```
logo

NTDB id 986930 MHB67 RS11360 WP 103131947.1  MYGIEYLRQELLEEACRMRASDVHIVPKEKEASVSFRVDSDLIQQRMIDKKSGERLIAHFKFLSSMDIGEKRRPQNGSLA   80
NTDB id 96 BSU 24730 NP 390353.1             MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT   80
consensus                                    !  !! *   !*!!!*  *!!!*!!!!*!**! **!!!!* !*  ! * !  * !!! !!!!!! !!!!!*!*!!!!!!

logo

NTDB id 986930 MHB67 RS11360 WP 103131947.1  VMLRKGQVFIRMSTLPTVNDESLVIRILPQDHVPKIKHLSLFPKASARLLSFLNHSHGLILFTGPTNSGKTTTLYSLIQF  160
NTDB id 96 BSU 24730 NP 390353.1             LKLKEGNVHLRMSTLPTINEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQY  160
consensus                                    * !* !*!**!!!!!!!*!*!!!!!!**!! **! ! *!!!!!!  ! !!!!! !!!!***!!!!! !!!!!!!!!!*!*

logo

NTDB id 986930 MHB67 RS11360 WP 103131947.1  AKKNFNRNIITLEDPVETRNEEVLQVQVNEKAGITYAAGLRAILRHDPDVIVLGEIRDAETARTAIRAALTGHLVLSTLH  240
NTDB id 96 BSU 24730 NP 390353.1             AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSLH  240
consensus                                    !!!*!!!!!*!!!!!!!!! !*!!!!!!!!!!!*!! !!!*!!!!!!!!*!*!!!!!!!!!  !*!!!*!!!!!!**!!

logo

NTDB id 986930 MHB67 RS11360 WP 103131947.1  AKNAKGALYRMLEFGVTMNELEQTMVAIAAQRLIELTCPFCGERCQ.LYCKLNRPIRRTNVFELLFGKELGECIKEAKGE  319
NTDB id 96 BSU 24730 NP 390353.1             TRDAKGAIYRLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN  320
consensus                                     * !!!!*!!*!!!!* !!!*!!!**!!!!!!!**! !!!!    ! **!!*  !* !!  !*!!!*!! !  !! !!!!

logo

NTDB id 986930 MHB67 RS11360 WP 103131947.1  YAHSSYETLQRLIRKGVALGYLSKNTYHRWVYEEASL  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