```
logo

NTDB id 390956 F9K07 RS20395 WP 159595164.1  MFSFFRKKPPAPP.AEPAPDLTEPPA.........................................PVPEAPT...   32
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                    !!!!!!*!** ***! * * * !**!*****************************************!*! !****

logo

NTDB id 390956 F9K07 RS20395 WP 159595164.1  ............PPTVAAPVAMAVAVAAPPERKGWLDKLKSGLRKTG....SSIATVFTGTQIDEALYEELEAALLM   95
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                    ***************!!* *!*         **** !!  *!! !! !* **** !*! !! ! !! ! !!!!!!  !*

logo

NTDB id 390956 F9K07 RS20395 WP 159595164.1  ADTGVKATTDLLVDLKRRVKENKTTDPKAVKGLLADAITDLLAPLQKPLVIGEH.KPTVVMVAGVNGAGKTTSIGKLTRH  174
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                    *! !* !!  !* !** !!      !*  **! ! *!* !!* !! !!!!**!** ! !*!*!!*!!!!!!!!!!!! **

logo

NTDB id 390956 F9K07 RS20395 WP 159595164.1  LADEGATVLLAAADTFRAAAREQLAVWADRNTVEIITQQGGDPAAVSFDAVTAGKARGRDVVLVDTAGRLPTQLHLMEEL  254
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                       ! *!!!!!*!!!!!!!!!!!  !* !! ! *!*!  !!*!!!*!!!! !*!!!! !*!! !!!!!!!!!!!!!!!*

logo

NTDB id 390956 F9K07 RS20395 WP 159595164.1  RKIKRVVQKAEASAPHEVLLVIDGNTGQNALNQVKAFDDALQLTGLIITKLDGTAKGGVLCAIAREKPVPVYFIGVGEKL  334
NTDB id 1118 NGFG RS11455 WP 003696286.1     KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                    *!*!!!*!!! * !!!!***!*!*! !!!!*!!!!!!!!!! !!!!!*!!!!!!!!!!*!*!*! **!!!! *!!!!! *

logo

NTDB id 390956 F9K07 RS20395 WP 159595164.1  EDLETFDAREFAQALLS   351
NTDB id 1118 NGFG RS11455 WP 003696286.1     DDLRPFDARAFVDALLD   417
consensus                                    *!! *!!!! !  !!!
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X non conserved

X similar

X ≥ 50% conserved