```
logo
                                                  ..........MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLIL
NTDB id 397 FSA28 RS09390 WP 002263438.1          ..........MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLIL   68
NTDB id 388 SMU RS09020 WP 002263438.1            ..........MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTLIL   68
NTDB id 432 SGO RS09405 WP 012130939.1            ...MVRTIVRLRQLPIKAFTVLESLLVLMISSFILLALSSSVQATFEQIQAKIFFLEFEHFYQESQKLSVSSQRKLVLEI   77
NTDB id 177425 A4H00 RS03445 WP 067087315.1       MNTLQKIQANLVRSTLKAFTLLESLLTLFIVSFLALALSGSVHSVFRQVEEQVFFMEFEQLYKSSQKLAVSSQQTISLDV   80
NTDB id 503 SM12261 RS00960 WP 029689647.1        ..MLNKI..PVKPLQIKAFTMLESLLVLGLVSILALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNL   76
NTDB id 531 SMSK321 RS09225 WP 000588032.1        ...............MIKAFTMLESLLVLGLVSIVALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSLNL   66
NTDB id 273 KZH43 RS09250 WP 000588026.1          ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNL   66
NTDB id 232 SPD RS09880 WP 000588026.1            ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNL   66
NTDB id 198 SPR RS09415 WP 000588026.1            ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNL   66
NTDB id 163 SP RS10365 WP 000588028.1             ...............MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSLNL   66
consensus                                                         **!!!!**!!!**!***!***!**!*!****! ******!!**!!**!***!!**!*******!**

logo
NTDB id 397 FSA28 RS09390 WP 002263438.1          KSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES...   134
NTDB id 388 SMU RS09020 WP 002263438.1            KSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES...   134
NTDB id 432 SGO RS09405 WP 012130939.1            SSQEISNGYARLPIPKGIQAPESTQIYFDKAGGNSSLSKVQFQTK.EGLVTYQLYIGNGKFKKTTN...   142
NTDB id 177425 A4H00 RS03445 WP 067087315.1       TPQQITNGYQSLYLPESVQVDNGQKIVFDKDGGNSSLAKLQFLTD.QKIVRYQLYLGSGRYKKTEE...   145
NTDB id 503 SM12261 RS00960 WP 029689647.1        DGKTISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKD   144
NTDB id 531 SMSK321 RS09225 WP 000588032.1        DGQTISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTG.KGTIRYQLYLGNGKIKRIKETKN   134
NTDB id 273 KZH43 RS09250 WP 000588026.1          DGQMISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 232 SPD RS09880 WP 000588026.1            DGQMISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 198 SPR RS09415 WP 000588026.1            DGQMISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
NTDB id 163 SP RS10365 WP 000588028.1             DGQTLSNGSQKLPVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTS.KGAIRYQLYLGNGKIKRIKETKN   134
consensus                                         *** **!!***!**!************!!**!!!!!!*!**!*!* *****!!!!*!*!**!*******
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X non conserved

X similar

X $\geq$ 50% conserved