```
logo
NTDB id 364783 FE661 RS13675 WP 009568264.1  MVFDWFKRNK.....................................................SL....   12
NTDB id 1118 NGFG RS11455 WP 003696286.1     .MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEA   79
consensus                                    **! *!*! !*************************************************************** *****

logo
NTDB id 364783 FE661 RS13675 WP 009568264.1  ............PAGETFAEEVNSPATVATIPEDTPKGLFSRLRQGLARSREQLAGGVGRLVLGKKVIDADLLEELEALL   80
NTDB id 1118 NGFG RS11455 WP 003696286.1     GEAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAG.VFGGGQIGEDLYEELETVL  158
consensus                                    ************* *!! *! !  *      *!*** !   !!*!!!!*!!* *!  ** *! !   !  !! !!!! *!

logo
NTDB id 364783 FE661 RS13675 WP 009568264.1  LQADLGSAATREVMDSVTERVRRNELTDPAALQAALRESLLDILRPRAQPWS.PEKGHSQVLLMVGINGAGKTTTIGKLA  159
NTDB id 1118 NGFG RS11455 WP 003696286.1     ITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLA  238
consensus                                    * *!*!  !!  *!  !  !!    ! !*  ! *!!*! ! !***!   !  *!!  ** !*** !!!!!!!!!*!!!!!

logo
NTDB id 364783 FE661 RS13675 WP 009568264.1  ARWKAEGFTVVLAAGDTFRAAAVEQLQGWGRRAQVPVVAQGTGADSASVIFDALTTTRARGSDLLIADTAGRLHTQGHLM  239
NTDB id 1118 NGFG RS11455 WP 003696286.1     KYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTG.DSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLM  317
consensus                                      * ! ! *!*!!!!!!!!!!! !!!!*!! ! *!*!* ! !!*!!! !*!!!*   *!!! !***!!!!!!! !! !!!

logo
NTDB id 364783 FE661 RS13675 WP 009568264.1  EELKKIRRVLGKQDPEAPQQIWLVLDAGTGQNALNQARQFHDAVGLTGICITKLDGTAKGGVVAAIAKALPIPIRYIGVG  319
NTDB id 1118 NGFG RS11455 WP 003696286.1     EEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVG  397
consensus                                    !!*!!**!!! !  ! !!* ! *!!!!  !!!!*!! * !*!!*!!!!***!!!!!!!!!!**!!*!   !*!*!!!!!!

logo
NTDB id 364783 FE661 RS13675 WP 009568264.1  EQVEDLRPFDPESFVDALFAESKS   343
NTDB id 1118 NGFG RS11455 WP 003696286.1     EGIDDLRPFDARAFVDALLD....   417
consensus                                    ! **!!!!!!*  !!!!!  ****
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X non conserved

X similar

X $\geq$ 50% conserved