```
logo
NTDB id 1170264 ACLV7A RS01475 WP 411863400.1  ..MKKKIRWPLYVIAALIVTFLAFVVPLPYYIEVPGGSEDIRQVLKINDTEDQAAGAYHFVTVGVQHATLAHMVYAWLTP   78
NTDB id 360 SMU RS02495 WP 002262039.1         MKTNKKFKWWLISGISLILLLMVFFFPLPYYLEMPGGAYDIRSVVTVNHKEDKEKGSYNFVAVSLSKATPIQVLYAWLTP   80
consensus                                      **  !! *!*!    !!*  * !  !!!!!*!*!!!  !!! !* *!* !!   ! !*!! ! * *!!* ***!!!!!!

logo
NTDB id 1170264 ACLV7A RS01475 WP 411863400.1  FTDIRSAQETTGGSSDVEFMRINQFYMQTSQNMAKYQGLKTAGKDIELKYFGVYVLNVTDNSTFKGILNISDTVTAVNDQ  158
NTDB id 360 SMU RS02495 WP 002262039.1         FTEITSAKETTGGVSDADYMRINQFYMETSQNGAIYKALSLAHKKASFNYMGVYVLQVSRNSTFKGVLNIADTVTGVNDK  160
consensus                                      !!*! !! !!!!! !! **!!!!!!!! !!!! ! ! *!  !*!     ! !!!!!*!* !!!!!!*!!! !!!!*!!!

logo
NTDB id 1170264 ACLV7A RS01475 WP 411863400.1  TFDSSKDLIDYVSSQKLGDSVKVTYEEDGQTKSAEGKIITLENGKNGIGIGLIDRTEVISNVPISFSTAGIGGPSAGLMF  238
NTDB id 360 SMU RS02495 WP 002262039.1         TFHSSKELIKYVSGLKLGSKVSVQYTSENKKKTAKGKVIKLSNGKNGIGIGLVDHTKVTSDDKIEFSTNGIGGPSAGLMF  240
consensus                                      !!*!!!*!! !!!  !!!  ! ! !  *   !*! !!*! ! !!!!!!!!!!*!*! ! !  *! !!! !!!!!!!!!!!

logo
NTDB id 1170264 ACLV7A RS01475 WP 411863400.1  SLAIYTQIAHPGLRNGRIVAGTGTIDRDGNVGDIGGIDKKVVASARSGAAIFFAPDNPVSEEEQKAHPDAKNNYQTALEA  318
NTDB id 360 SMU RS02495 WP 002262039.1         TLDIYDQLVKEDLRKGRIIAGTGTIGEDGSVGDIGGADMKVAAADRINADIFFVPNNPVSKATLKKNPKALTNYQEAKQA  320
consensus                                      *! !! !* ** !! !!!*!!!!!!  !! !!!!!! ! !! !  !  ! !!! ! !!!!    ! *! !  !!! !  !

logo
NTDB id 1170264 ACLV7A RS01475 WP 411863400.1  AKTIKTDMKIVPVKTLQDAIDYLKNNP  345
NTDB id 360 SMU RS02495 WP 002262039.1         AKKLGTKMKIVPVKNVQEAIDYLRKH.  346
consensus                                      !! * ! !!!!!!! *!*!!!!!* **
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X non conserved

X similar

X $\geq$ 50% conserved