```
logo
NTDB id 88591 K6J72 RS06135 WP 221279224.1  MKPLVKKALIFGGAMVAVFATSLILNKPK.....KAPEVLDLKDTNYPLADYLFTPPKETQPKENPELAHLQALLQENAKQIAQL   80
NTDB id 1218 CAA10657.1 2413..3543( )       MNKWLKGVIVFIGGFATITTISLVYHQKPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKDATIKALQEQLKAALKAL   85
consensus                                   ! * *!  **! !*   *   !!* * *******  ! *!   *  !!! !! !!*** * * ! *  !** !!       *  !

logo
NTDB id 88591 K6J72 RS06135 WP 221279224.1  KA.QLQEAKKSPLPQPPQEMEPEQEPQEQEIEPPEPTQEELEIEETLNGRIQGFEEPK.........EAEIDYGASHFENLKTHD  155
NTDB id 1218 CAA10657.1 2413..3543( )       NSKEMNYSKEETFKSPPI..DPNTT.......PPKK.DFSLKQLDLLAARITPFKQSPKNYEENLIFPVDNPKGIDGFTNLKEKD  160
consensus                                     * **   !  * * !! ***!   *******!! **   !   * ! *!! *!  ************ * * !  *! !!! *!

logo
NTDB id 88591 K6J72 RS06135 WP 221279224.1  LATHENRLLRTITADKMIPAFLVTPISSQLAGKVVAQVESDIFANMGQAVLIPKGSKAIGYYSNNNKIGEYRLDIVWSRIITPQG  240
NTDB id 1218 CAA10657.1 2413..3543( )       IATNENKLLRTITADKMIPAFLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYYSNNNKMGEYRLDIVWSRIITPHG  245
consensus                                   *!!*!!*!!!!!!!!!!!!!!!!*!!!!!!*!!!!*!!!!!!!!!*!! !!!!!!!!! !!!!!!!!!*!!!!!!!!!!!!!!!*!

logo
NTDB id 88591 K6J72 RS06135 WP 221279224.1  VNITLTNAKGADVKGYSGLVGQMITHNFQRYGMPLLVSTLSNGLLIGLTSALAN...KTGKNNLFGDFLLMQLTRQTGLGLNQII  322
NTDB id 1218 CAA10657.1 2413..3543( )       INIMLTNAKGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNKEGATNFFGDYLLMQLMRQSGMGINQVV  330
consensus                                   *!! !!!!!!!!*!!! !!!! *! *!!!!!!*!!!*!!*!!!!!!*!!!! !***! !  ! !!!*!!!!! !!*!*!*!!**

logo
NTDB id 88591 K6J72 RS06135 WP 221279224.1  AQILKDKSNLKPIIIVREGSRVFISPNLDIFFPTPQKGEVVAQFFKHSSP   372
NTDB id 1218 CAA10657.1 2413..3543( )       NQILRDKSKIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK  ...   376
consensus                                    !!!*!!! * !!***!!!!!!!!!!! !!!!! !   !!*! ! !****
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X non conserved

X similar

X ≥ 50% conserved