```
logo
NTDB id 121579 EG63 RS07200 WP 038418227.1  MKQSLLKKIGLGVGGFGALMALGLLANKG..SNNNNKEEIYSIEESKFPLSDYFFEEVKEKPKKEENPKTITTPQPTAPTTANVS   83
NTDB id 1218 CAA10657.1 2413..3543( )       .MNKWLKGVIVFIGGFATITTISLVYHQKPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKD.ATIKAL.........   74
consensus                                   * *  !! * * *!!!* *  * !* *  ** * !! * *   *! !*!! !! !   * **  !  * * ! * **********

logo
NTDB id 121579 EG63 RS07200 WP 038418227.1  TNPIITDYSHLNNNLNKQNDTITHLQQVLQS....PPKMKPKLKHTKEQLEFLNTRLKPLEPTKQA......IKPETEYGVDSFT   158
NTDB id 1218 CAA10657.1 2413..3543( )       QEQLKAALKALN....SKEMN.YSKEETFKSPPIDPNTTPPKKDFSLKQLDLLAARITPFKQSPKNYEENLIFPVDNPKGIDGFT   154
consensus                                     **     *!!****     * *      !****!*  *!!  **  !!* !  !* !  ***  ****** *** * !*! !!

logo
NTDB id 121579 EG63 RS07200 WP 038418227.1  NLKYKDIGTNEHKLLRTITADKMIPAFLITPISSQIAGKVTAQVESDIFASMGRAVLIPKGSKVIGYYNNNNQIGQYRLNIAWTR   243
NTDB id 1218 CAA10657.1 2413..3543( )       NLKEKDIATNENKLLRTITADKMIPAFLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYYSNNNKMGEYRLDIVWSR   239
consensus                                   !!! !!!*!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!*!!*!!!!!!!!!!!!!! !!! *! !!! ! !*!

logo
NTDB id 121579 EG63 RS07200 WP 038418227.1  IITPQGINIMLTNARGADVKGYNGLVGKYISRNFQRYGIPLLTNTLSNGLLIGITSALANRQNRKGDGNPFFGDYLLMQLTRNTG   328
NTDB id 1218 CAA10657.1 2413..3543( )       IITPHGINIMLTNAKGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNKEGA.TNFFGDYLLMQLMRQSG   323
consensus                                   !!!!*!!!!!!!!!*!!!*!!!!!!!!  ! !!!!!!!*!!!  !!*!!!!!!!!!!!! !! !* ! * *!!!!!!!!!! !**!

logo
NTDB id 121579 EG63 RS07200 WP 038418227.1  MGINQVVNQMLRQYGAQNPIIIIREGSRVFISPNLDIFIPKPKDGEVLAEFFKEKKPPIKTKEEINNEEEI   399
NTDB id 1218 CAA10657.1 2413..3543( )       MGINQVVNQILRDKSKIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK ................   376
consensus                                   !!!!!!*!!*!!      !!**!!!!!!!!!!!! !!! ! !** !!*!!! !*****************
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X non conserved

X similar

X ≥ 50% conserved