```
logo
NTDB id 280790 C7H73 RS04830 WP 106845604.1  MFSFFKKKTPDDIPAEAPLPPAPAAAPVPAPAPAAL..........A....TPAAPAAPESAGAMAWLRKSF.......G   59
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                    !!!!!**! * * !!  * ** * !!*!* * *  ***********!**** *! *! *   !!*  ** * *******!

logo
NTDB id 280790 C7H73 RS04830 WP 106845604.1  TG...APSAPPAVAE........PMDKPLPAGAARQHWVERLKTGLKKT....GTSITTVFTGTQINDALYEELEEALLM  124
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                     **** *!!**!!!********* * * *   ** *!  !!! !! !****** !*  !! ! !! * !!!!!!  !*

logo
NTDB id 280790 C7H73 RS04830 WP 106845604.1  ADTGVRATQHLLQDLKRRVKETKTTEPAAVKALLADALADLLRPLEKPLVIGEH.TPTVVMVAGVNGAGKTTSIGKLTRH  203
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                    *! !* !! *!* !** !!      **  *** ! *!! !!**!!!!!!!**!** ! !*!*!!*!!!!!!!!!!!! **

logo
NTDB id 280790 C7H73 RS04830 WP 106845604.1  LADHGQSVLLAAADTFRAAAREQLGVWATRNTVEIVSQEGGDPAAVSFDAVSAGRARGKDVVLVDTAGRLPTQLHLMQEL  283
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                       *! !!!!!!*!!!!!!!!!!!  !* !! ! **!!  !!*!!!*!!!! !**!!! !*!! !!!!!!!!!!!!! !*

logo
NTDB id 280790 C7H73 RS04830 WP 106845604.1  QKIRRVIAKADAGAPHEVLLVIDGNTGQNALAQVRAFDDALSLTGLIVTKLDGTAKGGVLAAIAQERPIPVYFIGVGERV  363
NTDB id 1118 NGFG RS11455 WP 003696286.1     KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                     !**!!* !! *!!!!!***!*!*! !!!!* !!*!!!!!! !!!!!!!!!!!!!!!!*!!!*! *!!*!! *!!!!! *

logo
NTDB id 280790 C7H73 RS04830 WP 106845604.1  EDLETFSAREFAQALLA   380
NTDB id 1118 NGFG RS11455 WP 003696286.1     DDLRPFDARAFVDALLD   417
consensus                                    *!! *! !! !  !!!
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X non conserved

X similar

X $\geq$ 50% conserved