```
logo
NTDB id 114298 R615 RS00675 WP 025264486.1  MFDFFKRKKDKNKSEKPESAE......LDSALEQTLEPISSQPEISEIEPQPVDVITEPEKAMIKPPV..VEE..MVVEPTPVEI   75
NTDB id 1118 NGFG RS11455 WP 003696286.1    MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKE..........DVESLAESVKGRAESAVETVSGAVEQVKETVAEM   75
consensus                                   !! !!*!!!    *  *!   ********! * ! * *!  ***********!* * !* !*   **!**!  **  ! *!* !*

logo
NTDB id 114298 R615 RS00675 WP 025264486.1  PPVVAPNELDI.TPSSAVAEINPD..VMVDVPVKAVAEEKKGFFGRIRSGLSKTRSGFTDGLASLLLGKKEIDDDLMDDLETQLI  157
NTDB id 1118 NGFG RS11455 WP 003696286.1    PSEAGEAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGV.FGGGQIGEDLYEELETVLI  159
consensus                                   !*  **    ****  !!!!  ****  !  *!    !*! !* *!** !! !*!      !! ** !   ! *!! **!!! !!

logo
NTDB id 114298 R615 RS00675 WP 025264486.1  MADVGVEATTEIIKRLTARVSRKELKDSDALYEALIDELQDMLAVSQKPLVIDSSKKPYVILMVGINGVGKTTTIGKLAKQFQKR  242
NTDB id 1118 NGFG RS11455 WP 003696286.1    TGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKYFQAQ  244
consensus                                    *!*!*!!!  **! * *!!! ! !!!   !  !! * ! !** *  !!!!** *! !*!!** !!!! !!!!*!!!!!! !!

logo
NTDB id 114298 R615 RS00675 WP 025264486.1  GKSVMLAAGDTFRAAAVEQLQVWGERNNVPVVAQHTGADSASVLFDALQAAQARKVDVLIADTAGRLHNKDNLMQELEKVVRVMQ  327
NTDB id 1118 NGFG RS11455 WP 003696286.1    GKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTG.DSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQ  328
consensus                                   !!!!*!!!!!!!!!!! !!!! !! !!!!*!* !*!!*!!! !*!!!*!!! !! *!***!!!!!!!    *!! !* !! !!*!

logo
NTDB id 114298 R615 RS00675 WP 025264486.1  KLDPSAPHEVMLVLDAGTGQNAINQTKHFLQSVGVTGLTLTKLDGTAKGGIIFALAKQFGLPVRYIGVGEGIDDLRPFNADEFTR  412
NTDB id 1118 NGFG RS11455 WP 003696286.1    KAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGIDDLRPFDARAFVD  413
consensus                                   !  ! !!!!***!!!!  !!!!*!! !*!   *!*!!! *!!!!!!!!!!!* !!!   **!!!!!!!!!!!!!!!!! !  !

logo
NTDB id 114298 R615 RS00675 WP 025264486.1  ALFRHDTKED   422
NTDB id 1118 NGFG RS11455 WP 003696286.1    ALLD......   417
consensus                                   !!  ******
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X non conserved

X similar

X ≥ 50% conserved