```
logo
NTDB id 1094361 ACMDW7 RS03710 WP 377305454.1  MFSFFRKKSPPP..AVQP.QVVESQAIAP.........................................QTAEAPA...   33
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      !!!!!!*! *****!* **!! !* !  ****************************************  !! ! ***

logo

NTDB id 1094361 ACMDW7 RS03710 WP 377305454.1  ...TEPVSAPAAVAAPAPAANAAPAEAPATEAAEKKSWLARLKQGLSRT....GSSIAGVFGYVKVDEALREQLEDALLM  106
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                      ***  * !!* !!! * * ! * * !!*!!   *!  ! !!!!!!! ******* !*!!!!!   * ! !*! !!  !*

logo

NTDB id 1094361 ACMDW7 RS03710 WP 377305454.1  ADCGVTATQKILAGLRARVKSERIEDAAAVRTALAKVLADHLRALERTPQVPAAADEKPWVVMIAGVNGAGKTTSIGKIT  186
NTDB id 1118 NGFG RS11455 WP 003696286.1       GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLP..ETKEPFVIMLAGINGAGKTTSIGKLA  238
consensus                                      *!*!* !!  **  *!*!!   * !*  *! !!   ! !****!!*** *!**   !*!*!*!!*!!!!!!!!!!!*

logo

NTDB id 1094361 ACMDW7 RS03710 WP 377305454.1  HHLSEGGHSVLLAAADTFRAAAREQLAVWGNRNQVSVVSQEGADPGAVVFDAVQSARAKGIDVVIADTAGRLPTQLHLME  266
NTDB id 1118 NGFG RS11455 WP 003696286.1       KYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLME  318
consensus                                      **    !*!!!!!!*!!!!!!!!!!!  !! !!*!*!*!!  *!**!!*!!!!! !*!*!!!*!*!!!!!!!!!!!!!!!

logo

NTDB id 1094361 ACMDW7 RS03710 WP 377305454.1  ELKKIARVCGKALDGAPHERWLVIDGNTGQNAVAQVKAFNEALQLTGLIVTKLDGTAKGGVLAAIASTCPVPVLFIGVGE  346
NTDB id 1118 NGFG RS11455 WP 003696286.1       EIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGE  398
consensus                                      !*!!* !!* !!**!!!!!  *!*!*! !!!!! !!!!! *!! !!!!!!!!!!!!!!!*!!!*!! *!!!! *!!!!!

logo

NTDB id 1094361 ACMDW7 RS03710 WP 377305454.1  QLQDLQPFSAEEFAGALTGA   366
NTDB id 1118 NGFG RS11455 WP 003696286.1       GIDDLRPFDARAFVDALLD.   417
consensus                                       * !! !! !  !  !!  *
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X non conserved

X similar

X ≥ 50% conserved