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logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1   MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFF..   78
NTDB id 320326 D4H18 RS00275 WP 139522392.1   MRCLTCLKLSFKPLCPNCLNDLP.LSLRIR.VLEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGKEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1      MRCLTCLKLSFKPLCPNCLNDLP.LSLKVR.VLEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
consensus                                     !!!**!********!**!***!* *!!**!  ******!!!!!*!*!!**!**!!****!******!*******!******

logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1   .DPRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
NTDB id 320326 D4H18 RS00275 WP 139522392.1   QGLNIPFYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGRLRANNVVSYAGKSLEFRANNPRNFTFKGDENLDYFLLD   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 320326 D4H18 RS00275 WP 139522392.1   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