```
logo
NTDB id 181367 BSA41 RS04070 WP 041089810.1  MKYQRLVMIFSFLLLLSACSQAPLKGKIEKVGLLVPDTINDQVWGTKGYKGLLSIQSTFGVDVYYKEGMVDKEKIVDAIE   80
NTDB id 94 BSU 11300 NP 389012.2             .MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIE   79
consensus                                    *    !!!!!!! !!!!!*! ! ! !!!!!!!!*! !!!!!! !!!!!!!!!!! !!! * !!!!!!!!*    ! !* !!!

logo
NTDB id 181367 BSA41 RS04070 WP 041089810.1  EFHKKGVNLIIGHGNEYSEIFNLISEDYPNIQFITVNGNKPQAGNVTNVTFKGEAMGFFGGMTAAHMSKTKKIGILATYD  160
NTDB id 94 BSU 11300 NP 389012.2             DFHKRGVNLLYGHGSEYAEVFNLVGEDYPDMEFVISN.AKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFT  158
consensus                                    *!!!*!!!!* !!! !! !*!!!* !!!! * !*  !* !* ! !!! !*! !!!!!!!!!!!!!!!!!!  *!**!**

logo
NTDB id 181367 BSA41 RS04070 WP 041089810.1  WQSEVDGFIKGAKYQDEHVQVLAEFVENWDDADKAVELYQKLKKQGVDVVYPAGDGYNIPVIEQIKADNLSAIGYVTDQS  240
NTDB id 94 BSU 11300 NP 389012.2             WQPEVDGFIKGAKYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQS  238
consensus                                    !!*!!!!!!!!!!!  *** !   * **!!!   !! !!!!*!  ! !!!!!!!!!!!*!!! !!! ! ! !!!!!!!!!

logo
NTDB id 181367 BSA41 RS04070 WP 041089810.1  NLGSHTVLTSTVQHVDKAYSIIAKKFNEGKLNEQGEYSFDFKEGVIEMGKFSSTIDDAFVKEIESDIASYKKTGKLPNEK  320
NTDB id 94 BSU 11300 NP 389012.2             DLGENTVLTSTVQNVDKAYEIIAEQFNKGTLEG.GDHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE  317
consensus                                     !! *!!!!!!!!*!!!!! !!!  !! ! !  *!** *!   !!*!!! !!* *!  !   !   ! *! !!! !!
```

X non conserved

X similar

X $\geq$ 50% conserved