

Accession Number	Description	Fold Enrichment	UniquePeptideCount
<b>P14316</b>	<b>Interferon regulatory factor 2</b>	<b>3.74</b>	<b>2</b>
P21462	fMet-Leu-Phe receptor	1.07	1
Q9UIF8	Bromodomain adjacent to zinc finger domain protein 2B	0.98	1
Q8WWZ4	ATP-binding cassette sub-family A member 10	0.98	1
Q9NZR2	Low-density lipoprotein receptor-related protein 1B	0.93	1
P54136	Arginine--tRNA ligase cytoplasmic	0.92	2
P31943	Heterogeneous nuclear ribonucleoprotein H	0.86	3
P07814	Bifunctional glutamateproline--tRNA ligase	0.84	4
P01709	Ig lambda chain V-II region MGC	0.80	1
P68104	Elongation factor 1-alpha 1	0.79	3
P02675	Fibrinogen beta chain	0.74	1
H7BY58	Protein-L-isoaspartate O-methyltransferase	0.71	4
P14618	Pyruvate kinase PKM	0.68	2
Q14847	LIM and SH3 domain protein 1	0.67	1
Q149N6	Dedicator of cytokinesis 4	0.66	5
O60701	UDP-glucose 6-dehydrogenase	0.62	2
Q02878	60S ribosomal protein L6	0.61	1
P07437	Tubulin beta chain	0.60	5
P60709	Actin cytoplasmic 1	0.59	3
P02769	Serum albumin	0.58	25
P49411	Elongation factor Tu mitochondrial	0.52	1
P04264	Keratin type II cytoskeletal 1	0.51	8
P08238	Heat shock protein HSP 90-beta	0.49	2
P25705	ATP synthase subunit alpha mitochondrial	0.43	1
P05783	Keratin type I cytoskeletal 18	0.41	3
P68363	Tubulin alpha-1B chain	0.41	2
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.36	1
P11021	78 kDa glucose-regulated protein	0.35	3
P68371	Tubulin beta-4B chain	0.35	2
P13645	Keratin type I cytoskeletal 10	0.34	4
Q86YZ3	Hornerin	0.26	1
P05787	Keratin type II cytoskeletal 8	0.26	2
P19474	E3 ubiquitin-protein ligase TRIM21	0.24	1
P35527	Keratin type I cytoskeletal 9	0.21	2
P51659	Peroxisomal multifunctional enzyme type 2	0.21	2
Q00610	Clathrin heavy chain 1	0.14	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.12	1