```
logo
NTDB id 42118 LCGT RS09420 WP 014025492.1  MKNEKIKKHKKLKLFFAIALPL.LIVVGLFFPLPYYIEQPGGTIPVNQMVDVAGKKDEHKGNFYLTTVEMVQANAASMLYSKSNS   84
NTDB id 360 SMU RS02495 WP 002262039.1     .....MKTNKKFKWWLISGISLILLLMVFFFPLPYYLEMPGGAYDIRSVVTVNHKEDKEKGSYNFVAVSLSKATPIQVLYAWLTP   80
consensus                                  ******! *!! ! *   ***!*!***  !!!!!!!*! !!!  **  *! ! *! ! *!! *    ! *  ! *  *!!    *

logo
NTDB id 42118 LCGT RS09420 WP 014025492.1  FATILSSEEMTGGMTNQQFDLVNQFYMQTAQNTAVYQAFKLAGKPYEMKYQGVYVLSITEDSTFKNDLQLSDTITAVNGHTFKSS  169
NTDB id 360 SMU RS02495 WP 002262039.1     FTEITSAKETTGGVSDADYMRINQFYMETSQNGAIYKALSLAHKKASFNYMGVYVLQVSRNSTFKGVLNIADTVTGVNDKTFHSS  165
consensus                                  !  ! !  ! !!!**   *  *!!!!! ! !! !*! !  !!*!*    ! !!!!! **  !!!!  !** !!*!*!! *!!*!!

logo
NTDB id 42118 LCGT RS09420 WP 014025492.1  TDMIDYVSQQKVGDEVTIKYTRMDGSNHEATGKYIKLSNGKTGIGIGLVDHTQVVTDPKVKIDAGSIGGPSAGMMFTLEIYSQIT  254
NTDB id 360 SMU RS02495 WP 002262039.1     KELIKYVSGLKLGSKVSVQYTSEN.KKKTAKGKVIKLSNGKNGIGIGLVDHTKVTSDDKIEFSTNGIGGPSAGLMFTLDIYDQLV  249
consensus                                   **! !!!   !*!   !** !!   *  * ! !! !!!!!!! !!!!!!!!!! ! *!*!*      !!!!!!!*!!!!*!! !*

logo
NTDB id 42118 LCGT RS09420 WP 014025492.1  GKDLRQGREIAGTGTINEDGSIGQIGGVDKKVATASNSGAKIFLCPDETEEQAK...ASGTTNNYTAALAAAKKLNTDMKIVPVK  336
NTDB id 360 SMU RS02495 WP 002262039.1     KEDLRKGRIIAGTGTIGEDGSVGDIGGADMKVAAADRINADIFFVPNNPVSKATLKKNPKALTNYQEAKQAAKKLGTKMKIVPVK  334
consensus                                    !!! !! !!!!!!! !!!!*! !!! ! !!! !    ! !! *!  *   ! *** *    !!  !  !!!!! ! !!!!!!!

logo
NTDB id 42118 LCGT RS09420 WP 014025492.1  TIQDALDYLEK.   347
NTDB id 360 SMU RS02495 WP 002262039.1     NVQEAIDYLRKH   346
consensus                                   *!*!*!!! !*
```

X non conserved

X similar

X ≥ 50% conserved