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logo                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                
NTDB id 432 SGO RS09405 WP 012130939.1      MVRTIVRLRQLPIKAFTVLESLLVLMISSFILLALSSSVQATFEQIQAKIFFLEFEHFYQESQKLSVSSQRKLVLEISSQEISNG   85
NTDB id 531 SMSK321 RS09225 WP 000588032.1  ...........MIKAFTMLESLLVLGLVSIVALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQTISNG   74
NTDB id 503 SM12261 RS00960 WP 029689647.1  .MLNKIPVKPLQIKAFTMLESLLVLGLVSILALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGKTISNG   84
NTDB id 163 SP RS10365 WP 000588028.1       ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQTLSNG   74
NTDB id 273 KZH43 RS09250 WP 000588026.1    ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQMISNG   74
NTDB id 232 SPD RS09880 WP 000588026.1      ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQMISNG   74
NTDB id 198 SPR RS09415 WP 000588026.1      ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQMISNG   74
NTDB id 22445 GBS RS01085 WP 000244259.1    .....MVLRKFQVKAFTVLESLIVLSVVAFMTLVFSTSFNNIFRQVEETIFFISFEHLYRDTQKLSAFGQKKQTLTISHNYLENT   80
NTDB id 397 FSA28 RS09390 WP 002263438.1    .........MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLILKSEYISNG   76
NTDB id 388 SMU RS09020 WP 002263438.1      .........MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLILKSEYISNG   76
consensus                                                 **!!!!**!!!**!*******!**!*!****! ******!!**!!**!***!!*!********!***** **!*
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logo                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                
NTDB id 432 SGO RS09405 WP 012130939.1      YARLPIPKGIQAPESTQIYFDKAGGNSSLSKVQFQTK.ECLVTYQLYIGNGKFKKTTN...   142
NTDB id 531 SMSK321 RS09225 WP 000588032.1  SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTG.KGTIRYQLYLGNGKIKRIKETKN   134
NTDB id 503 SM12261 RS00960 WP 029689647.1  SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKD   144
NTDB id 163 SP RS10365 WP 000588028.1       SQKLPVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 273 KZH43 RS09250 WP 000588026.1    SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 232 SPD RS09880 WP 000588026.1      SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 198 SPR RS09415 WP 000588026.1      SQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 22445 GBS RS01085 WP 000244259.1    YERLYLPKTVKVVKSDTLAFDANGGNSSLAKIQFECY.RKTVTYQLYIGSGNYRKKEN...   137
NTDB id 397 FSA28 RS09390 WP 002263438.1    VEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES...   134
NTDB id 388 SMU RS09020 WP 002263438.1      VEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES...   134
consensus                                   ***!**!!***********!!**!!!!!!*!**!*** *****!!!!*!*!**********
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X non conserved

X similar

X $\geq$ 50% conserved