```
logo
NTDB id 339259 EKL02 RS03785 WP 128900801.1  MFSFFKRKPKETAAPAA.APVEEIVAEKA..........QEPVAPV.............PLSPAV........PLPAAAT   48
NTDB id 1118 NGFG RS11455 WP 003696286.1     MFSFFRRKKKQETPALEEAQVQETAAKVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEAG   80
consensus                                    !!!!!*!!*!   **  *!*! !  !   ********** !*! *****************!*!!***********!  !

logo
NTDB id 339259 EKL02 RS03785 WP 128900801.1  PPAPV.......IFPPPPEAVQEVEIIPAPAPSAEQKRSWLTRLKAGLSKTS....ASLTTLFVGAKIDDDLYEELESAL  117
NTDB id 1118 NGFG RS11455 WP 003696286.1     EAAERVESAKEAVAETVGEAVG..QVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEELETVL  158
consensus                                    **!* ******** ****!!! ** * *!*!** !*!  !  !!! !! !* **** !!  *! !* ! *!!!!!!!* !

logo
NTDB id 339259 EKL02 RS03785 WP 128900801.1  LMADAGIDATQFLLDTLKKKVKDEKLTEAEQVKTALKALLIGLLTPLQKPLVLGR.HQPMVMMIAGVNGAGKTTTIGKLA  196
NTDB id 1118 NGFG RS11455 WP 003696286.1     ITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTSIGKLA  238
consensus                                    * *! !**!! *!*  ** *!    ! **  ** !!!  !  !* !! !!!!!* ** ! !*!*!!*!!!!!!!*!!!!!

logo
NTDB id 339259 EKL02 RS03785 WP 128900801.1  KHLQAHHQSVLLAAGDTFRAAAREQLAVWGERNNVTVIAQESGDPAAVAFDAVQSALARGTDVVMVDTAGRLPTQLHLME  276
NTDB id 1118 NGFG RS11455 WP 003696286.1     KYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQLHLME  318
consensus                                    !* !!** !!!!!!!!!!!!!!!!!!  !! !!!!!!! ! *!!*!!!*!!!!! ! !!! !*!* !!!!!!!!!!!!!!

logo
NTDB id 339259 EKL02 RS03785 WP 128900801.1  ELKKVKRVIGKSMAEAPHEVLLVIDGNTGQNALTQVKAFDDALQLTGLVVTKLDGTAKGGVLAAIARTRPVPVYFIGVGE  356
NTDB id 1118 NGFG RS11455 WP 003696286.1     EIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRYIGVGE  398
consensus                                    !*!!!!!!* ! ** !!!!***!*!*! !!!!* !!!!!!!!! !!!!*!!!!!!!!!!!*!!!*!  !!!!! *!!!!!

logo
NTDB id 339259 EKL02 RS03785 WP 128900801.1  GIEDLQPFNAQEFVDALLS   375
NTDB id 1118 NGFG RS11455 WP 003696286.1     GIDDLRPFDARAFVDALLD   417
consensus                                    !!*!! !! !  !!!!!!
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X non conserved

X similar

X ≥ 50% conserved