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logo

NTDB id 1011284 ABR335 RS07960 WP 017550534.1  .MSVEKIAELLIGQAVKNNVTDVHIVPKEKHYHVQFRQYGRLYPHRNLSAKAGERLISHLKFMSSMDISEKRKPQSGSFA   79
NTDB id 96 BSU 24730 NP 390353.1               MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT   80
consensus                                      * !*!!*   !!  !     *!*!!!!*!** ***!!  * ! **! *    * !!!!! !!*! !!! !*!!!! !!

logo

NTDB id 1011284 ABR335 RS07960 WP 017550534.1  MNVLQQTISLRISTLPTSLSKESLVIRILPHEEKFQLSQISLFPSSTKKLMALLNHSHGMLIFSGPTGSGKSTTMYTLVE  159
NTDB id 96 BSU 24730 NP 390353.1               LKLKEGNVHLRMSTLPT.INEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQ  159
consensus                                      * *    **!!*!!!!!**  !!!!!!**!*   * *  *!!!! *   !*  ! !!!!!!!!!*!!!!!!!*!!*!*!!

logo

NTDB id 1011284 ABR335 RS07960 WP 017550534.1  QCAKRFLRNVITLEDPVEKQSDSFLQVQVNEKAGITYSTGLKAILRHDPDIILVGEIRDAETARIAIRASLTGHLVLTTL  239
NTDB id 96 BSU 24730 NP 390353.1               YAKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSL  239
consensus                                       * !*! !!**!!!!!!!   *  !!!!!!!!!!*!!! !!!!!!!!!!!*!**!!!!!!!!! !!*!! *!!!!!!!*!

logo

NTDB id 1011284 ABR335 RS07960 WP 017550534.1  HTRDAKGAIYRLMEFGVSIHEMEQTLLAVSAQRLITLRCPVCGGIRCPHDSQDCPRSFEKRQTAVYELLYGKNLQRVLKE  319
NTDB id 96 BSU 24730 NP 390353.1               HTRDAKGAIYRLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENG...CSSVYCRQSFNTRRASVYELLYGKNLQQCIQE  316
consensus                                      !!!!!!!!!!!!*!!!* **!*!!!**!* !!!!* ! !! !   ***  !  !* !!  !   !!!!!!!!!!! ** !

logo

NTDB id 1011284 ABR335 RS07960 WP 017550534.1  AKGEQVYYHYPTLKEWLRKGIVLGYISEEEYHRWIAEEEEADKPQTAGALSYPGR  374
NTDB id 96 BSU 24730 NP 390353.1               AKGNHANYQYQTLRQIIRKGIALGYLTTNNYDRWVYHEKD...............  356
consensus                                      !!! *  !*!*!!*  *!!!! !!!**   !*!!* *! ***************
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X non conserved

X similar

X ≥ 50% conserved