```
logo
NTDB id 1002207 ABFO07 RS04420 WP 059566541.1  MFSFFKRFKKTQEPTPAESPSAAEPL.....................PDAQPASAPADVPAA.....STPQPVAQPPAPAQ   55
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      !!!!!*! !!    ! * ! *    * ********************  ** !*  !*  *****  * *!! *! *!

logo
NTDB id 1002207 ABFO07 RS04420 WP 059566541.1  PAAPAVVMTVTSAQNGRDEVVETVE....IV.PPPVQDAGAKKSWLARLKSGLAKTG....SNITGVFVNTKIDEDLYEE  126
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAER....VE...SAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEE  153
consensus                                      *!!* ****! *** *** ! !!! **** !****     *!  ! !!!! !!!!* ****  * !!!    ! !!!!!!

logo
NTDB id 1002207 ABFO07 RS04420 WP 059566541.1  LETALLMSDAGVDATEYLLGALREKVRAGRLTDPQQVKVALRDLLVELLQPLEKSLMLG.RAQPLVMMIAGVNGAGKTTS  205
NTDB id 1118 NGFG RS11455 WP 003696286.1       LETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTS  233
consensus                                      !!! !*  ! !**!!!!!*  *! *!    ! !** ** !!** ! *!* !!!!*!*!**   ! !*!*!!*!!!!!!!!

logo
NTDB id 1002207 ABFO07 RS04420 WP 059566541.1  IGKLAKHLQSFDQSVLLAAGDTFRAAAREQLTIWGERNNVTVVQQESGDPAAVIFDAVSAARARGIDVMMADTAGRLPTQ  285
NTDB id 1118 NGFG RS11455 WP 003696286.1       IGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQ  313
consensus                                      !!!!!!* !    !!!!!!!!!!!!!!!!!!  !! !!!!!!* ! *!!*!!!*!!!! !!*!!!!!***!!!!!!!!!!

logo
NTDB id 1002207 ABFO07 RS04420 WP 059566541.1  LHLMEELKKVKRVISKAHDGAPHEVLLVIDANTGQNALTQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYF  365
NTDB id 1118 NGFG RS11455 WP 003696286.1       LHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRY  393
consensus                                      !!!!!!*!!!!!!* !!**!!!!!***!*!!! !!!!* !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!  !!!!! *

logo
NTDB id 1002207 ABFO07 RS04420 WP 059566541.1  IGVGEKVEDLQPFSAEEFADALLG   389
NTDB id 1118 NGFG RS11455 WP 003696286.1       IGVGEGIDDLRPFDARAFVDALLD   417
consensus                                      !!!!! **!! !! !  ! !!!!
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X non conserved

X similar

X ≥ 50% conserved