```
logo
NTDB id 154112 DE10444 RS02245 WP 002229027.1  MKLKHLL.....................PLL...LSAALSAQAYALTEGEDYLVLDKPIPQEQSGKIEVLEFFGYFCVHCH   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 154112 DE10444 RS02245 WP 002229027.1  HFDPLLLKLGKALPSDAYLRTEHVVWQPEMLGLARMAAAVNLSGL..KYQANPAVFKAVYEQKIRLENRAVAGKWALSQK  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 154112 DE10444 RS02245 WP 002229027.1  GFDGKKLMRAYDSPEAAAAALKMQKLTEQYRIDSTPTVIVGGKYRVIFNNGFDGGVHTIKELVAKVREERKRQTPAVQK  214
NTDB id 1098 NMB RS01465 WP 002224844.1        AFDGKKVLAAYESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEFAD.WESGMNTIDLLADKVREEQKAAQ.....  232
NTDB id 1099 NMB RS01545 WP 002215704.1        AFDGKKVLAAFEAPESQARAAQMELTNKFQISGTPTVIVGGKYQVEFKD.WQSGMTTIDQLVDKVREEQKKPQ.....  231
consensus                                      *!!!!!***!***!!**!*! *!**!!* **!**!!!!!!!!!!*!*! * ***!* !!* !**!!!!!*!* *
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X non conserved

X similar

X $\geq$ 50% conserved