```
logo
NTDB id 108348 AACH39 RS09635 WP 338619764.1  MFSFFKKKSD.......................................................................  10
NTDB id 1118 NGFG RS11455 WP 003696286.1      MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG  80
consensus                                     !!!!!**!  ***********************************************************************

logo
NTDB id 108348 AACH39 RS09635 WP 338619764.1  ..............TDVSEVEAAPREPSDTPAIPQRSWTERLKSGLAKTRDKLGKNLASLFGGGQIDEALYEELETVLLT  76
NTDB id 1118 NGFG RS11455 WP 003696286.1      EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                     **************  !  !   * * !*  !*      !  !!! !!!!*!!!**! !! *!!!!!! ! !!!!!!!!!*!

logo
NTDB id 108348 AACH39 RS09635 WP 338619764.1  ADMGVDATEHLLREVRARVSLRGLKDSSELKQTLKDVLYELIAPLEVPLDTS.QGKPFVIMMAGVNGAGKTTSIGKLAKY  155
NTDB id 1118 NGFG RS11455 WP 003696286.1      GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                     *!!!**!!!*!***!!*!!!!*!!!!  !!*  !!* !!*!! !!! !!  **   !!!!!*!!*!!!!!!!!!!!!!!!!

logo
NTDB id 108348 AACH39 RS09635 WP 338619764.1  FQNQNKSVLLAAGDTFRAAAREQLMEWGERNNVTVITQQGGDSAAVCFDAINSAKARGIDIVLADTAGRLPTQLHLMEEI  235
NTDB id 1118 NGFG RS11455 WP 003696286.1      FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                     !! ! !!!!!!!!!!!!!!!!!!!  !! !!!!!!!*!  !!!!!!!!!!** !!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo
NTDB id 108348 AACH39 RS09635 WP 338619764.1  RKVKRVIQKALPEAPQEVVLVLDANIGQNALNQVKVFDEALGLTGLILTKLDGTAKGGIAAIAKQHPVPLRFIGVGEGL  315
NTDB id 1118 NGFG RS11455 WP 003696286.1      KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                     *!!!!!*!!!*! !!*!***!!!!!!!!!!*!!!! !!*!!!!!!!!*!!!!!!!!!!!!*!!*!  *!!!*!*!!!!!!*

logo
NTDB id 108348 AACH39 RS09635 WP 338619764.1  DDLRPFSARDYIDALFD  332
NTDB id 1118 NGFG RS11455 WP 003696286.1      DDLRPFDARAFVDALLD  417
consensus                                     !!!!!! !! **!!! !
```

X non conserved

X similar

X ≥ 50% conserved