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logo

NTDB id 128661 BF38 RS03020 WP 001013197.1  MNGIESFANMILKEACRVQASDLHIVPRQKDVVVQLRIGKDLMTKYCIEKEFGEKLVSHFKFLASMDIGERRKPQNGSLYLQMDG   85
NTDB id 96 BSU 24730 NP 390353.1            MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLTLKLKE   85
consensus                                   !  !!      ** !!*   !!!*!!!!! *! *** !* * !* ! ** !! * *!*!!!!!!  !!!!!!!!!!!!!! ! *

logo

NTDB id 128661 BF38 RS03020 WP 001013197.1  QEVYLRLSTLPTVYQESLVIRLHLQASIQPLSHLSLFPSTAKILLSFLRYSHGLLVFTGPTGSGKTTTMYALLEVIRKKKTRRIV   170
NTDB id 96 BSU 24730 NP 390353.1            GNVHLRMSTLPTINEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQYAKKHFNRNIV   170
consensus                                     !*!!*!!!!!*  !!!!!!***!  !*** *!!!!! !*  !!!!!**!!!*!*!!!!!!!!!!!!*! !*   *!*  ! !!

logo

NTDB id 128661 BF38 RS03020 WP 001013197.1  TLEDPVEKRNDDVLQIQINEKAGITYEAGLKAILRHDPDVILVGEIRDEETAKIAIRASLTGHLVMTTLHTNDARGAILRFMDFG   255
NTDB id 96 BSU 24730 NP 390353.1            TLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSLHTRDAKGAIYRLLEFG   255
consensus                                   !!!!!!! ! *!!!!*!*!!!!!*!! !!!!!!!!!!!!*!**!!!!! !!! !!*!! *!!!!!*!*!!! !!*!!! ! **!!

logo

NTDB id 128661 BF38 RS03020 WP 001013197.1  ITRQEIEQSLLAIAAQRLVELKCPFCKRKCST.LCKSMRQVRQASIYELLYGYELKQAIKEANGECVTYKHETLQSSIRKGYALG   339
NTDB id 96 BSU 24730 NP 390353.1            INMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGNHANYQYQTLRQIIRKGIALG   340
consensus                                   !  *!!!!***!!!!!!!!*! !!!!   !!** !*  !* ! !!*!!!!!!  ! !*! !! !    ! * !!   !!!! !!!

logo

NTDB id 128661 BF38 RS03020 WP 001013197.1  FLEEDVYV........   347
NTDB id 96 BSU 24730 NP 390353.1            YLTTNNYDRWVYHEKD   356
consensus                                   *!    ! ********
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X non conserved

X similar

X ≥ 50% conserved