```
logo
NTDB id 668294 MON40 RS02025 WP 003779857.1  MSDNKQNEVLTGYEQLKRRNRRRLVMASGLVAASTVLLGVALSTGPENKPEENAQTTTIQQNNANAEPANLADATVLEPS   80
NTDB id 1129 NGFG RS01435 WP 003687618.1     MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGA...TESQTANTAQ....TPA   73
consensus                                    !!*!!!!!!!!!!!!!!!!!!!!!! !! !!!!!**!!* !!!* !   ** *     !  ***     *!! ! **** !

logo

NTDB id 668294 MON40 RS02025 WP 003779857.1  TKEDLEAAAEAAKNADAEVADKPQETAAPEQTAQAEPQPDAPPAAPEDVGPPLVLINDTLSDSDIKGLEESEKIQKAEAA  160
NTDB id 1129 NGFG RS01435 WP 003687618.1     ....LKSA....AENGETAADKPQDLAGEDKP....SAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLQQAETA  141
consensus                                    ****!  !****       !!!!!* !*** ****** *! **  !! !!*!!!!!! ! !! !!!!!!!!!*! !! !

logo

NTDB id 668294 MON40 RS02025 WP 003779857.1  KREAAERRAEERRRNREEQRKAQQLQAQEAAEREANAAARKRALQAAKAEKAERAAAAERNKLAQLEKAEKAVAERKALK  240
NTDB id 1129 NGFG RS01435 WP 003687618.1     KTEPKQAKQRA..........AEKVSATADST.DTVAVEKPKRSAEPKPQKAERTAEAKP..............KAKETK  196
consensus                                    ! !*   *   **********!  * !     **  !  ***    *!* !!!! ! ! ***************  !  !

logo

NTDB id 668294 MON40 RS02025 WP 003779857.1  AKEEAAAKHKAAAEKAKETVTASASEPKERIRVDASKYEKIETANKKASAVREAEAKVAKAKTEQTAKAEKTATAERSSE  320
NTDB id 1129 NGFG RS01435 WP 003687618.1     TAEKVADKPKTAAEKTKPDTAKSDSAVKEAKKADK..AEGKKTAEK..........DRSDGKKHET..AQKTDKA..DKT  260
consensus                                      !  ! ! ! !!!! !*    ! ! *!!  * ! ** !   !! !**********    *! * !**! !!  !**

logo

NTDB id 668294 MON40 RS02025 WP 003779857.1  KAKTKTAEVKSGKKAAIQAGYAEKERALSLQRKMKAAGIDSTITEVMTDKGKVYRVKSGAYKNAYDAERDLNKLRVHGIA  400
NTDB id 1129 NGFG RS01435 WP 003687618.1     KTAEKEKSGKAGKKAAIQAGYAEKERALSLQRKMKAAGIDSTITEIMTDNGKVYRVKSSNYKNARDAERDLNKLRVHGIA  340
consensus                                    !   !    ! !!!!!!!!!!!!!!!!!!!!
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