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logo

NTDB id 232328 S100141 RS00440 WP 269768362.1  MKTIIEQMCLDADQIKNRMKSIIEQNIFNEHLLQLLGFVESRKYLTFAESALFHYDVFGGKDFKTAEKLAAGIEILTLS   80
NTDB id 93 BSU 31710 NP 391049.2               ...................MKEIVEQNIFNEDLSQLLYSFIDSKETFSFAESTILHYVVFGGENLDVATRLGAGIEILILS   62
consensus                                      ******************!! !*!!!!!!!*! !!!  !**!*   *!!!! * !! !!!!     ! *!*!!!!!! !!

logo

NTDB id 232328 S100141 RS00440 WP 269768362.1  ADIIDDLEDEDNNQALWMKIDRSEALSAAFSLYTAGLQGINSINMNPLILTYILKYTQDAMQGQHDDIINKPEKEDESLE   160
NTDB id 93 BSU 31710 NP 391049.2               SDIMDDLEDEDNHHALWMKINRSESLNAALSLYTVGLTSIYSLNNNPLIFKYVLKYVNEAMQGQHDDITNKSKTEDESLE   142
consensus                                       !!*!!!!!!!!**!!!!!! !!! ! !! !!!! !!  ! !*! !!!!  !*!!! **!!!!!!!!! !!*  !!!!!!

logo

NTDB id 232328 S100141 RS00440 WP 269768362.1  VIRLKCGSLFALSNVAGTMLATGKFSETVETYSYYKGMFEQIYSDYLALFSSKRSDILKDKHTLINLYLKRLFNEPSEKL   240
NTDB id 93 BSU 31710 NP 391049.2               VIRLKCGSLIALANVAGVLLATGEYNETVERYSYYKGIIAQISGDYYVLLSGNRSDIEKNKHTLIYLYLKRLFNDASEDL   222
consensus                                      !!!!!!!!! !! !!!! *!!!! * !!!! !!!!!!*  !!  !!  ! !  !!!! ! !!!!! !!!!!!!!**!! !

logo

NTDB id 232328 S100141 RS00440 WP 269768362.1  VRMFSHRETHLKFISDKNIFIQTLTDAGVTQYVSVLHQLYKQNVKML..............................   287
NTDB id 93 BSU 31710 NP 391049.2               LYLISHKDLYYKSLLDKEKFQEKLIKAGVTQYISVLLEIYKQKCISAIEQLNLDKEKKELIKECLLSYTKGDTRCKT   299
consensus                                      * * !!** * ! * !!  !   !  !!!!!!*!!!* *!!! *   ******************************
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X non conserved

X similar

X ≥ 50% conserved