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logo

NTDB id 432 SGO RS09405 WP 012130939.1              MVRTIVRLRQLPIKAFTVLESLLVLMISSFILLALSSSVQATFEQIQAKIFFLEFEHFYQESQKLSVSSQRKLVL    75
NTDB id 531 SMSK321 RS09225 WP 000588032.1          ...........MIKAFTMLESLLVLGLVSIVALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSL    64
NTDB id 503 SM12261 RS00960 WP 029689647.1          .MLNKIPVKPLQIKAFTMLESLLVLGLVSILALGLSGSVQSTFAAVEEQIFFMEFEELYRETQKRSVASQQKTSL    74
NTDB id 273 KZH43 RS09250 WP 000588026.1            ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSL    64
NTDB id 232 SPD RS09880 WP 000588026.1              ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSL    64
NTDB id 198 SPR RS09415 WP 000588026.1              ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSL    64
NTDB id 163 SP RS10365 WP 000588028.1               ...........MIKAFTMLESLLVLGLVSILALGLSGSVQSTFSAVEEQIFFMEFEELYRETQKRSVASQQKTSL    64
NTDB id 1111662 HMPREF9319 RS00400 WP 003062922.1   .....MKLKNKTVSAFTLLESLLTLGIACFIIMMLSGSLNGIFQNVEEKIFFLSFENLYCDTQKLANARQQSFTL    70
NTDB id 397 FSA28 RS09390 WP 002263438.1            .........MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTL    66
NTDB id 388 SMU RS09020 WP 002263438.1              .........MSRIKAFTLIESLVTLAITSFLILSFSGSITQTFAKVEERLFFLSFEHLYRDTQKLSVYQRQDMTL    66
consensus                                                   **!!!**!!!**!**********!*!****! ******!!**!!**!***!!**********!

logo

NTDB id 432 SGO RS09405 WP 012130939.1              EISSQEISNGYARLPIPKGIQAPESTQIYFDKAGGNSSLSKVQFQTKE.GLVTYQLYIGNGKFKKTTN......    142
NTDB id 531 SMSK321 RS09225 WP 000588032.1          NLDGQTISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTGK.GTIRYQLYLGNGKIKRIKETKN...    134
NTDB id 503 SM12261 RS00960 WP 029689647.1          NLDGKTISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKD...    144
NTDB id 273 KZH43 RS09250 WP 000588026.1            NLDGQMISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKN...    134
NTDB id 232 SPD RS09880 WP 000588026.1              NLDGQMISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKN...    134
NTDB id 198 SPR RS09415 WP 000588026.1              NLDGQMISNGSQKLTVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKN...    134
NTDB id 163 SP RS10365 WP 000588028.1               NLDGQTLSNGSQKLPVPKGIQAPSGQSITFDRAGGNSSLAKVEFQTSK.GAIRYQLYLGNGKIKRIKETKN...    134
NTDB id 1111662 HMPREF9319 RS00400 WP 003062922.1   TVSQEEVSTDFSAVKLPPSVKVDKTYQVNFDKAGGNSSLAKLAFQTSD.KEFNYQLYLGSGKYKKTETKSLHTP    143
NTDB id 397 FSA28 RS09390 WP 002263438.1            ILKSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES......    134
NTDB id 388 SMU RS09020 WP 002263438.1              ILKSEYISNGVEVLKIPKDVKLERNKTLHFDQAGGNSSLEKLVFQTSDEKRVTYQLYIGSGQYKKTES......    134
consensus                                           ***** *!********!***********!!*!!!!!!!*!**!!!** ****!!!!*!*!**!*******
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X non conserved

X similar

X $\geq$ 50% conserved