```
logo
NTDB id 91292 L6432 RS00740 WP 020450927.1  MVYRLVAIFFILLALSGCGQSPLKGKIEKVGLLVPDTINDQVWGTKGYKGLLRIQSVYNVDVYYKEGIDNDAAIKMAVEDFHKKG   85
NTDB id 94 BSU 11300 NP 389012.2            MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIEDFHKRG   85
consensus                                   !* !!! !! *!! !!!!!!*! !!!!!!!!*! !!!!!! !!!!!!!!!!! !!! !!!!!!!!!*  *  !  !*!!!!!*!

logo
NTDB id 91292 L6432 RS00740 WP 020450927.1  VNLIYGHGSEYEKAFNEICKDYPDIHFVLMNSKADHPNVTSISLEGEAMGFFGGMTAAHQSKTKKIGVLATYKWQPEVEGFSEGA   170
NTDB id 94 BSU 11300 NP 389012.2            VNLLYGHGSEYAEVFNLVGEDYPDMEFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFTWQPEVDGFIKGA   170
consensus                                   !!!*!!!!!!!   !! ** !!!!**!!* ! !! **!!!!** !!!!!!!!!!!!!! !!!  *!!*!** !!!!!*!!  !!

logo
NTDB id 91292 L6432 RS00740 WP 020450927.1  KYLDPDIEVMTEYVGQWDDSENAVKLYEKMKKAGADVVYPAGDGYNVPVIEQIKKDGLYAIGYVSDQSELGENVVLTSTIQHVDD   255
NTDB id 94 BSU 11300 NP 389012.2            KYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDLGENTVLTSTVQNVDK   255
consensus                                   !!  !!!!! ! !  *!!!   !!!!! !!!  !!!!!!!!!!!!!!!!! !!!!!!!!!!!!!*!!!*!!!! !!!!!*!*!!

logo
NTDB id 91292 L6432 RS00740 WP 020450927.1  AYELTADEFNKGTLKGGKRSFDIQDGVIEMGEFSPVVDQSFQEKMNRLIDEYKKTNKLPNEH   317
NTDB id 94 BSU 11300 NP 389012.2            AYEIIAEQFNKGTLEGGDHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE   317
consensus                                   !!!* !* !!!!!! !! * *!** !!*!!! !!!*!!! !! ** *!!  ! !!  !!  *
```

X non conserved

X similar

X $\geq$ 50% conserved