```
logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1      MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFD.   79
NTDB id 210685 HPYLPMSS1 RS07240 WP 077232506.1  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 210685 HPYLPMSS1 RS07240 WP 077232506.1  QGLNIPFYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRDFTFKGDESLDYFLLD   157
NTDB id 1221 C694 RS07590 WP 001203580.1          QGLNIPLYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD   157
consensus                                         **********!!*!!!** **!!!!!*!*********!!****!*!*!***!*!!!!!!!****!**************!*!

logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1      DIVTTGSSLLEAKKVLEENKISVLFALVLADAKV   191
NTDB id 210685 HPYLPMSS1 RS07240 WP 077232506.1  DIITTGTTLKEALKYLKALNIKAHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1          DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
consensus                                         !!*!!!**!*!!*!*!* **!***!!**!**!**
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X non conserved

X similar

X ≥ 50% conserved