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logo
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NTDB id 85019 RC16 RS04865 WP 002870583.1    MRCINCGAFALLCFCDLCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFD.    79
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1 MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFD.    79
NTDB id 1221 C694 RS07590 WP 001203580.1     MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE    78
consensus                                    !!!**!********!*!***!***!!*!!*******!!!!!*!*!!**!**!!****!******!*****!*!*****
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logo
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NTDB id 85019 RC16 RS04865 WP 002870583.1    ..PRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD    157
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1 ..PRMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD    157
NTDB id 1221 C694 RS07590 WP 001203580.1     QGLNIPLYGIATDDKIKS.FYSHSAALLKGFCQGNLKPTYGRLRANNAVSYAGKSLEFRANNPRNFTFKGDESLDYFLLD    157
consensus                                      *******!!*!!!*****!!!!!*!*********!!****!*!*!***!*!!!!!!!****!!************!*!
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logo
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NTDB id 85019 RC16 RS04865 WP 002870583.1    DIVTTGSSLLEAKKVLEENKISVLFALVLADAKV    191
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1 DIVTTGSSLLEAKKVLEENKISVLFALVLADAKV    191
NTDB id 1221 C694 RS07590 WP 001203580.1     DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE    191
consensus                                    !!*!!!**!*!!*!*!****!*!*!!**!**!**
```

X non conserved
X similar
X ≥ 50% conserved