```
logo
NTDB id 30639 SPCG RS10130 WP 000727272.1  ..MKKKIRWPLYVIAALIVTFLAFVVPLPYYIEVPGGSEDIRQVLKVNDTEDKEAGAYQFVTVGVQHATLAHMIYAWLTPFTDIR   83
NTDB id 360 SMU RS02495 WP 002262039.1     MKTNKKFKWWLISGISLILLLMVFFFPLPYYLEMPGGAYDIRSVVTVNHKEDKEKGSYNFVAVSLSKATPIQVLYAWLTPFTEIT   85
consensus                                  **  !! *!*!     !!*  * !  !!!!!*!*!!!  !!! !* !!* !!!! ! !*!! ! * *!!* ***!!!!!!!!*!

logo
NTDB id 30639 SPCG RS10130 WP 000727272.1  SAQETTGGSSDVEFLRINQFYMQTSQNMAKYQGLKTAGKDIELKYFGVYVLNVTDNSTFKGILNISDTVTAVNDQTFDSSKDLID  168
NTDB id 360 SMU RS02495 WP 002262039.1     SAKETTGGVSDADYMRINQFYMETSQNGAIYKALSLAHKKASFNYMGVYVLQVSRNSTFKGVLNIADTVTGVNDKTFHSSKELIK  170
consensus                                  !! !!!!! !! ***!!!!!!! !!!! ! ! *!  !*!     ! !!!!!*!* !!!!!!*!!! !!!!*!!! !!*!!!*!!

logo
NTDB id 30639 SPCG RS10130 WP 000727272.1  YVSSQKLGDSVKVTYEEDGQTKSAEGKIITLENGKNGIGIGLIDRTEVISNVPISFSTAGIGGPSAGLMFSLAIYTQIAHPDLRN  253
NTDB id 360 SMU RS02495 WP 002262039.1     YVSGLKLGSKVSVQYTSENKKKTAKGKVIKLSNGKNGIGIGLVDHTKVTSDDKIEFSTNGIGGPSAGLMFTLDIYDQLVKEDLRK  255
consensus                                  !!!  !!!  ! ! !  *   !*! !!*! ! !!!!!!!!!!*!*! ! !  *! !!! !!!!!!!!!!!*! !! !* **!!!

logo
NTDB id 30639 SPCG RS10130 WP 000727272.1  GRIVAGTGTIDRDGNVGDIGGIDKKVLASARAGAAIFFAPDNPVSEEEQKAHPDAKNNYQTALEAAKTIKTDMKIVPVKTLQDAI  338
NTDB id 360 SMU RS02495 WP 002262039.1     GRIIAGTGTIGEDGSVGDIGGADMKVAAADRINADIFFVPNNPVSKATLKKNPKALTNYQEAKQAAKKLGTKMKIVPVKNVQEAI  340
consensus                                  !!!*!!!!!!  !! !!!!!! ! !! !  !  ! !!! ! !!!!    ! *! !  !!! !  !!! * ! !!!!!!! *!*!!

logo
NTDB id 30639 SPCG RS10130 WP 000727272.1  DYLKNSP   345
NTDB id 360 SMU RS02495 WP 002262039.1     DYLRKH.   346
consensus                                  !!!* **
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X non conserved

X similar

X ≥ 50% conserved