

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q06547	GA-binding protein subunit beta-1	5.20	3
P0C629	Olfactory receptor 10J4	2.40	1
Q597H1	Transformation-related protein 14	2.40	1
Q5T5P2	Sickle tail protein homolog	2.40	1
Q7L190	Developmental pluripotency-associated protein 4	2.21	1
Q9H2U1	ATP-dependent RNA helicase DHX36	2.10	1
Q5T5A8	Late cornified envelope protein 3C	1.71	1
Q8TDW7	Protocadherin Fat 3	1.63	1
P11474	Steroid hormone receptor ERR1	1.34	1
H7BY58	Protein-L-isoaspartate O-methyltransferase	1.33	7
P01624	Ig kappa chain V-III region POM	1.23	1
Q6GYQ0	Ral GTPase-activating protein subunit alpha-1	1.22	1
Q13155	Aminoacyl tRNA synthase complex-interacting multifunctional protein 2	0.96	1
O94776	Metastasis-associated protein MTA2	0.92	1
Q16822	Phosphoenolpyruvate carboxykinase [GTP] mitochondrial	0.90	1
P23396	40S ribosomal protein S3	0.80	1
Q86Y82	Syntaxin-12	0.74	1
P07437	Tubulin beta chain	0.72	6
P02656	Apolipoprotein C-III	0.66	1
P13645	Keratin type I cytoskeletal 10	0.65	8
P68104	Elongation factor 1-alpha 1	0.61	2
O15090	Zinc finger protein 536	0.59	1
P68363	Tubulin alpha-1B chain	0.57	3
Q149N6	Dedicator of cytokinesis 4	0.55	4
A2NUT2	Lambda-chain (AA -20 to 215)	0.54	1
P49411	Elongation factor Tu mitochondrial	0.53	1
P05783	Keratin type I cytoskeletal 18	0.53	4
P04264	Keratin type II cytoskeletal 1	0.52	8
P19474	E3 ubiquitin-protein ligase TRIM21	0.52	3
P08238	Heat shock protein HSP 90-beta	0.49	2
P25705	ATP synthase subunit alpha mitochondrial	0.44	1
P60709	Actin cytoplasmic 1	0.44	2
P02769	Serum albumin	0.41	17
P35908	Keratin type II cytoskeletal 2 epidermal	0.41	5
O60701	UDP-glucose 6-dehydrogenase	0.40	1
P10809	60 kDa heat shock protein mitochondrial	0.39	1
P07814	Bifunctional glutamateproline--tRNA ligase	0.29	1
P11021	78 kDa glucose-regulated protein	0.26	2
P35527	Keratin type I cytoskeletal 9	0.22	2
P51659	Peroxisomal multifunctional enzyme type 2	0.21	2