```
logo
NTDB id 1007803 ABL838 RS26205 WP 418122633.1  MFSFFKKKPSAEAPDAPAPTPPAPAAPAPEPAPARSVF.......S.PSSWFGSKPAAEETPAPA.PVTTPASVPVPSPA   71
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVES.EVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEA   79
consensus                                      !!!!!**!*  ! !  * * ** *!!* * * *!  ! *******!** !  !  * ! !!* *!**   *   **!!*!

logo
NTDB id 1007803 ABL838 RS26205 WP 418122633.1  PAPAATPVVPSSPAPAPSATPAPAPTPVAAAPVAEPVIATERKGWFDKLKTGLRKTG....TGIQAVFVNAKIDEELYEE  147
NTDB id 1118 NGFG RS11455 WP 003696286.1       G..EAAERVESA....KEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEE  153
consensus                                      *** ! * !*! ***** ! * * * *!*  * ! *    ** !!  *!! !! !* ****  * *!!  * ! !*!!!!

logo
NTDB id 1007803 ABL838 RS26205 WP 418122633.1  LESALLMADTGVKATEYLLEDLRGRVKRQMATDASQVKALLAETVADLLKPLEKPLVIGQ.FTPTVIMVAGVNGAGKTTS  226
NTDB id 1118 NGFG RS11455 WP 003696286.1       LETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTS  233
consensus                                      !!* !* *! !* !!!!!* !*!!!!      !*  *** ! ! * !!*!!!!!!!!** *  ! !!!*!!*!!!!!!!!

logo
NTDB id 1007803 ABL838 RS26205 WP 418122633.1  IGKLTKHLADEGASVLLAAADTFRAAAREQLLVWADRNTVEIVSQEGGDPSAVSFDAVNAGKARGKDVVLVDTAGRLPTQ  306
NTDB id 1118 NGFG RS11455 WP 003696286.1       IGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQ  313
consensus                                      !!!! !*    ! !!!!!!*!!!!!!!!!!!  !* !! ! **!!  !!* !!*!!!!*!*!!!! !*!! !!!!!!!!

logo
NTDB id 1007803 ABL838 RS26205 WP 418122633.1  LHLMDELKKIKRVVTKADGTAPHEVLLVIDGNTGQNALAQVKAFDETLGLTGLVVTKLDGTAKGGVLCAIARERPIPVYF  386
NTDB id 1118 NGFG RS11455 WP 003696286.1       LHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRY  393
consensus                                      !!!!*!*!!*!!!* !! * !!!!***!*!*! !!!!* !!!!!!* !!!!!!*!!!!!!!!!!!!*!*!*! *!!*!! *

logo
NTDB id 1007803 ABL838 RS26205 WP 418122633.1  IGVGEKLEDLETFNAREFAQALLG  410
NTDB id 1118 NGFG RS11455 WP 003696286.1       IGVGEGIDDLRPFDARAFVDALLD  417
consensus                                      !!!!! **!! *! !! !  !!!
```

X non conserved

X similar

X $\geq$ 50% conserved