```
logo
NTDB id 1007785 ABL844 RS26695 WP 418129582.1  MFSFFKKKPPAEAPDAPAPTPPTPPTPAAPAPEPAPAR.....SVFS.PSSWFGSKPAAEETPAPAP.VTAPASVPVPSP   73
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAK..VESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSE   78
consensus                                      !!!!!**!** ! !  * * ** ** **** *!*!*  ***** ! !** !  !  * ! !!* *!**   *   **!!*

logo
NTDB id 1007785 ABL844 RS26695 WP 418129582.1  APAPVAAPVVPPSPAPAPSATPAPAPTPVAAAPVAEPVIATERKGWFDKLKTGLRKTG....TGIQAVFVNAKIDEELYE  149
NTDB id 1118 NGFG RS11455 WP 003696286.1       A..GEAAERVE....SAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYE  152
consensus                                      !*** !!* !******!* ! * * * *!*  * ! *    ** !!  *!! !! !* ****  * *!!  * ! !*!!!

logo
NTDB id 1007785 ABL844 RS26695 WP 418129582.1  ELESALLMADTGVKATEYLLDDLRGRVKRQMATDASQVKALLAETVADLLKPLEKPLVIGQ.FTPTVIMVAGVNGAGKTT  228
NTDB id 1118 NGFG RS11455 WP 003696286.1       ELETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTT  232
consensus                                      !!!* !* *! !* !!!!!* !*!!!!      !*  *** ! ! * !!*!!!!!!!!** *  ! !!!*!!*!!!!!!!

logo
NTDB id 1007785 ABL844 RS26695 WP 418129582.1  SIGKLTKHLADEGASVLLAAADTFRAAAREQLLVWADRNTVEIVSQEGGDPSAVSFDAVNAGRARGKDVVLVDTAGRLPT  308
NTDB id 1118 NGFG RS11455 WP 003696286.1       SIGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPT  312
consensus                                      !!!!! !*    ! !!!!!!*!!!!!!!!!!!  !* !! ! **!!  !!* !!*!!!!*!**!!! !*!! !!!!!!!!

logo
NTDB id 1007785 ABL844 RS26695 WP 418129582.1  QLHLMDELKKIKRVVTKADGTAPHEVLLVIDGNTGQNALAQVKAFDETLGLTGLVVTKLDGTAKGGVLCAIARERPIPVY  388
NTDB id 1118 NGFG RS11455 WP 003696286.1       QLHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVR  392
consensus                                      !!!!!*!*!!*!!!* !! * !!!!***!*!*! !!!!* !!!!!!* !!!!!!*!!!!!!!!!!!*!*!*! *!!*!!

logo
NTDB id 1007785 ABL844 RS26695 WP 418129582.1  FIGVGEKLEDLETFNAREFAQALLG  413
NTDB id 1118 NGFG RS11455 WP 003696286.1       YIGVGEGIDDLRPFDARAFVDALLD  417
consensus                                      *!!!!! **!! *! !! !  !!!
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X non conserved

X similar

X $\geq$ 50% conserved