```
logo
NTDB id 1122874 EL216 RS10685 WP 085390292.1  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 1122874 EL216 RS10685 WP 085390292.1  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 1122874 EL216 RS10685 WP 085390292.1  GFDGKKLIAAYDSFGNQAQAKRMEDLTNTYQISGTPTVIVGGKYQVKFTGDWQAGMKTIDELVAKVRTERGMKTPSAKPV  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 1122874 EL216 RS10685 WP 085390292.1  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