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| G1 | CYB5 MOUSE Mass: 15232 Score: 49 Expect: 0.2 Matches: 5 Cytochrome b5 OS=Mus musculus GN=Cyb5a PE=1 SV=2 (potvrzeno MS/MS) |
| G2 | MDHC MOUSE Mass: 36659 Score: 78 Expect: 0.00025 Matches: 13 Malate dehydrogenase, cytoplasmic OS=Mus musculus GN=Mdh1 PE=1 SV=3 |
| G3 | GSTM1 MOUSE Mass: 26067 Score: 242 Expect: 1.1e-20 Matches: 26 Glutathione S-transferase Mu 1 OS=Mus musculus GN=Gstm1 PE=1 SV=2 |
| G4 | HPPD MOUSE Mass: 45254 Score: 59 Expect: 0.02 Matches: 9 4-hydroxyphenylpyruvate dioxygenase OS=Mus musculus GN=Hpd PE=1 SV=3 |
| G5 | RGN MOUSE Mass: 33899 Score: 214 Expect: 6.7e-18 Matches: 25 Regucalcin OS=Mus musculus GN=Rgn PE=1 SV=1 |
| G6 | ENPL MOUSE Mass: 92703 Score: 249 Expect: 2.1e-21 Matches: 32 Endoplasmin OS=Mus musculus GN=Hsp90b1 PE=1 SV=2 |
| G7 | SODC MOUSE Mass: 16104 Score: 84 Expect: 6.4e-05 Matches: 8 Superoxide dismutase [Cu-Zn] OS=Mus musculus GN=Sod1 PE=1 SV=2 |
| G8 | SODM MOUSE Mass: 24816 Score: 93 Expect: 8.2e-06 Matches: 10 Superoxide dismutase [Mn], mitochondrial OS=Mus musculus GN=Sod2 PE=1 SV=3 |
| G9 | Mixture 1 Total score: 192 Expect: 1.1e-15 Matches: 28 Components (only one family member shown for each component): APOA1 MOUSE Mass: 30597 Score: 118 Expect: 2.7e-08 Matches: 16 Apolipoprotein A-I OS=Mus musculus GN=Apoa1 PE=1 SV=2 LGUL MOUSE Mass: 20967 Score: 105 Expect: 5.3e-07 Matches: 12 Lactoylglutathione lyase OS=Mus musculus GN=Glo1 PE=1 SV=3 |
| G10 | GSTA3 MOUSE Mass: 25401 Score: 126 Expect: 4.2e-09 Matches: 16 Glutathione S-transferase A3 OS=Mus musculus GN=Gsta3 PE=1 SV=2 |