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logo

NTDB id 34173 BGLU RS16435 WP 015877057.1   MFSFFKRLIGSKADDASQDA...PAADEQASAA............HAPDAAGHAAAAAS...........RPAPAPAHDAEPAAS    59
NTDB id 1118 NGFG RS11455 WP 003696286.1    MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAGEA...    82
consensus                                   !!!!!*!       *      ****!!      !*************!*   !*!  !  *********** *!* ! * *!****

logo

NTDB id 34173 BGLU RS16435 WP 015877057.1   PAAPVPAAPAPAAVPAAPA.VEVVPPPAPEPAAKKSWIARLRSGLSKTS....TSLTNVFVNTRIDDELYEELEAALLMSDAGID   139
NTDB id 1118 NGFG RS11455 WP 003696286.1    .AERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLITGDMGME   166
consensus                                   *! *!* !* * ! * **!*  ! ***!* * *!  ! !!!* !! !* **** !!  !!    ! **!!!!!!  !*  ! !**

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NTDB id 34173 BGLU RS16435 WP 015877057.1   ATEYLLGALREKVRAERLSDPLQVKAALRALLVDLLAPLEKSLMLGR.AQPLVMMIAGVNGAGKTTSIGKLAKHLQRFDQSVLLA   223
NTDB id 1118 NGFG RS11455 WP 003696286.1    ATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLA   251
consensus                                   !!!!!*  *!  *!    ! !*  ***!!*  ! !!* !!!!*!*!* *  ! !*!*!!*!!!!!!!!!!!!!!* !    !!!!!

logo

NTDB id 34173 BGLU RS16435 WP 015877057.1   AGDTFRAAAREQLAIWGERNNVTVVQQESGDPAAVIFDAVGAARARKIDVVMADTAGRLPTQLHLMEELRKVKRVIGKAQSDAPH   308
NTDB id 1118 NGFG RS11455 WP 003696286.1    AGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPH   336
consensus                                   !!!!!!!!!!!!!  !! !!!!!!* ! *!!*!!!*!!!! !!*!! !!*!*!!!!!!!!!!!!!!!!!!**!!!!!* !! * !!!

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NTDB id 34173 BGLU RS16435 WP 015877057.1   EVLLVIDANTGQNALAQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPIPVYFIGVGEKVEDLQPFNAEEFADALLG   389
NTDB id 1118 NGFG RS11455 WP 003696286.1    EIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD   417
consensus                                   !***!*!!! !!!!! !!!!!!!!!!!!!!!!!!!
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