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logo

NTDB id 1123982 EL323 RS09375 WP 002212479.1  MKLKHLLPLL.......................LSAVLSAQAYALTEREDYLVLDKPIPQEQSGKIEVLEFFGYFCVHCH   57
NTDB id 1098 NMB RS01465 WP 002224844.1       MKSRHLALGVAALFALAACDSKVQTSVPADSAPAA.SAAAAPAGLVEGQNYTVLANPIPQQQAGKVEVLEFFGYFCPHCA   79
NTDB id 1099 NMB RS01545 WP 002215704.1       MKLKTLALTSLTLLALAACSKQAETSVPADSAQSSSSAPAAPAELNEGVNYTVLSTPIPQQQAGKIEVLEFFGYFCPHCA   80
consensus                                     !!***!** *  * *****     ********  * ** !****! !* *!*!!  !!!!*!*!!*!!!!!!!!!!*!!*

logo

NTDB id 1123982 EL323 RS09375 WP 002212479.1  HFDPLLLKLGKALPSDAYLRTEHVVWQPEMLGLARMAAAVNLSGL..KYQANPAVFKAVYEQKIRLENRSVAGKWALSQK  135
NTDB id 1098 NMB RS01465 WP 002224844.1       HLEPVLSKHAKSFKDDMYLRTEHVVWQKEMLTLARLAAAVDMAAADSKDVANSHIFDAMVNQKIKLQNPEVLKKWLGEQT  159
NTDB id 1099 NMB RS01545 WP 002215704.1       HLEPVLSEHIKTFKDDTYMRREHVVWGDEMKPLARLAAAVEMAGE..SDKANSHIFDAMVNQKINLADTDTLKKWLSEQT  158
consensus                                     !**!*!***!****! !*!*!!!!!* !!* !!!*!!!!*****   ** !!***!*!***!!!*! * *****!!* *!*

logo

NTDB id 1123982 EL323 RS09375 WP 002212479.1  GFDGKKLMRAYDSPEAAAAALKMQKLTEQYGIDSTPTVIVGGKYRVIFNNGFDGGVHTIKELVAKVREERKRQTPAVQK  214
NTDB id 1098 NMB RS01465 WP 002224844.1       AFDGKKVLAAYESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEFAD.WESGMNTIDLLADKVREEQKAAQ.....  232
NTDB id 1099 NMB RS01545 WP 002215704.1       AFDGKKVLAAFEAPESQARAAQMEELTNKFQISGTPTVIVGGKYQVEFKD.WQSGMTTIDQLVDKVREEQKKPQ.....  231
consensus                                     *!!!!!***!***!!**!*! *!**!!* **!**!!!!!!!!!!*!*! * ***!* !!* !**!!!!!*!* *
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X non conserved

X similar

X $\geq$ 50% conserved