```
logo
NTDB id 1021297 AB1W24 RS02550 WP 039052211.1  MFSFFKKKTPPPAAPATPAPPPEPVAQPAEPVSPPSIAPAAPPVPAVAPEPPVAVPPAAAPPRPPVAAEPV..AVQTAAP   78
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVA.QIVGNIKEDVESLAE.........SVKGRAESAVETVSGAVEQVKE   70
consensus                                      !!!!!**! *** **  *!***!* ! *  *! ** ***  **!* *!**********  **!** ! !*!**!!    *

logo
NTDB id 1021297 AB1W24 RS02550 WP 039052211.1  ASPAMPAAPAPLVAEPV..AAEAVAALAPPSPPAAQAPAPAAPAPEVPAPEVPAPKKASWLSRLKQGLARTG....QSIG  152
NTDB id 1118 NGFG RS11455 WP 003696286.1       TVAEMPSEAGEA.AERVESAKEAVAE...........IVGEAVGQVQEAVATIEEHKLGWAARLKQGLAKSRDKMAKSLA  138
consensus                                        * !!  *** *!!*!**! !!!! ************ * !***  *!*  * **!  !  !!!!!!!** **** !**

logo
NTDB id 1021297 AB1W24 RS02550 WP 039052211.1  GIFVGVKVDENLFEELESALIMADAGIEATDKLLTALRARVKKERIEDPAQVKAALRQLLADHLKPLERQFDLK.RAQPL  231
NTDB id 1118 NGFG RS11455 WP 003696286.1       GVFGGGQIGEDLYEELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPF  218
consensus                                      !*! !  * ! !*!!!!* !! *! !*!!!* !*  *!*!!    * !*  ***!!*  ! !**!!!!**  !**   !

logo
NTDB id 1021297 AB1W24 RS02550 WP 039052211.1  VVMLAGVNGAGKTTSIGKLAHTFQRQGASVLLAAGDTFRAAAREQLVEWGSRNNVTVIAQDGGDPAAVAFDAVNAGRARA  311
NTDB id 1118 NGFG RS11455 WP 003696286.1       VIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARG  298
consensus                                      !*!*!!*!!!!!!!!!!!!!!* !! !! !!!!!!!!!!!!!!!!!!  !! !!!!!!! !  !!*!!!*!!!!*!**!!*

logo
NTDB id 1021297 AB1W24 RS02550 WP 039052211.1  AGVVMIDTAGRLPTQLHLMEELKKIRRVIGKADAAAPHEVLLVIDGNTGQNALAQIRAFDAAIQLTGLVVTKLDGTAKGG  391
NTDB id 1118 NGFG RS11455 WP 003696286.1       IDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGG  378
consensus                                        *!* !!!!!!!!!!!!!!!!*!!**!!* !! **!!!!***!*!*! !!!!* !**!!! !* !!!!*!!!!!!!!!!!

logo
NTDB id 1021297 AB1W24 RS02550 WP 039052211.1  TLAAVAAGSQGVRPVPVYWIGVGEGLEDLQPFVAEEFASALLAD   435
NTDB id 1118 NGFG RS11455 WP 003696286.1       ILAALA....SDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD.   417
consensus                                       !!!*!****  !!!!! *!!!!!!**!! !! !  !  !!! *
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X non conserved

X similar

X $\geq$ 50% conserved