```
logo
NTDB id 122987 BMMGA3 RS04395 WP 004433592.1  MLKRFGVILLCLLVLAGCEQPTKTGKLKKVGLLVPDTVNDQVWGTKGYKGMLKIQSRFNVDVYIKEGMNSELRVERAVKE   80
NTDB id 94 BSU 11300 NP 389012.2              MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIED   80
consensus                                     !* !  *!  *!!*! !! !**  !!* !!!*! !!!*!! !!!!!!!!!*! !!!**!!!!! !!!* *!  *  !* *

logo
NTDB id 122987 BMMGA3 RS04395 WP 004433592.1  FDQKGVNLIFGHGNEYAEYFNKISKNYPNIHFVSFNGDAKNKNTTSLNFQAYAMGFFGGMVASHMTKKNKVGIIAAYEWQ  160
NTDB id 94 BSU 11300 NP 389012.2              FHKRGVNLLYGHGSEYAEVFNLVGEDYPDMEFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFTWQ  160
consensus                                     !* *!!!!**!!! !!!! !! *   !! **!!  !* !!  ! !!**! * !!!!!!!! ! !!*! ! !!*!! * !!

logo
NTDB id 122987 BMMGA3 RS04395 WP 004433592.1  PEVEGFYEGAKFENKQVDVNIQYVGQWDDDQTAMTILDNMIKAGVDVVYPAGDGYNVPVIEKAKEQGLFAIGYISDESDL  240
NTDB id 94 BSU 11300 NP 389012.2              PEVDGFIKGAKYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDL  240
consensus                                     !!!*!!  !!!*!!* **!!  !  *!!!! !!* *   !   ! !!!!!!!!!!!!!!!   !  !!*!!!!**! !!!

logo
NTDB id 122987 BMMGA3 RS04395 WP 004433592.1  GKSTVLTSTIQHVDKLYELVAEQYNNGKLKSGNLHFDFKDGVISLGEFSPLIDDEFKAEINSYIEDYKKTGKLPNE.  316
NTDB id 94 BSU 11300 NP 389012.2              GENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE  317
consensus                                     !  !!!!!!*!*!!! !!**!!!*! ! !  ! ***!   !!* *! !!!!*! *!   !   !  ! !!! !!  *
```

X non conserved

X similar

X $\geq$ 50% conserved