```
logo
NTDB id 174656 B14 RS09055 WP 003183477.1  MYTIESLSGRLIDEAYRMKASDIHIVPGEKEAVVRFRIDDELFQKDGLTRNECSRLISHFKFLSSMDIGERRLPQSGALTLYINR   85
NTDB id 96 BSU 24730 NP 390353.1           MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLTLKLKE   85
consensus                                  ! *!! *!   !!*!!!  !!!!!!!!! !**!***!!*!* !  !  * * !! !!!!!!!!! !!!!!!! !! ! !!! *

logo
NTDB id 174656 B14 RS09055 WP 003183477.1  EPVHLRMSTLPTIHDESLVIRLLPKMSSKPLTKLSLFPSATFKLLSFLKHSHGLMLFTGPTGSGKTTTLYSLIDYAKRHFKRNII  170
NTDB id 96 BSU 24730 NP 390353.1           GNVHLRMSTLPTINEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQYAKKHFNRNIV  170
consensus                                   *!!!!!!!!!!!**!!!!!!**!    *** !!!!!!     !!!!!!!!!!***!!!!!!!!!!!!!!!!!* !!!*!! !!!*

logo
NTDB id 174656 B14 RS09055 WP 003183477.1  TLEDPVESRSENYLQVQVNEKAGMTYSAGLKAVLRHDPDMIILGEIRDAETAQIAVRAALTGHLVLSSMHAKNAKGAIYRLLELG  255
NTDB id 96 BSU 24730 NP 390353.1           TLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSLHTRDAKGAIYRLLEFG  255
consensus                                  !!!!!!!*! ! *!!!!!!!!!!*!!!!!!!!*!!!!!!!!!!!!!!!!!!! !!!!!!*!!!!!!*!*! * !!!!!!!!!! !

logo
NTDB id 174656 B14 RS09055 WP 003183477.1  VDMTEIEQTLVAVSAQRLVDIVCPFCGDSCT.LYCRLSRPVRRASVFELLYGKSLNLCIEEAKGRCGGIKTDTLKMLIRKGIALG  339
NTDB id 96 BSU 24730 NP 390353.1           INMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGNHANYQYQTLRQIIRKGIALG  340
consensus                                  * ! !!!!!**!* !!!!!!* !!!!   !***!!! !!* !!!!!*!!!!!! !* !! !!!!  *     !!* *!!!!!!!!

logo
NTDB id 174656 B14 RS09055 WP 003183477.1  YLPSKTYERWIGHED.   354
NTDB id 96 BSU 24730 NP 390353.1           YLTTNNYDRWVYHEKD   356
consensus                                  !!**  !*!!* !! *
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X non conserved

X similar

X ≥ 50% conserved