```
logo
NTDB id 1018861 ABXR98 RS04280 WP 374695168.1  MFGFFKRKKNTEPVAEQPVPAPH..QQENSE...QLQNIADDSVQTEVIPQPEPAISDPVAEKPEQAD...TAPVQPQTE   72
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDV..........ESLAESVKGRAESAVETVSGAVEQVKE   70
consensus                                      !! !!*!!!  !**!  * * ****   !!*** * !! *! *********** * **!  **! ! ******! *  !

logo
NTDB id 1018861 ABXR98 RS04280 WP 374695168.1  TEQDIIVPTAVDEPAAIAVTDDTASPVQTPVASENTVVIAAEPDTVSEQTEKPAKLSWTARLKQGLSKSRNQMAKSLASV  152
NTDB id 1118 NGFG RS11455 WP 003696286.1       TV..AEMPSEAGEAA......ERVESAKEAVA..ETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGV  140
consensus                                      ! **  *!*   !*!*******   *   *!!** !!  !  *     !  **!! ! !!!!!!! !!!  !!!!!! !

logo
NTDB id 1018861 ABXR98 RS04280 WP 374695168.1  FGGGKIDEDLYEELETVLLTSDMGIEATEKLLAQVRQRVTLKGLKNGDELRSALKDALYELLLPLDQPLILPDDKKPFVI  232
NTDB id 1118 NGFG RS11455 WP 003696286.1       FGGGQIGEDLYEELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVI  220
consensus                                      !!!! ! !!!!!!!!!!!!*! !!!*!!!! !*  !! !!*!!!!! ! !!! !!!*!!!*!* !!* !!*!!* ! !!!!

logo
NTDB id 1018861 ABXR98 RS04280 WP 374695168.1  MMAGVNGAGKTTSIGKLAKYFQSQGKSVLLAAGDTFRAAAREQLQQWGERNNVTVIAQASGDSAAVCYDALEAAKARGID  312
NTDB id 1118 NGFG RS11455 WP 003696286.1       MLAGINGAGKTTSIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGID  300
consensus                                      !*!!*!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!! !! !!!!!!! ! *!!!!!!!*!!* !!!!!!!!

logo
NTDB id 1018861 ABXR98 RS04280 WP 374695168.1  VVLADTAGRLPTQLHLMEEIKKVKRVLQKSLPDAPHEVMLVLDANVGQNAINQVVAFDDALGLTGLIVSKLDGTAKGGIL  392
NTDB id 1118 NGFG RS11455 WP 003696286.1       IVLADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGIL  380
consensus                                      *!!!!!!!!!!!!!!!!!!!!!!!!!!!! *!  !!!!***!!!!!*!!!!*!!! !!!!!!!!!!!!!*!!!!!!!!!!!

logo
NTDB id 1018861 ABXR98 RS04280 WP 374695168.1  AALAAKRPVPVRFIGVGEGIDDLRPFDARAFVDALIEDE   431
NTDB id 1118 NGFG RS11455 WP 003696286.1       AALASDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD..   417
consensus                                      !!!!  !!!!!!!*!!!!!!!!!!!!!!!!!!!!!!****
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X non conserved

X similar

X $\geq$ 50% conserved