```
logo

NTDB id 1013375 ABVJ71 RS06380 WP 353856134.1  MKYQRLVMIFSFLLLLSACSQAPLKGHIEKVGLLVPDTINDQVWGTKGYKGLLSIQSTFDVDVYYKEGMVDKEKIVDAIE   80
NTDB id 94 BSU 11300 NP 389012.2               .MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIE   79
consensus                                      *    !!!!!!! !!!!!*! ! ! !!*!!!!!*! !!!!!! !!!!!!!!!!! !!! * !!!!!!!!*    ! !* !!!

logo

NTDB id 1013375 ABVJ71 RS06380 WP 353856134.1  DFHKKGVNLIIGHGNEYSDIFNLISEDYPKIQFITVNGGKPRANNVTNVTFKGEAMGFFGGMTAAHMSKTKKIGILATYD  160
NTDB id 94 BSU 11300 NP 389012.2               DFHKRGVNLLYGHGSEYAEVFNLVGEDYPDMEFVISN.AKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFT  158
consensus                                      !!!!*!!!!* !!! !! **!!!* !!!! * !*  !**!**! !!! !*! !!!!!!!!!!!!!!!!!!  *!**!**

logo

NTDB id 1013375 ABVJ71 RS06380 WP 353856134.1  WQSEVDGFIKGAKYQDSHVQVLAEFVENWDDAAKAVKLYEKLKKQGADVVYPAGDGYNIPVIEQIKADNLSAIGYVTDQS  240
NTDB id 94 BSU 11300 NP 389012.2               WQPEVDGFIKGAKYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQS  238
consensus                                      !!*!!!!!!!!!!!  *** !   * **!!!   !!!!! !*!  !!!!!!!!!!!!!*!!! !!! ! ! !!!!!!!!!

logo

NTDB id 1013375 ABVJ71 RS06380 WP 353856134.1  NLGSHTVLTSTVQHVDKAYSIIAKKFNEGTLNEKSEYSFDFKEGVIEMGKFSTTIDRSFVKDIKNEITYYKKTGKLPNEK  320
NTDB id 94 BSU 11300 NP 389012.2               DLGENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDH.YYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE  317
consensus                                       !! *!!!!!!!!*!!!!! !!!  !! !!!    ** *!   !!*!!! !!* *!  !   !   !  ! !!! !!
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X non conserved

X similar

X ≥ 50% conserved