```
logo
NTDB id 988747 AAGF34 RS14560 WP 342616447.1  MFSFFKKKSPPAPVAAPAAVPDGSAAPV.....................................QPAAAG   34
NTDB id 1118 NGFG RS11455 WP 003696286.1      MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                     !!!!!**! ** **! * ! *  *!!*!******************************************** !  !!

logo
NTDB id 988747 AAGF34 RS14560 WP 342616447.1  TQIGSALAAPVP........VPDAIAAPVAPKPERQKWLDRLKAGLGK....TASSLSGVFGGNKVDEALFEELESALLM  102
NTDB id 1118 NGFG RS11455 WP 003696286.1      EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                              !* *********!*    !*  * ***  !  !!! !!*!**** ! !! !!!!!  * ! !*!!!!* !*

logo
NTDB id 988747 AAGF34 RS14560 WP 342616447.1  ADAGVSATQFLTDNLRKKAKSSNLTNPVALKNQLIADLTELLQPLQKPLVIGQ.QTPTVIMVAGVNGAGKTTSIGKLTRH  181
NTDB id 1118 NGFG RS11455 WP 003696286.1      GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                     *! !* !! *!   *! *     !  *  !*  !   ! *!* !! !!!!** *  ! !!!*!!*!!!!!!!!!!!! **

logo
NTDB id 988747 AAGF34 RS14560 WP 342616447.1  LSEEGASVLLAAADTFRAAAREQLSIWADRNTVEIISQAGADPAAVSFDAVSAGKARGKDVVLIDTAGRLPTQLHLMEEL  261
NTDB id 1118 NGFG RS11455 WP 003696286.1      FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                        ! !!!!!!*!!!!!!!!!!!  !* !! ! *!!!  *!*!!!*!!!! !*!!!! !*!! !!!!!!!!!!!!!!!*

logo
NTDB id 988747 AAGF34 RS14560 WP 342616447.1  KKIKRVIQKADVSAPHEVLLVIDGNTGQNAVAQVKAFDDALGLTGLIVTKLDGTAKGGVLAAIARDRPIPVYFIGVGEQL  341
NTDB id 1118 NGFG RS11455 WP 003696286.1      KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                     !!*!!!*!!! * !!!!***!*!*! !!!!! !!!!!!!!!!!!!!!
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