```
logo
NTDB id 1112226 GBSCOH1 RS04345 WP 000163017.1  MTKKQKQITASIVTLLFSLLAIGLGESKSLPNDHILKQVTDLIISNESGKNTSLKNISGTPSRELSESVLTSNVKKQLGT   80
NTDB id 262 KZH43 RS08770 WP 001036779.1        MNKKTRQTLIGLLV..LLLLSTGSYYIKQMPSAPN..........SPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKG   68
NTDB id 221 SPD RS09380 WP 001036779.1          MNKKTRQTLIGLLV..LLLLSTGSYYIKQMPSAPN..........SPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKG   68
NTDB id 187 SPR RS08930 WP 001036779.1          MNKKTRQTLIGLLV..LLLLSTGSYYIKQMPSAPN..........SPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKG   68
NTDB id 152 SP RS09870 WP 001036779.1           MNKKTRQTLIGLLV..LLLLSTGSYYIKQMPSAPN..........SPKTNLSQKKQASEAPSQALAESVLTDAVKSQIKG   68
consensus                                       !*!!**!*******  **!!**!****!**!****          *********!**!**!!**!*!!!!!**!!*!***

logo
NTDB id 1112226 GBSCOH1 RS04345 WP 000163017.1  NIAWNQSGAFIINQNKTDLNAKVSSAPYAINEIKKVNNQIVPTKANALLTKATRQYRNREETGNGRTYWKPAGWHQINGL  160
NTDB id 262 KZH43 RS08770 WP 001036779.1        SLEWNGSGAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPPGWHQVKNL  148
NTDB id 221 SPD RS09380 WP 001036779.1          SLEWNGSGAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPPGWHQVKNL  148
NTDB id 187 SPR RS08930 WP 001036779.1          SLEWNGSGAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPPGWHQVKNL  148
NTDB id 152 SP RS09870 WP 001036779.1           SLEWNGSGAFIVNGNKTNLDAKVSSKPYADNKTKTVGKETVPTVANALLSKATRQYKNRKETGNGSTSWTPPGWHQVKNL  148
consensus                                       ***!!*!!!!!*!*!!!*!*!!!!!*!!!*!**!*!****!!!*!!!!!*!!!!!!*!!*!!!!!*!*!*!*!!!!***!

logo
NTDB id 1112226 GBSCOH1 RS04345 WP 000163017.1  KGSYNHAVDRGHLIGYALVGSLRGFDASTSNPKNIATQAAWANQANSNQSTGQNYYETLVRKALDRHKTVRYRVTLIYDR  240
NTDB id 262 KZH43 RS08770 WP 001036779.1        KGSYTHAVDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRYRVTLYYAS  228
NTDB id 221 SPD RS09380 WP 001036779.1          KGSYTHAVDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRYRVTLYYAS  228
NTDB id 187 SPR RS08930 WP 001036779.1          KGSYTHAVDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRYRVTLYYAS  228
NTDB id 152 SP RS09870 WP 001036779.1           KGSYTHAVDRGHLLGYALIGGLDGFDASTSNPKNIAVQTAWANQAQAEYSTGQNYYESKVRKALDQNKRVRYRVTLYYAS  228
consensus                                       !!!!*!!!!!!!!*!!!!*!*!*!!!!!!!!!!!!!!*!*!!!!!!****!!!!!!!!**!!!!!!**!*!!!!!!!*!**

logo
NTDB id 1112226 GBSCOH1 RS04345 WP 000163017.1  .DNLLSSGSHIEAKSSDGSLEFNVFIPNVQSGLLFDYATGKVKQTK  285
NTDB id 262 KZH43 RS08770 WP 001036779.1        NEDLVPSASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ  274
NTDB id 221 SPD RS09380 WP 001036779.1          NEDLVPSASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ  274
NTDB id 187 SPR RS08930 WP 001036779.1          NEDLVPSASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ  274
NTDB id 152 SP RS09870 WP 001036779.1           NEDLVPSASQIEAKSSDGELEFNVLVPNVQKGLQLDYRTGEVTVTQ  274
consensus                                       ***!**!*!*!!!!!!!!!*!!!!!**!!!!*!!**!!*!!*!**!*
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X non conserved

X similar

X ≥ 50% conserved