```
logo
NTDB id 177452 A4H00 RS07900 WP 067089091.1  MARKKTKKQSMSLASLVITFLLVAGGYFWAEGTSTVSHRP...VTRVENAASRPSEELAHSVLTDTVRAQLGSDLVYNGA   77
NTDB id 262 KZH43 RS08770 WP 001036779.1     ....MNKKTRQTLIGLLVLLLLSTGSYYIKQMP.SAPNSPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKGSLEWNGS   75
NTDB id 221 SPD RS09380 WP 001036779.1       ....MNKKTRQTLIGLLVLLLLSTGSYYIKQMP.SAPNSPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKGSLEWNGS   75
NTDB id 187 SPR RS08930 WP 001036779.1       ....MNKKTRQTLIGLLVLLLLSTGSYYIKQMP.SAPNSPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKGSLEWNGS   75
NTDB id 152 SP RS09870 WP 001036779.1        ....MNKKTRQTLIGLLVLLLLSTGSYYIKQMP.SAPNSPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKGSLEWNGS   75
consensus                                        **!!****!**!****!!**!*!****** *****!*********!***!!**!!*!!!!!*!**!****!**!!*

logo
NTDB id 177452 A4H00 RS07900 WP 067089091.1  GAFVMNGNRTDLDAAVASVPYADNKTKIAQGKVVPTVANALLSKATRQYQKREETGNGSTSWKPAGWHQVRHLEGEYNHA  157
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 177452 A4H00 RS07900 WP 067089091.1  IDRGHLLAYSLVGGLKGYDASTSNPKNIAVQTAWSNQSNRPDATGQNYFETQVRRALDRNKRVRYRVTLIYQSQEDLVPV  237
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 177452 A4H00 RS07900 WP 067089091.1  GSHLEAKAADGSLEFNVFVPNVQQGLVIDYYSGKIHLNQ  276
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