```
logo
NTDB id 997761 KZZ19 RS20740 WP 237979491.1  MNGIEIFANTILKEACRVQASDLHIVPRQKDVVVQLRIGKDLMTKHCIEKEFGEKLVSHFKFLASMDIGEKRKPQNGSLY   80
NTDB id 96 BSU 24730 NP 390353.1             MDSIEKVSKNLIEEAYLTKASDIHIVPREKDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT   80
consensus                                    !  !!     ** !!*   !!!*!!!!! *! *** !* * !* !*** !! * *!*!!!!!!  !!!!!*!!!!!!!!

logo
NTDB id 997761 KZZ19 RS20740 WP 237979491.1  LQMDGQEVYLRLSTLPTVYQESLVIRLHLQASIQPLSHLSLFPSTAKKLLSFLSYSHGLLVFTGPTGSGKTTTMYALLEV  160
NTDB id 96 BSU 24730 NP 390353.1             LKLKEGNVHLRMSTLPTINEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQY  160
consensus                                    ! *    !*!!*!!!!!*  !!!!!!***!  !*** *!!!!! !*  !!!!! *!!!*!*!!!!!!!!!!!!*! !*

logo
NTDB id 997761 KZZ19 RS20740 WP 237979491.1  IRKKKTRRIVTLEDPVEKRNDDVLQIQINEKAGITYEAGLKAILRHDPDIILVGEIRDEETAKIAIRASLTGHLVMTTLH  240
NTDB id 96 BSU 24730 NP 390353.1             AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSLH  240
consensus                                     *!*  ! !!!!!!!!! ! *!!!!*!*!!!!!*!! !!!!!!!!!!!!*!**!!!!! !!! !!*!! *!!!!!*!*!!

logo
NTDB id 997761 KZZ19 RS20740 WP 237979491.1  TNDAKGAILRFMDFGITRQEIEQSLLAIAAQRLVELKCPFCQRMCST.LCKSMRQVRQASIYELLYGYELKQAIKEANGE  319
NTDB id 96 BSU 24730 NP 390353.1             TRDAKGAIYRLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN  320
consensus                                    ! !!!!!! ! **!!!  *!!!!***!!!!!!!!*! !!!!   !!** !*  !* ! !!*!!!!!!  ! !*! !! !

logo
NTDB id 997761 KZZ19 RS20740 WP 237979491.1  CVTYKHETLESLIRKGYALGFLEEDVYV........  347
NTDB id 96 BSU 24730 NP 390353.1             HANYQYQTLRQIIRKGIALGYLTTNNYDRWVYHEKD  356
consensus                                       ! * !!  *!!!! !!!*!    ! ********
```

X non conserved

X similar

X $\geq$ 50% conserved