```
logo
NTDB id 21661 NMA RS10560 WP 002220358.1  MKLKHLLPLL.......................LSAVLSAQAYALTEGEDYLVLDKPIPQKQSGKIEVLEFFGYFCVHCHHFDPL   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 21661 NMA RS10560 WP 002220358.1  LLKLGKALPSDAYLRTEHVVWQPEMLGLARMAAAVKLSGL..KYQANPAVFKAVYEQKIRLENRSVAEKWALSQKGFDGKKLMRA  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 21661 NMA RS10560 WP 002220358.1  YDSPAAAAAAASKMQQLTEQYRIDSTPTVVVGGKYRVIFNNGFDGGVHTIKELVAKVREERKRQTPAVQK  214
NTDB id 1098 NMB RS01465 WP 002224844.1   YESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEFA.DWESGMNTIDLLADKVREEQKAAQ.....  232
NTDB id 1099 NMB RS01545 WP 002215704.1   FEAPESQARAAQMEELTNKFQISGTPTVIVGGKYQVEFK.DWQSGMTTIDQLVDKVREEQKKPQ.....  231
consensus                                 ***!***!*! *!**!!* **!**!!!!*!!!!!*!*!  ****!* !!* !**!!!!!*!* *
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X non conserved

X similar

X $\geq$ 50% conserved