

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q21810	Zinc finger protein 160	4.36	2
Q9NVL6	RING finger protein 166	4.54	2
P09028	Malate dehydrogenase mitochondrial	1.46	1
Q9NVL3	Double-strand RNA-binding protein Staufen homolog 2	1.40	6
O95793	Double-strand RNA-binding protein Staufen homolog 1	1.29	8
E1P552	RNA-binding region (RNP1 RRM) containing 2 isoform CRA b	1.24	3
P17897	T-complex protein 1 subunit alpha	1.21	7
Q9Y581	Insulin-like peptide INSL6	1.18	2
Q9NH64	Olfactory receptor 51A7	1.11	2
Q9L210	Phb-interacting protein INBP-36	1.10	3
Q9L211	Phosphatase and actin regulator 4	1.08	2
O14647	Chromodomain-helicase-DNA-binding protein 2	1.08	1
Q9HFA3	Transmembrane protein 181A	1.03	2
Q9H3K6	BoLA-like protein 2	1.01	2
Q9BTV4	Transmembrane protein 43	0.99	1
P61174	Inosophosphate isomerase	0.97	1
P83731	60S ribosomal protein L24	0.97	2
P16150	Leukosialin	0.95	1
Q9RE38	Cell growth-inhibiting protein 34	0.95	1
P30837	Aldehyde dehydrogenase X mitochondrial	0.93	3
COJY72	Apolipoprotein B (including Aq(X) antigen)	0.92	2
Q9BRK6	RNA-binding protein 14	0.92	4
Q15436	Protein transport protein Sec23A	0.91	3
Q9WVA5	Choline transporter-like protein 2	0.91	1
Q14003	Potassium voltage-gated channel subfamily C member 3	0.88	1
P32269	60S ribosomal protein L9	0.87	2
ABNCG0	Maestro heat-like repeat-containing protein family member 6	0.84	1
Q9Y5V9	Sodium channel protein type 10 subunit alpha	0.83	3
P41252	Isoleucine--RNA ligase cytoplasmic	0.82	4
P95102	Methionine--RNA ligase cytoplasmic	0.82	3
Q13200	ZRS proteinase non-A1Pase regulatory subunit 2	0.80	3
Q701N2	Keratin-associated protein 5-5	0.80	1
P61976	Heterogeneous nuclear ribonucleoprotein K	0.79	7
P45564	Short-branched chain specific acyl-CoA dehydrogenase mitochondrial	0.79	3
P32528	60S ribosomal protein L22	0.78	1
Q9H1C5	Glycylglycyltransferase 8 domain-containing protein 2	0.77	2
P14959	Aspartate--RNA ligase cytoplasmic	0.76	4
P98164	Low-density lipoprotein receptor-related protein 2	0.75	1
Q04828	Aldehyde reductase family 1 member 1	0.73	7
O00571	ATP-dependent RNA helicase DDX3X	0.73	13
P27694	Replication protein A 70 kDa DNA-binding subunit	0.72	3
P92699	60S ribosomal protein L23	0.70	2
Q9BTCS	Metastasis-associated protein MTA3	0.70	2
Q9H479	Fructosamine-3-kinase	0.69	2
P55768	Formylglycine aminopeptidase	0.69	2
P19429	Troponin I cardiac muscle	0.69	1
P30101	Protein disulfide-isomerase A3	0.69	1
Q4YV74	Redoxin	0.69	1
P11586	C-1-tetrahydrofolate synthase cytoplasmic	0.68	1
P35637	RNA-binding protein FUS	0.67	2
Q9WUJ4	Programmed cell death-interacting protein	0.67	2
Q7L014	Probable ATP-dependent RNA helicase DDX46	0.67	2
P32119	Peroxiredoxin-2	0.66	2
Q9P202	Whirrin	0.65	2
J3QR09	Ribosomal protein L19	0.65	1
P09558	Phosphoglycerate kinase 1	0.63	1
P52507	Heterogeneous nuclear ribonucleoprotein F	0.62	4
P05388	60S acidic ribosomal protein P0	0.62	2
B4E0N9	Glutamate dehydrogenase	0.62	2
P11943	Heterogeneous nuclear ribonucleoprotein H	0.61	13
P62277	40S ribosomal protein S13	0.61	1
Q9JLQ0	Strain-interacting protein 2	0.60	1
P07747	Protein 1	0.60	2
P40227	T-complex protein 1 subunit zeta	0.60	1
Q00425	Insulin-like growth factor 2 mRNA-binding protein 3	0.60	1
P98311	Tubulin beta-8 chain	0.59	28
P23528	Cofilin-1	0.59	1
Q13266	Dehydrogenasereductase SDR family member 2 mitochondrial	0.58	14
P19338	Nuclein	0.57	3
Q9A2Z8	SLAM family member 9	0.57	1
P5044	Septin H1	0.56	1
Q9Y159	Zinc finger protein-like 1	0.56	1
Q9Y1M5	FK506-binding protein 15	0.56	2
P09651	Heterogeneous nuclear ribonucleoprotein A1	0.54	2
Q9B1G7	Retrotansposon-derived protein PEG10	0.53	1
Q06520	Bile salt sulfotransferase	0.52	2
Q13838	Salt-inducible RNA helicase DDX398	0.52	1
P15024	Desmoplakin	0.52	22
P51571	Translucan-associated protein subunit delta	0.51	1
P53261	Actin cytochrome 2	0.51	19
Q3LXA3	Bifunctional ATP-dependent dihydroxyacetone kinase/FAD-AMP lyase (cyclizing)	0.50	2
P07437	Tubulin beta chain	0.49	24
Q22995	Cytochrome b-c1 complex subunit 2 mitochondrial	0.49	1
Q9NE13	Transmembrane protein 145	0.49	1
Q9Y5V3	Melanoma-associated antigen D1	0.49	1
P16822	Phosphoenolpyruvate carboxylase (C1P) mitochondrial	0.49	1
Q9N2I8	Insulin-like growth factor 2 mRNA-binding protein 1	0.49	4
Q9BVJ8	Kinesin-like protein KIF-C3	0.48	4
P45990	Voltage-dependent anion-selective channel protein 2	0.46	2
Q9N6C5	Immunoglobulin superfamily member 1	0.46	1
P62701	40S ribosomal protein S4 X isoform	0.46	3
P11909	80 kDa heat shock protein mitochondrial	0.46	10
Q9WVW5	Choline transporter-like protein 1	0.46	1
Q9BEY1	DNA J homolog subfamily A member 3 mitochondrial	0.44	2
P93727	Keratin type I cytoskeletal 19	0.43	11
O60337	E3 ubiquitin-protein ligase MARCH6	0.43	1
P14818	Pyruvate kinase PKM	0.42	6
Q98530	Peroxidase-1	0.42	2
P47897	Glutamine--RNA ligase	0.41	1
Q96263	Troponin-2	0.40	2
Q98211	ATP-dependent RNA helicase A	0.40	1
Q9P2E9	Ribosome-binding protein 1	0.40	1
P36646	Slitase-70 protein mitochondrial	0.40	1
Q13151	Heterogeneous nuclear ribonucleoprotein A0	0.39	8
P05763	Keratin type I cytoskeletal 18	0.39	17
P46776	60S ribosomal protein L2/a	0.39	2
P27635	60S ribosomal protein L10	0.39	2
P14625	Endoplasmic	0.39	5
P18621	60S ribosomal protein L17	0.39	2
D3DPG0	Titin isoform CRA_a	0.38	3
Q12931	Heat shock protein 75 kDa mitochondrial	0.38	2
P48643	T-complex protein 1 subunit epsilon	0.38	3
AT1441	G protein-coupled receptor 128	0.36	1
Q9VTE0	Putative elongation factor 1-alpha-like 3	0.36	5
P98363	Tubulin alpha-1B chain	0.35	14
Q07955	Serinearginine-rich splicing factor 1	0.35	1
P15880	40S ribosomal protein S2	0.35	1
Q95765	Single-strand DNA-binding protein	0.34	1
P05787	Keratin type II cytoskeletal 8	0.34	23
P05141	ADP/ATP translocase 2	0.33	4
Q15094	Protein disulfide-isomerase A6	0.33	3
P08238	Heat shock protein HSP 90-beta	0.33	8
P07814	Bifunctional glutathione--RNA ligase	0.33	11
P92926	GTP-binding nuclear protein Ran	0.33	3
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.32	7
P62905	Histone H4	0.32	1
P04076	Fructose-bisphosphate aldolase A	0.30	2
P13639	Elongation factor 2	0.30	1
P60842	Eukaryotic initiation factor 4A-1	0.30	2
P95227	Fatty acid synthase	0.30	3
P25705	ATP synthase subunit alpha mitochondrial	0.30	4
P07237	Protein disulfide-isomerase	0.29	1
P19174	E3 ubiquitin-protein ligase TRIM21	0.29	10
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.26	2
P94886	Delta-L-pyrimidine-5-carboxylate synthase	0.26	2
Q15365	POW1-binding protein 1	0.25	2
P04406	Glycerolaldehyde-3-phosphate dehydrogenase	0.25	2
P92671	Fibrinogen alpha chain	0.25	7
P51659	Peroxisomal multifunctional enzyme type 2	0.24	17
Q14847	LIM and SH3 domain protein 1	0.24	1
Q14946	Deficator of cytokinesis 4	0.23	11
Q94776	Metastasis-associated protein MTA2	0.23	1
P22061	Protein-L-isoaspartate (D-aspartate) O-methyltransferase	0.22	8
P41136	Arginine--RNA ligase cytoplasmic	0.22	8
P06576	ATP synthase subunit beta mitochondrial	0.22	2
Q13330	Metastasis-associated protein MTA1	0.21	2
Q9JLQ0	Lincosaminidase myosin-1b	0.21	6
P02679	Fibrinogen gamma chain	0.20	3
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.19	8
P08107	Heat shock 70 kDa protein H1A18	0.18	1
P07900	Heat shock protein HSP 90-alpha	0.18	2
A2NUJ2	Lambda-chain (AA-20 to 215)	0.18	2
Q60701	UDP-glucose 6-dehydrogenase	0.17	2
P02769	Serum albumin	0.17	57
Q15058	Kinesin-like protein KIF 14	0.16	11
P11021	FS kDa glucose-regulated protein	0.16	9
P49411	Elongation factor Tu mitochondrial	0.15	1
P06748	Nucleophosmin	0.15	2
P05153	Alpha-amylase	0.15	2
Q5BKZ1	DBIRD complex subunit ZNF 326	0.15	1
P02768	Serum albumin	0.14	6
P11442	Heat shock cognate 71 kDa protein	0.14	2
P52272	Heterogeneous nuclear ribonucleoprotein M	0.13	1
P03038	L-lactate dehydrogenase A chain	0.12	2
P13647	Keratin type I cytoskeletal 5	0.11	2
Q92769	Histone deacetylase 2	0.11	2
P13645	Keratin type I cytoskeletal 10	0.07	5
P129N17	ATPase family AAA-domain-containing protein 3A	0.06	8
Q04264	Keratin type II cytoskeletal 1	0.06	5
Q00610	Clastrin heavy chain 1	0.06	2
P35527	Keratin type I cytoskeletal 9	0.04	1
P35508	Keratin type II cytoskeletal 2 epidermal	0.03	1