```
logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFF..   78
NTDB id 251208 CNX73 RS00280 WP 001203595.1  MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGKEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1     MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
consensus                                    !!!**!********!**!***!* *!!*!!* *****!!!!*!*!!**!**!!****!******!*******!******

logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  .DPRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
NTDB id 251208 CNX73 RS00280 WP 001203595.1  QGLNIPFYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGTLRATNAVSYAGKSLEFRANNPRDFTFKGDENLDYFLLD   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 251208 CNX73 RS00280 WP 001203595.1  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