```
logo
NTDB id 1094070 ACINK0 RS14065 WP 414657016.1  MPVYEYRARDREGKVLKSQMDAESEAQVRDALRARNLMILEIKAPKTGLNADVKIPGLTDRPPGLKQVAVFSKQLATLIN   80
NTDB id 1029 TT RS02230 WP 011228203.1         MPVYQYKARDRQGRLVEATIEAEDLRTAARLLRDRGLFVAEIKEPGKGLQAEVRIPALE.RGPGLKDLAIFSRQLATMLG   79
consensus                                      !!!! !*!!!! !***   **!!       !! ! ! * !!! !  !!*!*!*!!*! *!*!!!! *!*!!*!!!!**

logo
NTDB id 1094070 ACINK0 RS14065 WP 414657016.1  AGVPLVQSLAILQRQVDNKTFETVLKKVRGDVESGTPLSEALQQHPKVFSRLFINLVRAGETSGTLDTVLERIADFQEKQ  160
NTDB id 1029 TT RS02230 WP 011228203.1         AGLTLLQALAILERQTENRKFREILKQVRTDVEGGMAFSEALSKH.KIFSRLYVNLVRAGETSGGLDLILDRLASFLEKE  158
consensus                                      !!**!*! !!!! !! *!* !  *!! !! !!! ! * !!!!  !*!*!!!!**!!!!!!!!!! !! *!*!*! ! !!

logo
NTDB id 1094070 ACINK0 RS14065 WP 414657016.1  LALQGKIKSALTYPVVVLVFALGITYFLLTTIVPQFASILTQLNAPLPFITRMLMAVSDFLRSSGLFIFLIVGVIFFAYR  240
NTDB id 1029 TT RS02230 WP 011228203.1         LELRGKIRSAMTYPVIVFVFAVGVAYFLLTGIVPQFAQILTDLGSELPLLTRFLIAVSDLLRAATLPLLLLAVALFFAYR  238
consensus                                      ! ! !!!*!!*!!!!*! !!!*!* !!!!! !!!!!! !!! !  *!! *!! !*!!!! !!   !** !*   *!!!!!

logo
NTDB id 1094070 ACINK0 RS14065 WP 414657016.1  WYYRTPKGRYVIDDIKLRLPVFGNLIQKSAISSFARTFGLLISSGVNIIESLEITKGTANNAIVEETIDNAKNVVISGDQ  320
NTDB id 1029 TT RS02230 WP 011228203.1         WYYGTPQGRRVIDRLKLRLPVFGNLNRKTAVARFSRTLALLLSSGVNIVEALDITKGTAGNSVVEEIVEAAKLKIQQGDP  318
consensus                                      !!! !! !! !!! *!!!!!!!!!!  !*!*  ! !! *!!*!!!!!!*! !*!!!!!! ! *!!! ** !!  *  !!*

logo
NTDB id 1094070 ACINK0 RS14065 WP 414657016.1  MSSSLAT.SKVFPPMVVSMVAIGEETGSLDNMLGKVADFYDREVDEAVESLTAAIEPMMIVFLGGIVGLIVAGMFLPMFS  399
NTDB id 1029 TT RS02230 WP 011228203.1         LNLTLAQHPFVFPPMVSSMVAIGEETGALDTMLSKVADFYEREVDEAVASLTAAIEPLMIIFLGVIVGMIVAGMFLPLFK  398
consensus                                      *  *!! ** !!!!!! !!!!!!!!!! !! !! !!!!!!*!!!!!!! !!!!!!!!*!!*!!! !!!*!!!!!!!!*!

logo
NTDB id 1094070 ACINK0 RS14065 WP 414657016.1  IIGALSQ.   406
NTDB id 1029 TT RS02230 WP 011228203.1         IIGTLSVQ   406
consensus                                      !!! !! *
```

X non conserved

X similar

X $\geq$ 50% conserved