```
logo
NTDB id 1099879 ACNANZ RS17800 WP 416400512.1  MFSFFKKKSPSPAEPSSAP.APDAASALP...........ADA.....PTGGSA..........LGWLRKSFGAQD....   49
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      !!!!!**! * * **   ** ** !  ************ ! ******  ! !***********  ** * *  *****

logo
NTDB id 1099879 ACNANZ RS17800 WP 416400512.1  .......AAAPQAAPAADEAPPAVPEPAPKPEGARQGWMDRLKSGLRKTG....GSIATVFTGTQINDALYEELEDALLM  118
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                      ******* ! *  !*   !!** !*!* * *! ** !!  !!! !! !* **** !*! !! ! !! * !!!!!!  !*

logo
NTDB id 1099879 ACNANZ RS17800 WP 416400512.1  ADTGVKATQHLLADLRRRVKEAKATEPAAVKALLADALADLLRPLEKPLVIGE.HQPTVIMVAGVNGAGKTTSIGKLTRH  197
NTDB id 1118 NGFG RS11455 WP 003696286.1       GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                      *! !* !! *!* !*! !!      **  *** ! *!! !!**!!!!!!!**!** ! !!!*!!*!!!!!!!!!!!! **

logo
NTDB id 1099879 ACNANZ RS17800 WP 416400512.1  LADHGESVLLAAADTFRAAAREQLGVWATRNTVEIISQEGGDPAAVSFDAVTAGRARGKDVVLVDTAGRLPTQLHLMEEL  277
NTDB id 1118 NGFG RS11455 WP 003696286.1       FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                         *! !!!!!!*!!!!!!!!!!!  !* !! ! *!!!  !!*!!!*!!!! !**!!! !*!! !!!!!!!!!!!!!!!*

logo
NTDB id 1099879 ACNANZ RS17800 WP 416400512.1  RKIKRVVTKADATAPHEVLLVIDGNTGQNALAQVRAFDDALQLTGLIVTKLDGTAKGGVLAAIAQERPVPVYFIGVGEKV  357
NTDB id 1118 NGFG RS11455 WP 003696286.1       KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                      *!*!!!* !! * !!!!***!*!*! !!!!* !!*!!!!!! !!!!!!!!!!!!!!!!*!!!*! *!!!!! *!!!!! *

logo
NTDB id 1099879 ACNANZ RS17800 WP 416400512.1  EDLETFNAREFAQALLA   374
NTDB id 1118 NGFG RS11455 WP 003696286.1       DDLRPFDARAFVDALLD   417
consensus                                      *!! *! !! !  !!!
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X non conserved

X similar

X $\geq$ 50% conserved