Overview

Useful For
Aiding in the diagnosis of *Helicobacter pylori* infection and prediction of clarithromycin resistance or susceptibility directly from stool

Testing Algorithm
For more information see *Helicobacter pylori* Diagnostic Algorithm.

Special Instructions

- [Helicobacter pylori Diagnostic Algorithm]

Highlights
This test detects the *Helicobacter pylori* 23S ribosomal RNA gene and the 3 most common 23S ribosomal RNA gene single nucleotide variations (A2143G, A2142G, and A2142C) that lead to resistance to clarithromycin.

Method Name
Real-Time Polymerase Chain Reaction (PCR)

NY State Available
Yes

Specimen

Specimen Type
Fecal

Ordering Guidance
Confirmation of eradication testing should not be ordered until 4 or more weeks after cessation of treatment.

Specimen Required
The high sensitivity of amplification by polymerase chain reaction requires the specimen to be processed in an environment in which contamination of the specimen by *Helicobacter pylori* DNA is unlikely.

Patient Preparation: Proton pump inhibitors and bismuth compounds should be avoided for 2 weeks prior to testing; antibiotics should be avoided for 4 weeks prior to testing.

Supplies: Culture and Sensitivity Stool Transport Vial (T058)

Specimen Type: Preserved feces

Submission Container/Tube: Commercially available transport system specific for recovery of enteric pathogens from...
Test Definition: HPFRP
Helicobacter pylori with Clarithromycin Resistance Prediction, Molecular Detection, PCR, Feces

fecal specimens (15 mL of nonnutritive transport medium containing phenol red as a pH indicator, either Cary-Blair or Para-Pak C and S)

**Specimen Volume:** 5 mL

**Collection Instructions:**
1. Collect fresh fecal specimen and submit 1 gram or 5 mL in container with transport medium.
2. Place feces in preservative within 2 hours of collection.
3. Place vial in a sealed plastic bag and send ambient or refrigerated. **Specimens sent frozen will be rejected.**

**Forms**
If not ordering electronically, complete, print, and send one of the following with the specimen:
1. Microbiology Test Request (T244)
2. Gastroenterology and Hepatology Client Test Request (T728)

**Specimen Minimum Volume**
1 mL

**Reject Due To**

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<tr>
<th>Fecal swab</th>
<th>ESwab</th>
<th>transport medium Feces in gel transport medium</th>
<th>ECOFIX preservative formalin PVA fixative Unpreserved stool</th>
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**Specimen Stability Information**

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<td>Fecal</td>
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<tr>
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**Clinical & Interpretive**

**Clinical Information**

*Helicobacter pylori* is the main cause of peptic ulcer disease and, when left untreated, a risk factor for gastric cancer. Traditionally, *H pylori* diagnosis has included noninvasive tests (eg, urea breath test, fecal antigen test) or invasive tests...
Antimicrobial resistance in *H. pylori* is poorly studied but is rising, challenging its treatment; in 2012, an international clinical consortium study group recommended monitoring of clarithromycin resistance rates and ceasing its use at a threshold range of 15% to 20%. Local monitoring has been practically impossible as not all patients undergo invasive testing, which yields a culture isolate that can be subjected to susceptibility testing. Even if invasive testing is performed, the organism can be difficult to isolate in culture and is highly fastidious once isolated, oftentimes not being amenable to phenotypic susceptibility testing. Further, there are only a handful of specialized microbiology laboratories that perform *H. pylori* susceptibility testing. In an internal study of local and referred isolates published in 2016, clarithromycin resistance was observed to be most commonly due to A2143G (70/88 isolates, 79.6%), followed by A2142G (12/88 isolates, 13.6%) and A2142C (3/88 isolates, 3.4%) alterations in the 23S ribosomal RNA gene. Overall, one of these alterations was found in 97% of clarithromycin resistant *H. pylori* isolates studied.

**Reference Values**

Not detected

**Interpretation**

A detected result indicates the presence of *Helicobacter pylori* 23S ribosomal RNA gene; also indicated is whether or not one the 3 most common 23S ribosomal RNA gene single nucleotide variations (A2143G, A2142G, and A2142C) associated with clarithromycin resistance is detected.

A not detected result for *H. pylori* indicates the absence of detectable *H. pylori* DNA but does not negate the presence of the organism and may occur due to inhibition of the polymerase chain reaction (PCR), sequence variability underlying primers or probes, or the presence of *H. pylori* DNA in quantities less than the limit of detection of the assay.

**Cautions**

Proton pump inhibitors (PPI) and bismuth compounds should be avoided for 2 weeks prior to initial testing; antibiotics should be avoided for 4 weeks prior to initial testing. False-negative results may occur if testing occurs prior to these recommended timeframes. If testing is performed to determine response to therapy, testing should be performed at least 4 weeks after completion of antibiotic therapy and after PPI therapy has been withheld for 2 weeks. Histamine 2-receptor antagonists have been shown to slightly decrease the sensitivity of some *Helicobacter pylori* tests and, if possible, should be discontinued 2 weeks before testing. Antacids do not appear to impair test performance and may be taken until one day before testing.

Test results should be used as an aid in the diagnosis. The single assay should not be used as the only criterion to form a treatment decision; results of this test should be correlated with clinical presentation and results of other laboratory tests. A negative result does not negate the presence of the organism or active disease.

Potential cross-reactivity may occur with the following nonpylori *Helicobacter* species: *Helicobacter acinonychis*, *Helicobacter cetorum*, and *Helicobacter mustalae* (not been reported to cause disease in humans) and *Helicobacter canis*, *Helicobacter cinaedi*, *Helicobacter bizzozeronii*, and *Helicobacter heilmannii* (infrequently found in humans). This assay examines the three most common 23S ribosomal RNA single point variants associated with clarithromycin resistance. Other mechanisms of clarithromycin resistance are not assessed, nor are mechanisms of resistance to non-clarithromycin antimicrobial agents.
Supportive Data
During laboratory verification studies, 745 fecal samples previously tested with the Meridian Premier Platinum HpSA Plus fecal antigen test were assayed with this test. The assay detected *Helicobacter pylori* DNA in 306/335 antigen positive fecal samples (91% sensitivity [87.5-93.9%, 95% CI]). The *H pylori* with Clarithromycin Resistance Prediction (HPFRP) assay also detected *H pylori* DNA in 12/410 antigen negative fecal samples (97.1% specificity [94.9%-98.5%, 95% CI]). Positive and negative predictive values were 96.2% (93.5-98.0%, 95% CI) and 93.0% (90.1-95.2%, 95% CI), respectively. Simple Kappa Coefficient measurement of the performance of the assay against that of the antigen test was 0.89 (0.85-0.92, 95%CI), an almost perfect correlation.(3)

Assessment of clarithromycin resistance prediction was made by performing bidirectional Sanger sequencing on all HPFRP positive samples. All 76 samples with predicted clarithromycin susceptible *H pylori*, demonstrated wildtype 23S ribosomal RNA gene sequence at positions 2142 and 2143. All 37 samples with predicted clarithromycin resistant, *pylori* demonstrated single nucleotide polymorphisms of A2143G, A2142G, or A2142C in the detected *H pylori* 23S ribosomal RNA gene.

**Clinical Reference**

**Performance**

**Method Description**
Fecal samples (approximately 50-100 mg) are placed into 50% STAR buffer and extracted on a Hamilton Microlab STAR device using a Mayo Microlab Maxwell HT Fecal DNA Purification Kit. The polymerase chain reaction (PCR) assay
employs a target-specific detection system including primers as well TaqMan detection probes alongside a SimpleProbe for melt curve analysis-based genotyping targeting the 23S ribosomal RNA gene. The LightCycler 480 II instrument amplifies and monitors target nucleic acid sequences by fluorescence during PCR cycling. Detection of amplified product is based on the TaqMan probe principle. For PCR product detection, the TaqMan probe binds the complementary strand of amplified target. Specific PCR Taq polymerase with 5'-3' exonuclease activity degrades the probe, releasing the fluorophore and breaking the proximity to the quencher molecule, allowing fluorescence of the fluorophores. At the conclusion of PCR cycling, amplified product is thermally denatured and then cooled to allow for a fluorescein labeled SimpleProbe to anneal to an 18-base pair region of the amplified target that includes 2 positions mutations at which are known to confer clarithromycin resistance. The temperature is slowly raised while consistently monitoring fluorescence. The process is completed in a closed system to mitigate contamination. Further, contamination control is achieved through UNG enzymatic treatment and a master mix which includes dUTPs.(Chen D, Cunningham SA, Cole NC, Kohner PC, Mandrekar JN, Patel R: Phenotypic and molecular antimicrobial susceptibility of Helicobacter pylori. Antimicrob Agents Chemother. 2017 Mar 24;61(4):e02530-16. doi: 10.1128/AAC.02530-16)

PDF Report
No

Day(s) Performed
Monday through Friday

Report Available
4 to 6 days

Specimen Retention Time
7 days

Performing Laboratory Location
Rochester

Fees & Codes

Fees
- Authorized users can sign in to Test Prices for detailed fee information.
- Clients without access to Test Prices can contact Customer Service 24 hours a day, seven days a week.
- Prospective clients should contact their account representative. For assistance, contact Customer Service.

Test Classification
This test was developed, and its performance characteristics determined by Mayo Clinic in a manner consistent with CLIA requirements. This test has not been cleared or approved by the US Food and Drug Administration.

CPT Code Information
87798
Test Definition: HPFRP
Helicobacter pylori with Clarithromycin Resistance Prediction, Molecular Detection, PCR, Feces

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