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logo
NTDB id 589 KW2 RS10840 WP 096816401.1       MMITMTRIKMNSAISMIRAFTLLESLLVLLIVSFITLFFSAELTQTVHLFKGELFVLQFENLYKISQENAALQSSPENLG   80
NTDB id 432 SGO RS09405 WP 012130939.1       ....MVRTIVRLRQLPIKAFTVLESLLVLMISSFILLALSSSVQATFEQIQAKIFFLEFEHFYQESQKLSVSSQRKLVLE   76
NTDB id 531 SMSK321 RS09225 WP 000588032.1   ...............MIKAFTMLESLLVLGLVSIVALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLN   65
NTDB id 503 SM12261 RS00960 WP 029689647.1   .....MLNKIPVKPLQIKAFTMLESLLVLGLVSILALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLN   75
NTDB id 273 KZH43 RS09250 WP 000588026.1     ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLN   65
NTDB id 232 SPD RS09880 WP 000588026.1       ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLN   65
NTDB id 198 SPR RS09415 WP 000588026.1       ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLN   65
NTDB id 163 SP RS10365 WP 000588028.1        ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLN   65
NTDB id 185383 A9497 RS05170 WP 011226626.1  ....MRNMVVKAAQWPIRAFTLLESLMTLAVVVFLTLSLSGSVTGIFQQVEINLFYLRFEYLYRDSQRLASAEGSNVELQ   76
NTDB id 397 FSA28 RS09390 WP 002263438.1     .............MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLI   67
NTDB id 388 SMU RS09020 WP 002263438.1       .............MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLI   67
consensus                                                 *!*!!!**!!!**!*******!**!******* ******!***!!**!***!***********!*

logo
NTDB id 589 KW2 RS10840 WP 096816401.1       SKNGKLIYENKEIDIPKEVEMV.EFLIKFDEKGENSSLQKIKVYLPYEKKTILYQMEMGSGKYKKKIN...   147
NTDB id 432 SGO RS09405 WP 012130939.1       ISSQEISNGYARLPIPKGIQAPESTQIYFDKAGGNSSLSKVQFQ.T.KEGLVTYQLYIGNGKFKKTTN...   142
NTDB id 531 SMSK321 RS09225 WP 000588032.1   LDGQTISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQ.T.GKGTIRYQLYLGNGKIKRIKETKN   134
NTDB id 503 SM12261 RS00960 WP 029689647.1   LDGKTISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQ.T.SKGAIRYQLYLGNGKIKRIKETKD   144
NTDB id 273 KZH43 RS09250 WP 000588026.1     LDGQMISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQ.T.SKGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 232 SPD RS09880 WP 000588026.1       LDGQMISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQ.T.SKGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 198 SPR RS09415 WP 000588026.1       LDGQMISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQ.T.SKGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 163 SP RS10365 WP 000588028.1        LDGQTLSNGSQKLPVPKGIQAPSGQSITFDRAGGNSSLAKVEFQ.T.SKGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 185383 A9497 RS05170 WP 011226626.1  LTKDKVSNGRSSLLIPKSIHLDKGQTLVFDAKGGNSSLTKIRFS.S.DKEVVTYLLNMGSGKYKKTIS...   142
NTDB id 397 FSA28 RS09390 WP 002263438.1     LKSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQ.TSDEKRVTYQLYIGSGQYKKTES...   134
NTDB id 388 SMU RS09020 WP 002263438.1       LKSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQ.TSDEKRVTYQLYIGSGQYKKTES...   134
consensus                                    **** ******** *!!***********!!**!*!!!!*!**** *   ** **!****!*!**!******
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X non conserved

X similar

X $\geq$ 50% conserved