

***Helicobacter* sp., Strain Hp CPY6081 (deposited as *Helicobacter pylori*, Strain Hp CPY6081)**

**Catalog No. NR-43639**

**Product Description:**

*Helicobacter* sp., strain Hp CPY6081 (also referred to as CPY6081) was isolated from gastric biopsy homogenate from a gastric cancer patient in Yamaguchi Prefecture, Japan. NR-43639 was deposited to BEI Resources as *Helicobacter pylori*; however, digital DNA-DNA hybridization (dDDH) analysis performed at BEI Resources resulted in reclassification to *Helicobacter* sp. NR-43639 was produced by inoculation of BEI Resources seed lot 63734557 into Brucella broth, which was used to inoculate a Columbia agar with 7% defibrinated horse blood, 5 µg/mL trimethoprim, 5 µg/mL vancomycin, 10 µg/mL cefsulodin and 2.5 µg/mL amphotericin B plate, and both were grown for 3 days at 37°C in a microaerophilic atmosphere (~ 6-16% O<sub>2</sub> and 2-10% CO<sub>2</sub>). Colonies from the plate were suspended in Brucella broth and the growth mixture was added to Columbia agar with 7% defibrinated horse blood, 5 µg/mL trimethoprim, 5 µg/mL vancomycin, 10 µg/mL cefsulodin and 2.5 µg/mL amphotericin B kolles, which were grown for 3 days at 37°C in a microaerophilic atmosphere to produce this lot. Quality control testing was completed under propagation conditions unless otherwise noted.

**Lot: 70029280**

**Manufacturing Date: 03OCT2019**

TEST	SPECIFICATIONS	RESULTS
<b>Phenotypic Analysis</b> Cellular morphology Colony morphology  Motility (wet mount) Analytical profile index (API® CAMPY)	Gram-negative rods Report results  Report results <i>H. pylori</i> (≥ 90%)	Gram-negative rods Circular, low convex, entire, smooth and gray (Figure 1)  Motile <i>H. pylori</i> (99.9%)
<b>Genotypic Analysis</b> Sequencing of 16S ribosomal RNA (rRNA) gene (~ 1330 base pairs)	≥ 99% sequence identity to <i>H. pylori</i> , strain Hp CPY6081 (GenBank: AKNN01000009.1)	100% sequence identity to <i>H. pylori</i> , strain Hp CPY6081 (GenBank: AKNN01000009.1)
<b>Confirmation of <i>H. pylori</i> by PCR amplification of Extracted DNA</b> Positive control (16S rRNA gene) Negative control ( <i>H. acinonychis</i> ) <i>ureA</i> <i>ssaA</i>	Amplicon present No amplicon present Amplicon present Amplicon present	Amplicon present No amplicon present Amplicon present Amplicon present
<b>Purity (post-freeze)</b> Microaerophilic growth <sup>1</sup>  Aerobic growth <sup>2,3</sup>	Consistent with expected colony morphology Consistent with expected colony morphology	Consistent with expected colony morphology Consistent with expected colony morphology
<b>Viability (post-freeze)<sup>2</sup></b>	Growth	Growth

<sup>1</sup>Purity of this lot was assessed for 7 days on Columbia agar with 7% defibrinated horse blood, 5 µg/mL trimethoprim, 5 µg/mL vancomycin, 10 µg/mL cefsulodin and 2.5 µg/mL amphotericin B at 37°C in a microaerophilic atmosphere.

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

<sup>3</sup>*H. pylori* is known to show weak growth under aerobic conditions (Bury-Moné, S., et al. "Is *Helicobacter pylori* a True Microaerophile?" *Helicobacter* 11 (2006): 296-303. PubMed: 16882333.).

Figure 1: Colony Morphology



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