```
logo
NTDB id 1090437 ACLRDN RS21585 WP 413457938.1  MFSFFKRKPKPEAQPALPPQAEPAPQPAPAPSVAP..AAVEAP.......VP.........APAPIAPVEPVSAPEPE..   60
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKV.ESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEA   79
consensus                                      !!!!!*!!*!*! **  **!  * * * * * !!**** *  ********!********** **** *! *  !* ! **

logo
NTDB id 1090437 ACLRDN RS21585 WP 413457938.1  .....PV..ATEAP...AAEEV.EIVAAPEPAPEAKRSWLSRLKAGLSKTSS....NLTTLFVGARIDEDLYEELESALL  125
NTDB id 1118 NGFG RS11455 WP 003696286.1       GEAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLI  159
consensus                                      ******!**! !!**** *! !* *  !* * *!*!  !  !!! !! !*  **** !  *! !* ! !!!!!!!!* !*

logo
NTDB id 1090437 ACLRDN RS21585 WP 413457938.1  VSDAGVEATQWLLAELKKKVKNERLTEAAQVRTALRTLLIDLLQPLQRPLVLGR.DKPMVMMIAGVNGAGKTTTIGKLAL  204
NTDB id 1118 NGFG RS11455 WP 003696286.1       TGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAK  239
consensus                                        ! !*!!! *!* *** *!    ! **  *! !!*  ! !!* !! *!!!!* *  ! !*!*!!*!!!!!!!*!!!!!

logo
NTDB id 1090437 ACLRDN RS21585 WP 413457938.1  HLQAHGQSVLLAAGDTFRAAAREQLAVWGERNNVQVIAQESGDPAAVAYDSVHSAQARGTHVVMVDTAGRLPTQLHLMDE  284
NTDB id 1118 NGFG RS11455 WP 003696286.1       YFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEE  319
consensus                                      * !!*! !!!!!!!!!!!!!!!!!!  !! !!!! !! ! *!!*!!!**! !* ! !!! **!* !!!!!!!!!!!!!*!

logo
NTDB id 1090437 ACLRDN RS21585 WP 413457938.1  LKKIKRVIAKAMNSAPHEVLLVIDGNTGQNALAQVKAFDDALGLTGLVVTKLDGTAKGGILAAIAKTRPVPLYFIGVGEK  364
NTDB id 1118 NGFG RS11455 WP 003696286.1       IKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEG  399
consensus                                      *!!*!!!* !!** !!!!***!*!*! !!!!* !!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!*!  !!!!* *!!!!!

logo
NTDB id 1090437 ACLRDN RS21585 WP 413457938.1  IEDLQPFNATEFVDALLS   382
NTDB id 1118 NGFG RS11455 WP 003696286.1       IDDLRPFDARAFVDALLD   417
consensus                                      !*!! !! !  !!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved