```
logo
NTDB id 668333 MON37 RS01095 WP 039408913.1  MKNKIFLLTSILSLTGCTPAYDDLNEWMTKTRQDAKSHIIPFEAPVVNPPAVYNPPAYSGLNAFDFRRLDNAPKGNNAPN   80
NTDB id 1101 NMB RS09450 WP 002214522.1      .MKHYALLISFLALSACSQGSEDLNEWMAQTRREAKAEIIPFQAPTLPVAPVYSPPQLTGPNAFDFRRMETDKKGENAPD   79
consensus                                    *  *  !! ! ! !**!*** *!!!!!!  !! *!! *!!!! !! *****!! !!  *!*!!!!!!!**  *!! !!!

logo
NTDB id 668333 MON37 RS01095 WP 039408913.1  PNRPKETLEAFSLENMRYVGMLSNGSKTAGFIDVNGHVYTVYPGNYIGQNYGKIQSITEDLIVLTELVEDSSGNWIYRKA   160
NTDB id 1101 NMB RS09450 WP 002214522.1      TKRIKETLEKFSLENMRYVGILKSGQKVSGFIEAEGYVYTVGVGNYLGQNYGRIESITDDSIVLNELIEDSTGNWVSRKA   159
consensus                                    * !*!!!!! !!!!!!!!!!*!  ! !  !!!*  !*!!!! *!!!*!!!!!*! !!!*! !!! !!*!!!*!!!* !!!

logo
NTDB id 668333 MON37 RS01095 WP 039408913.1  ELSLSNNAENSSNDIGNASATNSN   184
NTDB id 1101 NMB RS09450 WP 002214522.1      ELLLNSSDKNTEQAAAPAAEQN..   181
consensus                                    !! !     !* *  **!   !**
```

X non conserved

X similar

X $\geq$ 50% conserved