```
logo
NTDB id 1057505 ACEQUB RS01560 WP 197331682.1  MFSFWKKRKAEPQPAAEPVPPAAAPEAVQAPA.....P....V...ASPAPA..............PAPVAVPDTPATPA   54
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      !!!!****!  * !! !* **   *  !  * **********!***! * ***************** *  !** !  **

logo
NTDB id 1057505 ACEQUB RS01560 WP 197331682.1  ............ALDMQADDIE.TVPTPPVVEQARKGWMSRLRSGLSKTS....KGLTTLFVGVKVDEALREELETALLM  117
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                      ************  *  ** * *  * **  ! ** !!  !!* !! !* ****! !  *! !  * ! !*!!!!! !*

logo
NTDB id 1057505 ACEQUB RS01560 WP 197331682.1  ADAGVDATEYLLGELRRRVKAQRIETAEGVKTALRDLLIELLHPLEKTMVLGR.EQPTVIMIAGVNGAGKTTSIGKLCKH  196
NTDB id 1118 NGFG RS11455 WP 003696286.1       GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                      *! !**!!!!!* **! !!    *  *  ** !!** ! *!**!!!!**!!* *   ! !!!*!!*!!!!!!!!!!!!*!*

logo
NTDB id 1057505 ACEQUB RS01560 WP 197331682.1  FQTYGQSVLLAAGDTFRAAAREQLVIWGQRNNVTVVAQESGDPAAVIFDAVNAARARGIDIVMADTAGRLPTQLHLMEEL  276
NTDB id 1118 NGFG RS11455 WP 003696286.1       FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                      !!   ! !!!!!!!!!!!!!!!!!!  !! !!!!!!* ! *!!*!!!*!!!!*!!*!!!!!!!*!!!!!!!!!!!!!!!!*

logo
NTDB id 1057505 ACEQUB RS01560 WP 197331682.1  KKVRRVIGKAMATAPHETLLVIDANTGQNALAQVKAFDDTLGLTGLIITKLDGTAKGGILAAIARQRPVPVYFIGVGEQV  356
NTDB id 1118 NGFG RS11455 WP 003696286.1       KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                      !!!*!!* !!** !!!! **!*!!! !!!!* !!!!!!! !!!!!!!*!!!!!!!!!!!!!!!*!  !!!!! *!!!!! *

logo
NTDB id 1057505 ACEQUB RS01560 WP 197331682.1  EDLQPFSAREFADALLG  373
NTDB id 1118 NGFG RS11455 WP 003696286.1       DDLRPFDARAFVDALLD  417
consensus                                      *!! !! !! ! !!!!
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X non conserved

X similar

X ≥ 50% conserved