```
logo
NTDB id 174460 A2G56 RS03815 WP 062709264.1  MAKPYYSKKSSKSAISLLAVILTLVLGLVAGNDKLPDDNPLKQLAQVVTNASNQEKARSNGSSANELTPSQELAQSVLTP   80
NTDB id 262 KZH43 RS08770 WP 001036779.1     .....MNKKTRQTLIGLLVLLL.LS....TGSYY......IKQM....PSAPNSPKTNLSQKKQASEAPSQALAESVLTD   60
NTDB id 221 SPD RS09380 WP 001036779.1       .....MNKKTRQTLIGLLVLLL.LS....TGSYY......IKQM....PSAPNSPKTNLSQKKQASEAPSQALAESVLTD   60
NTDB id 187 SPR RS08930 WP 001036779.1       .....MNKKTRQTLIGLLVLLL.LS....TGSYY......IKQM....PSAPNSPKTNLSQKKQASEAPSQALAESVLTD   60
NTDB id 152 SP RS09870 WP 001036779.1        .....MNKKTRQTLIGLLVLLL.LS....TGSYY......IKQM....PSAPNSPKTNLSQKKQASEAPSQALAESVLTD   60
consensus                                         **!!*****!*!!***! !*    *!***      *!!*    **!*!**!************!!!*!!*!!!!*

logo
NTDB id 174460 A2G56 RS03815 WP 062709264.1  SVKKLLGSNIKWNGAGAFIINDNKTNLDANVSSVPYVDNKTKQVQGEVVPTVANAMLSKQSRQYRDREATGNGSTSWKPA  160
NTDB id 262 KZH43 RS08770 WP 001036779.1     AVKSQIKGSLEWNGSGAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPP  140
NTDB id 221 SPD RS09380 WP 001036779.1       AVKSQIKGSLEWNGSGAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPP  140
NTDB id 187 SPR RS08930 WP 001036779.1       AVKSQIKGSLEWNGSGAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPP  140
NTDB id 152 SP RS09870 WP 001036779.1        AVKSQIKGSLEWNGSGAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPP  140
consensus                                    *!!********!!!*!!!!*!*!!!!!!!*!!!*!!*!!!!!*!**!*!!!!!!!*!!!**!!!**!**!!!!!!!!*!*

logo
NTDB id 174460 A2G56 RS03815 WP 062709264.1  GWHQVHGLPGGYDHAIDRGHLIGYALAGNIKGFDPSASNPKNIAVQTAWSNQARSHASTGQNYFETQIRRALDNNKRVRY  240
NTDB id 262 KZH43 RS08770 WP 001036779.1     GWHQVKNLKGSYTHAVDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRY  220
NTDB id 221 SPD RS09380 WP 001036779.1       GWHQVKNLKGSYTHAVDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRY  220
NTDB id 187 SPR RS08930 WP 001036779.1       GWHQVKNLKGSYTHAVDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRY  220
NTDB id 152 SP RS09870 WP 001036779.1        GWHQVKNLKGSYTHAVDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRY  220
consensus                                    !!!!!**!*!*!*!!*!!!!!*!!!!*!***!!!*!*!!!!!!!!!!!!!*!!****!!!!!!*!***!*!!!*!!!!!!

logo
NTDB id 174460 A2G56 RS03815 WP 062709264.1  RVTLVYDGD.NILASGSHLEAKSSDGSLEFNVFVPNVQSGITLNYRNGQVTVNNSFSD   297
NTDB id 262 KZH43 RS08770 WP 001036779.1     RVTLYYASNEDLVPSASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ....   274
NTDB id 221 SPD RS09380 WP 001036779.1       RVTLYYASNEDLVPSASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ....   274
NTDB id 187 SPR RS08930 WP 001036779.1       RVTLYYASNEDLVPSASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ....   274
NTDB id 152 SP RS09870 WP 001036779.1        RVTLYYASNEDLVPSASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ....   274
consensus                                    !!!!*!********!*!**!!!!!!!*!!!!!*!!!!!!*!**!*!!*!*!!!!**
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X non conserved

X similar

X ≥ 50% conserved