Tubulin beta chain	0.64	15
P60842	Eukaryotic initiation factor 4A-I	0.63	3
P49368	T-complex protein 1 subunit gamma	0.62	1
O75533	Splicing factor 3B subunit 1	0.60	3
Q07954	Prolow-density lipoprotein receptor-related protein 1	0.57	1
Q76KP1	N-acetyl-beta-glucosaminyl-glycoprotein 4-beta-N-acetylgalactosaminyltransferase 1	0.53	1
P48643	T-complex protein 1 subunit epsilon	0.53	1
P62826	GTP-binding nuclear protein Ran	0.49	1
P68363	Tubulin alpha-1B chain	0.46	9
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.46	9
P51659	Peroxisomal multifunctional enzyme type 2	0.45	16
P06576	ATP synthase subunit beta mitochondrial	0.43	2
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.42	4
Q06830	Peroxiredoxin-1	0.40	1
Q5D862	Filaggrin-2	0.39	1
P04075	Fructose-bisphosphate aldolase A	0.33	1
P49411	Elongation factor Tu mitochondrial	0.33	2
P25705	ATP synthase subunit alpha mitochondrial	0.32	2
P02769	Serum albumin	0.30	50
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.30	1
P07814	Bifunctional glutamate/proline--tRNA ligase	0.27	4
P08107	Heat shock 70 kDa protein 1A/1B	0.25	2
P01709	Ig lambda chain V-II region MGC	0.25	1
P06733	Alpha-enolase	0.24	2
P35580	Myosin-10	0.21	10
P68104	Elongation factor 1-alpha 1	0.20	2
O60701	UDP-glucose 6-dehydrogenase	0.20	1
P13647	Keratin type II cytoskeletal 5	0.18	2
P11142	Heat shock cognate 71 kDa protein	0.15	1
P00338	L-lactate dehydrogenase A chain	0.13	1
P13645	Keratin type I cytoskeletal 10	0.13	5
P02533	Keratin type I cytoskeletal 14	0.12	1
P35527	Keratin type I cytoskeletal 9	0.12	4
P35908	Keratin type II cytoskeletal 2 epidermal	0.11	4
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.09	1
Q00610	Clathrin heavy chain 1	0.08	2
H6VRG3	Keratin 1	0.07	3