```
logo
NTDB id 67012 TAO RS08125 WP 096527433.1  MVNFFKRWQKNPTQS.......................................................PDITSP   21
NTDB id 1118 NGFG RS11455 WP 003696286.1  MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAGEAAER   85
consensus                                 !  !!*!  !**!* ************************************************************   *

logo
NTDB id 67012 TAO RS08125 WP 096527433.1  TPIDTKVVPENSK.......EEI...TTPQKQFFSRLYDRLSRTRNHLSSGTYSLILGKKTIDDQLLEEIETQLLIADVGVETAQ   96
NTDB id 1118 NGFG RS11455 WP 003696286.1  VESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSL.AGVFGGGQIGEDLYEELETVLITGDMGMEATE  169
consensus                                  *     !*!   *******! ****      *  !!   ! **! **   **  * !   ! * ! !!*!! !* *!*!*!

logo
NTDB id 67012 TAO RS08125 WP 096527433.1  STIQDLTARVSRKELKDPERLMGALQENMQALLKPCHQPLVLPENLKPFTILMVGINGAGKTTTIGKLAKRFQAQNRSVMLAAGD  181
NTDB id 1118 NGFG RS11455 WP 003696286.1  YLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKYFQAQGKSVLLAAGD  254
consensus                                  ** !* *!!! ! !!!*  ! !!! ! *  !*!!** !!!!!!   !! !** !!!!!!!!!*!!!!!! !!!! *!!*!!!!!

logo
NTDB id 67012 TAO RS08125 WP 096527433.1  TFRAAAVEQLQVWGERNQIPVIAQHKGADSASVIFDAMQSAQARGIDVLIADTAGRLHTQANLMEELKKIKRVMTKADPTAPHEI  266
NTDB id 1118 NGFG RS11455 WP 003696286.1  TFRAAAREQLQAWGGRNNVTVISQTTG.DSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEIKKVKRVLQKAIPGAPHEI  338
consensus                                 !!!!!! !!!! !! !!***!! !* !*!!! !*!!!*! ! !!!!!***!!!!!!! !! *!!!!*!!*!!!* !! ! !!!!!

logo
NTDB id 67012 TAO RS08125 WP 096527433.1  LLTVDAGIGQNALAQTQQFHQAIGLTGIALTKLDGTAKGGIIFAIARKMALPIRFIGVGEGVDDLRPFDAEEFVEALLTLAPPKE  351
NTDB id 1118 NGFG RS11455 WP 003696286.1  IVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGIDDLRPFDARAFVDALLD......  417
consensus                                 ** *!! !!!!!* !   !* !*!!!!* *!!!!!!!!!!!* !*!   **!*!*!!!!!!*!!!!!!!!  !!*!!! ******

logo

NTDB id 67012 TAO RS08125 WP 096527433.1  SV   353
NTDB id 1118 NGFG RS11455 WP 003696286.1  ..   417
consensus                                 **
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X non conserved

X similar

X ≥ 50% conserved