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logo
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```
NTDB id 36661 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
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```
NTDB id 36661 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
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```
NTDB id 36661 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 ≥ 50% conserved