```
logo
NTDB id 241056 Sp14A RS07155 WP 115130480.1  MVKPYDVKKSSKSKFSVLVVIITLILGLVAGTDQLSDDNPLKQLAQVVTNGSNQEKVNSNGSSANESTPNQDLAQSVLTS   80
NTDB id 262 KZH43 RS08770 WP 001036779.1     ......MNKKTRQTLIGLLVLLLL....STGSYY......IKQM....PSAPNSPKTNLSQKKQASEAPSQALAESVLTD   60
NTDB id 221 SPD RS09380 WP 001036779.1       ......MNKKTRQTLIGLLVLLLL....STGSYY......IKQM....PSAPNSPKTNLSQKKQASEAPSQALAESVLTD   60
NTDB id 187 SPR RS08930 WP 001036779.1       ......MNKKTRQTLIGLLVLLLL....STGSYY......IKQM....PSAPNSPKTNLSQKKQASEAPSQALAESVLTD   60
NTDB id 152 SP RS09870 WP 001036779.1        ......MNKKTRQTLIGLLVLLLL....STGSYY......IKQM....PSAPNSPKTNLSQKKQASEAPSQALAESVLTD   60
consensus                                          **!********!*!***!    **!***      *!!*    ****!**!*!**********!*!*!!*!!!!*

logo
NTDB id 241056 Sp14A RS07155 WP 115130480.1  SVRKSLGDDIKWNGAGAFVINNNKTNLDASVSSMPYVDNKIKRVQGKLVPSVANAMLSKHSRQYRNRELTGNGSTSWKPA  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 241056 Sp14A RS07155 WP 115130480.1  GWHQVHDLPGEYDHAIDRGHLIGYAIAGNIDGFDPSASNPQNIAVQTAWSNQARDHDSTGQNYYETQIRRALDNNKRVRY  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 241056 Sp14A RS07155 WP 115130480.1  RVTLVYEGD.NIIASGSHLESKSSDGSLEFNVFVPNVQPGITLNYANGQVTVNH  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 $\geq$ 50% conserved