```
logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFDP   80
NTDB id 247200 C0052 RS07225 WP 096479194.1  MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1     MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
consensus                                    !!!**!********!**!***!* *!!*!!* *****!!!!*!*!!**!**!!****!******!*****!*!******
```

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logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  ...RMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
NTDB id 247200 C0052 RS07225 WP 096479194.1  KGLNAPLYGIAIDDKVKS.FYSHSAALLKGFCQGNLKPTYGRLRAINAVSYAGKSLEFRANNPRDFTFKGDKDLDYFLLD   157
NTDB id 1221 C694 RS07590 WP 001203580.1     QGLNIPLYGIAIDDKIKS.FYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD   157
consensus                                     *********!!*!!!*** *!!!!!*!*********!!****!*!*!***!*!!!!!!!****!*******  ****!*!
```

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logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  DIVTTGSSLLEAKKVLEENKISVLFALVLADAKV   191
NTDB id 247200 C0052 RS07225 WP 096479194.1  DIITTGTTLKEALKYLKALNIKAHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1     DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
consensus                                    !!*!!!**!*!!*!*!* **!***!!**!**!**
```

X non conserved

X similar

X $\geq$ 50% conserved