

logo

NTDB id 1016392 ABYM97 RS16050 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

AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAAILRHDPDMIILGEIRDAETAETAVRAAMTGHVLVLTSLH
AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAAILRHDPDMIILGEIRDAETAETAVRAAMTGHVLVLTSLH
AKKHFNRNIVTLEDPVETRDEDVLQVQVNEKAGVTYSAGLKAAILRHDPDMIILGEIRDAETAETAVRAAMTGHVLVLTSLH
!!

240
240

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consensus

TRDAKGAIYRLLLEFGINMNEIEQTVIAIAAQRLVDLACPFCEGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN
TRDAKGAIYRLLLEFGINMNEIEQTVIAIAAQRLVDLACPFCEGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN
TRDAKGAIYRLLLEFGINMNEIEQTVIAIAAQRLVDLACPFCEGCSSVYCRQSRNTRRASVYELLYGKNLQQCIQEAKGN
!!

320
320

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consensus

HANYQYQTLRQIIRKGIALGYLTTNMYDRWVYHEKD
HANYQYQTLRQIIRKGIALGYLTTNMYDRWVYHEKD
HANYQYQTLRQIIRKGIALGYLTTNMYDRWVYHEKD
!!

356
356

non conserved
 similar
 ≥ 50% conserved