```
logo
NTDB id 1123729 EL276 RS06405 WP 107879585.1  MKLKTVLLTAVAALA.......................LTANVYAATEGKDYTVLPKPIPQSQADKIEVLEFFGYFCVHCY   58
NTDB id 1099 NMB RS01545 WP 002215704.1       MKLKTLALTSLTLLALAACSKQAETSVPADSAQSSSSAPAAPAELNEGVNYTVLSTPIPQQQAGKIEVLEFFGYFCPHCA   80
NTDB id 1098 NMB RS01465 WP 002224844.1       MKSRHLALGVAALFALAACDSKVQTSVPADSAPAA.SAAAAPAGLVEGQNYTVLANPIPQQQAGKVEVLEFFGYFCPHCA   79
consensus                                     !!*****!* *****!****     ********    ** !*****  !! *!!!!  !!!!*!!*!*!!!!!!!!!!*!!*

logo
NTDB id 1123729 EL276 RS06405 WP 107879585.1  HLDPVLLKHAKTFPSDTYLRTEHVVWQPEMLGFARVAAAVNSSGL..KYQANQAVFRAVYEEKINLADSATFKQWAAAQK  136
NTDB id 1099 NMB RS01545 WP 002215704.1       HLEPVLSEHIKTFKDDTYMRREHVVWGDEMKPLARLAAAVEMAGE..SDKANSHIFDAMVNQKINLADTDTLKKWLSEQT  158
NTDB id 1098 NMB RS01465 WP 002224844.1       HLEPVLSKHAKSFKDDMYLRTEHVVWQKEMLTLARLAAAVDMAAADSKDVANSHIFDAMVNQKIKLQNPEVLKKWLGEQT  159
consensus                                     !!*!!!**!*!*!**!*!*!*!!!!!* !!* *!!*!!!!****    ** !!***!*!****!!*!**********!*!***!*

logo
NTDB id 1123729 EL276 RS06405 WP 107879585.1  SFDGKKLIAAYDSFGNQAQAKRMEDLTNTYQISGTPTVIVGGKYQVKFTGDWQAGMKTIDELVAKVRTERGMKTPSAKPV  216
NTDB id 1099 NMB RS01545 WP 002215704.1       AFDGKKVLAAFEAPESQARAAQMEELTNKFQISGTPTVIVGGKYQVEFK.DWQSGMTTIDQLVDKVREEQKKPQ......  231
NTDB id 1098 NMB RS01465 WP 002224844.1       AFDGKKVLAAYESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEFA.DWESGMNTIDLLADKVREEQKAAQ......  232
consensus                                     *!!!!!**!!******!!*! *!**!!***!!*!!!!!!!!!!!*!*!  !!**!! !!! !**!!!*!**  *

logo
NTDB id 1123729 EL276 RS06405 WP 107879585.1  AVAPALKSKGALLAKAANK   235
NTDB id 1099 NMB RS01545 WP 002215704.1       ....................   231
NTDB id 1098 NMB RS01465 WP 002224844.1       ....................   232
consensus
```

X non conserved

X similar

X ≥ 50% conserved