

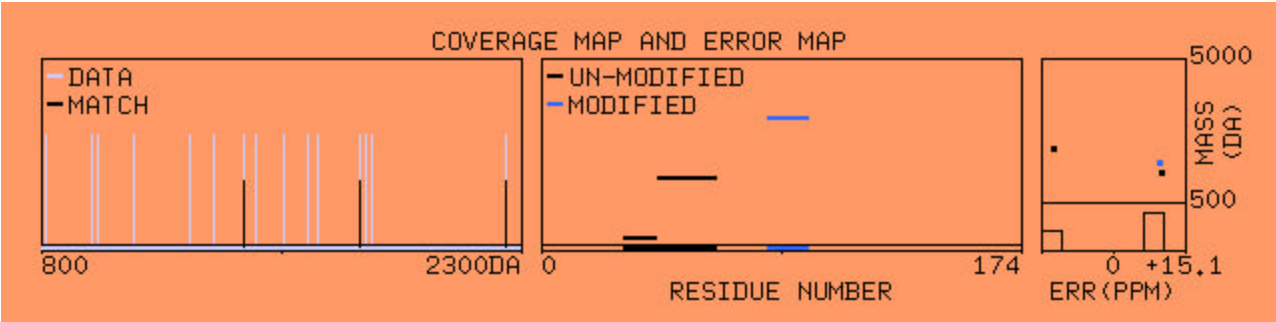
ProFound - Search Result Details


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Details for rank 1 candidate in search 20060404163403-1DDC-192168001107

1. FTTSGH0066 61794 62315 [+3 L= 522 r=-1.226] (FTT0061 61794 62315 +)
====>ref|YP_044976.1| membrane-bound ATP synthase , F1 sector,
delta-subunit[Acinetobacter sp. ADP1] Score = 125 bits (314), Expect =
6e-28====>ref|YP_169136.1| ATP synthase delta chain [Francisella tularensis subsp.
tularensisSchu 4] Score = 335 bits (858), Expect = 5e-91

Sample ID : Schu4 264 [Pass:1]
Measured peptides : 15
Matched peptides : 3
Min. sequence coverage: 28%



Note: click on the  symbol to change column format.

| Measured | Avg/ | Computed | Error |  Residues | Missed | | |
|----------|------|----------|-----------------------------------------------------------------------------------------|--------------------------------------------------------------------------------------------|--------|-----|------------------------|
| Mass(M) | Mono | Mass |  (ppm) | Start | To | Cut | Peptide sequence |
| 1433.827 | M | 1433.812 | 10 | 31 | 42 | 1 | LLQTFSELIKDK |
| 1795.890 | M | 1795.873 | 10 | 83 | 97 | 0 | MLIMPEIADQFESIK |
| | | | | | | | (2)+O@M; |
| 2254.249 | M | 2254.277 | -13 | 43 | 64 | 1 | SVAAIVSSPTISQIEVVDALKK |

Unmatched Monoisotopic Masses:

815.653 962.459 975.262 1088.630 1268.669 1338.848 1471.177 1562.860 1636.092
1667.991 1818.886 1835.632

Search again using unmatched masses:

Search again

 in

All taxa