```
logo
NTDB id 261867 CYK58 RS03715 WP 126131098.1  MKQSLLKKIGLVIGGFGALMALGLLANKG..SNNNNKEEIFQVNESKFPLSDYFFEEVKEEPKKEENPKT.ITTPQPTAP    77
NTDB id 1218 CAA10657.1 2413..3543( )        .MNKWLKGVIVFIGGFATITTISLVYHQKPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKDATIKALQ....    75
consensus                                    * *  !! * * !!!!* *  * !* *  ** * !! * * *  ! !*!! !! !   * **  !  *  *!  *!****

logo
NTDB id 261867 CYK58 RS03715 WP 126131098.1  TTTNVSTNPIITDYSHLNSNLNKQNDTITHLQQVLQS....PPKMKPKLKHTKEQLEFLNTRLKPLEPTKPAT......K   147
NTDB id 1218 CAA10657.1 2413..3543( )        .......EQLKAALKALNSKE....MN.YSKEETFKSPPIDPNTTPPKKDFSLKQLDLLAARITPFKQSPKNYEENLIFP   143
consensus                                    ******* **     *!!!  ****  * *      !****!*  *!!  **  !!* !  !* !  ****  *******

logo
NTDB id 261867 CYK58 RS03715 WP 126131098.1  PETEYGVDSFSNLRYKDIGTNEHKLLRTITADRMIPAFLITPISSQIAGKVTAQVESDVFASMGRAVLIPKGSKVIGYYN   227
NTDB id 1218 CAA10657.1 2413..3543( )        VDNPKGIDGFTNLKEKDIATNENKLLRTITADKMIPAFLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYYS   223
consensus                                    ** * !*! !*!!* !!!*!!!*!!!!!!!!!*!!!!!!!!!!!!!!!!!!! !!!!!!*!!*!!*!!!!!!!!!!!!!!

logo
NTDB id 261867 CYK58 RS03715 WP 126131098.1  NNNQIGQYRLNIAWTRIITPQGINIMLTNARGADVKGYNGLVGKYISRNFQRYGIPLLTNTLSNGLLIGITSALANQQNK   307
NTDB id 1218 CAA10657.1 2413..3543( )        NNNKMGEYRLDIVWSRIITPHGINIMLTNAKGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNK   303
consensus                                    !!! *! !!! ! !*!!!!!*!!!!!!!!!*!!!*!!!!!!!!  ! !!!!!!!*!!!  !!*!!!!!!!!!!! !  !!

logo
NTDB id 261867 CYK58 RS03715 WP 126131098.1  KGDGNPFFGDYLLMQLTRNTGMGINQVINQMLRQYGAQNPIIIREGSRVFISPNLDIFIPKPKDGEVLAEFFKEKKPLI   387
NTDB id 1218 CAA10657.1 2413..3543( )        EGA.TNFFGDYLLMQLMRQSGMGINQVVNQILRDKSKIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK .....   376
consensus                                     ! * *!!!!!!!!!! !**!!!!!!!*!!*!!      !!**!!!!!!!!!!!! !!! ! !** !!*!!! !******

logo
NTDB id 261867 CYK58 RS03715 WP 126131098.1  KKSNEEVNNEEEI   400
NTDB id 1218 CAA10657.1 2413..3543( )        .............   376
consensus                                    *************
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X non conserved

X similar

X $\geq$ 50% conserved