```
logo

NTDB id 311394 D0819 RS11725 WP 021480498.1  MKEIVKQNIFEDDLLKLLNLFIESRNFFPFAESAFLHYCVFGGKEFEIAIELGAGIEIITLSSDILDDIQDNDNLTAPWM   80
NTDB id 93 BSU 31710 NP 391049.2             MKEIVEQNIFNEDLSQLLYSFIDSKETFSFAESTILHYVVFGGENLDVATRLGAGIEILILSSDIMDDLEDEDNHHALWM   80
consensus                                    !!!!! !!!! *!!  !!  !!*!*  !*!!!!  !!!*!!!!   **!  !!!!!!!* !!!!!*!!* ! !!**!*!!

logo

NTDB id 311394 D0819 RS11725 WP 021480498.1  NIDQSVALSSVFSLYTVGLQSLKAVDPKSDIVNSILKYCLDAMQGQHRDIINKPQTEEDCLEAMKLKCGSLFALANVSGT  160
NTDB id 93 BSU 31710 NP 391049.2             KINRSESLNAALSLYTVGLTSIYSLNNNPLIFKYVLKYVNEAMQGQHDDITNKSKTEDESLEVIRLKCGSLIALANVAGV  160
consensus                                     !  !  !    !!!!!!! !*  * * * !   *!!!* *!!!!!! !! !!* !!***!! **!!!!!! !!!!! !

logo

NTDB id 311394 D0819 RS11725 WP 021480498.1  MLATGTYNKVVESYSYSKGIFEQIYGDYSALFNPNRSDILKDKHTLVNLYLKRLFNKDSEELLTIFKNKNFFYG.LLSKK  239
NTDB id 93 BSU 31710 NP 391049.2             LLATGEYNETVERYSYYKGIIAQISGDYYVLLSGNRSDIEKNKHTLIYLYLKRLFNDASEDLLYLISHKDLYYKSLLDKE  240
consensus                                    *!!!! !!  !! !!! !!!  !! !!!  !  *!!!!! ! !!!!* !!!!!!!!  !!*!! *  *!  *! *!! !

logo

NTDB id 311394 D0819 RS11725 WP 021480498.1  EFMEKLKKAGVTQYASVLHKIYKKKCETDIEKLNLDKHKIELIKKQLLK.........  288
NTDB id 93 BSU 31710 NP 391049.2             KFQEKLIKAGVTQYISVLLEIYKQKCISAIEQLNLDKEKKELIKECLLSYTKGDTRCKT  299
consensus                                     ! !!! !!!!!!! !!!* !!! !! * !! !!!!!*! !!!! *!! **********
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X non conserved

X similar

X ≥ 50% conserved