```
logo
NTDB id 47779 NMBB RS02245 WP 002218776.1  MKLKHLLPLL.......................LSAVLSAQAYALTEGEDYLVLDKPIPQEQSGKIEVLEFFGYFCVHCHHFDPL   62
NTDB id 1098 NMB RS01465 WP 002224844.1    MKSRHLALGVAALFALAACDSKVQTSVPADSAPAA.SAAAAPAGLVEGQNYTVLANPIPQQQAGKVEVLEFFGYFCPHCAHLEPV   84
NTDB id 1099 NMB RS01545 WP 002215704.1    MKLKTLALTSLTLLALAACSKQAETSVPADSAQSSSSAPAAPAELNEGVNYTVLSTPIPQQQAGKIEVLEFFGYFCPHCAHLEPV   85
consensus                                  !!***!** *  * *****     ********  * ** !***!! !! *!*!!  !!!!*!*!!*!!!!!!!!!!*!!*!**!*

logo
NTDB id 47779 NMBB RS02245 WP 002218776.1  LLKLGKALPSDAYLRTEHVVWQPEMLGLARMAAAVNLSGL..KYQANPAVFKAVYEQKIRLENRAVAGKWALSQKGFDGKKLMRA  145
NTDB id 1098 NMB RS01465 WP 002224844.1    LSKHAKSFKDDMYLRTEHVVWQKEMLTLARLAAAVDMAAADSKDVANSHIFDAMVNQKIKLQNPEVLKKWLGEQTAFDGKKVLAA  169
NTDB id 1099 NMB RS01545 WP 002215704.1    LSEHIKTFKDDTYMRREHVVWGDEMKPLARLAAAVEMAGE..SDKANSHIFDAMVNQKINLADTDTLKKWLSEQTAFDGKKVLAA  168
consensus                                  !****!****! !*!*!!!!!* !!* !!!*!!!!****   ** !!***!*!***!!!*! * ****!!* *!**!!!!!***!

logo
NTDB id 47779 NMBB RS02245 WP 002218776.1  YDSPEAAAAALKMQKLTEQYRIDSTPTVVVGGKYRVIFNNGFDGGVHTIKELVAKVRAERKLSNHPVQK   214
NTDB id 1098 NMB RS01465 WP 002224844.1    YESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEFAD.WESGMNTIDLLADKVREEQKAAQ.....   232
NTDB id 1099 NMB RS01545 WP 002215704.1    FEAPESQARAAQMEELTNKFQISGTPTVIVGGKYQVEFKD.WQSGMTTIDQLVDKVREEQKKPQ.....   231
consensus                                  ***!!**!*! *!**!!* **!**!!!!*!!!!!*!*! * ***!* !!* !**!!!*!*!  *
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X non conserved

X similar

X $\geq$ 50% conserved