```
logo
NTDB id 1069769 RSP597 RS08285 WP 247546103.1  MFSFWKKRKAEPQPAAEPAPPAAAPEAAQAQVPAPVASPAPAPAPVASPVA......V.PVASP.....VAVPDTPAAP.   67
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      !!!!****!  * !! !*!**   *      !* *!* * * * **! *! ******!**!  ******  !** !  **

logo
NTDB id 1069769 RSP597 RS08285 WP 247546103.1  .A.........ALDMQADDIE.TVPTPPVMEQARKGWMSRLRSGLSKTS....KSLTTLFVGVKVDEALREELETALLM  131
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                      *!**********  *  ** * *  * **  ! ** !!  !!* !! !* ****!!!  *! !  * ! !*!!!!! !*

logo
NTDB id 1069769 RSP597 RS08285 WP 247546103.1  ADAGVDATEYLLNELRRRVKAQRIETAESVKTALRDLLIELLHPLEKTMVLG.RDQPMVIMIAGVNGAGKTTSIGKLCKH  210
NTDB id 1118 NGFG RS11455 WP 003696286.1       GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                      *! !**!!!!!* **! !!    *  *  ** !!** ! *!**!!!!**!!**   ! !!!*!!*!!!!!!!!!!!!*!*

logo
NTDB id 1069769 RSP597 RS08285 WP 247546103.1  FQTYGQSVLLAAGDTFRAAAREQLVIWGQRNNVTVVAQESGDPAAVIFDAVNAARARGIDIVMADTAGRLPTQLHLMEEL  290
NTDB id 1118 NGFG RS11455 WP 003696286.1       FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                      !!   ! !!!!!!!!!!!!!!!!!!  !! !!!!!!* ! *!!*!!!*!!!!*!!*!!!!!!!*!!!!!!!!!!!!!!!!*

logo
NTDB id 1069769 RSP597 RS08285 WP 247546103.1  KKVRRVIGKAMVTAPHETLLVIDANTGQNALAQVKAFDDTLGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEQV  370
NTDB id 1118 NGFG RS11455 WP 003696286.1       KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                      !!!*!!* !!** !!!! **!*!!! !!!!* !!!!!!! !!!!!!!!!!!!!!!!!!!!!!*!  !!!!! *!!!!! *

logo
NTDB id 1069769 RSP597 RS08285 WP 247546103.1  EDLQPFSAREFADALLG   387
NTDB id 1118 NGFG RS11455 WP 003696286.1       DDLRPFDARAFVDALLD   417
consensus                                      *!! !! !! ! !!!!
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X non conserved

X similar

X ≥ 50% conserved