```
logo
NTDB id 1146411 CS885 RS07815 WP 089086745.1  MKQSLLKKIGLGVGGFVVLMALGLLANKG..SNNNNKEEIFQVSESKFPLSDYFFEEVKEEPKKAKETPEVTTQEPTTQT   78
NTDB id 1218 CAA10657.1 2413..3543( )         .MNKWLKGVIVFIGGFATITTISLVYHQKPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTES.SKDATIKALQEQL   78
consensus                                     * *  !! * * *!!!  *  * !* *  ** * !! * * *  ! !*!! !! !   * **    **** !   *  !

logo
NTDB id 1146411 CS885 RS07815 WP 089086745.1  SPNPSVSANPNIKDYSHLNNNLNQQNNTITHLQQVLQSPSKIKPKPKHTKEQLEFLNTRLKPLEPTKPATK......PEM  152
NTDB id 1218 CAA10657.1 2413..3543( )         KA.ALKALNSKEMNYSK..........EETFKSPPIDPNTTPPKKDFSLKQLDLLAARITPFKQSPKNYEENLIFPVDN  146
consensus                                      ***    !*    !!************  *   **  !   *!!* **  !!* !  !* !  ****   ********

logo
NTDB id 1146411 CS885 RS07815 WP 089086745.1  EYGVDSFTNLKYKDIGTNEHKLLRTITADRMIPAFLITPISSQIAGKVTAQVESDIFASMGRAVLIPKGSKVIGYYNNNN  232
NTDB id 1218 CAA10657.1 2413..3543( )         PKGIDGFTNLKEKDIATNENKLLRTITADKMIPAFLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYYSNNN  226
consensus                                     * !*! !!!!! !!!*!!!*!!!!!!!!!*!!!!!!!!!!!!!!!!!! !!!!!!!!!*!!*!!!!!!!!!!!!!! !!!

logo
NTDB id 1146411 CS885 RS07815 WP 089086745.1  QIGQYRLNIAWTRIITPQGINIMLTNARGADVKGYNGLVGKYISRNFQRYGIPLLTNTLSNGLLIGITSALANRQNRKGD  312
NTDB id 1218 CAA10657.1 2413..3543( )         KMGEYRLDIVWSRIITPHGINIMLTNAKGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNKEGA  306
consensus                                      *! !!! ! !*!!!!!*!!!!!!!!!!*!!!*!!!!!!!!  ! !!!!!!!*!!!  !!*!!!!!!!!!!! !! !* !

logo
NTDB id 1146411 CS885 RS07815 WP 089086745.1  GNPFFGDYLLMQLTRNTGMGINQVINQMLRQYGAQNPIIIIREGSRVFISPNLDIFIPKPKDGEVLAEFFKEKKPLIKTK  392
NTDB id 1218 CAA10657.1 2413..3543( )         .TNFFGDYLLMQLMRQSGMGINQVVNQILRDKSKIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK........  376
consensus                                     * *!!!!!!!!!! !**!!!!!!!*!!*!!      !!**!!!!!!!!!!!! !!! ! !** !!*!!! !*********

logo
NTDB id 1146411 CS885 RS07815 WP 089086745.1  EEINNEEEI   401
NTDB id 1218 CAA10657.1 2413..3543( )         .........   376
consensus                                     *********
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X non conserved

X similar

X ≥50% conserved