```
logo

NTDB id 421679 G3M68 RS01125 WP 163497690.1   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

NTDB id 421679 G3M68 RS01125 WP 163497690.1   KGLNIPLYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGRLRAMNTIKYAGKSLEFRTNHPRDFTFKGDKNLDYFLLD   157
NTDB id 1221 C694 RS07590 WP 001203580.1      QGLNIPLYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD   157
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1 ..PRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
consensus                                      ********!!*!!!** **!!!!!*!*********!!****!*!*! **!*!!!!!!! ***!*******  ****!*!

logo

NTDB id 421679 G3M68 RS01125 WP 163497690.1   DIITTGTTIKEALKYLKALNIKAHFAIALCSADE   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