```
logo
NTDB id 340661 ETK69 RS16560 WP 269321599.1  MKTIIEQMCLDANQIKNRMKSIIERNIFNEHLMQLLLRFIEPKKNFSLAESAVLHYVIFDGENQETAVNLGAAIEILTLT   80
NTDB id 93 BSU 31710 NP 391049.2             ...................MKEIVEQNIFNEDLSQLLYSFIDSKETFSFAESTILHYVVFGGENLDVATRLGAGIEILILS   62
consensus                                    ******************!! !*! !!!!!*! !!!  !!**!  !! !!! *!!!!*! !!! * !  !!!*!!!! !*

logo
NTDB id 340661 ETK69 RS16560 WP 269321599.1  SDIIDDLEDEDNRDAPWMKIDRSLALSAAFSLYTAGLQSINSISTNPLVIKYVLKYIQDAMQGQHDDLINKPETEDECLE  160
NTDB id 93 BSU 31710 NP 391049.2             SDIMDDLEDEDNHHALWMKINRSESLNAALSLYTVGLTSIYSLNNNPLIFKYVLKYVNEAMQGQHDDITNKSKTEDESLE  142
consensus                                    !!!*!!!!!!!!**!*!!!! !!  ! !! !!!! !! !! !*  !!!* !!!!!!***!!!!!!!!* !!* !!!!*!!

logo
NTDB id 340661 ETK69 RS16560 WP 269321599.1  VIKLKCGSLFALANVTGTLLATGEFNKSVESYSYYKGIFEQISGDYHAIFKDNTSDILKDKHTLINLYLKRLFNKSSQEL  240
NTDB id 93 BSU 31710 NP 391049.2             VIRLKCGSLIALANVAGVLLATGEYNETVERYSYYKGIIAQISGDYYVLLSGNRSDIEKNKHTLIYLYLKRLFNDASEDL  222
consensus                                    !!*!!!!!! !!!!! ! !!!!!!*! *!! !!!!!!!  !!!!!!* *   ! !!! ! !!!!! !!!!!!!!  ! *!

logo
NTDB id 340661 ETK69 RS16560 WP 269321599.1  LRIFSDQKLYSQLRLNKKEFIQKLYDSGASQYVSVLYQLYMHKCGTEVENLNLDIKKIKIIKKHLLNNKNNKKDDEDAEF  320
NTDB id 93 BSU 31710 NP 391049.2             LYLISHKDLYYKSLLDKEKFQEKLIKAGVTQYISVLLEIYKQKCISAIEQLNLDKEKKELIKECLLSYTKGDTRCKT...  299
consensus                                    ! * !*  !!    ! !  !  !!   ! *!!*!!!  *! *!! * *!*!!!!  !  *!!  !!        *  ***

logo
NTDB id 340661 ETK69 RS16560 WP 269321599.1  NKLFFELS   328
NTDB id 93 BSU 31710 NP 391049.2             ........   299
consensus                                    ********
```

X non conserved

X similar

X $\geq$ 50% conserved