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logo

NTDB id 1099 NMB RS01545 WP 002215704.1          MKLKTLALTSLTLLALAACSKQAETSVPADSAQSSSSAPAAPAELNEGVNYTVLSTPIPQQQAGKIEVLEFFGYFCPHCA   80
NTDB id 1099512 ACM67D RS09195 WP 118997788.1    MKSRHLALGVAALFALAACDSKVQTSVPADSAPAA.SAAAAPAGLVEGQNYTVLSNPIPQQQAGKVEVLEFFGYFCPHCA   79
NTDB id 1098 NMB RS01465 WP 002224844.1          MKSRHLALGVAALFALAACDSKVQTSVPADSAPAA.SAAAAPAGLVEGQNYTVLANPIPQQQAGKVEVLEFFGYFCPHCA   79
consensus                                        !!***!!!****!*!!!!!*****!!!!!!!*** !!*!!!!*!*!!*!!!!!**!!!!!!!!!*!!!!!!!!!!!!!!!

logo

NTDB id 1099 NMB RS01545 WP 002215704.1          HLEPVLSEHIKTFKDDTYMRREHVVWGDEMKPLARLAAAVEMAGE..SDKANSHIFDAMVNQKINLADTDTLKKWLSEQT  158
NTDB id 1099512 ACM67D RS09195 WP 118997788.1    HLEPVLSKHAKSFKDDMYLRTEHVVWQKEMLPLARLAAAVDMAAAESKDVANSHIFDAMVNQKIKLQNPEVLKKWLGEQT  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 1099512 ACM67D RS09195 WP 118997788.1    AFDGKKVLAAYESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEFADWESGMNTIDLLADKVREEQKAAQ  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