```
logo
NTDB id 174681 B14 RS15650 WP 009328848.1  MVYRLVAIFFILLALSGCGQSPLKGKIEKVGLLVPDTISDQVWGTKGYKGLLRIQSVYNVDVYYKEGIDNDAAIKMAVEDFHKKG   85
NTDB id 94 BSU 11300 NP 389012.2           MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIEDFHKRG   85
consensus                                  !* !!! !! *!! !!!!!!*! !!!!!!!!*! !!!! ! !!!!!!!!!!! !!! !!!!!!!!!!*  *  !  !*!!!!!*!

logo
NTDB id 174681 B14 RS15650 WP 009328848.1  VNLIYGHGSEYEKAFNEICKDYPDIHFVLMNSKAEHQNVISISLEGEAMGFFGGMTAAHQSKTKKIGVLATYKWQPEVEGFAEGA  170
NTDB id 94 BSU 11300 NP 389012.2           VNLLYGHGSEYAEVFNLVGEDYPDMEFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFTWQPEVDGFIKGA  170
consensus                                  !!!*!!!!!!!   !! ** !!!!**!!* ! !! * !! !**  !!!!!!!!!!!!!! !!!  *!!*!** !!!!!*!!  !!

logo
NTDB id 174681 B14 RS15650 WP 009328848.1  KYLDPDIEVMTEYVGQWDDSENAVKLYEKMKKAGADVVYPAGDGYNVPVIEQIKKDGLYAIGYVSDQSELGENVVLTSTIQHVDD  255
NTDB id 94 BSU 11300 NP 389012.2           KYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDLGENTVLTSTVQNVDK  255
consensus                                  !!  !!!!! ! !  *!!!   !!!!! !!!  !!!!!!!!!!!!!!!!! !!!!!!!!!!!!!*!!!*!!!! !!!!!*!*!!

logo
NTDB id 174681 B14 RS15650 WP 009328848.1  AYELTADEFNKGTLKGGKRSFDIQEGVIEMGKFSPVVDPSFQEEMKRLIDEYKKTNKLPNEH  317
NTDB id 94 BSU 11300 NP 389012.2           AYEIIAEQFNKGTLEGGDHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE  317
consensus                                  !!!* !* !!!!!! !! * *!** !!*!!! !!!*!!* !!  * *!!  ! !!  !!  *
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X non conserved

X similar

X ≥ 50% conserved