```
logo
NTDB id 20714 AYT26 RS14590 WP 050768984.1  ......................MPTDFELMVGLLVEDTIDDQGWNRKAYEGLLNIHSNLDVDVVLEEGVNSEQKAHRRIKELVDGG   64
NTDB id 94 BSU 11300 NP 389012.2            MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNIQSKYNVDVYYKEGVKTEEDIINAIEDFHKRG   85
consensus                                   ***********************  *   !!*! *!!! !  !  !*! !!!!!*!   !!!   !!! *!   *  ! * *  !

logo
NTDB id 20714 AYT26 RS14590 WP 050768984.1  VNLIFGHGHAFAEYFSTIHNQYPDVHFVSFNGEVKGENITSLHFEGYAMGYFGGMVAASMSETHKVGVIAAFPWQPEVEGFVDGA  149
NTDB id 94 BSU 11300 NP 389012.2            VNLLYGHGSEYAEVFNLVGEDYPDMEFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFTWQPEVDGFIKGA  170
consensus                                   !!!**!!!* *!! !  **  !!!**!!  !*  !**!*!!*!! ! !!!*!!!! !!*!! !* !!!!! !*!!!!!*!!* !!

logo
NTDB id 20714 AYT26 RS14590 WP 050768984.1  KYMNES.EAFVRYVGEWTDADKALELFQELQKEQVDVFYPAGDGYHVPVVEAIKDQGDFAIGYVGDQADLGGSTILTSTVQHVDD  233
NTDB id 94 BSU 11300 NP 389012.2            KYENPDIEVNTKYTDHWDDDTTAVKLYQKMKNEGADVVYPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDLGENTVLTSTVQNVDK  255
consensus                                   !! !* *!   *!  *! !   !* !*! *  !  !! !!!!!!!*!!!*  !!  ! *!!!!! !! !!!  !*!!!!!!*!!

logo
NTDB id 20714 AYT26 RS14590 WP 050768984.1  LYVLVAKRFQEGKLESGNLYYDFQDGVVSLGEFSSVVPDFVREQITDAISTYIQTGQFPHEEER  297
NTDB id 94 BSU 11300 NP 389012.2            AYEIIAEQFNKGTLEGGDHYYDLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE..  317
consensus                                    ! **!  !* ! !! ! *!!! * !!! *! !!**!* *    !   ! !!  !!  !* !**
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X non conserved

X similar

X $\geq$50% conserved