```
logo
NTDB id 1010563 ABRA92 RS21355 WP 350225956.1  MCTKNETKPLPSFIEERLNFHIQDLIKSNENQKHLVLGKRPSENAVVMQSNDYLSLSHNELIQKAHRDAISERDDNVVMS   80
NTDB id 1155 A1552VC RS17030 WP 001039912.1    ....MNKPQLPDFIQNKIDHYIENYFDINKNGKHLVLGKQASPDDIILQSNDYLALANHPLIKARLAKSLLEEQQSLFMS   76
consensus                                      ****   **!! !!  ** **!      ! ! !!!!!!! *!*  ***!!!!!! ! ***!!   *   * !    * !!

logo
NTDB id 1010563 ABRA92 RS21355 WP 350225956.1  AIFLQDDQSKPAFEHQLATFVGMESCLLSQSGWAANIGLLQTICAPNVPVYIDFFAHMSLWEGARTAGAQIHPFMHNNMN  160
NTDB id 1155 A1552VC RS17030 WP 001039912.1    ASFLQNDYDKPMIEKRLAKFTGFDECLLSQSGWNANVGLLQTICQPNTNVYIDFFAHMSLWEGARYANAQAHPFMHNNCD  156
consensus                                      ! !!! !  !!  !* !! ! ! * !!!!!!!! !!*!!!!!!! !! *!!!!!!!!!!!!!!!! ! !! !!!!!!!*

logo
NTDB id 1010563 ABRA92 RS21355 WP 350225956.1  HLRKQIQRHGAGIIVVDSVYSTIGTIAPLRAIYEMAKEFDCGLVVDESHSLGTHGPKGSGLLQELGLTQMVDFVTVSLAK  240
NTDB id 1155 A1552VC RS17030 WP 001039912.1    HLRMLIQRHGPGIIVVDSIYSTLGTIAPLAELVNISKEFGCALLVDESHSLGTHGPNGAGLLAELGLTREVHFMTASLAK  236
consensus                                      !!!  !!!!!*!!!!!!!*!!!*!!!!!!  *  * !!! !*!*!!!!!!!!!!!! ! !!! !!!!!  !*!*! !!!!

logo
NTDB id 1010563 ABRA92 RS21355 WP 350225956.1  TFAYRAGAILGPNKLAQSLPFVAYPAIFSSTVLPQEVVRLEKTLEVIKEADDKRECLFKRAKELAIGLKRIGFNIRSESQ  320
NTDB id 1155 A1552VC RS17030 WP 001039912.1    TFAYRAGAIWCNNEVNRCVPFISYPAIFSSTLLPYEAAGLETTLEIIESADNRRQHLDRMARKLRIGLSQLGLTIRSESQ  316
consensus                                      !!!!!!!!! **! *  **!!* !!!!!!!!*!! !   !! !!!*!  !! *!  ! * !* ! !!!  *!  !!!!!!

logo
NTDB id 1010563 ABRA92 RS21355 WP 350225956.1  IIALECGSERNTERVRDFLEERGVFGAVFCRPATGRNKNIIRFSVNADMTAQQVDHVLSACQEAFDHPELEFV  393
NTDB id 1155 A1552VC RS17030 WP 001039912.1    IIGLETGDERNTEKVRDYLESNGVFGSVFCRPATSKNKNIIRLSLNSDVNDEQIAKIIEVCSDAVNYGDFYFR  389
consensus                                      !!*!!*! !!!!!*!!!*!!  !!!! !!!!!!! *!!!!!! !*! !*   !* ***  ! *!  ***  !
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X non conserved

X similar

X $\geq$ 50% conserved