```
logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFDP   80
NTDB id 239201 CGC32 RS00270 WP 089482153.1  MRCLTCLKLSFKPLCKNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYVLIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1     MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
consensus                                    !!!**!********! *!***!* *!!*!!* *****!!!!*!*!!**!**!!* **!******!*****!*!******

logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  ...RMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
NTDB id 239201 CGC32 RS00270 WP 089482153.1  KGLTTPLYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGCLRATNAVSYAGKSLEFRANNPRNFTFKGDETLDYFLLD   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 239201 CGC32 RS00270 WP 089482153.1  DIITTGTTLKEALKYLKTLNIKAHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1     DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
consensus                                    !!*!!!**!*!!*!*!****!***!!**!**!**
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X non conserved

X similar

X ≥ 50% conserved