```
logo
NTDB id 406784 GO998 RS15425 WP 211904044.1  MFSFWKKRKAEPQPAAEPVPPAAAPEAVQAPA.....P....V...ASPAPA..............PAPVAVPDTPATPA   54
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                    !!!!****!  * !! !* **   *  !  * **********!***! * ***************** *  !** !  **

logo
NTDB id 406784 GO998 RS15425 WP 211904044.1  ............ALDMQADDIE.AVPTPPVVEQARKGWMSRLRSGLSKTS....KGLTTLFVGVKVDEALREELETALLM  117
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                    ************  *  ** * *  * **  ! ** !!  !!* !! !* ****! !  *! !  * ! !*!!!!! !*

logo
NTDB id 406784 GO998 RS15425 WP 211904044.1  ADAGVDATEYLLGELRRRVKAQRIETAEGVKTALRDLLIELLHPLEKTMVLGR.EQPTVIMIAGVNGAGKTTSIGKLCKH  196
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                    *! !**!!!!!* **! !!    *  *  ** !!** ! *!**!!!!**!!* *  ! !!!*!!*!!!!!!!!!!!!*!*

logo
NTDB id 406784 GO998 RS15425 WP 211904044.1  FQTYGQSVLLAAGDTFRAAAREQLVIWGQRNNVTVVAQESGDPAAVIFDAVNAARARGIDIVMADTAGRLPTQLHLMEEL  276
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                    !!   ! !!!!!!!!!!!!!!!!!!  !! !!!!!!* ! *!!*!!!*!!!!*!!*!!!!!!!*!!!!!!!!!!!!!!!!*

logo
NTDB id 406784 GO998 RS15425 WP 211904044.1  KKVRRVIGKAMATAPHETLLVIDANTGQNALAQVKAFDDTLGLTGLIITKLDGTAKGGILAAIARQRPVPVYFIGVGEQV  356
NTDB id 1118 NGFG RS11455 WP 003696286.1     KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                    !!!*!!* !!** !!!! **!*!!! !!!!* !!!!!!! !!!!!!!*!!!!!!!!!!!!!!!*!  !!!!! *!!!!! *

logo
NTDB id 406784 GO998 RS15425 WP 211904044.1  EDLQPFSAREFADALLG   373
NTDB id 1118 NGFG RS11455 WP 003696286.1     DDLRPFDARAFVDALLD   417
consensus                                    *!! !! !! ! !!!!
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X non conserved

X similar

X $\geq$ 50% conserved