```
logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFD.   79
NTDB id 316265 D4J00 RS00260 WP 139519086.1  MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1      MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
consensus                                     !!!**!********!**!***!* *!!*!!* *****!!!!*!*!!**!**!!****!******!*****!*!******

logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  ..PRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
NTDB id 316265 D4J00 RS00260 WP 139519086.1  QGLNIPFYGIAIDDKIKS.FYSHSAALLKGFCQGNLKATYGRLRANNAVSYAGKSLEFRTNNPRDFTFKGDENLDYFLLD   157
NTDB id 1221 C694 RS07590 WP 001203580.1      QGLNIPLYGIAIDDKIKS.FYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD   157
consensus                                     ********!!*!!!*** *!!!!!*!*********!*****!*!*!***!*!!!!!!! ***!******** ****!*!

logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  DIVTTGSSLLEAKKVLEENKISVLFALVLADAKV   191
NTDB id 316265 D4J00 RS00260 WP 139519086.1  DIITTGTTLKEALKYLKTLNIKTHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1      DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
consensus                                     !!*!!!**!*!!*!*!****!***!!**!**!**
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X non conserved

X similar

X ≥ 50% conserved