```
logo
NTDB id 965276 NSQ36 RS05580 WP 050945301.1  MKYQRLVMIFSFLLLLSACSQAPLKGQIEKVGLLVPDTINDQVWGTKGYKGLLNIQSTFGVDVYYKEGMVDKEKIVDAIE   80
NTDB id 94 BSU 11300 NP 389012.2             .MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIE   79
consensus                                    *   !!!!!!! !!!!!*! ! ! !! !!!!!*! !!!!!! !!!!!!!!!!!!!!! * !!!!!!!!*   ! !* !!!

logo
NTDB id 965276 NSQ36 RS05580 WP 050945301.1  EFHKKGVNLIIGHGSEYSEIFNIISEDYPKTQFITVNGNKPQADNVANVTFKGEAMGFFGGMTAAHMSKSKKIGILATYD  160
NTDB id 94 BSU 11300 NP 389012.2             DFHKRGVNLLYGHGSEYAEVFNLVGEDYPDMEFVISN.AKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFT  158
consensus                                    *!!!*!!!!* !!!!!! !*!!** !!!!   !*  !* !* !!!!  !*! !!!!!!!!!!!!!!!!!*  *!**!**

logo
NTDB id 965276 NSQ36 RS05580 WP 050945301.1  WQSEVDGFIKGAKYQDEHVQVLAEFVENWDDADKAVELYRKLEKQGVDVVYPAGDGYNIPVIEQIKADNLSAIGYVTDQS  240
NTDB id 94 BSU 11300 NP 389012.2             WQPEVDGFIKGAKYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQS  238
consensus                                    !!*!!!!!!!!!!!  *** !   * **!!!   !! !! !*   ! !!!!!!!!!!!*!!! !!! ! ! !!!!!!!!!

logo
NTDB id 965276 NSQ36 RS05580 WP 050945301.1  NLGSHTVLTSTVQHVDKAYSIIAKKFNEGKLNAQGEYSFDFKEGVIEMGKFSSTIDRAFLKDIESDIASYKKTGKLPNEK  320
NTDB id 94 BSU 11300 NP 389012.2             DLGENTVLTSTVQNVDKAYEIIAEQFNKGTLEGG.DHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE  317
consensus                                     !! *!!!!!!!!*!!!!! !!!  !! ! ! * *** *!   !!*!!! !!* *!  !   !   ! *! !!! !!
```

X non conserved

X similar

X $\geq$ 50% conserved