```
logo
NTDB id 231792 CCD82 RS08165 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 231792 CCD82 RS08165 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 231792 CCD82 RS08165 WP 002229027.1  GFDGKKLMRAYDSPEAAAAALKMQKLTEQYRIDSTPTVIVGGKYRVIFNNGFDGGVHTIKELVAKVREERKRQTPAVQK  214
NTDB id 1098 NMB RS01465 WP 002224844.1      AFDGKKVLAAYESPESQARADKMQELTETFQIDGTPTVIVGGKYKVEFAD.WESGMNTIDLLADKVREEQKAAQ.....  232
NTDB id 1099 NMB RS01545 WP 002215704.1      AFDGKKVLAAFEAPESQARAAQMEELTNKFQISGTPTVIVGGKYQVEFKD.WQSGMTTIDQLVDKVREEQKKPQ.....  231
consensus                                    *!!!!!***!***!!**!*! *!**!!* **!**!!!!!!!!!!*!*! * ***!* !!* !**!!!!!*!* *
```

X non conserved

X similar

X $\geq$ 50% conserved