```
logo

NTDB id 1144963 BGLY RS06425 WP 046131230.1  MVYRLVVIFLMLLAMCGCGQASLNGKIEKVGLLVPDTISDQVWGTKGYKGLLRIQSVYGVDVYYKEGIDNEAAIKMAVDD   80
NTDB id 94 BSU 11300 NP 389012.2             MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIED   80
consensus                                    !* !!!*!! *!! **!!!! *  !!!!!!!*! !!!! ! !!!!!!!!!!! !!! ! !!!!!!!!*  !  !  !**!

logo

NTDB id 1144963 BGLY RS06425 WP 046131230.1  FQKKGVNLIFGHGSEYEKVFNRISGDYPGIHFVLVNSKAEHKNVTSIDFDGRAMGFFGGMTAAHQSKSGKLGILATHSWQ  160
NTDB id 94 BSU 11300 NP 389012.2             FHKRGVNLLYGHGSEYAEVFNLVGEDYPDMEFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFTWQ  160
consensus                                    !*!*!!!!**!!!!!!  !!! *  !!! **!!* ! !! * !!!!**! ! !!!!!!!!!!!! !!*  *!**!***!!

logo

NTDB id 1144963 BGLY RS06425 WP 046131230.1  PEVKGFIEGAKYVNPDISVMTEYVGQWDDSEKAVQKYEKMKKAGADVVYPAGDGYNVPVIEQIKKDGLYAIGYVSDQSEL  240
NTDB id 94 BSU 11300 NP 389012.2             PEVDGFIKGAKYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDL  240
consensus                                    !!! !!! !!!! !!!! ! ! !  *!!!   !!  ! !!!  !!!!!!!!!!!!!!!!! !!!!!!!!!!!!!*!!!*!

logo

NTDB id 1144963 BGLY RS06425 WP 046131230.1  GENVVLTSTIQHVDEAYELIAGEFNKGTLKGGRRYVFDIQDGVIEMGEFSPVVDQAFQKKIKGLIEQYKKTGKLPDENVR  319
NTDB id 94 BSU 11300 NP 389012.2             GENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE..  317
consensus                                    !!! !!!!!*!*!! !!!*!!  !!!!!! !! * *!** !!*!!! !!!*!!! !! *!  !!  ! !!! !!   **
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X non conserved

X similar

X $\geq$ 50% conserved