```
logo
NTDB id 221645 BLMD RS13525 WP 080131442.1  MYTIESLSGRLIDEAYRMKASDIHIVPGEKEAVVRFRIDDELFQKDRLTRNECSRLISHFKFLSSMDIGKRRLPQSGALTLYINR   85
NTDB id 96 BSU 24730 NP 390353.1            MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLTLKLKE   85
consensus                                   ! *!! *!  !!*!!!  !!!!!!!!! !**!***!!*!* !  !  * * !! !!!!!!!!! !!!! !! !! ! !!! *

logo
NTDB id 221645 BLMD RS13525 WP 080131442.1  QPVHLRMSTLPTIHDESLVIRLLPKMSSKPLTKLSLFPSATFKLLSFLKHSHGLILFTGPTGSGKTTTLYSLIEYAKRHFKRNII  170
NTDB id 96 BSU 24730 NP 390353.1            GNVHLRMSTLPTINEESLVIRVMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQYAKKHFNRNIV  170
consensus                                    *!!!!!!!!!!!**!!!!!!**!    *** !!!!!!     !!!!!!!!!!***!!!!!!!!!!!!!!!!!* !!!*!! !!!*

logo
NTDB id 221645 BLMD RS13525 WP 080131442.1  TLEDPVESRSENILQVQVNEKAGMTYSAGLKAVLRHDPDMIILGEIRDAETAQIAVRAALTGHLVLSSMHAKNAKGAIYRLLEFG  255
NTDB id 96 BSU 24730 NP 390353.1            TLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAEIAVRAAMTGHLVLTSLHTRDAKGAIYRLLEFG  255
consensus                                   !!!!!!!*! ! *!!!!!!!!!!*!!!!!!!!*!!!!!!!!!!!!!!!!!!!! !!!!!!*!!!!!!*!*! * !!!!!!!!!!!!

logo
NTDB id 221645 BLMD RS13525 WP 080131442.1  VDMTEIEQTLVAISAQRLVGLVCPFCGDSCS.LYCRLSRPVRRASVFELLYGKSLNLCIEEAKGRCGDIKTETLKTLIQKGIALG  339
NTDB id 96 BSU 24730 NP 390353.1            INMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGNHANYQYQTLRQIIRKGIALG  340
consensus                                   * ! !!!!!**!! !!!!! ! !!!!   !!**!!! !!* !!!!!*!!!!!! !* !! !!!!  *     !!* *! !!!!!!

logo
NTDB id 221645 BLMD RS13525 WP 080131442.1  YLPSNTYERWIGHED.   354
NTDB id 96 BSU 24730 NP 390353.1            YLTTNNYDRWVYHEKD   356
consensus                                   !!**! !*!!* !! *
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X non conserved

X similar

X ≥ 50% conserved