```
logo
NTDB id 1007911 ABL841 RS19505 WP 418148816.1  MFSFFKKKPPAESPAAPAPSPAAEPAAAPEPAPAPAAAPA.RSVFSP.SSWFGSKPAAEEAAP...PPAAPPVAPPPPPP   75
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAA.KVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSE.   78
consensus                                      !!!!!**!** !*!! * * *   *!* *!* *!*  ** *  ! !** !  !  * ! !  ******  **!!**!***

logo
NTDB id 1007911 ABL841 RS19505 WP 418148816.1  TVAPPAAPVPAPAPAPAPAPAPVPA.PAPAVLTEAAMPAERKGWFDKLKTGLRKTG....TGIQAVFVNAQIDEALYEEL  150
NTDB id 1118 NGFG RS11455 WP 003696286.1       .AGEAAERVESAKEA...VAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEEL  154
consensus                                      * ***! *!* * *!*** * *!* ****     !  * ** !!  *!! !! !* ****  * *!!  *!! ! !!!!!

logo
NTDB id 1007911 ABL841 RS19505 WP 418148816.1  ESALLMADAGVKATEFLLDDLRGRVKRQMATDAAAQVKRLLADAITDLLQPLEKPLVIGQ.FTPTVIMVAGVNGAGKTTSI  229
NTDB id 1118 NGFG RS11455 WP 003696286.1       ETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSI  234
consensus                                      !* !* *! !* !!!*!* !*!!!!      !*  **  ! *!* !!* !!!!!!!** *  ! !!!*!!*!!!!!!!!!

logo
NTDB id 1007911 ABL841 RS19505 WP 418148816.1  GKLTKHLASEGASVLLAAADTFRAAAREQLLVWADRNTVEIVSQEGGDPSAVSFDAVTAGKARGKDVVLVDTAGRLPTQL  309
NTDB id 1118 NGFG RS11455 WP 003696286.1       GKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQL  314
consensus                                      !!! !*    ! !!!!!!*!!!!!!!!!!!  !* !! ! **!!  !!* !!*!!!! !*!!!! !*!! !!!!!!!!!!

logo
NTDB id 1007911 ABL841 RS19505 WP 418148816.1  HLMDELKKIKRVVTKADATAPHEVLLVIDGNTGQNALAQVRAFDETLGLTGLVVTKLDGTAKGGVLCAIARERPIPVYFI  389
NTDB id 1118 NGFG RS11455 WP 003696286.1       HLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYI  394
consensus                                      !!!*!*!!*!!!* !! * !!!!***!*!*! !!!!* !!*!!!* !!!!!!*!!!!!!!!!!!!*!*!*! *!!*!! *!

logo
NTDB id 1007911 ABL841 RS19505 WP 418148816.1  GVGEKLEDLETFNAREFALALLG   412
NTDB id 1118 NGFG RS11455 WP 003696286.1       GVGEGIDDLRPFDARAFVDALLD   417
consensus                                      !!!! **!! *! !! !  !!!
```

X non conserved

X similar

X $\geq$ 50% conserved