```
logo

NTDB id 1100986 ACNI65 RS01380 WP 416760760.1  MFSFLKKKLGFGSSEPAPPPAPETAPAVEPVQAPAPAAPAPQI.PIPATVAAPA...............PV..YVPAPTP   62
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      !!!! **!     ** * *** *!!!* !!*  !* ** * * ***** *! **!*************** ** *! * *

logo

NTDB id 1100986 ACNI65 RS01380 WP 416760760.1  APAPVVAPAPAPVAPP....APIPVAAPEPEPAPRKSWMDKLRQGLRKTG....SSISQVFTGTQIDDALYEELEAALLM  134
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                       *!* ! *!* *!!****** * *  !* * *  *  !  *!*!!! !* **** !*  !! ! !! * !!!!!!  !*

logo

NTDB id 1100986 ACNI65 RS01380 WP 416760760.1  ADAGVKATEFLLEDLKRRVKDAKVTDPRAVRQLLVDALAELLQPLQRQLVIGE.QSPMVMMVVGVNGAGKTTSIGKLTRH  213
NTDB id 1118 NGFG RS11455 WP 003696286.1       GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                      *! !* !!!*!* !** !!    * !*  *!  ! *!! *!* !! **!!**!*  ! !*!* !*!!!!!!!!!!!! **

logo

NTDB id 1100986 ACNI65 RS01380 WP 416760760.1  LADAEANVLLAAADTFRAAAREQLAVWADRNRVQIVSQEGGDPSAVTFDAVVAGKARGCDVVIADTAGRLPTQLHLMEEL  293
NTDB id 1118 NGFG RS11455 WP 003696286.1       FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                            !!!!!*!!!!!!!!!!!  !* !! ! **!!  !!* !!*!!!! !*!!!!*!*!*!!!!!!!!!!!!!!!!*

logo

NTDB id 1100986 ACNI65 RS01380 WP 416760760.1  KKIKRVIGKAQETAPHEVLLVVDGNTGQNALMQVKSFDDALGLTGLIVTKLDGTAKGGVLAAIARERPIPVYFIGVGEKL  373
NTDB id 1118 NGFG RS11455 WP 003696286.1       KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                      !!*!!!* !! * !!!!***!*!*! !!!!* !!! !!!!!!!!!!!!!!!!!!!!!!!!*!!!*! *!!*!! *!!!!! *

logo

NTDB id 1100986 ACNI65 RS01380 WP 416760760.1  EDLQTFDAREFAQALLE   390
NTDB id 1118 NGFG RS11455 WP 003696286.1       DDLRPFDARAFVDALLD   417
consensus                                      *!! *!!!! !  !!!*
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X non conserved

X similar

X ≥ 50% conserved