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logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1   MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFDP   80
NTDB id 667317 MPG57 RS00265 WP 245081158.1   MRCLTCLKLSFKPLCSNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEVEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   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 667317 MPG57 RS00265 WP 245081158.1   KGLNTPLYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGRLRANNTISYAGKSLEFRANNPRNFTFKGDETLDYFLLD   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 667317 MPG57 RS00265 WP 245081158.1   DIITTGTTLKEALKYLKTLNIKAHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1      DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
consensus                                     !!*!!!**!*!!*!*!****!***!!**!**!**
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X non conserved

X similar

X $\geq$ 50% conserved