```
logo
NTDB id 1114595 A66 RS08850 WP 053039757.1  ..MKKKIRWPLYVIAALIVTFLAFVVPLPYYIEVPGGSEDIRQVLKVNDTEDKEAGAYQFVTVGVQHATLAHMIYAWLTPFTDIR   83
NTDB id 360 SMU RS02495 WP 002262039.1      MKTNKKFKWWLISGISLILLLMVFFFPLPYYLEMPGGAYDIRSVVTVNHKEDKEKGSYNFVAVSLSKATPIQVLYAWLTPFTEIT   85
consensus                                   **  !! *!*!     !!*  * !  !!!!!*!*!!!  !!! !* !!* !!!! ! !*!! ! * *!!* ***!!!!!!!!*!

logo
NTDB id 1114595 A66 RS08850 WP 053039757.1  SAQETTGGSSDVEFMRINQFYMQTSQNMAKYQGLKTAGKDIELKYFGVYVLNVTDNSTFKGILNISDTVTAVNDQTFDSSKDLID  168
NTDB id 360 SMU RS02495 WP 002262039.1      SAKETTGGVSDADYMRINQFYMETSQNGAIYKALSLAHKKASFNYMGVYVLQVSRNSTFKGVLNIADTVTGVNDKTFHSSKELIK  170
consensus                                   !! !!!!! !! **!!!!!!!! !!!! ! ! *!  !*!     ! !!!!!*!* !!!!!!*!!! !!!!*!!! !!*!!!*!!

logo
NTDB id 1114595 A66 RS08850 WP 053039757.1  YVSSQKLGDYVSSQKLGDSVKVTYEEDGQTKSAEGKIITLENGKNGIGIGLIDRTEVISNVPISFSTAGIGGPSAGLMFSLAIYT  253
NTDB id 360 SMU RS02495 WP 002262039.1      YVSGLKLG.........SKVSVQYTSENKKKTAKGKVIKLSNGKNGIGIGLVDHTKVTSDDKIEFSTNGIGGPSAGLMFTLDIYD  246
consensus                                   !!!  !!!*********  ! ! !  *   !*! !!*! ! !!!!!!!!!!*!*! ! !  *! !!! !!!!!!!!!!!*! !!

logo
NTDB id 1114595 A66 RS08850 WP 053039757.1  QIAHPDLRNGRIVAGTGTIDRDGNVGDIGGIDKKVVASARAGAAIFFAPDNPVSEEEQKAHPDAKNNYQTALEAAKTIKTDMKIV  338
NTDB id 360 SMU RS02495 WP 002262039.1      QLVKEDLRKGRIIAGTGTIGEDGSVGDIGGADMKVAAADRINADIFFVPNNPVSKATLKKNPKALTNYQEAKQAAKKLGTKMKIV  331
consensus                                   !* **!!! !!!*!!!!!!  !! !!!!!! ! !! !  !  ! !!! ! !!!!    ! *! !  !!! !  !!! * ! !!!!

logo
NTDB id 1114595 A66 RS08850 WP 053039757.1  PVKTLQDAIDYLKNNP   354
NTDB id 360 SMU RS02495 WP 002262039.1      PVKNVQEAIDYLRKH.   346
consensus                                   !!! *!*!!!!!* **
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X non conserved

X similar

X ≥ 50% conserved