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logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1   MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFD.   79
NTDB id 320137 D4K20 RS00280 WP 139544739.1    MRCLTCLKLSFKPLCPNCLNDLP.LNLKVR.VLESVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1       MRCLTCLKLSFKPLCPNCLNDLP.LSLKVR.VLEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
consensus                                      !!!**!********!**!***!* **!*!! *** **!!!!!*!*!!**!**!!****!******!*****!*!******

logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1   ..PRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
NTDB id 320137 D4K20 RS00280 WP 139544739.1    QGLNIPFYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRTNNPRNFKFKGDKNLDYFLLD   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 320137 D4K20 RS00280 WP 139544739.1    DIITTGTTLKEALKYLKTLNIKTHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1       DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
consensus                                      !!*!!!**!*!!*!*!****!***!!**!**!**
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X non conserved

X similar

X ≥ 50% conserved