```
logo
NTDB id 1009357 ABNN95 RS30135 WP 039346088.1  MFSFFKRFKKTQEPDPAESQSADAQQTDEPSDAPAVE.....APPVADVPQAPAQPAAP.......AVV..MTVTPTND.   65
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      !!!!!*! !!    ! * ! !  *     !*  !* ! ***** ***!* * **! *! ******** ! ***   !*  *

logo
NTDB id 1009357 ABNN95 RS30135 WP 039346088.1  ........GRDEVVETVEIV.....PPPLQDATAKKSWLARLKTGLAKTG....SSITGVFVNTKIDEDLYEELETALLM  128
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                      *********** ! !!!   ********     *!  ! !!!! !!!!* **** !* !!!    ! !!!!!!!!! !*

logo
NTDB id 1009357 ABNN95 RS30135 WP 039346088.1  SDAGVDATEYLLGALREKVRTGRLTDPQQVKAALHDLLVELLTPLEKSLMLGR.AQPLVMMITGVNGAGKTTSIGKLAKH  207
NTDB id 1118 NGFG RS11455 WP 003696286.1       GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                       ! !**!!!!!*  *! *!    ! !** ***!** ! *!* !!!!*!*!* *  ! !*!* !*!!!!!!!!!!!!!!*

logo
NTDB id 1009357 ABNN95 RS30135 WP 039346088.1  LQSFDQSVLLAAGDTFRAAAREQLAVWGERNNVTVVQQESGDPAAVIFDAVSAARARKIDVMMADTAGRLPTQLHLMEEL  287
NTDB id 1118 NGFG RS11455 WP 003696286.1       FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                       !    !!!!!!!!!!!!!!!!!!  !! !!!!!!* ! *!!*!!!*!!!! !!*!! !!***!!!!!!!!!!!!!!!!!*

logo
NTDB id 1009357 ABNN95 RS30135 WP 039346088.1  KKVKRVISKAHDGAPHEVLLVIDANTGQNALTQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEKV  367
NTDB id 1118 NGFG RS11455 WP 003696286.1       KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                      !!!!!!* !!**!!!!!***!*!!! !!!!* !!!!!!!!!!!!!!!!!!
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