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logo

NTDB id 1119841 EL086 RS08560 WP 371212477.1  MFLKQLLRKWRPSVAKENASKSILGKLWRIKWWLLSIFTILFLLFALFFPLNNYYVELPGGAFDTKEVLTVDKKADDSKG   80
NTDB id 360 SMU RS02495 WP 002262039.1        .......................MKTNKKFKWWLISGISLILLMVFFFPL.PYYLEMPGGAYDIRSVVTVNHKEDKEKG   56
consensus                                     ************************    * !!!!*!  *** !!   !!!!**!!*!*!!!!*! * !*!! *! !  !!

logo

NTDB id 1119841 EL086 RS08560 WP 371212477.1  SYNFVAVAQTKATLALMLYAQLNDFAKLQTAEEATGNYSDEDFMRINQFYMETSQNQAIYQGLTLAGKEASLEYMGVYVL   160
NTDB id 360 SMU RS02495 WP 002262039.1        SYNFVAVSLSKATPIQVLYAWLTPFTEITSAKETTGGVSDADYMRINQFYMETSQNGAIYKALSLAHKKASFNYMGVYVL   136
consensus                                     !!!!!!!  *!!!*  *!!! ! *!  * *! ! !!  !! !*!!!!!!!!!!!!! !!! *!*!!*! !!  !!!!!!!

logo

NTDB id 1119841 EL086 RS08560 WP 371212477.1  QVADDSSFKGVLNIADTVTAVNGKSFDNSTDLIKYVQGLKLGSKVKVTYTTDDKEKTATGKIIKIANGKNGIGIGLTDHT   240
NTDB id 360 SMU RS02495 WP 002262039.1        QVSRNSTFKGVLNIADTVTGVNDKTFHSSKELIKYVSGLKLGSKVSVQYTSENKKKTAKGKVIKLSNGKNGIGIGLVDHT   216
consensus                                     !!   !*!!!!!!!!!!!!!*!! !*!* ! *!!!!! !!!!!!!! ! !!** ! !!! !!*!!* !!!!!!!!!! !!!

logo

NTDB id 1119841 EL086 RS08560 WP 371212477.1  EVKSPEDVKFKLDGVGGPSAGLMFTLAIYDQVSGQDLKAGRKIAGTGTIEKDGSVGDIGGAYLKVKSAADSGADIFFVPN   320
NTDB id 360 SMU RS02495 WP 002262039.1        KVTSDDKIEFSTNGIGGPSAGLMFTLDIYDQLVKEDLRKGRIIAGTGTIGEDGSVGDIGGADMKVAAADRINADIFFVPN   296
consensus                                      ! !** * !   !*!!!!!!!!!!! !!!!*   !!* !! !!!!!!!  !!!!!!!!!! *!!  !    !!!!!!!!

logo

NTDB id 1119841 EL086 RS08560 WP 371212477.1  NPVTKEMKKADPDAKTNYQEAKEAAKKLGTKMKIVPVKTAQEAIDYLKKTK   371
NTDB id 360 SMU RS02495 WP 002262039.1        NPVSKATLKKNPKALTNYQEAKQAAKKLGTKMKIVPVKNVQEAIDYLRKH.   346
consensus                                     !!!*!   !  ! ! !!!!!!! !!!!!!!!!!!!!!!  !!!!!!!*!**
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X non conserved

X similar

X ≥ 50% conserved