```
logo
NTDB id 1167939 ACLIIK RS24420 WP 410367420.1  ..................................................................MAPTPAPVVPAPAPVS   16
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      ************************************************************* * * *! * ! *

logo

NTDB id 1167939 ACLIIK RS24420 WP 410367420.1  APAP.V.ST.PAPVAAPAPVIAAVPEPAPAEAPKKASWLSRLKQGLSRT....GQSIGGIFVGVKVDENLREELESALIM   89
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                       *!**!*! **!*   *** ** !*!* *   **!  !  !!!!!! ******* !**!*! !  * ! !*!!!!* !!

logo

NTDB id 1167939 ACLIIK RS24420 WP 410367420.1  ADAGLEATEKLLTALRARVKKERIEDPAKVKTALRQLLADHLRPLERAFDLKR.TQPLVVMIAGVNGAGKTTSIGKLAHT  168
NTDB id 1118 NGFG RS11455 WP 003696286.1       GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                      *! !*!!!! !*  *!*!!    * !*  ** !!*  ! !***!!!**  !* *  ! !*!*!!*!!!!!!!!!!!!!!*

logo

FQRQGASVLLAAGDTFRAAAREQLVEWGSRNNVSVISQDGGDPAAVAFDSVNAGRARGMGVVMVDTAGRLPTQLHLMEEL  248
FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                      !! !! !!!!!!!!!!!!!!!!!!  !! !!!!*!!!!  !!*!!!*!! !*!**!!!* *!* !!!!!!!!!!!!!!!*

logo

NTDB id 1167939 ACLIIK RS24420 WP 410367420.1  KKIRRVIGKADASAPHEVLLVVDGNTGQNALAQIRAFDAAINLTGLVVTKLDGTAKGGTLAAVAAGSQGVRPVPVYWIGV  328
NTDB id 1118 NGFG RS11455 WP 003696286.1       KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALA....SDRPVPVRYIGV  396
consensus                                      !!**!!* !! * !!!!***!*!*! !!!!* !**!!! !* !!!!*!!!!!!!!!!!! !!!*!****  !!!!! *!!!

logo

NTDB id 1167939 ACLIIK RS24420 WP 410367420.1  GESLEDLQPFVADEFAAALLAD   350
NTDB id 1118 NGFG RS11455 WP 003696286.1       GEGIDDLRPFDARAFVDALLD.   417
consensus                                      !! **!! !! !  !  !!! *
```

X non conserved

X similar

X ≥ 50% conserved