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logo
NTDB id 432 SGO RS09405 WP 012130939.1          MVRTIVRLRQLPIKAFTVLESLLVLMISSFILLALSSSVQATFEQIQAKIFFLEFEHFYQESQKLSVSSQRKLVLEISSQ   80
NTDB id 531 SMSK321 RS09225 WP 000588032.1      ...........MIKAFTMLESLLVLGLVSIVALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQ   69
NTDB id 503 SM12261 RS00960 WP 029689647.1      .MLNKIPVKPLQIKAFTMLESLLVLGLVSILALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGK   79
NTDB id 163 SP RS10365 WP 000588028.1           ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQ   69
NTDB id 273 KZH43 RS09250 WP 000588026.1        ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQ   69
NTDB id 232 SPD RS09880 WP 000588026.1          ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQ   69
NTDB id 198 SPR RS09415 WP 000588026.1          ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQ   69
NTDB id 1159070 QOR62 RS01115 WP 000244261.1    .....MVLRKFQVKAFTVLESLIVLSVVAFMTLVFSTSFNNIFRQVEETIFFISFEHLYRDTQKLSAFGQKKQTLTISHN   75
NTDB id 397 FSA28 RS09390 WP 002263438.1        .........MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLILKSE   71
NTDB id 388 SMU RS09020 WP 002263438.1          .........MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLILKSE   71
consensus                                                **!!!!**!!!**!*******!**!*!****! ******!!**!!**!**!!*!********!*****

logo
NTDB id 432 SGO RS09405 WP 012130939.1          EISNGYARLPIPKGIQAPESTQIYFDKAGGNSSLSKVQFQTK.EGLVTYQLYIGNGKFKKTTN...   142
NTDB id 531 SMSK321 RS09225 WP 000588032.1      TISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTG.KGTIRYQLYLGNGKIKRIKETKN   134
NTDB id 503 SM12261 RS00960 WP 029689647.1      TISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKD   144
NTDB id 163 SP RS10365 WP 000588028.1           TLSNGSQKLPVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 273 KZH43 RS09250 WP 000588026.1        MISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 232 SPD RS09880 WP 000588026.1          MISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 198 SPR RS09415 WP 000588026.1          MISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 1159070 QOR62 RS01115 WP 000244261.1    YLENTYERLYLPKTVKVVKSDTLAFDTNGGNSSLAKIQFECY.RKTVTYQLYIGSGNYRKKEN...   137
NTDB id 397 FSA28 RS09390 WP 002263438.1        YISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES...   134
NTDB id 388 SMU RS09020 WP 002263438.1          YISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES...   134
consensus                                         **!****!**!!***********!!**!!!!!!*!**!*** *****!!!!*!*!**********
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X non conserved

X similar

X ≥ 50% conserved