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logo

NTDB id 1142437 DQN84 RS08380 WP 066137453.1  .MSIEKIADLLLKQAIQLTATDIHITPRKHNYHIQFRLHGLLTTIQSIPFQAGERLISHLKFMSSIDISEKRKPQSGSFE   79
NTDB id 96 BSU 24730 NP 390353.1              MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT   80
consensus                                     * !!!!*   !*  !    !*!!!! !! *  *!*!!*** !     **   * !!!!! !!*! *!! !*!!!! !!

logo

NTDB id 1142437 DQN84 RS08380 WP 066137453.1  LNLNNQLIALRISTLPTTLAKESLVIRILPQDHSFILEKMSLFPSTSPILKSFMLHAHGMLIFTGPTGSGKSTTMYAVAE   159
NTDB id 96 BSU 24730 NP 390353.1              LKLKEGNVHLRMSTLPT.INEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQ   159
consensus                                     ! !    **!!*!!!!!**  !!!!!!**!! * * **!*!!!! ! * ! !!* ! !!!!!!!!!!!!!!*!!*! *

logo

NTDB id 1142437 DQN84 RS08380 WP 066137453.1  HCAGTLNRRVVTLEDPVEKQSDMLLQVQLNDKAGITYSTGLKAILRHDPDVIIIGEIRDAETAHIAIRAALTGHLVLTSL   239
NTDB id 96 BSU 24730 NP 390353.1              YAKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSL   239
consensus                                     **  * !! *!!!!!!!!   * *!!!!*!*!!!*!!! !!!!!!!!!!!*!!*!!!!!!!!!*!!*!!!*!!!!!!!!!

logo

NTDB id 1142437 DQN84 RS08380 WP 066137453.1  HTRDAKGAIYRLLEFGVKLQDIEQTLIGIMAQRLIALLCPLCGESCSKFCTR..RVAKRTGVYEILYGQALVGALEESKG   317
NTDB id 96 BSU 24730 NP 390353.1              HTRDAKGAIYRLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKG   319
consensus                                     !!!!!!!!!!!!!!!!* ***!!!!*!*! !!!!* ! !! !   !!  **!**!  *!  !!!*!!!  !  ** ! !!

logo

NTDB id 1142437 DQN84 RS08380 WP 066137453.1  GKEVYHYPLIKDLIRKGIALGYVPVEEYKHWVLEE..   352
NTDB id 96 BSU 24730 NP 390353.1              NHANYQYQTLRQIIRKGIALGYLTTNNYDRWVYHEKD   356
consensus                                      *  !*!* ** *!!!!!!!!!**   ! *!! *!**
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X non conserved

X similar

X ≥ 50% conserved