```
logo
NTDB id 1027659 AB3Z07 RS04615 WP 251684364.1  MYLKLGSVLIILLSLFGCGQPVASGELEKVGLLVTETIDEKVWGTKGYKGLLKIQSELNVDVYYKEGMNDEAGIREAIEE   80
NTDB id 94 BSU 11300 NP 389012.2               MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIED   80
consensus                                      !  *!  *  *!! ! !!!!**  ! *!!!!*! **!! * !!!!!!!!!!! !!!  !!!!!!!!!*  !  !  !!!*

logo
NTDB id 1027659 AB3Z07 RS04615 WP 251684364.1  FHNNGVNLIFGHGSEFANYFNEISSEYPDIHFVVFNGEANGENVTSLNFDSNAMGFFGGMVAGEMTKTNKVAVLAAFEWQ  160
NTDB id 94 BSU 11300 NP 389012.2               FHKRGVNLLYGHGSEYAEVFNLVGEDYPDMEFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFTWQ  160
consensus                                      !!  !!!!**!!!!!*!  !! *  *!!!**!!* !* ! **!!!!**!   !!!!!!!! !**!*!!! !*!*! ! !!

logo
NTDB id 1027659 AB3Z07 RS04615 WP 251684364.1  PEVDGFFEGAYYQNENVEVDIQYVQDWNNTDIALTLLDQMIEDGVDIVYVAGDGYNIAVIEKLKEEGLYAIGFVSDQSDL  240
NTDB id 94 BSU 11300 NP 389012.2               PEVDGFIKGAKYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDL  240
consensus                                      !!!!!!  !! ! !* *!!   !  *!     !* !   !  *! !*!!*!!!!!!**!!  *! *!!!!!!*!*!!!!!

logo
NTDB id 1027659 AB3Z07 RS04615 WP 251684364.1  GKGTVLTSTVQHVDTLYEVVAKEFDNGKLENGELFYDFKDGVISLGTFSPLVDEEFKSTVMESIEEYKETGLLPNQL  317
NTDB id 94 BSU 11300 NP 389012.2               GENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE  317
consensus                                      !  !!!!!!!!*!!  !!**!  !  ! !! !***!!   !!* *!!!!!!!! *!   *   !  !  !! !! *
```

X non conserved

X similar

X $\geq$ 50% conserved