```
logo                                                  MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE
                                                      MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE
NTDB id 35684 C695 RS07595 WP 001203580.1             MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1              MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1          MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFD.   79
consensus                                             !!!**!********!**!***!* *!!*!!* *****!!!!!*!*!!**!**!!****!******!*****!*!******

logo                                                  QGLNIPLYGIAIDDKIKS.FYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD
                                                      QGLNIPLYGIAIDDKIKS.FYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD
NTDB id 35684 C695 RS07595 WP 001203580.1             QGLNIPLYGIAIDDKIKS.FYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD   157
NTDB id 1221 C694 RS07590 WP 001203580.1              QGLNIPLYGIAIDDKIKS.FYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD   157
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1          ..PRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
consensus                                             *********!!*!!!*** *!!!!!*!*********!!****!*!*!***!*!!!!!!!****!!************!*!

logo                                                  DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE
                                                      DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE
NTDB id 35684 C695 RS07595 WP 001203580.1             DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1              DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1          DIVTTGSSLLEAKKVLEENKISVLFALVLADAKV   191
consensus                                             !!*!!!**!*!!*!*!****!*!*!!**!**!**
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X non conserved
X similar
X $\geq$ 50% conserved