

| Accession Number | Description | Fold Enrichment | UniquePeptideCount |
|------------------|---|-----------------|--------------------|
| Q09472 | Histone acetyltransferase p300 | 15.94 | 8 |
| P62913 | 60S ribosomal protein L11 | 1.53 | 2 |
| P60842 | Eukaryotic initiation factor 4A-I | 1.38 | 3 |
| P35555 | Fibrillin-1 | 1.34 | 2 |
| A7MBN2 | TNC protein | 1.17 | 1 |
| P00338 | L-lactate dehydrogenase A chain | 0.95 | 6 |
| Q02543 | 60S ribosomal protein L18a | 0.88 | 2 |
| P35556 | Fibrillin-2 | 0.82 | 1 |
| P49327 | Fatty acid synthase | 0.80 | 2 |
| P60709 | Actin cytoplasmic 1 | 0.76 | 8 |
| O60701 | UDP-glucose 6-dehydrogenase | 0.73 | 3 |
| P04406 | Glyceraldehyde-3-phosphate dehydrogenase | 0.72 | 2 |
| P02768 | Serum albumin | 0.71 | 8 |
| P19474 | E3 ubiquitin-protein ligase TRIM21 | 0.71 | 6 |
| P08238 | Heat shock protein HSP 90-beta | 0.70 | 4 |
| P01709 | Ig lambda chain V-II region MGC | 0.65 | 2 |
| P35527 | Keratin type I cytoskeletal 9 | 0.64 | 12 |
| Q9H222 | ATP-binding cassette sub-family G member 5 | 0.62 | 1 |
| P68363 | Tubulin alpha-1B chain | 0.57 | 5 |
| P05783 | Keratin type I cytoskeletal 18 | 0.54 | 5 |
| D3DPG0 | Titin isoform CRA_a | 0.51 | 1 |
| P02769 | Serum albumin | 0.50 | 37 |
| P68104 | Elongation factor 1-alpha 1 | 0.36 | 2 |
| A2NUT2 | Lambda-chain (AA -20 to 215) | 0.31 | 1 |
| H6VRG3 | Keratin 1 | 0.30 | 7 |
| Q13268 | Dehydrogenase/reductase SDR family member 2 mitochondrial | 0.26 | 1 |
| P68371 | Tubulin beta-4B chain | 0.24 | 2 |
| P07437 | Tubulin beta chain | 0.23 | 2 |
| P13645 | Keratin type I cytoskeletal 10 | 0.21 | 4 |
| P15924 | Desmoplakin | 0.20 | 1 |
| P02671 | Fibrinogen alpha chain | 0.20 | 1 |
| Q9ULV0 | Unconventional myosin-Vb | 0.15 | 1 |
| P11021 | 78 kDa glucose-regulated protein | 0.10 | 1 |
| Q15058 | Kinesin-like protein KIF14 | 0.09 | 1 |