

Accession Number	Description	Fold Enrichment	UniquePeptideCount
P36873			
P68032	Actin alpha cardiac muscle 1	10.81	37
P61158	Actin-related protein 3	4.31	8
P59998	Actin-related protein 23 complex subunit 4	3.44	5
P61160	Actin-related protein 2	2.42	5
P09327	Villin-1	1.95	26
Q43707	Alpha-actinin-4	1.90	32
P12814	Alpha-actinin-1	1.83	20
Q15144	Actin-related protein 23 complex subunit 2	1.74	4
<b>Q14978</b>	<b>Zinc finger protein 263</b>	<b>1.49</b>	<b>1</b>
P35579	Myosin-9	1.37	44
Q15143	Actin-related protein 23 complex subunit 1B	1.31	1
Q15145	Actin-related protein 23 complex subunit 3	1.31	1
P36873	Serine/threonine-protein phosphatase PP1-gamma catalytic subunit	1.20	1
Q13724	Mannosyl-oligosaccharide glucosidase	1.16	1
Q0IIN1	Keratin 77	1.07	1
Q3B7A7	Phosphoribosylglycinamide formyltransferase phosphoribosylglycinamide synthetase	1.05	1
D3DR22	Hydroxysteroid (17-beta) dehydrogenase 12 isoform CRA_a	0.98	1
O15050	TPR and ankyrin repeat-containing protein 1	0.95	1
Q01082	Spectrin beta chain non-erythrocytic 1	0.93	6
P60709	Actin cytoplasmic 1	0.89	50
P32969	60S ribosomal protein L9	0.89	2
Q75477	Erlin-1	0.89	2
Q40763	Signal transducer and activator of transcription 3	0.88	1
P49327	Fatty acid synthase	0.85	12
P40227	T-complex protein 1 subunit zeta	0.83	2
O00410	Importin-5	0.77	1
P62241	40S ribosomal protein S8	0.75	1
P48735	Isocitrate dehydrogenase [NADP] mitochondrial	0.75	1
Q9ULV0	Unconventional myosin-Vb	0.74	32
Q00325	Phosphate carrier protein mitochondrial	0.71	1
P55786	Purromycin-sensitive aminopeptidase	0.70	1
Q15393	Splicing factor 3B subunit 3	0.70	1
P61247	40S ribosomal protein S3a	0.70	2
P0CG38	POTE ankyrin domain family member 1	0.68	6
P46779	60S ribosomal protein L28	0.67	1
P62273	40S ribosomal protein S29	0.66	2
P04843	Dolichyl-diphosphooligosaccharide--protein glycosyltransferase subunit 1	0.64	1
P61626	Lysozyme C	0.63	1
P26373	60S ribosomal protein L13	0.62	1
P53618	Coatomer subunit beta	0.61	1
P50225	Sulfotransferase 1A1	0.60	1
P01859	Ig gamma-2 chain C region	0.56	2
P35232	Prohibitin	0.56	1
P62266	40S ribosomal protein S23	0.55	1
P33993	DNA replication licensing factor MCM7	0.55	1
Q13509	Tubulin beta-3 chain	0.54	10
P62805	Histone H4	0.53	3
Q16647	Prostacyclin synthase	0.53	1
P06576	ATP synthase subunit beta mitochondrial	0.51	7
P62244	40S ribosomal protein S15a	0.51	1
Q13813	Spectrin alpha chain non-erythrocytic 1	0.50	1
P56192	Methionine--tRNA ligase cytoplasmic	0.50	1
P27635	60S ribosomal protein L10	0.50	3
P45954	Shortbranched chain specific acyl-CoA dehydrogenase mitochondrial	0.49	1
Q43795	Unconventional myosin-1b	0.49	1
P48643	T-complex protein 1 subunit epsilon	0.49	2
P61353	60S ribosomal protein L27	0.48	1
P60748	Ig lambda chain V-III region LOI	0.46	2
Q6IPI7	RPL14 protein	0.46	1
P62917	60S ribosomal protein L8	0.45	1
P62826	GTP-binding nuclear protein Ran	0.44	2
P18085	ADP-ribosylation factor 4	0.43	1
Q08211	ATP-dependent RNA helicase A	0.41	1
P46776	60S ribosomal protein L27a	0.40	2
P17987	T-complex protein 1 subunit alpha	0.39	1
Q3LXA3	Bifunctional ATP-dependent dihydroxyacetone kinaseFAD-AMP lyase (cyclizing)	0.38	1
P07437	Tubulin beta chain	0.38	17
P78527	DNA-dependent protein kinase catalytic subunit	0.38	2
AZNU72	Lambda-chain (AA-20 to 215)	0.36	6
P68371	Tubulin beta-4B chain	0.35	15
P15880	40S ribosomal protein S2	0.35	1
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.35	1
P62913	60S ribosomal protein L11	0.34	1
P68363	Tubulin alpha-1B chain	0.34	13
P36542	ATP synthase subunit gamma mitochondrial	0.34	1
P51659	Peroxisomal multifunctional enzyme type 2	0.32	22
Q99623	Prohibitin-2	0.31	1
P49411	Elongation factor Tu mitochondrial	0.31	4
P08238	Heat shock protein HSP 90-beta	0.31	7
P25705	ATP synthase subunit alpha mitochondrial	0.31	4
P62263	40S ribosomal protein S14	0.30	1
P54136	Arginine--tRNA ligase cytoplasmic	0.30	2
Q4W4Y1	Dopamine receptor interacting protein 4	0.30	1
P18621	60S ribosomal protein L17	0.30	1
Q79533	Splicing factor 3B subunit 1	0.30	2
P62701	40S ribosomal protein S4 X isoform	0.29	1
P18124	60S ribosomal protein L7	0.28	1
P02656	Apolipoprotein C-III	0.28	2
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.27	2
P01617	Ig kappa chain V-II region TEW	0.27	2
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.27	1
Q15365	Poly(rC)-binding protein 1	0.26	2
O60701	UDP-glucose 6-dehydrogenase	0.26	4
P31943	Heterogeneous nuclear ribonucleoprotein H	0.26	4
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.26	5
P68104	Elongation factor 1-alpha 1	0.26	6
Q15084	Protein disulfide-isomerase A6	0.26	1
Q00571	ATP-dependent RNA helicase DDX3X	0.26	3
Q9NZI8	Insulin-like growth factor 2 mRNA-binding protein 1	0.25	1
P38646	Stress-70 protein mitochondrial	0.25	1
Q02543	60S ribosomal protein L18a	0.25	2
Q9Y4L1	Hypoxia up-regulated protein 1	0.25	1
P10809	60 kDa heat shock protein mitochondrial	0.24	4
Q14847	LIM and SH3 domain protein 1	0.24	1
P01700	Ig lambda chain V-I region HA	0.23	2
P35580	Myosin-10	0.21	21
P11142	Heat shock cognate 71 kDa protein	0.17	3
P08727	Keratin type I cytoskeletal 19	0.17	3
P05141	ADP-ATP translocase 2	0.17	1
P02671	Fibrinogen alpha chain n	0.17	4
P02768	Serum albumin	0.17	7
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.15	2
Q02769	Serum albumin	0.15	49
Q15058	Kinesin-like protein KIF14	0.14	9
H6VRC3	Keratin 1	0.13	13
P05783	Keratin type I cytoskeletal 18	0.13	4
P06748	Nucleophosmin	0.12	1
P05787	Keratin type II cytoskeletal 8	0.11	6
P19474	E3 ubiquitin-protein ligase TRIM21	0.10	2
P22061	Protein-L-isoaspartate(D-aspartate) O-methyltransferase	0.09	2
P02533	Keratin type I cytoskeletal 14	0.09	1
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1 n	0.08	2
P35908	Keratin type II cytoskeletal 2 epidermal	0.07	5
P35527	Keratin type I cytoskeletal 9	0.07	4
P13645	Keratin type I cytoskeletal 10	0.06	4
P11021	78 kDa glucose-regulated protein	0.04	1