SUPPORTING INFECTIOUS DISEASE RESEARCH

### Helicobacter pylori, Strain CPY6081

#### Catalog No. NR-43639

#### **Product Description:**

*Helicobacter pylori* (*H. pylori*), strain CPY6081 was isolated from the gastric biopsy homogenate from a gastric cancer patient in Yamaguchi Prefecture, Japan. The strain designation on the vial label is incorrect. The correct strain designation is CPY6081. NR-43639 was produced by the inoculation of the BEI Resources seed lot 63734557 into Brucella broth and grown for 3 days at 37°C in a microaerophilic atmosphere (~ 6-16% O<sub>2</sub> and 2-10% CO<sub>2</sub>). Broth inoculum was added to Columbia agar with 7% defibrinated horse blood, 5  $\mu$ g/mL trimethoprim, 5  $\mu$ g/mL vancomycin, 10  $\mu$ g/mL cefsulodin and 2.5  $\mu$ g/mL amphotericin B kolles, which were grown for 3 days at 37°C in a microaerophilic atmosphere to produce this lot.

#### Lot: 70053916

## Manufacturing Date: 14JUL2022

| TEST   | SPECIFICATIONS  | RESULTS   |
|--|---|---|
| Phenotypic Analysis  |   |   |
| Cellular morphology  | Gram-negative rods  | Gram-negative rods  |
| Colony morphology  | Report results  | Circular, convex, entire, smooth and gray   |
| Motility (wet mount)   | Report results  | Motile  |
| VITEK <sup>®</sup> MS (MALDI-TOF)                                      | H. pylori   | H. pylori (99.9%)   |
| Genotypic Analysis   |   |   |
| Sequencing of 16S ribosomal RNA gene<br>(~ 1400 base pairs)            | ≥ 99% sequence identity to<br><i>H. pylori,</i> strain CPY6081<br>(GenBank: AKNN01000009.1) | 99.9% sequence identity to<br><i>H. pylori,</i> strain CPY6081<br>(GenBank: AKNN01000009.1)       |
| Digital DNA-DNA hybridization (dDDH) <sup>1</sup>                      | > 70% dDDH value for identity to<br>Helicobacter pylori type species                        | < 70% dDDH value for identity to<br>any <i>Helicobacter pylori</i> type<br>species <sup>1,2</sup> |
| Purity (post-freeze)   |   |   |
| Microaerophilic  | Growth consistent with expected   | Growth consistent with expected   |
| 7 days at 37°C on Tryptic Soy agar with 5%<br>defibrinated sheep blood | colony morphology   | colony morphology   |
| Aerobic with 5% CO <sub>2</sub>  | Growth consistent with expected   | Growth consistent with expected   |
| 7 days at 37°C on Tryptic Soy agar with 5%<br>defibrinated sheep blood | colony morphology   | colony morphology <sup>3</sup>  |
| Viability (post-freeze)  | Growth  | Growth  |

<sup>1</sup>Relatedness between bacterial strains has traditionally been determined using DDH. For additional information, refer to Auch, A. F., et al. "Digital DNA-DNA Hybridization for Microbial Species Delineation by Means of Genome-to-Genome Sequence Comparison." <u>Stand. Genomic Sci.</u> 2 (2010):117-134. PubMed: 21304684. dDDH analysis was performed using the Type (Strain) Genome Server.

<sup>2</sup>TYGS was unable to identify a matching type strain for this item. The closest match is to *Mangrovihabitans endophyticus* CGMCC 4.7299 with a dDDH value of 69.1. This suggests NR-43639 may belong to a novel species closely related to *Mangrovihabitans endophyticus* likely in the genus *Mangrovihabitans*.

<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?" <u>Helicobacter</u> 11 (2006): 296-303. PubMed: 16882333)

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# **Certificate of Analysis for NR-43639**

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## /Sonia Bjorum Brower/

Sonia Bjorum Brower

Technical Manager or designee, ATCC Federal Solutions

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