```
logo
NTDB id 366814 FGY92 RS00200 WP 235063213.1  MTQLKKKRIGLYARVSTEMQ.TDGYSIQGQLNQLTEYCQFQGYEVVNEYTDRGISGKTTQRPELQRMLKDANDGKLDCIM   79
NTDB id 11 SA RS00430 WP 001186602.1         MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRILKDAKEGYIDCVM   80
consensus                                    ! !!! !!*!*! !!!!!!!*!*!!!! !!*!!* !!! !***!!   ! !!!!!!!* *!!!!!!*!!!! *! *!!*!

logo
NTDB id 366814 FGY92 RS00200 WP 235063213.1  VYKTNRLARNTSDLLNIVEELYKINVEFFSLTEKIEIASSTGKLMLQILASFSEFERNTIVENVYNGQRQRAIEGYYQGN  159
NTDB id 11 SA RS00430 WP 001186602.1         VYKTNRLARNTSDLLKIVEDLHKQNVEFFSLSERMEVNTSSGKLMLQILASFSEFERNNIVENVFMGQTRRAQEGYYQGN  160
consensus                                    !!!!!!!!!!!!!!! !!!*!*! !!!!!!!*!**!* *!*!!!!!!!!!!!!!!!! !!!!!* !!  !! !!!!!!!

logo
NTDB id 366814 FGY92 RS00200 WP 235063213.1  LPLGYDKVPDSKKELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSISSITYIISNPFYIGKIQFAK  239
NTDB id 11 SA RS00430 WP 001186602.1         LPLGYDKIPNSKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSISSITYILANPFYIGKIQFAK  240
consensus                                    !!!!!!!*! !!*!!!!!!!!!!!!!!!
```

X non conserved
X similar
X $\geq$ 50% conserved