```
logo
NTDB id 1105954 V3C22 RS06875 WP 002264825.1  MTRKTNKSQRKQQKLCLSLLAALSVFVCTYIATSDHIADSNPIKQAAQLLTGKNKDILSNGSMSHQATPSQELANTVMTD   80
NTDB id 262 KZH43 RS08770 WP 001036779.1      MNKK.......TRQTLIGLLVLLLLSTGSYYIKQMPSAPNSPK............TNL.SQKKQASEAPSQALAESVLTD   60
NTDB id 221 SPD RS09380 WP 001036779.1        MNKK.......TRQTLIGLLVLLLLSTGSYYIKQMPSAPNSPK............TNL.SQKKQASEAPSQALAESVLTD   60
NTDB id 187 SPR RS08930 WP 001036779.1        MNKK.......TRQTLIGLLVLLLLSTGSYYIKQMPSAPNSPK............TNL.SQKKQASEAPSQALAESVLTD   60
NTDB id 152 SP RS09870 WP 001036779.1         MNKK.......TRQTLIGLLVLLLLSTGSYYIKQMPSAPNSPK............TNL.SQKKQASEAPSQALAESVLTD   60
consensus                                     !**!       *******!!**!******!*******!***!*           **! *********!!!*!!**!*!!

logo
NTDB id 1105954 V3C22 RS06875 WP 002264825.1  AVKNALGRQIEWSGAGSFIINHNKTDLNADVASQPYANNQTKIVQGQTVPTVANALLSKATRQYRNRQETGNGRTNWQPA  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 1105954 V3C22 RS06875 WP 002264825.1  GWHQLYDLPGDYDHAVDRGHLLAYALVGGLKRFDASTSNPQNVATQAAWANEANSSNSRGQNYYETLIRKALDSHKRVRY  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 1105954 V3C22 RS06875 WP 002264825.1  RVTLIYDDNNDILASGSHLEAKSSDGSLEFNVFIPNVQSGLTFDYATGQVKAN.  293
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