```
logo
NTDB id 1015662 ABV523 RS01775 WP 369608688.1  MFGFFKRKKNTEPEQ.EQASVNPAQPESVEQPDTAAETHTQPEPAASHQSVEEHTSNAAVSESTATETTEDII..VPAAV   77
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAA.KVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEA   79
consensus                                      !! !!*!!!  !** *! ! !**  ** !! *     *  ***   * !!  *   !  * ! ! !   * ****!

logo
NTDB id 1015662 ABV523 RS01775 WP 369608688.1  DEPAAIATSTESEVVQAPASENTIVIPVETTETPSETDKPARLGWAARLKQGLSKSRNQMAKSLASVFGGGKIDEDLYEE  157
NTDB id 1118 NGFG RS11455 WP 003696286.1       GEAAERVESAK......EAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEE  153
consensus                                       !*!     !  *******!   !*  *!     *  !  ***!!!!!!!!!!! !!!  !!!!!! !!!!! ! !!!!!!

logo
NTDB id 1015662 ABV523 RS01775 WP 369608688.1  LETVLLTSDMGIEATEKLLAQVRQRVTLKGLKNGNELRSALKEALYELLQPLQQPLSIPIDKKPFVIMMAGVNGAGKTTS  237
NTDB id 1118 NGFG RS11455 WP 003696286.1       LETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTS  233
consensus                                      !!!!!*! !!!*!!!! !*  !! !!*!!!!! !!!!! !!!!!!!*!* !!  !! *!  ! !!!!!*!!*!!!!!!!!

logo
NTDB id 1015662 ABV523 RS01775 WP 369608688.1  IGKLAKYFQSQGKSVLLAAGDTFRAAAREQLQQWGERNNVTVIAQESGDSAAVCFDALEAAKARGIDIVLADTAGRLPTQ  317
NTDB id 1118 NGFG RS11455 WP 003696286.1       IGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQ  313
consensus                                      !!!!!!!!! !!!!!!!!!!!!!!!!!!!!!! !! !!!!!!! ! *!!!!!!!!!!* !!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 1015662 ABV523 RS01775 WP 369608688.1  LHLMEEIKKVKRVLQKSLPEAPHEVMLVLDANVGQNAINQVVAFDDALGLTGLIVSKLDGTAKGGILAALAAKRPIPVRF  397
NTDB id 1118 NGFG RS11455 WP 003696286.1       LHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRY  393
consensus                                      !!!!!!!!!!!!!!!! *! !!!!***!!!!!*!!!!*!!! !!!!!!!!!!!!!*!!!!!!!!!!!!!!!  !!*!!!*

logo
NTDB id 1015662 ABV523 RS01775 WP 369608688.1  IGVGEGIDDLRPFDAKAFVDALIEDEE   424
NTDB id 1118 NGFG RS11455 WP 003696286.1       IGVGEGIDDLRPFDARAFVDALLD...   417
consensus                                      !!!!!!!!!!!!!!!*!!!!!!*****
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X non conserved

X similar

X $\geq$ 50% conserved