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logo
NTDB id 256527 CVS37 RS17120 WP 122946308.1   MFSFFKRFKKTQEPDPAESQSADAQQTDEPSDAPAVE.....APPVAEVPRAPAQPAAPA.......VV..MTVTPTND.   65
NTDB id 1118 NGFG RS11455 WP 003696286.1      MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                     !!!!!*! !!   ! * ! !  *     !*  !* ! ***** ***!! ****! *! * *******! ***   !*  *

logo
NTDB id 256527 CVS37 RS17120 WP 122946308.1   ........GRDEVVETVEIV.....PPPLQDATAKKSWLARLKTGLAKTG....SSITGVFVNTKIDEDLYEELETALLM   128
NTDB id 1118 NGFG RS11455 WP 003696286.1      EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT   160
consensus                                     *********** ! !!!   ********     *!  ! !!!! !!!!* **** !* !!!    ! !!!!!!!!! !*

logo
NTDB id 256527 CVS37 RS17120 WP 122946308.1   SDAGVDATEYLLGALREKVRAGRLTDPQQVKAALHDLLVELLSPLEKSLMLGR.AQPLVMMITGVNGAGKTTSIGKLAKH   207
NTDB id 1118 NGFG RS11455 WP 003696286.1      GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY   240
consensus                                      ! !**!!!!!*  *! *!    ! !** ***!!** ! *!* !!!!*!*!* *  ! !*!* !*!!!!!!!!!!!!!!*

logo
NTDB id 256527 CVS37 RS17120 WP 122946308.1   LQSFDQSVLLAAGDTFRAAAREQLAVWGERNNVTVVQQESGDPAAVIFDAVSAARARKIDVMMADTAGRLPTQLHLMEEL   287
NTDB id 1118 NGFG RS11455 WP 003696286.1      FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI   320
consensus                                      !    !!!!!!!!!!!!!!!!!!  !! !!!!!!* ! *!!*!!!*!!!! !!*!! !!***!!!!!!!!!!!!!!!!*

logo
NTDB id 256527 CVS37 RS17120 WP 122946308.1   KKVKRVISKAHDGAPHEVLLVIDANTGQNALTQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEKV   367
NTDB id 1118 NGFG RS11455 WP 003696286.1      KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI   400
consensus                                     !!!!!!* !!**!!!!!***!*!!! !!!!* !!!!!!!!!!!!!!!!!!
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