```
logo
NTDB id 1057947 ACFJI0 RS17270 WP 382321390.1  MFSFFRKKAPTPPTPALPEAPAP.......................................................SPA   26
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQE.TPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEA   79
consensus                                      !!!!!!*! * **!!!!*!!* ***************************************************!*!

logo
NTDB id 1057947 ACFJI0 RS17270 WP 382321390.1  AP................APAAAAPTPAAPAAAPRQGWLDKLKAGLRKTG....SGIATVFTGTQIDDALYEELESALL   85
NTDB id 1118 NGFG RS11455 WP 003696286.1       GEAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLI  159
consensus                                      ******************* * *  * * !*    * !!  *!! !! !* ****  *! !! ! !! * !!!!!!* !*

logo
NTDB id 1057947 ACFJI0 RS17270 WP 382321390.1  MADTGVKATTHLLSDLKRRVKESKTTDPAAVKGLLADAIADLLAPLQKPLVIGEH.KPTVIMVAGVNGAGKTTSIGKLTR  164
NTDB id 1118 NGFG RS11455 WP 003696286.1       TGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAK  239
consensus                                       *! !* !! *!* !** !!      !*  **! ! *!* !!* !! !!!!**!** ! !!!*!!*!!!!!!!!!!!! *

logo
NTDB id 1057947 ACFJI0 RS17270 WP 382321390.1  HLADSGASVLLAAADTFRAAAREQLAVWADRNTVEIITQQGGDPAAVSFDAVTAGKARGRDVVLVDTAGRLPTQLHLMEE  244
NTDB id 1118 NGFG RS11455 WP 003696286.1       YFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEE  319
consensus                                      *    ! !!!!!!*!!!!!!!!!!!  !* !! ! *!*!  !!*!!!*!!!! !*!!!! !*!! !!!!!!!!!!!!!!!

logo
NTDB id 1057947 ACFJI0 RS17270 WP 382321390.1  LRKIKRVVQKADAGAPHEVLLVIDGNTGQNALMQVKAFDEALQLTGLVITKLDGTAKGGVLCAIAREKPVPVYFIGVGEK  324
NTDB id 1118 NGFG RS11455 WP 003696286.1       IKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEG  399
consensus                                      **!*!!!*!!! *!!!!!***!*!*! !!!!* !!!!!!*!! !!!!**!!!!!!!!!!*!*!*! **!!!! *!!!!!

logo
NTDB id 1057947 ACFJI0 RS17270 WP 382321390.1  LEDLETFDAREFAQALLA   342
NTDB id 1118 NGFG RS11455 WP 003696286.1       IDDLRPFDARAFVDALLD   417
consensus                                      **!! *!!!! !  !!!
```

X non conserved

X similar

X $\geq$ 50% conserved