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logo

NTDB id 223322 A6J88 RS27720 WP 080614426.1  MSDNKQNEVLTGYEQLKRRNRRRLVMASGLVAASTVLLGVALSTGPENKPEEENAQTTTIQQNNANAEPANLADATVLEP   80
NTDB id 1129 NGFG RS01435 WP 003687618.1     MSENKQNEVLTGYEQLKRRNRRRLVTASSLVAASCILLAAALSSDPADSNPAPQAGETGATE....SQTANTAQT....P   72
consensus                                    !!*!!!!!!!!!!!!!!!!!!!!!! !! !!!!!**!!* !!!* !   ** **!   !    ****  *!! !  ****!

logo

NTDB id 223322 A6J88 RS27720 WP 080614426.1  STKEDLEAAAEAAKNADAEVADKPQETAAPEQTAQAEPQPDAPPAAPEDVGPPLVLINDTLSDSDIKGLEESEKIQKAEA  160
NTDB id 1129 NGFG RS01435 WP 003687618.1     A....LKSA....AENGETAADKPQDLAGEDKP....SAADSEISEPENVGAPLVLINDRLEDSNIKGLEESEKLQQAET  140
consensus                                     ****!  !****       !!!!!* !*** ****** *! **  !! !!*!!!!!!! ! !! !!!!!!!!!*! !!

logo

NTDB id 223322 A6J88 RS27720 WP 080614426.1  AKREAAERRAEERRRNREEQRKAQQLQAQEAAEREANAAARKRALQAAKAEKAERAAAAERNKLAQLEKAEKAVAERKAL  240
NTDB id 1129 NGFG RS01435 WP 003687618.1     AKTEPKQAKQRA..........AEKVSATADST.DTVAVEKPKRSAEPKPQKAERTAEAKP..............KAKET  195
consensus                                    !! !*   *   **********!  * !     **  !  ***    *!* !!!! ! ! ***************  !

logo

NTDB id 223322 A6J88 RS27720 WP 080614426.1  KAKEEAAAKHKAAAEKAKEAVTASASEPKERIRVDASKYEKIETANKKASAVREAEAKVAKAKTEQTAKAEKTATAERSS  320
NTDB id 1129 NGFG RS01435 WP 003687618.1     KTAEKVADKPKTAAEKTKPDTAKSDSAVKEAKKADK..AEGKKTAEK..........DRSDGKKHET..AQKTDKA..DK  259
consensus                                    !  !  ! ! ! !!!! !*    ! ! *!!  * ! ** !   !! !**********    *! * !**! !!  !**

logo

NTDB id 223322 A6J88 RS27720 WP 080614426.1  EKAKTKTAEVKSGKKAAIQAGYAEKERALSLQRKMKAAGIDSTITEVMTDKGKVYRVKSGAYKNAYDAERDLNKLRVHGI  400
NTDB id 1129 NGFG RS01435 WP 003687618.1     TKTAEKEKSGKAGKKAAIQAGYAEKERALSLQRKMKAAGIDSTITEIMTDNGKVYRVKSSNYKNARDAERDLNKLRVHGI  339
consensus                                     !   !    ! !!!!!!!!!!!!!!!!!!!!
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