```
logo
NTDB id 1099 NMB RS01545 WP 002215704.1      MKLKTLALTSLTLLALAACSKQAETSVPADSAQSSSSAPAAPAELNEGVNYTVLSTPIPQQQAGKIEVLEFFGYFCPHCA   80
NTDB id 668290 MON40 RS01385 WP 003780094.1 MKLKTLTL.ATALLALTACDNKVQTSVPADSAPAASAASAAPVALVEGLNYTTLSTPIPQQQAGKVEVLEFFGYFCPHCA   79
NTDB id 1098 NMB RS01465 WP 002224844.1     MKSRHLALGVAALFALAACDSKVQTSVPADSAPAASAA.AAPAGLVEGQNYTVLANPIPQQQAGKVEVLEFFGYFCPHCA   79
consensus                                   !!***!*!    *!*!!*!!* ***!!!!!!!***!*! !!!*!*!!*!!!*!**!!!!!!!!!*!!!!!!!!!!!!!!!

logo
NTDB id 1099 NMB RS01545 WP 002215704.1      HLEPVLSEHIKTFKDDTYMRREHVVWGDEMKPLARLAAAVEMAG..ESDKANSHIFDAMVNQKINLADTDTLKKWLSEQT  158
NTDB id 668290 MON40 RS01385 WP 003780094.1 HLEPVLSKHVKTFKDDTYLRTEHVVWGDEMKPLARLAAAVDMAVADTKDIANSHIFDAMVNQKVKLQDPETLKKWLAEQT  159
NTDB id 1098 NMB RS01465 WP 002224844.1     HLEPVLSKHAKSFKDDMYLRTEHVVWQKEMLTLARLAAAVDMAAADSKDVANSHIFDAMVNQKIKLQNPEVLKKWLGEQT  159
consensus                                   !!!!!!!*!*!*!!!!*!*!*!!!!!**!!**!!!!!!!!*!!*****!*!!!!!!!!!!!!!!**!*****!!!!!*!!!

logo
NTDB id 1099 NMB RS01545 WP 002215704.1      AFDGKKVLAAFEAPESQARAAQMEELTNKFQISGTPTVIVGGKYQVEFKDWQSGMTTIDQLVDKVREEQKKPQ  231
NTDB id 668290 MON40 RS01385 WP 003780094.1 AFDGKKVLAAYESPESQTRADKMAELTNTYKIDGTPTVIVGGKYKVEFADWESGMKTIDLLADKVREEQKTAK  232
NTDB id 1098 NMB RS01465 WP 002224844.1     AFDGKKVLAAYESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEFADWESGMNTIDLLADKVREEQKAAQ  232
consensus                                   !!!!!!!!!!!*!*!!!!*!!**! !!!****!*!!!!!!!!!!!!*!!!*!!*!!! !!!*!*!!!!!!!! **
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X non conserved

X similar

X $\geq$ 50% conserved