```
logo
NTDB id 401305 GKC22 RS05420 WP 034663299.1  MKYQRLVMIFSFLLLLSACSQAPLKGQIEKVGLLVPDTINDQVWGTKGYKGLLNIQSTFGVDVYYKEGMVDKEKIVDAIE   80
NTDB id 94 BSU 11300 NP 389012.2             .MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIE   79
consensus                                    *   !!!!!!! !!!!!*! ! ! !! !!!!!*! !!!!!! !!!!!!!!!!!!!!! * !!!!!!!!*   ! !* !!!

logo
NTDB id 401305 GKC22 RS05420 WP 034663299.1  EFHKKGVNLIIGHGNEYSEIFNLISEDYPKTQFITVNGNKPQADNVANVTFKGEAMGFFGGMTAAHMSKSKKIGILATYD  160
NTDB id 94 BSU 11300 NP 389012.2             DFHKRGVNLLYGHGSEYAEVFNLVGEDYPDMEFVISN.AKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFT  158
consensus                                    *!!!*!!!!* !!! !! !*!!!* !!!!   !*  !* !* !!!!  !*! !!!!!!!!!!!!!!!!!*  *!**!**

logo
NTDB id 401305 GKC22 RS05420 WP 034663299.1  WQSEVDGFIKGAKYQNEHVQVLAEFVENWDDADKAVELYQKLKKQGVDVVYPAGDGYNIPVIEQIKADNLSAIGYVTDQS  240
NTDB id 94 BSU 11300 NP 389012.2             WQPEVDGFIKGAKYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQS  238
consensus                                    !!*!!!!!!!!!!!  *** !   * **!!!   !! !!!!*!  ! !!!!!!!!!!!*!!! !!! ! ! !!!!!!!!!

logo
NTDB id 401305 GKC22 RS05420 WP 034663299.1  NLGSHTVLTSTVQHVDKAYTIIAKKFNEGKLNEQGDYSFDFKEGVIEMGKFSSTIDRAFVKDIESDIATYKKTGKLPNEK  320
NTDB id 94 BSU 11300 NP 389012.2             DLGENTVLTSTVQNVDKAYEIIAEQFNKGTLE.GGDHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE  317
consensus                                     !! *!!!!!!!!*!!!!! !!!  !! ! ! * !!* *!   !!*!!! !!* *!  !   !   ! !! !!! !!
```

X non conserved

X similar

X $\geq$ 50% conserved