```
logo
NTDB id 339123 ESP47 RS05960 WP 014098021.1  .MSVEKIAELLIGQAVKNNVTDVHIVPKEKHYHVQFRQYGRLYPHRNLSAKAGERLISHLKFMSSMDISEKRKPQSGSFA   79
NTDB id 96 BSU 24730 NP 390353.1             MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT   80
consensus                                    * !*!!*   !!  !     *!*!!!!*!** ***!!  * ! **! *    * !!!!! !!*! !!! !*!!!! !!

logo
NTDB id 339123 ESP47 RS05960 WP 014098021.1  MNVLQQTISLRISTLPTSLSKESLVIRILPHEEKFQLSQISLFPSSTKKLMALLNHSHGMLIFSGPTGSGKSTTMYTLVE  159
NTDB id 96 BSU 24730 NP 390353.1             LKLKEGNVHLRMSTLPT.INEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQ  159
consensus                                    * *    **!!*!!!!!**  !!!!!!**!*   * *  *!!!! *   !*  ! !!!!!!!!*!!!!!!!*!!*!*!!

logo
NTDB id 339123 ESP47 RS05960 WP 014098021.1  HCAKRFLRNVITLEDPVEKQSDSFLQVQVNEKAGITYSTGLKAILRHDPDIILVGEIRDAETARIAIRASLTGHLVLTTL  239
NTDB id 96 BSU 24730 NP 390353.1             YAKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSL  239
consensus                                    ** !*! !!**!!!!!!!   *  !!!!!!!!!!*!!! !!!!!!!!!!!*!**!!!!!!!!! !!*!! *!!!!!!!*!

logo
NTDB id 339123 ESP47 RS05960 WP 014098021.1  HTRDAKGAIYRLMEFGVSIHEMEQTLLAVSAQRLITLRCPVCGGIRCPHDSQDCPRSFEKRQTAVYELLYGKNLQRVLKE  319
NTDB id 96 BSU 24730 NP 390353.1             HTRDAKGAIYRLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENG...CSSVYCRQSFNTRRASVYELLYGKNLQQCIQE  316
consensus                                    !!!!!!!!!!!!*!!!* **!*!!**!* !!!!* ! !! !   ***  !  !* !!  !   !!!!!!!!!!! ** !

logo
NTDB id 339123 ESP47 RS05960 WP 014098021.1  AKGEQVYYHYPTLKEWLRKGIVLGYISEEEYHRWIAEEEEADKPQTAGALSYPGR  374
NTDB id 96 BSU 24730 NP 390353.1             AKGNHANYQYQTLRQIIRKGIALGYLTTNNYDRWVYHEKD...............  356
consensus                                    !!! *  !*!*!!*  *!!!! !!!**   !*!!* *! ***************
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X non conserved

X similar

X $\geq$ 50% conserved