```
logo
NTDB id 106809 ACAM56 RS21325 WP 369648861.1  MFSFFKKKPPAESPAAPAPSRAVEPVPAPEPAPAPAAAP.ARSVFS.PSSWFGSKPAAEEAVPPPAAPDMAPPVAPPPAP   78
NTDB id 1118 NGFG RS11455 WP 003696286.1      MFSFFRRKKKQETPALEEAQVQETAAK.VESEVAQIVGNIKEDVESLAESVKGRAFSAVETVSG.AVEQVKETVAEMPSE   78
consensus                                     !!!!!**!** !*!! * *     * ***!* *!*  ***   ! !** !  !  * ! ! !***! * * **!!**! *

logo
NTDB id 106809 ACAM56 RS21325 WP 369648861.1  ..AAAPHLAPVPAAPVPTPAVPTVPPPPPAL.ATEVAMPAERKGWFDKLKTGLRKTG....TGIQAVFVNAQIDEALYEE  151
NTDB id 1118 NGFG RS11455 WP 003696286.1      AGEAAERVESAKE.AV....AETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEE  153
consensus                                     ** !!*** * * **!**** *!!***** **      * ** !!  *!! !! !* ****  * *!!  *!! ! !!!!

logo
NTDB id 106809 ACAM56 RS21325 WP 369648861.1  LESALLMADAGVKATEYLLEDLRGRVKRQMATDAAQVKRLLADAITDLLQPLEKPLVIGQ.FTPTVIMVAGVNGAGKTTS  230
NTDB id 1118 NGFG RS11455 WP 003696286.1      LETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTS  233
consensus                                     !!* !* *! !* !!!!!* !*!!!!      !*  **  ! *!* !!* !!!!!!** *  ! !!!*!!*!!!!!!!!

logo
NTDB id 106809 ACAM56 RS21325 WP 369648861.1  IGKLTKHLASEGASVLLAAADTFRAAAREQLLVWADRNTVEIVSQEGGDPSAVSFDAVTAGKARGKDVVLVDTAGRLPTQ  310
NTDB id 1118 NGFG RS11455 WP 003696286.1      IGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQ  313
consensus                                     !!!! !*    ! !!!!!!*!!!!!!!!!!!  !* !! ! **!!  !!* !!*!!!! !*!!!! !*!! !!!!!!!!!

logo
NTDB id 106809 ACAM56 RS21325 WP 369648861.1  LHLMDELKKIKRVVTKADASAPHEVLLVIDGNTGQNALAQVRAFDETLGLTGLVVTKLDGTAKGGVLCAIARERPIPVYF  390
NTDB id 1118 NGFG RS11455 WP 003696286.1      LHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRY  393
consensus                                     !!!!*!*!!*!!!* !! * !!!!***!*!*! !!!!* !!*!!!* !!!!!!*!!!!!!!!!!!*!*!*! *!!*!! *

logo
NTDB id 106809 ACAM56 RS21325 WP 369648861.1  IGVGEKLEDLETFNAREFAMALLG   414
NTDB id 1118 NGFG RS11455 WP 003696286.1      IGVGEGIDDLRPFDARAFVDALLD   417
consensus                                     !!!!! **!! *! !! !  !!!
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X non conserved

X similar

X ≥ 50% conserved