

Accession Number	Description	Fold Enrichment	UniquePeptideCount
<b>O43829</b>	<b>Zinc finger and BTB domain-containing protein 14</b>	<b>8.21</b>	<b>15</b>
Q75319	RNAi-PCR complex-1-interacting phosphatase	1.43	1
P15927	Replication protein A 32 kDa subunit	1.31	2
Q96PK6	RNA-binding protein 14	1.16	5
Q14627	Interleukin-13 receptor subunit alpha-2	1.13	1
P28907	ADP-ribosyl cyclase/cyclic ADP-ribose hydrolase 1	1.13	2
Q96G43	Protein LTV1 homolog	1.07	1
Q9H3K6	BoiA-like protein 2	1.04	2
Q95789	Zinc finger MYM-type protein 6	1.01	1
Q9B1V4	Transmembrane protein 43	1.00	1
Q96S38	Cell growth-inhibiting protein 34	0.98	1
Q9P291	Armadillo repeat-containing X-linked protein 1	0.95	1
Q8IWA5	Choline transporter-like protein 2	0.92	1
Q9NUL3	Double-stranded RNA-binding protein Staufin homolog 2	0.92	3
Q02841	Voltage-dependent L-type calcium channel subunit beta-1	0.91	1
QRUKT9	Alysdolase domain-containing protein 15	0.91	1
P39656	Dolichyl-diphosphooligosaccharide-protein glycosyltransferase 48 kDa subunit	0.90	1
Q8NEQ5	Transmembrane protein C1orf162	0.88	1
Q9Y5L3	Ectonucleoside triphosphate diphosphohydrolase 2	0.88	1
P62829	60S ribosomal protein L23	0.88	4
P18621	60S ribosomal protein L17	0.87	7
Q5VTE0	Putative elongation factor 1-alpha-like 3	0.85	14
P52597	Heterogeneous nuclear ribonucleoprotein F	0.84	6
P42877	40S ribosomal protein S27	0.82	2
Q99823	Prohibitin-2	0.82	6
Q13200	26S proteasome non-ATPase regulatory subunit 2	0.81	1
P61204	ADP-ribosylation factor 3	0.81	2
Q701N2	Keratin-associated protein 5-5	0.81	1
A8K122	KIAA1946 isoform CRA_b	0.80	1
Q5TCH4	Cytochrome P450 4A22	0.79	2
Q9Y277	Voltage-dependent anion-selective channel protein 3	0.79	1
Q8NH64	Olfactory receptor 51A7	0.79	1
P14868	Aspartate--tRNA ligase cytoplasmic	0.78	4
Q9H9L7	Isthmin-2	0.77	2
Q13609	Tubulin beta-3 chain	0.76	15
Q16552	Interleukin-17A	0.75	1
Q07912	Activated CDC42 kinase 1	0.75	1
Q86W48	Lon protease homolog 2, peroxisomal	0.75	1
PA8643	T-complex protein 1 subunit epsilon	0.72	4
Q9HCC0	Methylcrotonoyl-CoA carboxylase beta chain mitochondrial	0.72	1
Q6GMR7	Fatty acid amide hydrolase 2	0.71	1
Q5TCS8	Adenylyl kinase 9	0.71	1
Q9H479	Fructosamine-3-kinase	0.71	2
P19429	Troponin 1 cardiac muscle	0.70	1
D4YW74	Polytrophin	0.70	1
P61247	40S ribosomal protein S3a	0.69	2
P35606	Coatomer subunit beta	0.69	2
Q8WUM4	Programmed cell death 6-interacting protein	0.69	3
P35537	RNA-binding protein FL5	0.68	2
P60842	Eukaryotic initiation factor 4A-1	0.68	7
Q8IYW2	Cilia- and flagella-associated protein 46	0.67	1
Q5QIP1	CCR5 chemokine receptor	0.67	1
Q75477	Erlin-1	0.66	1
Q9Y5V3	Melanoma-associated antigen D1	0.65	2
Q9HBT6	Cadherin-20	0.65	1
P23396	40S ribosomal protein S3	0.65	5
P50991	T-complex protein 1 subunit delta	0.63	2
P07437	Tubulin beta chain	0.63	30
P05388	60S acidic ribosomal protein P0	0.63	2
K9N2Q6	Interleukin 15 receptor alpha isoform IC3	0.62	1
Q9ULQ0	Striatin-interacting protein 2	0.61	1
Q00571	ATP-dependent RNA helicase DDX3X	0.60	10
P58371	Tubulin beta-4b chain	0.59	27
Q95793	Double-stranded RNA-binding protein Staufin homolog 1	0.58	3
P50454	Serpin H1	0.57	1
Q15436	Protein transport protein Sec23A	0.56	1
P35232	Prohibitin	0.55	1
P49368	T-complex protein 1 subunit gamma	0.55	2
P62266	40S ribosomal protein S23	0.55	1
P62081	40S ribosomal protein S7	0.54	2
Q75369	Filamin-B	0.53	1
Q16847	Prostacyclin synthase	0.53	2
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.52	4
Q13838	Spliceosome RNA helicase DDX39B	0.52	1
P17987	T-complex protein 1 subunit alpha	0.52	2
P53261	Actin cytoplasmic 2	0.50	18
Q02679	Fibrinogen gamma chain	0.49	10
P68363	Tubulin alpha-1B chain	0.47	19
Q14847	LIM and SH3 domain protein 1	0.47	4
P31943	Heterogeneous nuclear ribonucleoprotein H	0.47	9
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.46	2
P62633	Cellular nucleic acid-binding protein	0.46	1
P61978	Heterogeneous nuclear ribonucleoprotein K	0.45	3
P05141	ADP/ATP translocase 2	0.45	6
P54136	Arginine--tRNA ligase cytoplasmic	0.45	4
P68238	Heat shock protein HSP 90-beta	0.44	11
P62826	GTP-binding nuclear protein Ran	0.44	2
P14618	Pyruvate kinase PKM	0.43	6
Q5T1M5	FK506-binding protein 15	0.43	1
Q06830	Peroxisome-1	0.43	3
P51659	Peroxisomal multifunctional enzyme type 2	0.41	30
P22061	Protein-L-isoaspartate(D-aspartate) O-methyltransferase	0.41	16
P47897	Glutamine--tRNA ligase	0.41	1
PA9411	Elongation factor Tu mitochondrial	0.41	6
P46776	60S ribosomal protein L27a	0.40	2
P06576	ATP synthase subunit beta mitochondrial	0.39	5
Q92841	Probable ATP-dependent RNA helicase DDX17	0.39	2
Q15365	Poly(rC)-binding protein 1	0.39	4
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.38	8
P12004	Proliferating cell nuclear antigen	0.38	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.36	13
P17844	Probable ATP-dependent RNA helicase DDX5	0.36	1
P45880	Voltage-dependent anion-selective channel protein 2	0.36	1
Q36796	Single-stranded DNA-binding protein	0.35	1
P08727	Keratin type I cytoskeletal 19	0.34	8
P05783	Keratin type I cytoskeletal 18	0.34	14
P36542	ATP synthase subunit gamma mitochondrial	0.34	1
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.34	1
P05167	Keratin type II cytoskeletal 9	0.33	22
Q9NZI8	Insulin-like growth factor 2 mRNA-binding protein 1	0.33	2
Q9BVG8	Kinesin-like protein KIFC3	0.33	2
P62249	40S ribosomal protein S16	0.32	2
P13639	Elongation factor 2	0.30	1
Q16822	Phosphoenolpyruvate carboxykinase [GTP] mitochondrial	0.30	1
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.30	13
P62701	40S ribosomal protein S4 X isoform	0.29	1
Q02675	Fibrinogen beta chain	0.28	1
P02768	Serum albumin	0.26	12
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.26	2
P38646	Stress-70 protein mitochondrial	0.25	1
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.24	4
P14625	Endoplasmic	0.23	2
P02671	Fibrinogen alpha chain	0.23	6
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.23	1
Q60701	UDP-glucose 6-dehydrogenase	0.21	3
P25705	ATP synthase subunit alpha mitochondrial	0.20	2
Q15058	Kinesin-like protein KIF14	0.20	14
P01617	Ig kappa chain V-II region TEW	0.20	1
P08107	Heat shock 70 kDa protein 1A1B	0.20	3
Q149N6	Dedicator of cytokinesis 4	0.18	8
P49327	Fatty acid synthase	0.18	1
Q00639	Heterogeneous nuclear ribonucleoprotein U	0.18	3
A2NUT2	Lambda-chain (AA-20 to 215)	0.18	2
P10809	60 kDa heat shock protein mitochondrial	0.16	2
P11021	78 kDa glucose-regulated protein	0.16	9
P07814	Bifunctional glutamylproline--tRNA ligase	0.16	4
Q02769	Serum albumin	0.15	50
P06733	Alpha-enolase	0.15	2
P15924	Desmoplakin	0.13	4
P35908	Keratin type II cytoskeletal 2 epidermal	0.13	11
Q04264	Keratin type II cytoskeletal 1	0.11	10
P13645	Keratin type I cytoskeletal 10	0.07	5
P35527	Keratin type I cytoskeletal 9	0.07	4
Q00610	Clathrin heavy chain 1	0.04	1