

logo

NTDB id 1088295 ACLII2 RS08765 WP 004399124.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT
MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT
MDSIEKVSKNLIEEAYLTKASDIHIVPRERDAIHFRVDHALLKKRDMKKEECVRLISHFKFLSAMDIGERRKPQNGSLT
!!

80
80

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

LKLKEGNVHLMSTLPTINEESLVRMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQY
LKLKEGNVHLMSTLPTINEESLVRMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQY
LKLKEGNVHLMSTLPTINEESLVRMPQYNIPSIDKLSLFPKTGATLLSFLKHSHGMLIFTGPTGSGKTTTLYSLVQY
!!

160
160

logo

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consensus

AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAETAVRAAMTGHVLVLSLH
AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAETAVRAAMTGHVLVLSLH
AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAILRHDPDMIILGEIRDAETAETAVRAAMTGHVLVLSLH
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240
240

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consensus

TRDAKGAIYRLLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN
TRDAKGAIYRLLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN
TRDAKGAIYRLLLEFGINMNEIEQTVIAIAAQRLVDLACPFCENGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN
!!

320
320

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consensus

HANYQYQTLRQIIRKGIALGYLTTNMYDRWVYHEKD
HANYQYQTLRQIIRKGIALGYLTTNMYDRWVYHEKD
HANYQYQTLRQIIRKGIALGYLTTNMYDRWVYHEKD
!!

356
356

- non conserved
- similar
- ≥ 50% conserved