```
logo
NTDB id 1099918 ACNFWO RS03900 WP 132648645.1  MFSFFKKKVTPAPAAPVEPTP....AEA.SDQAPVLSDD..ELE........RA.................ALAAAPE...   46
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      !!!!!**!  * **!* !* *****!  *!* !***   ****!********!!*************** *!  ! ***

logo
NTDB id 1099918 ACNFWO RS03900 WP 132648645.1  .........PEALPEPAPLPLDAAPVAEVPAAAEPRSRWLDRLKLGLRKT....GSSLTQVFTGTKIDDALYEDLESALL  113
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAV.GQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLI  159
consensus                                      **********!!**!* * **** *   !*   ! *  !  !!! !! !****** !!  !! !  ! * !!!*!!* !*

logo
NTDB id 1099918 ACNFWO RS03900 WP 132648645.1  VADAGVGATEYLLQDLKRRVKEAKATEPAVVKALLADAVAELLAPLQKPLAIGQ.AMPTVVMVAGVNGAGKTTTIGKLTR  192
NTDB id 1118 NGFG RS11455 WP 003696286.1       TGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAK  239
consensus                                       *! !* !!!!!* !** !!      **  *** ! *!* *!* !! !!! ** *  ! !*!*!!*!!!!!!!*!!!! *

logo
NTDB id 1099918 ACNFWO RS03900 WP 132648645.1  HLAESDRRVLLAAADTFRAAAREQLAVWADRNRVEIVSQQGGDPAAVSFDAVTAGRARGCDVVIADTAGRLPTQLHLMDE  272
NTDB id 1118 NGFG RS11455 WP 003696286.1       YFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEE  319
consensus                                      *     * !!!!!*!!!!!!!!!!!  !* !! ! **!!  !!*!!!*!!!! !**!!!*!*!*!!!!!!!!!!!!!!*!

logo
NTDB id 1099918 ACNFWO RS03900 WP 132648645.1  LKKIKRTIAKAQDGAPHEVLLVIDGNTGQNALAQVKAFDAALGLTGLIVTKLDGTAKGGVLAAIARWCAEQGRVVPVYFI  352
NTDB id 1118 NGFG RS11455 WP 003696286.1       IKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALA.....SDRPVPVRYI  394
consensus                                      *!!*!! * !! *!!!!!***!*!*! !!!!* !!!!!! !!!!!!!!!!!!!!!!!!!*!!!*!*****  !*!!! *!

logo
NTDB id 1099918 ACNFWO RS03900 WP 132648645.1  GVGERLEDLQTFDAREFANALLG   375
NTDB id 1118 NGFG RS11455 WP 003696286.1       GVGEGIDDLRPFDARAFVDALLD   417
consensus                                      !!!! **!! *!!!! !  !!!
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X non conserved

X similar

X $\geq$ 50% conserved