```
logo
NTDB id 39726 TSC RS10145 WP 015717937.1  MKALLARLATAWRNLPQSTKLLLAGLLLVSAVALWYVGFYLPAQAPMEVQPTPGQPQVPSQGAEAPKAIEAPPIPPLAETAPQDQ   85
NTDB id 1026 TT RS05140 WP 011173435.1    MKNAWQRLKEAWANLPRSTKLLLAALLLVGGVALWYVGLYLPAQVAEAPTPAP...........STQVIEAPPIPPLASQEEAKP   74
consensus                                 !!    !!  !! !!! !!!!!!!*!!!! *!!!!!!! !!!!! *  * ! !*********** *  !!!!!!!!!!   *  *

logo
NTDB id 39726 TSC RS10145 WP 015717937.1  AAK.PSPE...ASPSQKPVLTTALPIPKTQQEAPLPNPFVPLVVEAPPSPPVPSPAPAPRPTPVPTGAPVRVTQGTPLPTPSVQA   166
NTDB id 1026 TT RS05140 WP 011173435.1    QAEAPAQEEAQAPASSQEEAQAPPPAPVARAEPPPPNPFVPLVVETPPPPPA....SASRPTPVPECPAVRVQT..PAQAPQAAP   153
consensus                                  ! *! *!***!**!  *    **! !    !*!*!!!!!!!!!! !!*!! *****!*!!!!!! !**!!!  **! * !   *

logo
NTDB id 39726 TSC RS10145 WP 015717937.1  SPRPLPGSQGALPAPKVLTPAFQVETPKAQVDTPLLLTPPAGLVEAPLPRAPQTPEGGKTEPSPATSPKTPLQALVEEKGIKLAG   251
NTDB id 1026 TT RS05140 WP 011173435.1    ATRPIPGTSGALPAPKILSPALSAPLPQARETPP.RVAVPTALVEAPLPGPEEK......GAEKAPTPASPLERLVAEKGLRLSG   231
consensus                                 *!!*!!* !!!!!!!*!*!!   * ! !   *!* * *! *!!!!!!! **  ****** * *!**! *!!  !! !!!**! !

logo
NTDB id 39726 TSC RS10145 WP 015717937.1  TLLGPVSVAILETKEGYLVLPVGSLLPGSEAVLRRIESDRVVLALKDESLEITLEKMQAGGGQ   314
NTDB id 1026 TT RS05140 WP 011173435.1    TLLGPVSVAILESKEGYLVVPAGSPIPGTEAVVRQVEEGSVTLALKEETLNLSL..VQAGGGQ   292
consensus                                 !!!!!!!!!!!!*!!!!!!*! !!**!!*!!!*! *!   ! !!!!*!*! **!***!!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved