```
logo
NTDB id 290770 C3363 RS11145 WP 005711167.1  MKYIDAKTQHIRNLLYPIIGETKAKTYFSYYGIMKDKAMFALYKD..DKFYLHIPNHCLEENIA.KNLSLLLDSQTGI.N   76
NTDB id 1391 A4U84 RS05295 WP 021111957.1    MKYIDVKTQHIRNLLYPIIGETQAKTYFSYYGIMKDGALFALYKN..SKFYLHISGNCLHEVLTNPNISQLSDSKYGI.N   77
NTDB id 1451 ASU2 RS08740 WP 015674037.1     MKYTTKELERLKALLS.PLGEVTFKPYFSFLGIFKNDTMFALYKD..QHLYVRKSAQYLEEITQTISIHFLVDRHINC.R   76
NTDB id 1376 D1108 RS09055 WP 005609196.1    MAISPKKFQYLKEIFS.PLGEINFKSYFSYLGIFKDDTMFALYDHKNDRLYLRKSAQFYPDIIRTIPIHFLIDRRIGKQQ   79
consensus                                    !************** ***!!** !*!!!* !! !** *!!!!**  ** !** ********* ******!*! **** *

logo
NTDB id 290770 C3363 RS11145 WP 005711167.1  LKSFYLIPPDLLNDLQQVSHWVVESVKDILHTKQNQYMEKKKCIRTLPNMTIQLERTLKKLGVYSIDDLVDRGEIDIFVN   156
NTDB id 1391 A4U84 RS05295 WP 021111957.1    SKSFYLIPTEILYDLKPISHWITNGIQEIINTKHTQYIKKKRCIRTLPNMTIQLERTLKKLGINSIDELIDRGEVDIFVN   157
NTDB id 1451 ASU2 RS08740 WP 015674037.1     SKIFYLLPSSILNNLQTYSHWITSAIHEYQQSKAKIENQNQNKIRMLPNLNISIERLLAKVGIYTVTDLKATGAINVFVR   156
NTDB id 1376 D1108 RS09055 WP 005609196.1    SHIFYLIPSSIIHNLHLYTHWILSAIEEYQTAKAKLISQNKNKIRLLPNLNINIERLLARIEIYTVDDLKNVGVINAFVK   159
consensus                                    ** !!!*!***** !* **!!***** * ***!****** ***!!*!!!* !**!! ! *********!***!** *!!*

logo
NTDB id 290770 C3363 RS11145 WP 005711167.1  LLKIGVDADQALLFRLHGAINRQYIYTISDKTKQNLLNDADNALYAAGLRKRFNTR...   212
NTDB id 1391 A4U84 RS05295 WP 021111957.1    LLKMGIDADQAILFRLYGAINRQYIYTISDKTKQDLLNDANHALYEAGLRKRFNVK...   213
NTDB id 1451 ASU2 RS08740 WP 015674037.1     LIQQGLEATALLLFKLYSAIQYKYIYMLTEQEKQNLLIEADNALYQAGLRKRFMLCVK.   214
NTDB id 1376 D1108 RS09055 WP 005609196.1    LIMLGLEVTELLLFKLYAALEHKYIYMLSKQEKQSLLIEADLSLYNAGLRKRFAISQAN   218
consensus                                    !***!*** * *!!*!**!*** !!! ***  !!*!! *!***!!*!!!!!!!***
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X non conserved

X similar

X $\geq$ 50% conserved