```
logo
NTDB id 246965 CNX70 RS05560 WP 096233884.1  MFSFFKKKPTAPEAAPAEPLVLPA.......................................ATPP...   28
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                    !!!!!**!*  * *!* !* ! * ********************************************  !****

logo

NTDB id 246965 CNX70 RS05560 WP 096233884.1  ........APQPAARPLSGAPAELVPVIVAAEPEKKSWMTRLKAGLSKTSN....TLSVLFVGAKIDDALYEELEAALLM   96
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVLIT  160
consensus                                    ********!* * ! **  !** * * *   !**!  !  !!! !! !*  *****!  *! !* ! * !!!!!!  !*

logo

NTDB id 246965 CNX70 RS05560 WP 096233884.1  SDAGIDATEFLLTELKKKVKEDKLLDAAAVKAALKVLLIDLLSPLEKRFELGR.HKPLVMMISGVNGAGKTTTIGKLAKH  175
NTDB id 1118 NGFG RS11455 WP 003696286.1     GDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLAKY  240
consensus                                     ! !**!!!*!* *** *!    ! !*  ***!!!  ! !!* !!!!*  !* ** ! !*!* !*!!!!!!!*!!!!!!*

logo

NTDB id 246965 CNX70 RS05560 WP 096233884.1  MQSNKQSVLLAAGDTFRAAAREQLMVWGERNNVTVISQESGDPAAVAYDSVQSAKARGTDVVMVDTAGRLPTQLHLMEEL  255
NTDB id 1118 NGFG RS11455 WP 003696286.1     FQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLMEEI  320
consensus                                     ! *  !!!!!!!!!!!!!!!!!!  !! !!!!!!!!! *!!*!!**! !! !!!!! !*!* !!!!!!!!!!!!!!!*

logo

NTDB id 246965 CNX70 RS05560 WP 096233884.1  KKIKRVIGKGMEGAPHETLLVIDGNTGQNALTQVKAFDDALQLTGLVITKLDGTAKGGVLAAIARVRPVPVYFIGIGEKI  335
NTDB id 1118 NGFG RS11455 WP 003696286.1     KKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGEGI  400
consensus                                    !!*!!!* !***!!!!! **!*!*! !!!!* !!!!!!!!! !!!!**!!!!!!!!!!!*!!!*!  !!!!! *!!*!! !

logo

NTDB id 246965 CNX70 RS05560 WP 096233884.1  EDLQPFVAAEFVEALLG   352
NTDB id 1118 NGFG RS11455 WP 003696286.1     DDLRPFDARAFVDALLD   417
consensus                                    *!! !! !  !!*!!!
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X non conserved

X similar

X ≥ 50% conserved