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logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1   MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFF..   78
NTDB id 421671 G3M69 RS04460 WP 163563095.1   MRCLTCLKLSFRPLCPNCLNDLP.LSLRVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGVEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1      MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
consensus                                     !!!**!********!**!***!* *!!*!!* *****!!!!!*!*!!**!**!!****!******!*******!******

logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1   .DPRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKEKRNYKLLKTIHEPVILVD   157
NTDB id 421671 G3M69 RS04460 WP 163563095.1   KGLNIPLYGIAIDDKIKS.FYSHSAALLKGFCQGNLKPTYGTLRATNAVSYAGKSLEFRANHPRDFTFKGDKNLDYFLLD   157
NTDB id 1221 C694 RS07590 WP 001203580.1      QGLNIPLYGIAIDDKIKS.FYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD   157
consensus                                      ********!!*!!!*** *!!!!!*!*********!!*** !*!*!***!*!!!!!!!****!*******  ****!*!

logo

NTDB id 1233 CJJ81176 RS04995 WP 002853564.1   DIVTTGSSLLEAKKVLEENKISVLFALVLADAKV   191
NTDB id 421671 G3M69 RS04460 WP 163563095.1   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