```
logo
NTDB id 325148 EGX58 RS04525 WP 002486185.1  MDKMKKKLVGGYIRVSTER.QVEGYSIEGQITQIEQYCQFNGYELVDIYADRGISGKSMNRPELQRMLNDAKNGKLDCVM   79
NTDB id 11 SA RS00430 WP 001186602.1         MQQLKTKRVGIYVRVSTEMQSTEGYSIDGQINQIKEYCDFHHFEVKDIYADRGISGKSMNRPELQRILKDAKEGYIDCVM   80
consensus                                    !  *! ! !! !*!!!!! *  !!!!!*!!! !!  !! !***!* !!!!!!!!!!!!!!!!!!!!*! !!! ! *!!!!

logo
NTDB id 325148 EGX58 RS04525 WP 002486185.1  VYKTNRLARNTSDLLTIVEELHRQNVEFFSLSERMEVKNSTGKLMLQILASFSEFERNTILENIYTGQRQRALEGYYQGN  159
NTDB id 11 SA RS00430 WP 001186602.1         VYKTNRLARNTSDLLKIVEDLHKQNVEFFSLSERMEVNTSSGKLMLQILASFSEFERNNIVENVFMGQTRRAQEGYYQGN  160
consensus                                    !!!!!!!!!!!!!!! !!!*!!*!!!!!!!!!!!!!!  !*!!!!!!!!!!!!!!!!!! !*!!** !!  !! !!!!!!!

logo
NTDB id 325148 EGX58 RS04525 WP 002486185.1  LPLGYNNIPDNKKELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGNPFSISAVTYILSNPFYIGKIQFAK  239
NTDB id 11 SA RS00430 WP 001186602.1         LPLGYDKIPNSKHELMINQHEANIVKYIFESYAKGHGYRKIANALNHKGYVTKKGKPFSISSITYILANPFYIGKIQFAK  240
consensus                                    !!!!!  !!  !*!!!!!!!!!!!!!!!!!!!
```

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