```
logo

NTDB id 316285 D4I21 RS00280 WP 139542625.1   MRCLTCLKLSFKPLCKNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGKEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1      MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1 MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFF..   78
consensus                                     !!!**!********! *!***!* *!!*!!* *****!!!!*!*!!**!**!!****!******!*******!******

logo

NTDB id 316285 D4I21 RS00280 WP 139542625.1   QGLNIPFYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGRLRATNAVSYAGKSLEFRANNPRNFKFKGDESLDYFLLD   157
NTDB id 1221 C694 RS07590 WP 001203580.1      QGLNIPLYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD   157
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1 .DPRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
consensus                                     ********!!*!!!** **!!!!!*!*********!!****!*!*!***!*!!!!!!!****!!************!*!

logo

NTDB id 316285 D4I21 RS00280 WP 139542625.1   DIITTGTTLKEALKYLKTLNIKTHFAIALCSADE   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