

Accession Number	Description	Fold Enrichment	UniquePeptideCount
Q96F45	Zinc finger protein 503	4.19	3
P62854	40S ribosomal protein S26	1.30	2
P14868	Aspartate--tRNA ligase cytoplasmic	1.23	3
P30041	Peroxioredoxin-6	1.20	2
P13639	Elongation factor 2	1.15	4
P62249	40S ribosomal protein S16	1.13	5
P62273	40S ribosomal protein S29	1.12	2
P11586	C-1-tetrahydrofolate synthase cytoplasmic	0.99	1
Q13907	Isopentenyl-diphosphate Delta-isomerase 1	0.95	1
P46779	60S ribosomal protein L28	0.94	1
P62847	40S ribosomal protein S24	0.93	1
P32969	60S ribosomal protein L9	0.93	1
P61626	Lysozyme C	0.91	1
P27635	60S ribosomal protein L10	0.88	3
P62081	40S ribosomal protein S7	0.86	2
P62987	Ubiquitin-60S ribosomal protein L40	0.86	2
Q1KMD3	Heterogeneous nuclear ribonucleoprotein U-like protein 2	0.80	4
Q06210	n=3 Tax=Homininae RepID=GFPT1_HUMAN	0.79	1
P48643	T-complex protein 1 subunit epsilon	0.77	2
P05141	ADP/ATP translocase 2	0.76	5
P63244	Guanine nucleotide-binding protein subunit beta-2-like 1	0.75	3
P31943	Heterogeneous nuclear ribonucleoprotein H	0.73	7
O43175	D-3-phosphoglycerate dehydrogenase	0.73	1
Q06830	Peroxioredoxin-1	0.73	3
P62244	40S ribosomal protein S15a	0.70	1
Q9BVG8	Kinesin-like protein KIFC3	0.70	3
P68363	Tubulin alpha-1B chain	0.69	13
O43795	Unconventional myosin-Ib	0.68	1
P26641	Elongation factor 1-gamma	0.67	1
P62805	Histone H4	0.67	2
P61353	60S ribosomal protein L27	0.66	1
Q99623	Prohibitin-2	0.65	2
P49327	Fatty acid synthase	0.65	4
Q6IPH7	RPL14 protein	0.63	1
P68371	Tubulin beta-4B chain	0.63	13
P62829	60S ribosomal protein L23	0.60	1
P62701	40S ribosomal protein S4 X isoform	0.60	2
P07437	Tubulin beta chain	0.60	13
P18124	60S ribosomal protein L7	0.59	2
P01859	Ig gamma-2 chain C region	0.59	1
P49368	T-complex protein 1 subunit gamma	0.57	1
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.57	3
P02768	Serum albumin	0.56	13
P17844	Probable ATP-dependent RNA helicase DDX5	0.50	1
Q149N5	DOCK4 protein	0.49	10
P15880	40S ribosomal protein S2	0.49	1
P06576	ATP synthase subunit beta mitochondrial	0.48	3
P68104	Elongation factor 1-alpha 1	0.48	6
Q14103	Heterogeneous nuclear ribonucleoprotein D0	0.48	1
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.47	5
Q96EY1	DnaJ homolog subfamily A member 3 mitochondrial	0.47	1
P14618	Pyruvate kinase PKM	0.46	3
P62826	GTP-binding nuclear protein Ran	0.46	1
O60701	UDP-glucose 6-dehydrogenase	0.46	4
P02679	Fibrinogen gamma chain	0.44	4
P60709	Actin cytoplasmic 1	0.43	11
P25705	ATP synthase subunit alpha mitochondrial	0.43	3
P46776	60S ribosomal protein L27a	0.42	1
P18621	60S ribosomal protein L17	0.41	1
P23396	40S ribosomal protein S3	0.39	1
A2NUT2	Lambda-chain (AA -20 to 215)	0.39	3
Q6IB91	PCK2 protein	0.39	1
P01700	Ig lambda chain V-I region HA	0.36	2
Q15084	Protein disulfide-isomerase A6	0.35	1
P49411	Elongation factor Tu mitochondrial	0.33	2
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.32	3
P60842	Eukaryotic initiation factor 4A-I	0.32	1
P02671	Fibrinogen alpha chain	0.31	4
P08238	Heat shock protein HSP 90-beta	0.29	3
P01617	Ig kappa chain V-II region TEW	0.28	1
Q15365	Poly(rC)-binding protein 1	0.27	1
P04406	Glyceraldehyde-3-phosphate dehydrogenase	0.27	1
P19474	E3 ubiquitin-protein ligase TRIM21	0.26	4
Q02543	60S ribosomal protein L18a	0.26	1
P08107	Heat shock 70 kDa protein 1A1B	0.25	2
P22061	Protein-L-isoaspartate(D-aspartate) O-methyltransferase	0.25	4
P02769	Serum albumin	0.24	37
P14625	Endoplasmic	0.24	1
P05783	Keratin type I cytoskeletal 18	0.23	4
P51659	Peroxisomal multifunctional enzyme type 2	0.23	7
P11142	Heat shock cognate 71 kDa protein	0.22	2
O00571	ATP-dependent RNA helicase DDX3X	0.21	1
Q15058	Kinesin-like protein KIF14	0.19	6
Q9BUJ2	Heterogeneous nuclear ribonucleoprotein U-like protein 1	0.17	3
P06733	Alpha-enolase	0.16	1
P08727	Keratin type I cytoskeletal 19	0.15	1
P00338	L-lactate dehydrogenase A chain	0.13	1
P05787	Keratin type II cytoskeletal 8	0.12	3
Q00610	Claithrin heavy chain 1	0.11	3
H6VRG3	Keratin 1	0.07	3
P11021	78 kDa glucose-regulated protein	0.06	1
P35908	Keratin type II cytoskeletal 2 epidermal	0.04	1
P13645	Keratin type I cytoskeletal 10	0.04	1