```
logo
NTDB id 80778 HXS57 RS06790 WP 006564967.1  MCLRIWLLLFLGCLLV......ADSTAQPEEEEEEPPIYSMQDLHAIQSSFFKREREGDLDNTLFMDYRLGQTPKLRLRYAMITTL   79
NTDB id 1217 CAA10656.1 1440..2420( )       ..MRKVLYALMGFLLAFSALKADDFLEEANETAPANLNHPMQDLNAIQGSFFDKNRSKMSNTLNIDYFQGQTYKIRLRYAMATLL   83
consensus                                   ***!  !   *!*!! ****** !    * !  * ** **!!!!*!!! !!! * !  * !!! *!!  !!!*!*!!!!!! ! !

logo
NTDB id 80778 HXS57 RS06790 WP 006564967.1  VFSEPIREVILGDSVGFSTKMLGHERHQNSNVLLIKPLQIGIDSNLSVVGKSGKIYTFYIFSTTFTSPKHPVLNVYVSNKHFFQS  164
NTDB id 1217 CAA10656.1 1440..2420( )       FFSKPISDFVLGDKVGFDAKILES....NDRILLIKPLQIGVDSNISVIDSEGKIFSFYVFSTTFTSSKHPNLQVFIEDKNYYSN  164
consensus                                    !! !! * *!!! !!!  !*! *****!  *!!!!!!!!!*!!!*!!*   !!!**!!*!!!!!!!*!!! !*!**  !***

logo
NTDB id 80778 HXS57 RS06790 WP 006564967.1  SPDQPLAKFEEKEIL......KSNVLESK..KDTPDSTRYLKIGDASNAMWIDKTQIKRNYKVLGSRKRAWFCLWLCRIDTAKLI  241
NTDB id 1217 CAA10656.1 1440..2420( )       AFLKPQNKENVLENALENTPTNNKPLKEKKEETKEKEEETITIGDNTNAMKIVKKDIQKGYKALKSSQRKWYCLGICSKKSKLSL  249
consensus                                    *  !  !    !  ******   *!  !**   *     * !!! *!!! ! !  ! * !! ! !  ! !*!! *!   *   *

   <0
logo
NTDB id 80778 HXS57 RS06790 WP 006564967.1  KPLDIFNDAHFTYFKFDRKASQVKFPVAYKVVDGYDNPINTRIVGDYLIAEDIANKWTLREGKLHACIRRTRP.....  314
NTDB id 1217 CAA10656.1 1440..2420( )       MPKEIFNDKQFTYFKFDKRLALSKFPVIYKVVDGYDNPVNTRIVGDYIIAEDVSTKWTLRLGKDYLCIRFVKKGKDE*  326
consensus                                    ! *!!!! *!!!!!!!**    !!!! !!!!!!!!!!*!!!!!!!!*!!!!*  !!!!! !! * !!!  ******
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X non conserved

X similar

X ≥ 50% conserved