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logo
NTDB id 667414 MPF95 RS06290 WP 245066660.1  MKQSLLKKIGLVIGGFGALMALGLLANKGS..NNNNKEEIYSIEESKFPLSDYFFEEVKEEPKKEENPKTITTPQPTAPT   78
NTDB id 1218 CAA10657.1 2413..3543( )        .MNKWLKGVIVFIGGFATITTISLVYHQKPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKD.ATIKA.....   73
consensus                                    * *  !! * * !!!!* *  * !* *  **** !! * *   *! !*!! !! !   * **  !  * * ! * *****

logo
NTDB id 667414 MPF95 RS06290 WP 245066660.1  TANVSTNPIITDYSHLNSNLNKQNDTIM..HLQQVLQS....PPKMKPKLKHIKEQLEFLNTRLKPLEPTKQA......I   146
NTDB id 1218 CAA10657.1 2413..3543( )        ..........LQEQLKAALKALNSKEMNYSKEETFKSPPIDPNTTPPKKDFSLKQLDLLAARITPFKQSPKNYEENLIF   142
consensus                                    ***********   *!   !   !   !***      !****!*  *!!  **  !!* !  !* !  ***  ******

logo
NTDB id 667414 MPF95 RS06290 WP 245066660.1  KPETEYGVDSFSNLRYKDIGTNEHKLLRTITADRMIPAFLITPISSQIAGKVTAQVESDIFASMGRAVLIPKGSKVIGYY   226
NTDB id 1218 CAA10657.1 2413..3543( )        PVDNPKGIDGFTNLKEKDIATNENKLLRTITADKMIPAFLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYY   222
consensus                                    *** * !*! !*!!* !!!*!!!*!!!!!!!!!*!!!!!!!!!!!!!!!!!!! !!!!!!!!!*!!*!!!!!!!!!!!!!!

logo
NTDB id 667414 MPF95 RS06290 WP 245066660.1  NNNNQIGQYRLNIAWTRIITPQGINIMLTNARGADVKGYNGLVGKYISRNFQRYGIPLLTNTLSNGLLIGVASALANQQN   306
NTDB id 1218 CAA10657.1 2413..3543( )        SNNNKMGEYRLDIVWSRIITPHGINIMLTNAKGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGN   302
consensus                                     !!! *! !!! ! !*!!!!!*!!!!!!!!!*!!!*!!!!!!!  ! !!!!!!!*!!!  !!*!!!!!!* !!! !  !

logo
NTDB id 667414 MPF95 RS06290 WP 245066660.1  KKCDGNPFFGDYLLMQLTRNTGMGINQVINQMLRQYGAQNPIIIIREGSRVFITPNLDIFIPKPKDGEVLAEFFKEKKPL   386
NTDB id 1218 CAA10657.1 2413..3543( )        KEGA.TNFFGDYLLMQLMRQSGMGINQVVNQILRDKSKIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK ....   376
consensus                                    ! ! * *!!!!!!!!!! !**!!!!!!!*!!*!!       !!**!!!!!!!!!*!! !!! ! !** !!*!!! !*****

logo
NTDB id 667414 MPF95 RS06290 WP 245066660.1  IKKSNEEVNNEEEI   400
NTDB id 1218 CAA10657.1 2413..3543( )        ..............   376
consensus                                    **************
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X non conserved

X similar

X ≥ 50% conserved