```
logo
NTDB id 107627 AABA48 RS08975 WP 286350165.1  ..............................................MTEERKGFFKGLWGKVTGGDRD.............  22
NTDB id 1118 NGFG RS11455 WP 003696286.1      MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG  80
consensus                                     ********************************************** !  !!        !*!*   *************

logo
NTDB id 107627 AABA48 RS08975 WP 286350165.1  ........QEEAVDETT.SAVAD.VAAPPEQEDRRPGFFERLKQGLSKTRDSLVGRIDRLVLGKKEIDADTLEELEEILI  92
NTDB id 1118 NGFG RS11455 WP 003696286.1      EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAG.VFGGGQIGEDLYEELETVLI  159
consensus                                     ********  !!! !! * !!* *  !**  !****!*  !!!!!! !*!! *   *  *! !   !  !  !!!! *!!

logo
NTDB id 107627 AABA48 RS08975 WP 286350165.1  TADLGVQTTVELIRGLEQRLSRNELKDGEALREALKEDIHGRLARDAHQL.DVTGASPFVIMVIGVNGVGKTTTIGKLAA  171
NTDB id 1118 NGFG RS11455 WP 003696286.1      TGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAK  239
consensus                                     !*!*!*  !  !** *  !*!   !!!!  !! !!!! **  * *  **!* *    !!!!!* !*!! !!!!*!!!!!

logo
NTDB id 107627 AABA48 RS08975 WP 286350165.1  RFTAQGKRVILAAGDTFRAAAAEQLQIWGERTGVDVIRHKEGADPSAVVFDSIKAAVARGADILIVDTAGRLHTKVNLME  251
NTDB id 1118 NGFG RS11455 WP 003696286.1      YFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTG.DSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLME  318
consensus                                      ! !!!! !*!!!!!!!!!!! !!!! !! !  ! !! *  !*!* !!*!! * !! !!! !!** !!!!!! ! **!!!

logo
NTDB id 107627 AABA48 RS08975 WP 286350165.1  ELKKVRRIMSREIPGAPHETLLVLDAATGQNALSQAKLFKEAAQVTGIALTKLDGTAKGGIVVAICNEFRIPVRYIGVGE  331
NTDB id 1118 NGFG RS11455 WP 003696286.1      EIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDAALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGE  398
consensus                                     !*!!!*!** * !!!!!!! **!!!!  !!!!* ! ! ! *!  *!!* *!!!!!!!!!!!* !** * **!!!!!!!!!!

logo
NTDB id 107627 AABA48 RS08975 WP 286350165.1  GVDDLRDFDPSQFVDALFQ  350
NTDB id 1118 NGFG RS11455 WP 003696286.1      GIDDLRPFDARAFVDALLD  417
consensus                                     !*!!!!*!!*  !!!!!
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X non conserved

X similar

X ≥ 50% conserved