

| Accession Number | Description   | Fold Enrichment | UniquePeptideCount |
|------------------|---|-----------------|--------------------|
| P68371           | Tubulin beta-4B chain   | 14.72           | 12                 |
| B3KQT2           | cDNA PSEC0148 fis clone PLACE1007202 highly similar to Protein disulfide-isomerase A3 | 10.15           | 8                  |
| O43237           | Cytoplasmic dynein 1 light intermediate chain 2                                       | 5.57            | 4                  |
| Q13838           | Spliceosome RNA helicase DDX39B   | 4.71            | 5                  |
| O00330           | Pyruvate dehydrogenase protein X component mitochondrial                              | 4.43            | 3                  |
| P26196           | Probable ATP-dependent RNA helicase DDX6  | 4.43            | 3                  |
| P26640           | Valine--tRNA ligase   | 4.43            | 3                  |
| P62495           | Eukaryotic peptide chain release factor subunit 1                                     | 4.43            | 3                  |
| Q13509           | Tubulin beta-3 chain  | 4.43            | 3                  |
| Q86UY0           | TXNDC5 protein  | 4.43            | 3                  |
| Q96KP4           | Cytosolic non-specific dipeptidase  | 4.43            | 3                  |
| P68363           | Tubulin alpha-1B chain  | 3.86            | 9                  |
| P02679           | Fibrinogen gamma chain  | 3.65            | 3                  |
| Q10713           | Mitochondrial-processing peptidase subunit alpha                                      | 3.65            | 3                  |
| E9PM69           | 26S protease regulatory subunit 6A  | 3.29            | 2                  |
| Q15460           | Prolyl 4-hydroxylase subunit alpha-2  | 3.29            | 2                  |
| P01859           | Ig gamma-2 chain C region   | 3.29            | 2                  |
| P60228           | Eukaryotic translation initiation factor 3 subunit E                                  | 3.29            | 2                  |
| Q02318           | Sterol 26-hydroxylase mitochondrial   | 3.29            | 2                  |
| Q13418           | Integrin-linked protein kinase  | 3.29            | 2                  |
| Q16877           | 6-phosphofructo-2-kinase/fructose-26-bisphosphatase 4                                 | 3.29            | 2                  |
| Q6AZ91           | RAB6B protein   | 3.29            | 2                  |
| Q8TD30           | Alanine aminotransferase 2  | 3.29            | 2                  |
| Q9GZL7           | Ribosome biogenesis protein WDR12   | 3.29            | 2                  |
| Q9Y4K3           | TNF receptor-associated factor 6  | 3.29            | 2                  |
| Q9Y512           | Sorting and assembly machinery component 50 homolog                                   | 3.29            | 2                  |
| Q01581           | Hydroxymethylglutaryl-CoA synthase cytoplasmic  | 3.27            | 3                  |
| Q14498           | RNA-binding protein 39  | 3.27            | 3                  |
| F5H7U0           | 6-phosphogluconate dehydrogenase decarboxylating                                      | 3.02            | 4                  |
| Q16851           | UTP--glucose-1-phosphate uridylyltransferase  | 2.83            | 4                  |
| P34897           | Serine hydroxymethyltransferase mitochondrial   | 2.71            | 2                  |
| Q13885           | Tubulin beta-2A chain   | 2.71            | 2                  |
| Q9BS26           | Endoplasmic reticulum resident protein 44   | 2.71            | 2                  |
| P41091           | Eukaryotic translation initiation factor 2 subunit 3                                  | 2.68            | 6                  |
| Q9NUQ8           | ATP-binding cassette sub-family F member 3  | 2.52            | 3                  |
| P05091           | Aldehyde dehydrogenase mitochondrial  | 2.40            | 3                  |
| Q9UMS4           | Pre-mRNA-processing factor 19   | 2.31            | 2                  |
| P01019           | Angiotensinogen   | 2.14            | 2                  |
| E7EWF1           | 60S ribosomal protein L4  | 1.99            | 2                  |
| P32189           | Glycerol kinase   | 1.99            | 2                  |
| Q15084           | Protein disulfide-isomerase A6  | 1.95            | 3                  |
| Q9Y230           | RuvB-like 2   | 1.93            | 5                  |
| P51812           | Ribosomal protein S6 kinase alpha-3   | 1.86            | 4                  |
| P13639           | Elongation factor 2   | 1.81            | 4                  |
| P14868           | Aspartate--tRNA ligase cytoplasmic  | 1.79            | 7                  |
| P38919           | Eukaryotic initiation factor 4A-III   | 1.70            | 5                  |
| O95573           | Long-chain-fatty-acid--CoA ligase 3   | 1.64            | 2                  |
| Q6YN16           | Hydroxysteroid dehydrogenase-like protein 2   | 1.63            | 4                  |
| P28838           | Cytosol aminopeptidase  | 1.57            | 4                  |
| P49257           | Protein ERGIC-53  | 1.51            | 4                  |
| Q14254           | Flotillin-2   | 1.45            | 2                  |
| P26641           | Elongation factor 1-gamma   | 1.42            | 7                  |
| Q9Y262           | Eukaryotic translation initiation factor 3 subunit L                                  | 1.41            | 2                  |
| P19474           | E3 ubiquitin-protein ligase TRIM21  | 1.40            | 8                  |
| P39656           | Dolichyl-diphosphooligosaccharide--protein glycosyltransferase 48 kDa subunit         | 1.37            | 3                  |
| P19338           | Nucleolin   | 1.36            | 2                  |
| P27708           | CAD protein   | 1.29            | 2                  |
| P60842           | Eukaryotic initiation factor 4A-I   | 1.19            | 6                  |
| Q3LXA3           | Bifunctional ATP-dependent dihydroxyacetone kinase/FAD-AMP lyase (cyclizing)          | 1.17            | 3                  |
| P78371           | T-complex protein 1 subunit beta  | 1.12            | 2                  |
| P17844           | Probable ATP-dependent RNA helicase DDX5  | 1.08            | 3                  |
| Q9Y265           | RuvB-like 1   | 1.03            | 2                  |
| P02769           | Serum albumin   | 1.00            | 18                 |
| <b>Q01664</b>    | <b>Transcription factor AP-4</b>  | <b>1.00</b>     | <b>3</b>           |
| P50991           | T-complex protein 1 subunit delta   | 0.99            | 2                  |
| P09327           | Villin-1  | 0.99            | 2                  |
| P54886           | Delta-1-pyrroline-5-carboxylate synthase  | 0.98            | 8                  |
| Q9NZI8           | Insulin-like growth factor 2 mRNA-binding protein 1                                   | 0.95            | 3                  |
| P25705           | ATP synthase subunit alpha mitochondrial  | 0.94            | 9                  |
| P17987           | T-complex protein 1 subunit alpha   | 0.93            | 4                  |
| P07900           | Heat shock protein HSP 90-alpha   | 0.85            | 2                  |
| P14625           | Endoplasmic   | 0.85            | 4                  |
| P49368           | T-complex protein 1 subunit gamma   | 0.81            | 3                  |
| P02768           | Serum albumin   | 0.80            | 3                  |
| P11142           | Heat shock cognate 71 kDa protein   | 0.75            | 3                  |
| P06733           | Alpha-enolase   | 0.75            | 9                  |
| P06576           | ATP synthase subunit beta mitochondrial   | 0.72            | 8                  |
| B5BTY4           | ATP-dependent RNA helicase DDX3X  | 0.65            | 4                  |
| P08238           | Heat shock protein HSP 90-beta  | 0.64            | 8                  |
| O60701           | UDP-glucose 6-dehydrogenase   | 0.52            | 2                  |
| P14618           | Pyruvate kinase isozymes M1/M2  | 0.50            | 5                  |
| P00367           | Glutamate dehydrogenase 1 mitochondrial   | 0.42            | 3                  |
| P15924           | Desmoplakin   | 0.40            | 2                  |
| B4E335           | Actin cytoplasmic 1 N-terminally processed  | 0.34            | 4                  |
| P10809           | 60 kDa heat shock protein mitochondrial   | 0.21            | 3                  |