```
logo
NTDB id 1121823 EL140 RS01275 WP 000730772.1  ..MKKKTRWPLYVILALVLTFLAFVVPLPYYIEVPGGAEDIRRVLKVNETEDTEAGAYQFVTVGIRHATLSHLVYAWLTP   78
NTDB id 360 SMU RS02495 WP 002262039.1        MKTNKKFKWWLISGISLILLLMVFFFPLPYYLEMPGGAYDIRSVVTVNHKEDKEKGSYNFVAVSLSKATPIQVLYAWLTP   80
consensus                                     **  !! *!*!   * !*!  * !  !!!!!*!*!!!! !!! !* !!* !! ! ! !*!! ! * *!!* ***!!!!!!

logo
NTDB id 1121823 EL140 RS01275 WP 000730772.1  FTDIRSAKETTGGSTDAEFMRINQFYMQTSQNMAKYQGLKTAGKDIELKYLGVYVLTVTDNSTFKGILNIADTVTAVNDK  158
NTDB id 360 SMU RS02495 WP 002262039.1        FTEITSAKETTGGVSDADYMRINQFYMETSQNGAIYKALSLAHKKASFNYMGVYVLQVSRNSTFKGVLNIADTVTGVNDK  160
consensus                                     !!*! !!!!!!!! *!!**!!!!!!!! !!!! ! ! *!  !*!     !*!!!!! !* !!!!!!*!!!!!!!!*!!!!

logo
NTDB id 1121823 EL140 RS01275 WP 000730772.1  TFDSSKDLVDYVNSQQLGDTVKVTYEEDGKVKSAEGKIITLENGKNGIGIGLIDRTEVTSDVPIRFSTAGIGGPSAGLMF  238
NTDB id 360 SMU RS02495 WP 002262039.1        TFHSSKELIKYVSGLKLGSKVSVQYTSENKKKTAKGKVIKLSNGKNGIGIGLVDHTKVTSDDKIEFSTNGIGGPSAGLMF  240
consensus                                     !!*!!!*!* !!    !!  ! ! !  * ! !*! !!*! ! !!!!!!!!!!*!*! !!!! *! !!! !!!!!!!!!!!

logo
NTDB id 1121823 EL140 RS01275 WP 000730772.1  SLAIYTQIADPGLRNGRIVAGTGTIDRDGNVGDIGGIDKKVVAASRQGANIFFAPDNPVTEEAKKADPNAKSNYETALEA  318
NTDB id 360 SMU RS02495 WP 002262039.1        TLDIYDQLVKEDLRKGRIIAGTGTIGEDGSVGDIGGADMKVAAADRINADIFFVPNNPVSKATLKKNPKALTNYQEAKQA  320
consensus                                     *! !! !*  * !! !!!*!!!!!!  !! !!!!!! ! !! !! !  ! !!! ! !!!*    !  ! ! *!!  !  !

logo
NTDB id 1121823 EL140 RS01275 WP 000730772.1  AKTIKTEMKIVPVKTLQDAIDYLKNNP   345
NTDB id 360 SMU RS02495 WP 002262039.1        AKKLGTKMKIVPVKNVQEAIDYLRKH.   346
consensus                                     !! * ! !!!!!!! *!*!!!!!* **
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X non conserved

X similar

X ≥ 50% conserved