```
logo
NTDB id 269963 C3L41 RS09570 WP 014025492.1  MKNEKIKKHKKLKLFFAIALPL.LIVVGLFFPLPYYIEQPGGTIPVNQMVDVAGKKDEHKGNFYLTTVEMVQANAASMLY   79
NTDB id 360 SMU RS02495 WP 002262039.1       .....MKTNKKFKWWLISGISLILLLMVFFFPLPYYLEMPGGAYDIRSVVTVNHKEDKEKGSYNFVAVSLSKATPIQVLY   75
consensus                                    ******! *!! ! *   ***!*!***  !!!!!!!*! !!!  **  *! ! *! ! *!! *    ! *  ! *  *!!

logo
NTDB id 269963 C3L41 RS09570 WP 014025492.1  SKSNSFATILSSEEMTGGMTNQQFDLVNQFYMQTAQNTAVYQAFKLAGKPYEMKYQGVYVLSITEDSTFKNDLQLSDTIT  159
NTDB id 360 SMU RS02495 WP 002262039.1       AWLTPFTEITSAKETTGGVSDADYMRINQFYMETSQNGAIYKALSLAHKKASFNYMGVYVLQVSRNSTFKGVLNIADTVT  155
consensus                                        *!  ! !  ! !!!**   *  *!!!!! ! !! !*! !  !!*!*    ! !!!!! **  !!!!  !** !!*!

logo
NTDB id 269963 C3L41 RS09570 WP 014025492.1  AVNGHTFKSSTDMIDYVSQQKVGDEVTIKYTRMDGSNHEATGKYIKLSNGKTGIGIGLVDHTQVVTDPKVKIDAGSIGGP  239
NTDB id 360 SMU RS02495 WP 002262039.1       GVNDKTFHSSKELIKYVSGLKLGSKVSVQYTSEN.KKKTAKGKVIKLSNGKNGIGIGLVDHTKVTSDDKIEFSTNGIGGP  234
consensus                                    *!! *!!*!! **! !!!  !*!  !** !!   *  * ! !! !!!!!!! !!!!!!!!!! ! *!*!*      !!!!

logo
NTDB id 269963 C3L41 RS09570 WP 014025492.1  SAGMMFTLEIYSQITGKDLRQGREIAGTGTINEDGSIGQIGGVDKKVATASNSGAKIFLCPDETEEQAK...ASGTTNNY  316
NTDB id 360 SMU RS02495 WP 002262039.1       SAGLMFTLDIYDQLVKEDLRKGRIIAGTGTIGEDGSVGDIGGADMKVAAADRINADIFFVPNNPVSKATLKKNPKALTNY  314
consensus                                    !!!*!!!!*!! !*   !!! !! !!!!!!! !!!!*! !!! ! !!! !    ! !! *!  *   ! *** *    !!

logo
NTDB id 269963 C3L41 RS09570 WP 014025492.1  TAALAAAKKLNTDMKIVPVKTIQDALDYLEK.   347
NTDB id 360 SMU RS02495 WP 002262039.1       QEAKQAAKKLGTKMKIVPVKNVQEAIDYLRKH   346
consensus                                      !  !!!!! ! !!!!!!! *!*!*!!! !*
```

X non conserved

X similar

X $\geq$ 50% conserved