```
logo
NTDB id 56633 HPOK113 RS07155 WP 015428359.1  MRCLTCLKLSFKPLCLNCLNDL.PLSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1      MRCLTCLKLSFKPLCPNCLNDL.PLSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFDP   80
consensus                                     !!!**!********! *!***! **!!*!!* *****!!!!*!*!!**!**!!****!******!*****!*!******
```

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logo
NTDB id 56633 HPOK113 RS07155 WP 015428359.1  KGLNAPLYGIAIDDKVKS.FYSHSAALLKGFCQGNLKPTYGRLRASNAVSYAGKSLEFRANNPRNFTFKGDKNLDYFLLD   157
NTDB id 1221 C694 RS07590 WP 001203580.1      QGLNIPLYGIAIDDKIKS.FYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD   157
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  ...RMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
consensus                                      *********!!*!!!*** *!!!!!*!*********!!****!*!*!***!*!!!!!!!****!!******  ****!*!
```

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logo
NTDB id 56633 HPOK113 RS07155 WP 015428359.1  DIITTGTTLKEALKYLKTLNIKAHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1      DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  DIVTTGSSLLEAKKVLEENKISVLFALVLADAKV   191
consensus                                     !!*!!!**!*!!*!*!****!***!!**!**!**
```

X non conserved

X similar

X $\geq$ 50% conserved