```
logo

NTDB id 1044764 ACCB35 RS07660 WP 374056417.1  MSKRFYTIFLIVFLIGISGCSPG.KSAEIEKVGLLVPETVDDGVWGTKGYKGLLKIQSDYNADVFYKESIKNDTSIKQAV   79
NTDB id 94 BSU 11300 NP 389012.2               MITRLVMIFSVLLL..LSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAI   78
consensus                                      !  !   !! ** !***!!! * *  * !!!!!*! !*!* ! !!!!!!!!!!! !!! !! !!*!!! *! *  ! *!*

logo

NTDB id 1044764 ACCB35 RS07660 WP 374056417.1  KAYNDDKVTLVFGHGSEYSGIFNTLSSEYPNMHFVSFNGNATEENTTSLQFKGYAMGFFGGMTAAFQSKTNKIGVIAAFD   159
NTDB id 94 BSU 11300 NP 389012.2               EDFHKRGVNLLYGHGSEYAEVFNLVGEDYPDMEFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFT   158
consensus                                        **   ! !**!!!!!!  *!! *  *!! !*!!  !* !  *! !!**! ! !!!!!!!!!!!* !!!! *!!!! !

logo

NTDB id 1044764 ACCB35 RS07660 WP 374056417.1  WQPEVQGFIDGAKFQKSESELIVEYTNHWDDQTIAFEKLEGLMNQGVDVVYPAGDGYNVSVIEKLKEKGLYAIGYVSDQS   239
NTDB id 94 BSU 11300 NP 389012.2               WQPEVDGFIKGAKYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQS   238
consensus                                      !!!!! !!! !!!*  ** !*   !! !!!! ! !      * ! ! !!!!!!!!!!!!*!!  *!  !!!!!!!!*!!!

logo

NTDB id 1044764 ACCB35 RS07660 WP 374056417.1  DLGESTVLTSTVQHADKLYSVVAERFANDELDSGNLEFDFEDGVISMGKYSPVIPKEFQAEMKQHVKAYIETGKLPNGKE   319
NTDB id 94 BSU 11300 NP 389012.2               DLGENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE.   317
consensus                                      !!!! !!!!!!!!* !! ! **!! !    !* ! * *!   !!* !! *!!*** *!!  *  **! !  !! !!   *

logo

NTDB id 1044764 ACCB35 RS07660 WP 374056417.1  SKQ   322
NTDB id 94 BSU 11300 NP 389012.2               ...   317
consensus                                      ***
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X non conserved

X similar

X ≥ 50% conserved