```
logo
NTDB id 1092568 ACLSSQ RS16940 WP 416243321.1  ..............................................................MFGFLKKSDK..........   10
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                      ************************************************************ ! *     **********

logo
NTDB id 1092568 ACLSSQ RS16940 WP 416243321.1  ........DQD.....ATPATAAAVAPVPEPLDKPLSWRERLFAGLSRTRSQLGGKLKSVFARGKVDDELLEELETLLLT   77
NTDB id 1118 NGFG RS11455 WP 003696286.1       EAAERVESAKEAVAETVGEAVGQVQEAVATIEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                      ********  ******  *! *    *!* * ***! !  !!  !! **!  **  !  !!* ! * **! !!!!!*!*!

logo
NTDB id 1092568 ACLSSQ RS16940 WP 416243321.1  SDVGMDATMHLLDELKRRAKQDKLEGAAAIQQALADTLAQMLAPLEAPLQV.DSHKPFVIMIAGVNGAGKTTSIGKLAKY  156
NTDB id 1118 NGFG RS11455 WP 003696286.1       GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                       !*!!*!! *!* *** !     !  *  *  !! * !  ** !!! !! ***** !!!!!*!!*!!!!!!!!!!!!!!!

logo
NTDB id 1092568 ACLSSQ RS16940 WP 416243321.1  FQGQGKSVLLAAGDTFRAAAREQLMTWGERNNVTVIAQESGDPAAVVFDAIAAAKARGIDVVLADTAGRLPTQLHLMEEI  236
NTDB id 1118 NGFG RS11455 WP 003696286.1       FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                      !!*!!!!!!!!!!!!!!!!!!!!!  !! !!!!!!! ! *!!*!!!*!!!* !!!!!!!!*!!!!!!!!!!!!!!!!!!!

logo
NTDB id 1092568 ACLSSQ RS16940 WP 416243321.1  AKVRRVIQKADPTGPHEVMLVLDANIGQNALAQVKAFDKSINVTGLVLTKLDGTAKGGVIAAIARQCPKPVRFIGVGEGI  316
NTDB id 1118 NGFG RS11455 WP 003696286.1       KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                       !!*!!*!!! ! *!!!***!!!!!!!!!!* !!!!!!  * *!!!**!!!!!!!!!!!**!!*!  *! !!!*!!!!!!!

logo
NTDB id 1092568 ACLSSQ RS16940 WP 416243321.1  DDLRPFVARDFVDALFGD   334
NTDB id 1118 NGFG RS11455 WP 003696286.1       DDLRPFDARAFVDALLD.   417
consensus                                      !!!!!! !! !!!!!  *
```

X non conserved

X similar

X $\geq$ 50% conserved