```
logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFF..   78
NTDB id 38590 HPSJM RS07255 WP 001203654.1    MRCLTCLKLSFKPLCSNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGKEFVKILQE   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 38590 HPSJM RS07255 WP 001203654.1    QGLNIPFYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGTLRATNAVSYAGKSLEFRANHPRDFTFKGDKNLDYFLLD   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 38590 HPSJM RS07255 WP 001203654.1    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