```
logo
NTDB id 668335 MON37 RS01480 WP 039408843.1  MKWTTKPA..AVLFSL.................ALAFSASAQAAVEGVDYTVLNKPIPQRDASKIEVLEFFGYFCIHCQ   60
NTDB id 1099 NMB RS01545 WP 002215704.1      MKLKTLALTSLTLLALAACSKQAETSVPADSAQSSSSAPAAPAELNEGVNYTVLSTPIPQQQAGKIEVLEFFGYFCPHCA   80
NTDB id 1098 NMB RS01465 WP 002224844.1      MKSRHLALGVAALFALAACDSKVQTSVPADSAPAA.SAAAAPAGLVEGQNYTVLANPIPQQQAGKVEVLEFFGYFCPHCA   79
consensus                                    !! *****  * !**!***     ********  * ** !****+**!!**!!!!  !!!!**!*!*!!!!!!!!!!*!!*

logo
NTDB id 668335 MON37 RS01480 WP 039408843.1  NFDPVLLSYSKTFPKDVYLRTEHVVWMPEMLGLARVAAAVNASGL..KYQANPAIFKAVHEQKINLADTATFKSWAAAQK  138
NTDB id 1099 NMB RS01545 WP 002215704.1      HLEPVLSEHIKTFKDDTYMRREHVVWGDEMKPLARLAAAVEMAGE..SDKANSHIFDAMVNQKINLADTDTLKKWLSEQT  158
NTDB id 1098 NMB RS01465 WP 002224844.1      HLEPVLSKHAKSFKDDMYLRTEHVVWQKEMLTLARLAAAVDMAAADSKDVANSHIFDAMVNQKIKLQNPEVLKKWLGEQT  159
consensus                                    ***!!!* * !*!**!*!*!*!!!!!  !!* !!!*!!!!****   ** !!**!!*!***!!!*!*****!*!***!*

logo
NTDB id 668335 MON37 RS01480 WP 039408843.1  SFDSKKLIAAYDAPASLAAAKKMQSLTETYRIDGTPDVIVGGKYRVIFSSDWANGQKIIGELINKVRQERSGKAAR  214
NTDB id 1099 NMB RS01545 WP 002215704.1      AFDGKKVLAAFEAPESQARAAQMEELTNKFQISGTPTVIVGGKYQVEF.KDWQSGMTTIDQLVDKVREEQKKPQ..  231
NTDB id 1098 NMB RS01465 WP 002224844.1      AFDGKKVLAAYESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEF.ADWESGMNTIDLLADKVREEQKAAQ..  232
consensus                                    *!!*!!**!!***!*!*!*! *!**!!****!*!!!*!!!!!!!*!*!   !! *!* *!* !**!!!*!***  *
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X non conserved

X similar

X $\geq$ 50% conserved