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logo

NTDB id 193650 BARD7 RS13820 WP 269465883.1  MKTIIEQMCLDADQIKNKMKSIIEQNIFNEHLLQLLFGFVESRKYLTFAESALLHYEVFGGKDFKTAEKLAAGIEILTLS   80
NTDB id 93 BSU 31710 NP 391049.2             ...................MKEIVEQNIFNEDLSQLLYSFIDSKETFSFAESTILHYVVFGGENLDVATRLGAGIEILILS   62
consensus                                    ******************!! !*!!!!!!!*! !!!* !**!*   *!!!! *!!! !!!!      ! *!*!!!!!! !!

logo

NTDB id 193650 BARD7 RS13820 WP 269465883.1  ADIIDDLEDEDNNQALWMKIDHSEALNAAFSLYTAGVQGINSINMNPLILTYILKYTQDAMQGQHDDIINKAEKEDESLE  160
NTDB id 93 BSU 31710 NP 391049.2             SDIMDDLEDEDNHHALWMKINRSESLNAALSLYTVGLTSIYSLNNNPLIFKYVLKYVNEAMQGQHDDITNKSKTEDESLE  142
consensus                                     !!*!!!!!!!!**!!!!!! *!! !!!! !!!! !*  ! !*! !!!!  !*!!! **!!!!!!!!! !!   !!!!!!

logo

NTDB id 193650 BARD7 RS13820 WP 269465883.1  VIRLKCGSLFALSNVAGTMLATGNYNGTVETYSYYKGMFEQIYSDYLALFSTKRSDILKEKHTLINLYLKRQFNEPSEKL  240
NTDB id 93 BSU 31710 NP 391049.2             VIRLKCGSLIALANVAGVLLATGEYNETVERYSYYKGIIAQISGDYYVLLSGNRSDIEKNKHTLIYLYLKRLFNDASEDL  222
consensus                                    !!!!!!!!! !! !!!! *!!!! !! !!! !!!!!!*  !!  !!  ! !  !!!! ! !!!!! !!!!! !!**!! !

logo

NTDB id 193650 BARD7 RS13820 WP 269465883.1  VRMFSHRETHLKFISDKNIFIQTLTDAGVTQYVSVLHQLFKQKCEKAIEGLNLENNKIDKLKKYVVNRGE.......  310
NTDB id 93 BSU 31710 NP 391049.2             LYLISHKDLYYKSLLDKEKFQEKLIKAGVTQYISVLLEIYKQKCISAIEQLNLDKEKKELIKECLLSYTKGDTRCKT  299
consensus                                    * * !!** * ! * !!  !   !  !!!!!!*!!!* **!!!!  !!! !!!*  ! * *! ***    *******
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X non conserved

X similar

X $\geq$ 50% conserved