

Accession Number	Description	Fold Enrichment	UniquePeptideCount
<b>Q9UIS9</b>	<b>Methyl-CpG-binding domain protein 1</b>	<b>4.25</b>	<b>2</b>
P01605	Ig kappa chain V-I region Lay	1.30	2
A2NUT2	Lambda-chain AA -20 to 215	1.05	4
P02656	Apolipoprotein C-III	0.96	2
P45880	Voltage-dependent anion-selective channel protein 2	0.91	1
P01617	Ig kappa chain V-II region TEW	0.88	2
P01024	Complement C3	0.80	1
P31943	Heterogeneous nuclear ribonucleoprotein H	0.79	3
P04264	Keratin type II cytoskeletal 1	0.71	14
P13645	Keratin type I cytoskeletal 10	0.69	12
P25705	ATP synthase subunit alpha mitochondrial	0.65	2
P02769	Serum albumin	0.63	38
P63261	Actin cytoplasmic 2	0.60	5
P08107	Heat shock 70 kDa protein 1A1B	0.51	2
P31942	Heterogeneous nuclear ribonucleoprotein H3	0.48	1
P06733	Alpha-enolase	0.48	2
P11142	Heat shock cognate 71 kDa protein	0.44	2
P35908	Keratin type II cytoskeletal 2 epidermal	0.42	7
P35527	Keratin type I cytoskeletal 9	0.39	6
P22061	Protein-L-isoaspartateD-aspartate O-methyltransferase	0.29	2
Q149N5	DOCK4 protein	0.26	2
Q13268	Dehydrogenasereductase SDR family member 2 mitochondrial	0.26	1
Q00839	Heterogeneous nuclear ribonucleoprotein U	0.26	1
P02671	Fibrinogen alpha chain	0.17	1
P68363	Tubulin alpha-1B chain	0.16	1
P51659	Peroxisomal multifunctional enzyme type 2	0.15	2
P05787	Keratin type II cytoskeletal 8	0.08	1