```
logo
NTDB id 667710 MPF98 RS02420 WP 245049743.1  MKQSLLKKIGLGVGGFVVLMALGLLANK..GSNNNNKEEIFQVSESKFPLSDYFFEEVKEEPKKEESPKTITTPQPTAPT   78
NTDB id 1218 CAA10657.1 2413..3543( )        .MNKWLKGVIVFIGGFATITTISLVYHQKPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKDA.TIKALQE..   76
consensus                                    * *  !! * * *!!!  *  * !* * **  * !! * * *  ! !*!! !! !   * **  ! !*  *! * *  **

logo
NTDB id 667710 MPF98 RS02420 WP 245049743.1  TANFSTNPIITDYSHLNSNLNKQNDTITHLQQVLQSPPKMKPKLKHTKEQLEFLNTRLKPLEPTKPA......IKPETEY  152
NTDB id 1218 CAA10657.1 2413..3543( )        .......QLKAALKALNSKEMNYS.KEETFKSPPIDPNTTPPKKDFSLKQLDLLAARITPFKQSPKNYEENLIFPVDNPK  148
consensus                                    *********     *!!!      *   *   **  !*  *!!  **  !!* !  !* !  **** ****** *** *

logo
NTDB id 667710 MPF98 RS02420 WP 245049743.1  GVDSFSNLRYKDIGTNEHKLLRTITADRMIPAFLITPISSQIAGKVTAQVESDIFASMGRAVLIPKGSKVIGYYNNNNQI  232
NTDB id 1218 CAA10657.1 2413..3543( )        GIDGFTNLKEKDIATNENKLLRTITADKMIPAFLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYYSNNNKM  228
consensus                                    !*! !*!!* !!!*!!!*!!!!!!!!!*!!!!!!!!!!!!!!!!!! !!!!!!!!!*!!*!!!!!!!!!!!!!! !!! *

logo
NTDB id 667710 MPF98 RS02420 WP 245049743.1  GQYRLNIAWTRIITPQGINIMLTNARGADVKGYNGLVGKYISRNFQRYGIPLLTNTLSNGLLIGVTSALANQQNRKGDGN  312
NTDB id 1218 CAA10657.1 2413..3543( )        GEYRLDIVWSRIITPHGINIMLTNAKGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNKEGA.T  307
consensus                                    ! !!! ! !*!!!!!*!!!!!!!!!*!!!*!!!!!!!!  ! !!!!!!!*!!!  !!*!!!!!!*!!!! !  !* ! *

logo
NTDB id 667710 MPF98 RS02420 WP 245049743.1  PFLGDYLLMQLTRNTGMGINQVINQMLRQYGAQNPIIIIREGSRVFITPNLDIFTPKPKDGEVLAEFFKEKKPLIKKSNE  392
NTDB id 1218 CAA10657.1 2413..3543( )        NFFGDYLLMQLMRQSGMGINQVVNQILRDKSKIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK ..........  376
consensus                                    *! !!!!!!!! !**!!!!!!!*!!*!!      !!**!!!!!!!!!*!! !!! ! !** !!*!!! !***********

logo
NTDB id 667710 MPF98 RS02420 WP 245049743.1  EVNNEEEI   400
NTDB id 1218 CAA10657.1 2413..3543( )        ........   376
consensus                                    ********
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X non conserved

X similar

X ≥ 50% conserved