

***Clostridium difficile*, Isolate 20110979**

**Catalog No. NR-49285**

**Product Description:** *Clostridium difficile* (*C. difficile*; also referred to as *Peptoclostridium difficile*), isolate 20110979 was obtained from the stool of an elderly adult female patient with a community-associated (CA) *C. difficile* infection in midwestern USA in 2011. Isolate 20110979 was deposited as PCR ribotype 027, North American pulsed-field gel electrophoresis type 1 (NAP1), containing *tcdA*, *tcdB* and *tcdC* (with 18 base pair deletion) of the PaLoc operon as well as the *C. difficile* binary toxin (CDT).

**Lot<sup>1</sup>: 63719790**

**Manufacturing Date: 12SEP2015**

| TEST  | SPECIFICATIONS   | RESULTS  |
|---|--|--|
| <b>Phenotypic Analysis</b><br>Cellular morphology<br>Colony morphology <sup>2</sup><br><br>Hemolysis <sup>2</sup><br>Motility (wet mount)<br>Biochemical tests:<br>Esculin hydrolysis <sup>3</sup><br>Gelatin hydrolysis <sup>3</sup><br>VITEK <sup>®</sup> MS (MALDI-TOF)          | Gram-positive rods<br>Report results<br><br>Non-hemolytic<br>Report results<br><br>Positive<br>Positive<br>Consistent with <i>C. difficile</i> | Gram-positive rods<br>Irregular, slight peaked, undulate, opaque, rough, and gray (Figure 1)<br>Non-hemolytic<br>Motile<br><br>Positive<br>Positive<br><i>C. difficile</i> (99.9%) |
| <b>Genotypic Analysis</b><br>Sequencing of 16S ribosomal RNA gene (~ 1390 base pairs)   | > 99% sequence identity to <i>C. difficile</i> type strain   | 99.9% sequence identity to CP011968.1  |
| <b>PCR Assay of Extracted DNA</b><br>Presence of <i>C. difficile</i> -specific genes <sup>4</sup><br>Triose phosphate isomerase ( <i>tpi</i> )<br>Presence of toxin genes <sup>4,5</sup><br><i>cdtB</i><br><i>tcdA</i> (wild type)<br><i>tcdA</i> (partial deletion)<br><i>tcdB</i> | ~ 230 base pair amplicon<br><br>~ 510 base pair amplicon<br>~ 370 base pair amplicon<br>No amplicon<br>~ 160 base pair amplicon                | ~ 230 base pair amplicon<br><br>~ 510 base pair amplicon<br>~ 370 base pair amplicon<br>No amplicon<br>~ 160 base pair amplicon  |
| <b>Purity (post-freeze)</b><br>Anaerobic growth <sup>6</sup><br><br>Aerobic growth <sup>7</sup>   | Growth consistent with expected morphology<br>No growth  | Growth consistent with expected morphology<br>No growth  |
| <b>Viability (post-freeze)<sup>2</sup></b>  | Growth   | Growth   |

<sup>1</sup>NR-49285 was produced by inoculation of the deposited material into Modified Reinforced Clostridial medium and incubated for 1 day at 37°C in an anaerobic atmosphere (< 0.5% O<sub>2</sub>; Remel™ Anaero Pack-Anaero™ R681001). The material from the initial growth was passaged in Modified Reinforced Clostridial medium for 1 day at 37°C in an anaerobic atmosphere to produce this lot.

<sup>2</sup>1 day at 37°C in an anaerobic atmosphere on Tryptic Soy agar with 5% defibrinated sheep blood

<sup>3</sup>Tests were assessed after 7 days at 37°C in an anaerobic atmosphere. The gelatin tube was placed at 4°C for one hour prior to result determination.

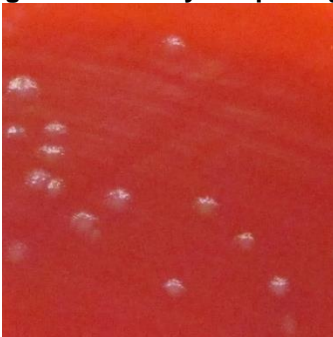
<sup>4</sup>Lemee, L., et al. "Multiplex PCR Targeting *tpi* (Triose Phosphate Isomerase), *tcdA* (Toxin A), and *tcdB* (Toxin B) Genes for Toxigenic Culture of *Clostridium difficile*." *J. Clin. Microbiol.* 42 (2004): 5710-5714. PubMed: 15583303.

<sup>5</sup>Antikainen, J., et al. "Detection of Virulence Genes of *Clostridium difficile* by Multiplex PCR." *APMIS*. 117 (2009): 607-613. PubMed: 19664132.

<sup>6</sup>Purity of this lot was assessed for 10 days at 37°C in an anaerobic atmosphere on Tryptic Soy agar with 5% defibrinated sheep blood.

<sup>7</sup>Purity of this lot was assessed for 10 days at 37°C in an aerobic atmosphere with 5% CO<sub>2</sub> on Tryptic Soy agar with 5% defibrinated sheep blood.

Figure 1: Colony Morphology



Date: 23 FEB 2016

Signature:

A handwritten signature in black ink, appearing to read "David C. Archer".

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