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logo
NTDB id 42081 LCGT RS00740 WP 014024177.1   .........MKNVMYRIKALTLLESLLVLFLCSFVLFLFSGSVKQSVRIARAEIFILQFERLYKDTQRMAGLKGQQQTLSASNGA   76
NTDB id 589 KW2 RS10840 WP 096816401.1      MMITMTRIKMNSAISMIRAFTLLESLLVLLIVSFITLFFSAELTQTVHLFKGELFVLQFENLYKISQENAALQSSPENLGSKNGK   85
NTDB id 397 FSA28 RS09390 WP 002263438.1    .............MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLILKSEY   72
NTDB id 388 SMU RS09020 WP 002263438.1      .............MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLILKSEY   72
NTDB id 503 SM12261 RS00960 WP 029689647.1  .....MLNKIPVKPLQIKAFTMLESLLVLGLVSILALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGKT   80
NTDB id 531 SMSK321 RS09225 WP 000588032.1  ...............MIKAFTMLESLLVLGLVSIVALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQT   70
NTDB id 163 SP RS10365 WP 000588028.1       ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQT   70
NTDB id 273 KZH43 RS09250 WP 000588026.1    ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQM   70
NTDB id 232 SPD RS09880 WP 000588026.1      ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQM   70
NTDB id 198 SPR RS09415 WP 000588026.1      ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNLDGQM   70
consensus                                                   *!*!*!**!!!**!***!******!******* ******!***!!*!!***!*********!*****

logo
NTDB id 42081 LCGT RS00740 WP 014024177.1   LYSQEEMLKVPEEAEIN.DFSITFDQKSGNSSLQKITIFLPYQKKTISYQLEIGSGKYKKRIT...   138
NTDB id 589 KW2 RS10840 WP 096816401.1      LIYENKEIDIPKEVEMV.EFLIKFDEKGENSSLQKIKVYLPYEKKTILYQMEMGSGKYKKKIN...   147
NTDB id 397 FSA28 RS09390 WP 002263438.1    ISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLV.FQTSDEKRVTYQLYIGSGQYKKTES...   134
NTDB id 388 SMU RS09020 WP 002263438.1      ISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLV.FQTSDEKRVTYQLYIGSGQYKKTES...   134
NTDB id 503 SM12261 RS00960 WP 029689647.1  ISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVE.FQTSK.GAIRYQLYLGNGKIKRIKETKD   144
NTDB id 531 SMSK321 RS09225 WP 000588032.1  ISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVE.FQTGK.GTIRYQLYLGNGKIKRIKETKN   134
NTDB id 163 SP RS10365 WP 000588028.1       LSNGSQKLPVPKGIQAPSGQSITFDRAGGNSSLAKVE.FQTSK.GAIRYQLYLGNGKIKRIKETKN   134
NTDB id 273 KZH43 RS09250 WP 000588026.1    ISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVE.FQTSK.GAIRYQLYLGNGKIKRIKETKN   134
NTDB id 232 SPD RS09880 WP 000588026.1      ISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVE.FQTSK.GAIRYQLYLGNGKIKRIKETKN   134
NTDB id 198 SPR RS09415 WP 000588026.1      ISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVE.FQTSK.GAIRYQLYLGNGKIKRIKETKN   134
consensus                                   **********!************!!****!!!!*!** ***** ****!!***!*!**!*******
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X non conserved

X similar

X ≥ 50% conserved