```
logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1   MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFF..   78
NTDB id 1123745 EL256 RS00300 WP 126442292.1   MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGKEFVRILQE   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 1123745 EL256 RS00300 WP 126442292.1   QGLNIPFYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGTLRATNAVSYAGKSLEFRTNHPRDFTFKGDENLDYFLLD   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 1123745 EL256 RS00300 WP 126442292.1   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