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|--------|--|--------|---|---|---|---|----------|---|---|---|---|--------|--------|-------|--------|------|-------|------|
| P07996 | Thrombospondin-1 OS=Homo sapiens GN=THBS1 PE=1 SV=2 - [TSP1_HUMAN]   | 1.03%  | 1 | 1 | 1 | 2 | 8.63E-05 | 1 | 1 | 1 | 1 | 1.03%  | 1.03%  | 2.86  | 27.25  | 1170 | 129.3 | 4.94 |
| P16646 | Stress-70 protein, mitochondrial OS=Homo sapiens GN=HSPA9 PE=1 SV=2 - [GRP75_HUMAN]                                | 1.33%  | 1 | 1 | 1 | 2 | 8.49E-05 | 1 | 1 | 1 | 1 | 1.33%  | 1.33%  | 2.77  | 58.04  | 679  | 73.6  | 6.16 |
| Q91014 | Vascular protein sorting-associated protein 7B OS=Homo sapiens GN=VPS7B PE=1 SV=1 - [VPS7B_HUMAN]                  | 5.96%  | 1 | 1 | 1 | 4 | 8.48E-05 | 1 | 1 | 2 | 2 | 5.96%  | 5.96%  | 6.28  | 36.85  | 285  | 31.3  | 7.34 |
| Q9Y696 | Chloride intracellular channel protein 4 OS=Homo sapiens GN=CLIC4 PE=1 SV=4 - [CLIC4_HUMAN]                        | 16.60% | 1 | 3 | 3 | 8 | 8.44E-05 | 3 | 3 | 4 | 4 | 16.60% | 16.60% | 11.93 | 84.47  | 253  | 28.8  | 5.59 |
| Q98277 | Charged multivesicular body protein 6 OS=Homo sapiens GN=CHMP6 PE=1 SV=3 - [CHMP6_HUMAN]                           | 6.47%  | 1 | 1 | 1 | 2 | 7.93E-05 | 1 | 1 | 1 | 1 | 6.47%  | 6.47%  | 2.72  | 52.77  | 201  | 23.5  | 5.31 |
| P73802 | Transglutinin-2 OS=Homo sapiens GN=TG2 PE=1 SV=3 - [TAGLN2_HUMAN]  | 7.04%  | 1 | 1 | 1 | 1 | 7.90E-05 | 1 | 1 | 1 | 1 | 7.04%  | 0.00%  | 2.25  | 199    | 22.4 | 8.25  |      |
| A6ND89 | Paranemin-3 OS=Homo sapiens GN=PALM3 PE=1 SV=2 - [PALM3_HUMAN]   | 4.46%  | 1 | 2 | 2 | 4 | 7.61E-05 | 2 | 2 | 2 | 2 | 4.46%  | 4.46%  | 5.96  | 66.61  | 673  | 71.7  | 4.53 |
| P08995 | Annexin A11 OS=Homo sapiens GN=ANXA11 PE=1 SV=1 - [ANXA11_HUMAN]   | 2.18%  | 1 | 1 | 1 | 1 | 7.31E-05 | 1 | 1 | 1 | 1 | 2.18%  | 2.18%  | 2.46  | 50.5   | 54.4 | 7.65  |      |
| P09651 | Heterogeneous nuclear ribonucleoprotein A1 OS=Homo sapiens GN=HNRNP_A1 PE=1 SV=5 - [ROA1_HUMAN]                    | 9.95%  | 2 | 2 | 3 | 6 | 7.22E-05 | 3 | 3 | 3 | 3 | 9.95%  | 9.95%  | 8.54  | 106.5  | 372  | 38.7  | 9.13 |
| O75781 | Paranemin-1 OS=Homo sapiens GN=PALM1 PE=1 SV=2 - [PALM_HUMAN]  | 4.91%  | 1 | 1 | 1 | 2 | 6.97E-05 | 1 | 1 | 1 | 1 | 4.91%  | 4.91%  | 2.91  | 35.72  | 387  | 42.1  | 5    |
| P61088 | Ubiquitin-conjugating enzyme E2 N OS=Homo sapiens GN=UBE2N PE=1 SV=1 - [UBE2N_HUMAN]                               | 7.24%  | 2 | 1 | 1 | 2 | 6.65E-05 | 1 | 1 | 1 | 1 | 7.24%  | 7.24%  | 3.98  | 45.07  | 152  | 17.1  | 6.57 |
| P05386 | 60S acidic ribosomal protein P1 OS=Homo sapiens GN=RPL1 PE=1 SV=1 - [RPL1_HUMAN]                                   | 14.00% | 1 | 1 | 1 | 2 | 6.40E-05 | 1 | 1 | 2 | 2 | 14.00% | 14.00% | 5.07  | 114    | 11.5 | 4.32  |      |
| P22626 | Heterogeneous nuclear ribonucleoprotein A2/B1 OS=Homo sapiens GN=HNRNP_A2B1 PE=1 SV=2 - [ROA2_HUMAN]               | 5.38%  | 1 | 1 | 2 | 4 | 6.32E-05 | 2 | 2 | 2 | 2 | 5.38%  | 5.38%  | 5.42  | 58.92  | 353  | 37.4  | 8.95 |
| Q15599 | Na(+)/H(+) exchange regulatory cofactor NHE-RF2 OS=Homo sapiens GN=SLC9A3R2 PE=1 SV=2 - [NHRF2_HUMAN]              | 2.67%  | 1 | 1 | 1 | 2 | 5.85E-05 | 1 | 1 | 1 | 1 | 2.67%  | 2.67%  | 2.44  | 49.1   | 337  | 37.4  | 7.93 |
| P10301 | Ras-related protein R-Ras OS=Homo sapiens GN=RRAS PE=1 SV=1 - [RRAS_HUMAN]   | 12.84% | 2 | 2 | 2 | 6 | 5.33E-05 | 2 | 2 | 3 | 3 | 12.84% | 12.84% | 11.3  | 118.15 | 218  | 23.5  | 6.93 |
| Q15247 | Tubulin-specific chaperone A OS=Homo sapiens GN=TRCA PE=1 SV=3 - [TRCA_HUMAN]                                      | 8.33%  | 1 | 1 | 1 | 1 | 4.71E-05 | 1 | 1 | 1 | 1 | 8.33%  | 8.33%  | 2.04  | 108    | 12.8 | 5.29  |      |
| Q99961 | Endophilin-A2 OS=Homo sapiens GN=SH3GL1 PE=1 SV=1 - [SH3GL1_HUMAN]   | 5.43%  | 1 | 1 | 1 | 2 | 4.71E-05 | 1 | 1 | 1 | 1 | 5.43%  | 5.43%  | 1.9   | 51.17  | 368  | 41.5  | 5.43 |
| P63218 | Guanine nucleotide-binding protein G(I)/G(S)/G(O) subunit gamma 5 OS=Homo sapiens GN=GNG5 PE=1 SV=3 - [GBG5_HUMAN] | 13.24% | 1 | 1 | 1 | 2 | 4.43E-05 | 1 | 1 | 1 | 1 | 13.24% | 13.24% | 2.64  | 49.75  | 68   | 7.3   | 9.85 |
| P10073 | Annexin A7 OS=Homo sapiens GN=ANXA7 PE=1 SV=3 - [ANXA7_HUMAN]  | 8.61%  | 1 | 3 | 3 | 6 | 4.31E-05 | 3 | 3 | 3 | 3 | 8.61%  | 8.61%  | 8.18  | 43.49  | 488  | 52.7  | 5.68 |
| Q91489 | EH domain-containing protein 1 OS=Homo sapiens GN=EHD1 PE=1 SV=2 - [EHD1_HUMAN]                                    | 2.25%  | 1 | 1 | 1 | 2 | 4.07E-05 | 1 | 1 | 1 | 1 | 2.25%  | 2.25%  | 2.6   | 22.35  | 534  | 60.6  | 6.83 |
| Q15365 | Poly(C)-binding protein 1 OS=Homo sapiens GN=PCBP1 PE=1 SV=2 - [PCBP1_HUMAN]                                       | 3.09%  | 3 | 1 | 1 | 1 | 3.94E-05 | 1 | 1 | 1 | 1 | 3.09%  | 3.09%  | 2.72  | 35.6   | 37.5 | 7.09  |      |
| Q53E24 | Centromeres protein of 55 kDa OS=Homo sapiens GN=CEP55 PE=1 SV=3 - [CEP55_HUMAN]                                   | 2.59%  | 1 | 1 | 1 | 1 | 3.90E-05 | 1 | 1 | 1 | 1 | 0.00%  | 2.59%  | 35.09 | 464    | 54.1 | 7.01  |      |
| A5DN16 | Vascular protein sorting-associated protein 37C OS=Homo sapiens GN=VPS37C PE=1 SV=2 - [VPS37C_HUMAN]               | 3.10%  | 1 | 1 | 1 | 2 | 3.67E-05 | 1 | 1 | 1 | 1 | 3.10%  | 3.10%  | 3.05  | 57.6   | 355  | 38.6  | 5.21 |
| Q98X67 | Junctional adhesion molecule C OS=Homo sapiens GN=JAM3 PE=1 SV=1 - [JAM3_HUMAN]                                    | 2.58%  | 1 | 1 | 1 | 2 | 3.52E-05 | 1 | 1 | 1 | 1 | 2.58%  | 2.58%  | 2.41  | 26.51  | 310  | 35    | 7.59 |
| Q14108 | Lysosome membrane protein 2 OS=Homo sapiens GN=SCARB2 PE=1 SV=2 - [SCRB2_HUMAN]                                    | 2.72%  | 1 | 1 | 1 | 1 | 3.16E-05 | 1 | 1 | 1 | 1 | 2.72%  | 2.72%  | 2.42  | 47.8   | 54.3 | 5.14  |      |
| P5399  | EST1 homolog OS=Homo sapiens GN=EST1 PE=1 SV=1 - [EST1_HUMAN]  | 2.75%  | 1 | 1 | 1 | 2 | 3.15E-05 | 1 | 1 | 1 | 1 | 2.75%  | 2.75%  | 2.68  | 33.97  | 364  | 39.7  | 5.35 |
| Q9C012 | Protein trexy homolog 3 OS=Homo sapiens GN=TTYH3 PE=1 SV=3 - [TTYH3_HUMAN]   | 3.06%  | 1 | 2 | 2 | 4 | 2.37E-05 | 2 | 2 | 2 | 2 | 3.06%  | 3.06%  | 4.98  | 38.26  | 523  | 57.5  | 5.39 |
| P6081  | Dextrin OS=Homo sapiens GN=DSTN PE=1 SV=3 - [DSTN_HUMAN]   | 7.88%  | 1 | 1 | 1 | 2 | 2.07E-05 | 1 | 1 | 1 | 1 | 7.88%  | 7.88%  | 3.12  | 34.28  | 165  | 18.5  | 7.85 |
| P14550 | Alcohol dehydrogenase [NAD(P)+] OS=Homo sapiens GN=AKR1A1 PE=1 SV=3 - [AKR1A1_HUMAN]                               | 5.83%  | 1 | 1 | 1 | 2 | 1.97E-05 | 1 | 1 | 1 | 1 | 5.83%  | 5.83%  | 1.75  | 20.39  | 325  | 36.5  | 6.79 |
| P09607 | Amyloid beta 44 protein OS=Homo sapiens GN=APP PE=1 SV=3 - [A4_HUMAN]  | 3.12%  | 1 | 2 | 2 | 6 | 8.52E-04 | 2 | 2 | 3 | 3 | 3.12%  | 3.12%  | 4.47  | 64.61  | 770  | 86.9  | 4.82 |
| P27105 | Erythrocyte band 7 integral membrane protein OS=Homo sapiens GN=STOM PE=1 SV=3 - [STOM_HUMAN]                      | 8.33%  | 1 | 2 | 2 | 4 | 0.00E+00 | 2 | 2 | 2 | 2 | 8.33%  | 8.33%  | 6.29  | 70.73  | 288  | 31.7  | 7.88 |