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logo

NTDB id 421394 G0D92 RS13925 WP 006416170.1  MFSFFKRFKKTQEPGSAESQPADAPQPDAPSDA.....HDADVPPAP....APAEPAAP.......AV..VMTVTPSDNG   62
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                    !!!!!*! !!   !*  ! !* * * *  *  !******  !!** *******!!*! ******** !** *   !!* !

logo

NTDB id 421394 G0D92 RS13925 WP 006416170.1  .........RDEVVETVEIV.....PPPVQDASAKKSWLARLKTGLAKTG....SSITGVFVNTKIDDALYEELETALLM  124
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                    *********** ! !!!   ********     *!  ! !!!! !!!!* **** !* !!!    ! * !!!!!!! !*

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NTDB id 421394 G0D92 RS13925 WP 006416170.1  SDAGVDATEYLLGALREKVRVGRLTDPQQVKSALRDLLVELLTPLEKSLVLGR.AQPLVMMIAGVNGAGKTTSIGKLAKH  203
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                     ! !**!!!!!*  *! *! *  ! !** ** !!** ! *!* !!!!*!!!* *  ! !*!*!!*!!!!!!!!!!!!!!*

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NTDB id 421394 G0D92 RS13925 WP 006416170.1  LQSFDQSVLLAAGDTFRAAAREQLTVWGERNNVTVVQQESGDPAAVIFDAVSAARARKIDVMMADTAGRLPTQLHLMEEL  283
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                     !    !!!!!!!!!!!!!!!!!!  !! !!!!!!* ! *!!*!!!*!!!! !!*!! !!***!!!!!!!!!!!!!!!!!*

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NTDB id 421394 G0D92 RS13925 WP 006416170.1  KKVKRVIAKAHDGAPHEVLLVIDANTGQNALTQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEKV  363
NTDB id 1118 NGFG RS11455 WP 003696286.1     KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                    !!!!!!* !!**!!!!!***!*!!! !!!!* !!!!!!!!!!!!!!!!!!
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