```
logo
NTDB id 58740 K751 RS01495 WP 015643444.1  MSKSLLKKIGLVIGGFGALMTLGLLANKGS..NNNNKEEIYSIEESKFPLSDYFFEEVKEEPKKEENPKTITTPQPAAPTTANVS   83
NTDB id 1218 CAA10657.1 2413..3543( )      .MNKWLKGVIVFIGGFATITTISLVYHQKPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKD.ATIKA.........L   74
consensus                                  *    !! * * !!!!* * !* !* *  **** !! * *   *! !*!! !! !   * **  !  * * ! * *********

logo
NTDB id 58740 K751 RS01495 WP 015643444.1  TNPIITDYSHLNSNLNKQNDTITHLQQVLQSPPKMKPKLKHTKEQLEFLDTRLKPLEPTKP......ATKLETEYGVDSFSNLKY   162
NTDB id 1218 CAA10657.1 2413..3543( )      QEQLKAALKALNSKEMNYS.KEETFKSPPIDPNTTPKKDFSLKQLDLLAARITPFKQSPKNYEENLIFPVDNPKGIDGFTNLKE   158
consensus                                    **     *!!!      *   *   **  !*  *!!  **  !!* !  !* !  **********  *** * !*! !*!!!

logo
NTDB id 58740 K751 RS01495 WP 015643444.1  KDIGTNEHKLLRTITADRMIPAFLITPISSQIAGKVTAQVESDIFASMGRAVLIPKGSKVIGYYDNNNQIGQYRLNIAWTRIITP   247
NTDB id 1218 CAA10657.1 2413..3543( )      KDIATNENKLLRTITADKMIPAFLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYYSNNNKMGEYRLDIVWSRIITP   243
consensus                                  !!!*!!!*!!!!!!!!!*!!!!!!!!!!!!!!!!!! !!!!!!!!!*!!*!!!!!!!!!!!!!! !!!  *!  !!!  !  !*!!!!!

logo
NTDB id 58740 K751 RS01495 WP 015643444.1  QGINIMLTNARGADVKGYNGLVGKYISRNFQRYGIPLLTNTLSNGLLIGVTSALANQQNKKGDGNPFFGDYLLMQFTRNTGMGIN   332
NTDB id 1218 CAA10657.1 2413..3543( )      HGINIMLTNAKGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNKEGA.TNFFGDYLLMQLMRQSGMGIN   327
consensus                                  *!!!!!!!!!*!!!*!!!!!!!  ! !!!!!!!*!!!  !!*!!!!!!*!!!! !  !! ! * *!!!!!!!!!  !**!!!!!

logo
NTDB id 58740 K751 RS01495 WP 015643444.1  QVINQMLRQYGAQNPIIIIREGSRVFITPNLDIFIPKPKDGEVLAEFFKEKKPLIKKSNEEVNNEEEI   400
NTDB id 1218 CAA10657.1 2413..3543( )      QVVNQILRDKSKIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK ...................   376
consensus                                  !!*!!*!!      !!**!!!!!!!!!*!! !!! ! !** !!*!!! !******************
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X non conserved

X similar

X ≥ 50% conserved