```
logo

NTDB id 235522 CEG11 RS14845 WP 050560672.1  MKSIIERNIFNEHLMQLLLRFIEPKKNFSLAESAVLHYVIFDGESQETAVNLGAAIEILTLTSDIIDDLEDEDNRDAPWM   80
NTDB id 93 BSU 31710 NP 391049.2             MKEIVEQNIFNEDLSQLLYSFIDSKETFSFAESTILHYVVFGGENLDVATRLGAGIEILILSSDIMDDLEDEDNHHALWM   80
consensus                                    !! !*! !!!!!*! !!!  !!**!  !! !!! *!!!!*! !!  * !  !!!*!!!! !*!!!*!!!!!!!!**!*!!

logo

NTDB id 235522 CEG11 RS14845 WP 050560672.1  KIDRSLALSAAFSLYTAGLQSINSISTNPLVIKYVLKYIQDAMQGQHDDLINKPETEDECLEVIKLKCGSLFALANVTGT  160
NTDB id 93 BSU 31710 NP 391049.2             KINRSESLNAALSLYTVGLTSIYSLNNNPLIFKYVLKYVNEAMQGQHDDITNKSKTEDESLEVIRLKCGSLIALANVAGV  160
consensus                                    !! !!  ! !! !!!! !! !! !*  !!!* !!!!!!***!!!!!!!!* !!* !!!!*!!!!*!!!!!! !!!!! !

logo

NTDB id 235522 CEG11 RS14845 WP 050560672.1  LLATGEFNKSVESYSYYKGIFEQISGDYHAIFKDNTSDILKDKHTLINLYLKRLFNKSSQELLRIFSDQKLYSQLRLNKK  240
NTDB id 93 BSU 31710 NP 391049.2             LLATGEYNETVERYSYYKGIIAQISGDYYVLLSGNRSDIEKNKHTLIYLYLKRLFNDASEDLLYLISHKDLYYKSLLDKE  240
consensus                                    !!!!!!*! *!! !!!!!!!  !!!!!!* *   ! !!! ! !!!!! !!!!!!!!  ! *!! * !*  !!    ! !

logo

NTDB id 235522 CEG11 RS14845 WP 050560672.1  EFIQKLYDSGASQYVSVLYQLYMHKCGTEVENLNLDIKKIKIIKKHLLNNKNNKKDDEDAEFNKLFFELS  310
NTDB id 93 BSU 31710 NP 391049.2             KFQEKLIKAGVTQYISVLLEIYKQKCISAIEQLNLDKEKKELIKECLLSYTKGDTRCKT..........  299
consensus                                     !  !!   ! *!!*!!!  *! *!! * *!*!!!!  !  *!!  !!        *  ***********
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X non conserved

X similar

X ≥ 50% conserved