```
logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  MRCINCGAFALLCFCELCELELSEFSLNVRKLDNNFKVYSFYKYHEIQHLLHSKHYFYGYFVYKMLAKLSFAKFKDFFDP   80
NTDB id 146246 AA976 RS07300 WP 033746068.1   MRCLTCLKLSFKPLCPNCLNDLP.LSLRVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVRILQE   78
NTDB id 1221 C694 RS07590 WP 001203580.1      MRCLTCLKLSFKPLCPNCLNDLP.LSLKVRV.LEGVSVYSFYAYSEIEELIKSKYALIGSRILPLLSQKAGAEFVKILQE   78
consensus                                     !!!**!********!**!***!* *!!*!!* *****!!!!*!*!!**!**!!****!******!*****!*!******

logo
NTDB id 1233 CJJ81176 RS04995 WP 002853564.1  ...RMTINVIALDDKVEDMLYSHSAILARYLKTKFVKPVFNVLKAQNHIKYSGKSLEFRQKHKRNYKLLKTIHEPVILVD   157
NTDB id 146246 AA976 RS07300 WP 033746068.1   KGLNTPLYGIAIDDKIK.SFYSHSAALLKGFCQGNLKPTYGTLRATNTISYAGKSLEFRANNPRDFTFKGDKNLDYFLLD   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 146246 AA976 RS07300 WP 033746068.1   DIITTGTTLKEALKHLKALNIKAHFAIALCSADE   191
NTDB id 1221 C694 RS07590 WP 001203580.1      DIITTGTTLKEALKYLKTLNIKVHFAIALCSADE   191
consensus                                     !!*!!!**!*!!*! !* **!***!!**!**!**
```

X non conserved

X similar

X $\geq$ 50% conserved