```
logo
NTDB id 973040 WN985 RS15385 WP 342307843.1  MFSFFKRFKKTQEPDPAESQSADASQTDEPSDAPAVDA.....PPAAEVP....QAPVQPA.G..PAVVM..TVTPTND.   65
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                    !!!!!*! !!   ! * ! !  *     !*  !* !  ******* !! *****  *! * *!*** !  **   !*  *

logo
NTDB id 973040 WN985 RS15385 WP 342307843.1  ........GRDEVVETVEI....V.PPPLQDASAKKSWLARLKTGLAKTG....SSITGVFVNTKIDEDLYEELETALLM  128
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                    *********** ! !!!  ****!****     *!  ! !!!! !!!!* **** !* !!!    ! !!!!!!!!! !*

logo
NTDB id 973040 WN985 RS15385 WP 342307843.1  SDAGVDATEYLLDALREKVRTGRLTDPQQVKSALHDLLVELLKPLEKSLMLGR.AQPLVMMITGVNGAGKTTSIGKLAKH  207
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                     ! !**!!!!!*  *! *!    ! !** ** !!** ! *!*!!!!!*!*!* *  ! !*!* !*!!!!!!!!!!!!!!*

logo
NTDB id 973040 WN985 RS15385 WP 342307843.1  LQSFDQSVLLAAGDTFRAAAREQLAVWGERNNVTVVQQESGDPAAVIFDAVSAARARKIDVMMADTAGRLPTQLHLMEEL  287
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                     !    !!!!!!!!!!!!!!!!!!  !! !!!!!!* ! *!!*!!!*!!!! !!*!! !!***!!!!!!!!!!!!!!!!*

logo
NTDB id 973040 WN985 RS15385 WP 342307843.1  KKVKRVISKAHDGSPHEVLLVIDANTGQNALTQVKAFDDALGLTGLIVTKLDGTAKGGILAAIARQRPVPVYFIGVGEKV  367
NTDB id 1118 NGFG RS11455 WP 003696286.1     KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                    !!!!!!* !!**! !!!***!*!!! !!!!* !!!!!!!!!!!!!!!!!!
```