```
logo

NTDB id 995308 AAVC46 RS06855 WP 034661296.1  MYGIEYLGQELLEEACRMRASDVHIVPKEKEASVSFRVDSDLIQQRTIDKKSGERLIAHFKFLSSMDIGEKRRPQNGSLA   80
NTDB id 96 BSU 24730 NP 390353.1              MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT   80
consensus                                     !  !! *   !*!!!*  *!!!*!!!!*!**! **!!!!* !*  ! * !  * !!! !!!!!! !!!!!*!*!!!!!!

logo

NTDB id 995308 AAVC46 RS06855 WP 034661296.1  VMLRSGQVFIRMSTLPTVNDESLVIRILPQDHVPKIKHLSLFPKASTRLLSFLNHSHGLILFTGPTNSGKTTTLYSLIQF  160
NTDB id 96 BSU 24730 NP 390353.1              LKLKEGNVHLRMSTLPTINEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQY  160
consensus                                     * !* !*!**!!!!!!!*!*!!!!!**!! **! ! *!!!!!!    !!!!! !!!!***!!!!! !!!!!!!!!!*!*

logo

NTDB id 995308 AAVC46 RS06855 WP 034661296.1  AKKNFNRNIITLEDPVETRNEEVLQVQVNEKAGITYAAGLRAILRHDPDLIVLGEIRDAETARTAIRAALTGHLVMSTLH  240
NTDB id 96 BSU 24730 NP 390353.1              AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSLH  240
consensus                                     !!!*!!!!!*!!!!!!!!!! !*!!!!!!!!!!!*!! !!!*!!!!!!!!*!*!!!!!!!!!!  !*!!!*!!!!!***!!

logo

NTDB id 995308 AAVC46 RS06855 WP 034661296.1  AKNAKGALYRMLEFGVTMNELEQTMVAIAAQRLIELTCPFCGETCQ.LYCKLNRPVRRTNVFELLFGKELGECIKEAKGE  319
NTDB id 96 BSU 24730 NP 390353.1              TRDAKGAIYRLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN  320
consensus                                      * !!!!*!!*!!!!* !!!*!!!**!!!!!!!**! !!!!   ! **!!*  !* !!  !*!!!*!! !  !! !!!!

logo

NTDB id 995308 AAVC46 RS06855 WP 034661296.1  YAHSSYETLQRLIRKGVALGYLSKNTYHRWVYEEASL  356
NTDB id 96 BSU 24730 NP 390353.1              HANYQYQTLRQIIRKGIALGYLTTNNYDRWVYHEKD.  356
consensus                                     *!*  ! !!  *!!!!*!!!!!* ! !*!!!!*!  *
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X non conserved

X similar

X ≥ 50% conserved