```
logo
NTDB id 73926 FOY54 RS02555 WP 176485116.1  MNSLVKKLIIGLGMGVVIMASSLWWHKRNEQP....EAVLEVEESNYPLSDYLFVPKQQQKEDPKDS.EDYKRLEQILEENAKQI   80
NTDB id 1218 CAA10657.1 2413..3543( )       MNKWLKGVIVFIGGFATITTISLVYHQKPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKDATIKALQEQLKAALKAL   85
consensus                                   !!  *! *!* *!    !  !! *! **  !***** *!  *!  !!! !! ! * *!*  *   *   ! !   !    ! *

logo
NTDB id 73926 FOY54 RS02555 WP 176485116.1  TALKAQLK..QASQKPP.PQITPPPPQEIDSTQQDILASRIKAFEISQDEPKEQEPIS.VLDYGVDHFDNLKAKDVATHENRLLR  161
NTDB id 1218 CAA10657.1 2413..3543( )       NSKEMNYSKEETFKSPPIDPNTTPPKKDFSLKQLDLLAARITPFKQSPKNYEENLIFPVDNPKGIDGFTNLKEKDIATNENKLLR  170
consensus                                        *  **      !!*** !*!* *    ! !*!! !! *!  !*  * !* * **  * !*!*! !!! !!*!!*!!*!!!

logo
NTDB id 73926 FOY54 RS02555 WP 176485116.1  TITADKMIPAFLITPISSQLGGKVIAQVESDVFANMGRAILIPKGSKVIGYYNNNNKIGEYRLDIVWTRILTPQGVNIMLTNAKG  246
NTDB id 1218 CAA10657.1 2413..3543( )       TITADKMIPAFLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYYSNNNKMGEYRLDIVWSRITPHGINIMLTNAKG  255
consensus                                   !!!!!!!!!!!!!!!!!!!!**!!!!!!!!!!*!!*!*!*!!!!!!!!!!!! !!!!*!!!!!!!!!*!!*!!*!*!!!!!!!!!

logo
NTDB id 73926 FOY54 RS02555 WP 176485116.1  ADVKGYSGLVGEMITHNFQRYGMPLLVSTISNGLLIALTSAIAN...KTGKNNFFGDYLLMQMTRQTGMGLNQIVAQILRDKSNL  328
NTDB id 1218 CAA10657.1 2413..3543( )       ADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNKEGATNFFGDYLLMQLMRQSGMGINQVVNQILRDKSKI  340
consensus                                   !!*!!! !!!!!*! *!!!!!!*!!!*!!**!!!!!**!!!* !***! !  !!!!!!!!!!* !!*!!!*!!*! !!!!!!! *

logo
NTDB id 73926 FOY54 RS02555 WP 176485116.1  KPIIIIREGSRVFIAPNLDIFFPVPKEGEILAQFFTHSLP   368
NTDB id 1218 CAA10657.1 2413..3543( )       APIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK ...   376
consensus                                    !!**!!!!!!!!! !! !!!!!*!*! !**! !  ****
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X non conserved

X similar

X ≥ 50% conserved