```
logo
NTDB id 1155450 QNH65 RS01690 WP 024417501.1  MAKKNVRKQAMSLLSLLATIVVVACGYLLSDNKSSNVADYSYYVNQSKASEGIPSQELASSVLTEDVKKQLGNSIEWNGA   80
NTDB id 262 KZH43 RS08770 WP 001036779.1      ....MNKKTRQTLIGLLVLLLLSTGSYYIKQMPSAPN.SPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKGSLEWNGS   75
NTDB id 221 SPD RS09380 WP 001036779.1        ....MNKKTRQTLIGLLVLLLLSTGSYYIKQMPSAPN.SPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKGSLEWNGS   75
NTDB id 187 SPR RS08930 WP 001036779.1        ....MNKKTRQTLIGLLVLLLLSTGSYYIKQMPSAPN.SPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKGSLEWNGS   75
NTDB id 152 SP RS09870 WP 001036779.1         ....MNKKTRQTLIGLLVLLLLSTGSYYIKQMPSAPN.SPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKGSLEWNGS   75
consensus                                         ***!****!**!!*******!*!******!*** *******!*!*****!!!*!!*!!!!**!!*!***!*!!!!*

logo
NTDB id 1155450 QNH65 RS01690 WP 024417501.1  GAFIINGNQTDLDASVASVPYGDNKTKTVQGETVPTVANALLAKSTRQYKDREETGNGRTSWTPAGWHQIHDLTGEYDHA  160
NTDB id 262 KZH43 RS08770 WP 001036779.1      GAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPPGWHQVKNLKGSYTHA  155
NTDB id 221 SPD RS09380 WP 001036779.1        GAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPPGWHQVKNLKGSYTHA  155
NTDB id 187 SPR RS08930 WP 001036779.1        GAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPPGWHQVKNLKGSYTHA  155
NTDB id 152 SP RS09870 WP 001036779.1         GAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPPGWHQVKNLKGSYTHA  155
consensus                                     !!!!*!!!*!*!!!*!*!*!!*!!!!!!!**!!!!!!!!!!!*!*!!!!!*!*!!!!!*!!!!!*!!!!***!*!*!*!!

logo
NTDB id 1155450 QNH65 RS01690 WP 024417501.1  VDRGHLLGYALIGGLSGFDASTSNPKNIAVQTAWSNQANDQTSTGQNYFESQIRRALDKNKRIRYRVTLVYAESEDLVPV  240
NTDB id 262 KZH43 RS08770 WP 001036779.1      VDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRYRVTLYYASNEDLVPS  235
NTDB id 221 SPD RS09380 WP 001036779.1        VDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRYRVTLYYASNEDLVPS  235
NTDB id 187 SPR RS08930 WP 001036779.1        VDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRYRVTLYYASNEDLVPS  235
NTDB id 152 SP RS09870 WP 001036779.1         VDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRYRVTLYYASNEDLVPS  235
consensus                                     !!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!*!!!****!!!!!!*!!**!*!!!*!!!*!!!!!!*!!**!!!!!*

logo
NTDB id 1155450 QNH65 RS01690 WP 024417501.1  GSHLEAKAADGSLEFNVFVPNVQSGLSINYHSGEIAVNQ  279
NTDB id 262 KZH43 RS08770 WP 001036779.1      ASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ  274
NTDB id 221 SPD RS09380 WP 001036779.1        ASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ  274
NTDB id 187 SPR RS08930 WP 001036779.1        ASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ  274
NTDB id 152 SP RS09870 WP 001036779.1         ASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ  274
consensus                                     *!**!!!**!!*!!!!!*!!!!!*!!***!**!!**!*!
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X non conserved

X similar

X ≥ 50% conserved