```
logo
NTDB id 275061 C5695 RS12360 WP 117730990.1  MYGIEYLGQELLEEACRMRASDVHIVPKEKEALVSFRVDSDLMQQRIIDKKSGERLIAHFKFLSSMDIGEKRRPQNGSLA   80
NTDB id 96 BSU 24730 NP 390353.1             MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT   80
consensus                                    !  !! *   !*!!!*  *!!!*!!!!*!**!***!!!!* !*  ! * !  * !!! !!!!!! !!!!!*!*!!!!!!

logo
NTDB id 275061 C5695 RS12360 WP 117730990.1  VMLRSGQVYIRMSTLPTVNDESLVIRILPQNHVPTINHLSLFPKASARLLSFLNHSHGLILFTGPTNSGKTTTLYSLIQF  160
NTDB id 96 BSU 24730 NP 390353.1             LKLKEGNVHLRMSTLPTINEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQY  160
consensus                                    * !* !*!**!!!!!!!*!*!!!!!!**!! **!*! *!!!!!!  ! !!!!! !!!!***!!!!! !!!!!!!!!!*!*

logo
NTDB id 275061 C5695 RS12360 WP 117730990.1  AKKNFNRNIITLEDPVETRNEEVLQVQVNEKAGITYAAGLRAILRHDPDMIVLGEIRDAETARTAIRAALTGHLVMSTLH  240
NTDB id 96 BSU 24730 NP 390353.1             AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSLH  240
consensus                                    !!!*!!!!!*!!!!!!!!! !*!!!!!!!!!!!*!! !!!*!!!!!!!!!!!*!!!!!!!!!  !*!!!*!!!!!***!!

logo
NTDB id 275061 C5695 RS12360 WP 117730990.1  AKNAKGALYRMLEFGVTMNELEQTMVAIAAQRLIELTCPFCGEACQ.LYCKLNRQVRRTNVFELLFGKELGECIKEAKGE  319
NTDB id 96 BSU 24730 NP 390353.1             TRDAKGAIYRLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN  320
consensus                                     * !!!!*!!*!!!!* !!!*!!!**!!!!!!!**! !!!!  *! **!!*  !* !!  !*!!!*!! !  !! !!!!

logo
NTDB id 275061 C5695 RS12360 WP 117730990.1  YAHSSYETLQRLIRKGVALGYLSKSTYHRWVYEEASL  356
NTDB id 96 BSU 24730 NP 390353.1             HANYQYQTLRQIIRKGIALGYLTTNNYDRWVYHEKD.  356
consensus                                    *!*  ! !!  *!!!!*!!!!!*   !*!!!!*!  *
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X non conserved

X similar

X $\geq$ 50% conserved