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logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFD.   79
NTDB id 329349 EAL91 RS01790 WP 121053967.1  MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1     MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
consensus                                    !!!**!********!**!***!* *!!*!!* *****!!!!*!*!!**!**!****!******!*****!*!******

logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  ..PRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
NTDB id 329349 EAL91 RS01790 WP 121053967.1  QGLNIPLYGIAIDDKIK.SFYSHAAVLLKGFCQGNLKPTYGTLRATNTVSYAGKSLEFRTNNPRNFTFKGDENLDYFLLD   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 329349 EAL91 RS01790 WP 121053967.1  DIITTGTTLKEALKYLKTLNTKVHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1     DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
consensus                                    !!*!!!**!*!!*!*!******!*!!**!**!**
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X non conserved

X similar

X ≥ 50% conserved