```
logo
NTDB id 127135 JL26 RS09225 WP 005711167.1  MKYIDAKTQHIRNLLYPIIGETKAKTYFSYYGIMKDKAMFALYKD..DKFYLHIPNHCLEENIA.KNLSLLLDSQTGI.NLKSFY   81
NTDB id 1391 A4U84 RS05295 WP 021111957.1   MKYIDVKTQHIRNLLYPIIGETQAKTYFSYYGIMKDGALFALYKN..SKFYLHISGNCLHEVLTNPNISQLSDSKYGI.NSKSFY   82
NTDB id 1451 ASU2 RS08740 WP 015674037.1   MKYTTKELERLKALLS.PLGEVTFKPYFSFLGIFKNDTMFALYKD..QHLYVRKSAQYLEEITQTISIHFLVDRHINC.RSKIFY   81
NTDB id 1376 D1108 RS09055 WP 005609196.1  MAISPKKFQYLKEIFS.PLGEINFKSYFSYLGIFKDDTMFALYDHKNDRLYLRKSAQFYPDIIRTIPIHFLIDRRIGKQQSHIFY   84
consensus                                  !************* ***!!** !*!!!* !! !** *!!!!**  ** !** ********* ******!*! **** *** !!

logo
NTDB id 127135 JL26 RS09225 WP 005711167.1  LIPPDLLNDLQQVSHWVVESVKDILHTKQNQYMEKKKCIRTLPNMTIQLERTLKKLGVYSIDDLVDRGEIDIFVNLLKIGVDADQ   166
NTDB id 1391 A4U84 RS05295 WP 021111957.1   LIPTEILYDLKPISHWITNGIQEIINTKHTQYIKKKRCIRTLPNMTIQLERTLKKLGINSIDELIDRGEVDIFVNLLKMGIDADQ   167
NTDB id 1451 ASU2 RS08740 WP 015674037.1   LLPSSILNNLQTYSHWITSAIHEYQQSKAKIENQNQNKIRMLPNLNISIERLLAKVGIYTVTDLKATGAINVFVRLIQQGLEATA   166
NTDB id 1376 D1108 RS09055 WP 005609196.1  LIPSSIIHNLHLYTHWILSAIEEYQTAKAKLISQNKNKIRLLPNLNINIERLLARIEIYTVDDLKNVGVINAFVKLIMLGLEVTE   169
consensus                                  !*!***** !* **!!***** * ***!****** ***!!*!!!* !**!! ! *********!***!** *!!*!***!*** *

logo
NTDB id 127135 JL26 RS09225 WP 005711167.1  ALLFRLHGAINRQYIYTISDKTKQNLLNDADNALYAAGLRKRFNTR...   212
NTDB id 1391 A4U84 RS05295 WP 021111957.1   AILFRLYGAINRQYIYTISDKTKQDLLNDANHALYEAGLRKRFNVK...   213
NTDB id 1451 ASU2 RS08740 WP 015674037.1   LLLFKLYSAIQYKYIYMLTEQEKQNLLIEADNALYQAGLRKRFMLCVK.   214
NTDB id 1376 D1108 RS09055 WP 005609196.1  LLLFKLYAALEHKYIYMLSKQEKQSLLIEADLSLYNAGLRKRFAISQAN   218
consensus                                   *!!*!**!*** !!! ***  !!*!! *!***!!*!!!!!!!***
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X non conserved

X similar

X $\geq$ 50% conserved