```
logo
NTDB id 1099 NMB RS01545 WP 002215704.1        ..MKLKTLALTSLTLLALAACSKQAETSVPADSAQSSSSAPAAPAELNEGVNYTVLSTPIPQQQAGKIEVLEFFGYFCPH   78
NTDB id 1144510 C7S04 RS10365 WP 010951331.1   MKSRHLALALGVAALFALAACDSKVQTSVPADSAPAA.SAAAAPAGLVEGQNYTVLANPIPQQQAGKVEVLEFFGYFCPH   79
NTDB id 1098 NMB RS01465 WP 002224844.1        MKSRH..LALGVAALFALAACDSKVQTSVPADSAPAA.SAAAAPAGLVEGQNYTVLANPIPQQQAGKVEVLEFFGYFCPH   77
consensus                                      *****  !!!****!*!!!!!*****!!!!!!!!*** !!*!!!!*!*!!*!!!!!**!!!!!!!!!!*!!!!!!!!!!!!

logo
NTDB id 1099 NMB RS01545 WP 002215704.1        CAHLEPVLSEHIKTFKDDTYMRREHVVWGDEMKPLARLAAAVEMAGE..SDKANSHIFDAMVNQKINLADTDTLKKWLSE   156
NTDB id 1144510 C7S04 RS10365 WP 010951331.1   CARLEPVLSKHAKSFKDDMYLRTEHVVWQKEMLPLARLAAAVDMAAAESKDVANSHIFDAMVNQKIKLQEPEVLKKWLGE   159
NTDB id 1098 NMB RS01465 WP 002224844.1        CAHLEPVLSKHAKSFKDDMYLRTEHVVWQKEMLTLARLAAAVDMAAADSKDVANSHIFDAMVNQKIKLQNPEVLKKWLGE   157
consensus                                      !!*!!!!!!*!*!*!!!!*!*!*!!!!!**!!**!!!!!!!!*!!**********!*!!!!!!!!!!!!!!!!*!*****!!!!!*!

logo
NTDB id 1099 NMB RS01545 WP 002215704.1        QTAFDGKKVLAAFEAPESQARAAQMEELTNKFQISGTPTVIVGGKYQVEFKDWQSGMTTIDQLVDKVREEQKKPQ   231
NTDB id 1144510 C7S04 RS10365 WP 010951331.1   QTAFDGKKVLAAYESPESQARAGKMQELTETFQIDGTPTVIVGGKYKVEFADWESGMNTIDLLADKVREEQKAAQ   234
NTDB id 1098 NMB RS01465 WP 002224844.1        QTAFDGKKVLAAYESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEFADWESGMNTIDLLADKVREEQKAAQ   232
consensus                                      !!!!!!!!!!!!!!*!*!!!!!!!**!*!!!**!!!*!!!!!!!!!!!!!!*!!!*!!*!!!*!!!!*!*!!!!!!!!!**!
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X non conserved

X similar

X $\geq$ 50% conserved