```
logo
NTDB id 63469 TEL01S RS01175 WP 028843511.1  ..............................................................................  0
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG  80
consensus                                    ******************************************************************************

logo
NTDB id 63469 TEL01S RS01175 WP 028843511.1  ...................................MGFFERLKTGLSRARKALFEGLSQILKGSKISDEILEELEERLLM  45
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                    ************************************!*  !!! !! * !  *   !  *  !  ! *** !!!!  !*

logo
NTDB id 63469 TEL01S RS01175 WP 028843511.1  ADVGYPTTEYIIEQLKQSKTNDAL.......ETLKEIMIDLLS.FDSK.LNSPDQPPFVITVVGVNGTGKTTTVAKLAAY  116
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                    *!*! * !!!**  **   *  *!*******  !!! * !!* * * *** **  *!!!! * !*!! !!!!***!!! !

logo
NTDB id 63469 TEL01S RS01175 WP 028843511.1  FQSTGKTVVLGAADTFRAAAIEQLREWGERTGSTVIYHSEGADSAAVAFDAVSHAKAREKDVVIIDTAGRLHTKKNLMEE  196
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTT.GDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEE  319
consensus                                    !!  !!*!*!*!*!!!!!!! !!!  !! !   !!! ** **!!!!!*!!!! *!!!!  !*!* !!!!!! !  *!!!!

logo
NTDB id 63469 TEL01S RS01175 WP 028843511.1  LKKVHRVIGKVVPGAPHEVLLVIDATTGQNGLVQAKVFKEMVDVTGIVITKLDGTAKGGISLAIKKELNLPIKFIGIGEG  276
NTDB id 1118 NGFG RS11455 WP 003696286.1     IKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEG  399
consensus                                    *!!!*!!* ! *!!!!!!***!*!!  !!!** ! ! ! * * *!!***!!!!!!!!!!!  !*  * **!***!!*!!!

logo
NTDB id 63469 TEL01S RS01175 WP 028843511.1  AEDLKPFDAKEFVEALIY  294
NTDB id 1118 NGFG RS11455 WP 003696286.1     IDDLRPFDARAFVDALLD  417
consensus                                     *!!*!!!!* !!*!!*
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X non conserved

X similar

X ≥ 50% conserved