```
logo
NTDB id 140098 TO73 RS02690 WP 003044366.1  MKEALARLQAAWKNLPQSTKLLLAGLLLVGAVTLWYVGLYLPAQTPLEAQPPAPAVPETPKALEAPPIPPLAEPAPAEAPKAAPQ   85
NTDB id 1026 TT RS05140 WP 011173435.1      MKNAWQRLKEAWANLPRSTKLLLAALLLVGGVALWYVGLYLPAQVAEAPTP.....APSTQVIEAPPIPPLASGEEAKPQAEAPA   80
consensus                                   !! !  !!  !! !!! !!!!!!!*!!!!!*! !!!!!!!!!!! *  * !*********  *!!!!!!!!! * *! **  !!

logo
NTDB id 140098 TO73 RS02690 WP 003044366.1  APEK.AEAAQTTPQAANLPMPKAKPEAPPPNPFVPLIVPAEAAPPAPVPTPPAPTVRPTPVPTGTPVRVSGGTPLPTPSVQGAPR   169
NTDB id 1026 TT RS05140 WP 011173435.1      QEEAQAPASSQEEAQAPPPAPVARAEPPPPNPFVPLVVETPP.......PPPASASRPTPVPEGPAVRVQT..PAQAPQAAPATR   156
consensus                                    *! *!*!    *  !**! ! !**!*!!!!!!!!!*!* **********!!!*  !!!!!! !**!!!  **! * !   *!*!

logo
NTDB id 140098 TO73 RS02690 WP 003044366.1  PLPGTAGALPAPQVLRPTPKVEVPQARLEPPVEVAAPGGLVETPLPQEEPPKEAQAPAPPAPKSPLAALVAEKGLRLSGTLMGPI   254
NTDB id 1026 TT RS05140 WP 011173435.1      PIPGTSGALPAPKILSPALSAPLPQARETPP.RVAVPTALVEAPLPGPEEK...GAEKAPTPASPLERLVAEKGLRLSGTLLGPV   237
consensus                                   !*!!! !!!!!! *! ! *  **!!!!  !!* !! ! *!!! !!! *!***** !* *! ! !!!  !!!!!!!!!!!!!*!!*

logo
NTDB id 140098 TO73 RS02690 WP 003044366.1  SVAILESKEGYLVLPAGSPIPGSEAVVKRIEGDRIVLALKEETLETPLEQAGGER   309
NTDB id 1026 TT RS05140 WP 011173435.1      SVAILESKEGYLVVPAGSPIPGTEAVVRQVEEGSVTLALKEETLNLSLVQAGGGQ   292
consensus                                   !!!!!!!!!!!!!*!!!!!!!!!*!!!!* *!    * !!!!!!!! **! !!!!
```

X non conserved

X similar

X $\geq$ 50% conserved