

# INTERIM CULTURE METHOD FOR THE DUODENOSCOPE – DISTAL END AND INSTRUMENT CHANNEL

CDC Disclaimer: This protocol has not been validated. The protocol is still being developed and evaluated for the major duodenoscope types. This is an interim protocol and will be updated accordingly.

## **Purpose**

This method is to culture bacteria from reprocessed duodenoscopes (after drying) specifically from the distal end and instrument channel. A laboratory will need to decide whether to process the samples with a <u>Culture Method A - Presence/ Absence by Enrichment</u> method or <u>Culture Method B - Quantitative</u>. The quantitative method also incorporates enriching the remainder of the sample to capture lower levels of contamination.

Sample Types: Instrument channel flush (50 ml) Distal end and elevator mechanism, sampled by a channel-opening brush (submerged in 50 ml)

# Materials and Reagents

- Vortex
- Incubator 35°C to 37°C
- Conical/ centrifugation tubes of various sizes tubes (50-cc, 1.5-cc)
- Sterile 0.01M phosphate buffered saline (PBS) with 0.02% Tween<sup>®</sup>-80 solution (PBST) (one example Teknova, #P3875)
- Blood agar plates
- Selective agar (suggest MacConkey II agar plates for the detection of enteric pathogens)
- Tryptic soy broth (5 mL) (one example Hardy Diagnostics, K89)
- Pipets and pipette tips



# Culture Method A – Presence/ Absence by Enrichment

Note: Process irrigation water and PBST negative controls using the same protocol as the samples

- 1. Vortex the sample for 2 minutes in 10 second bursts
- 2. Aseptically, remove the channel-opening brush
- 3. Transfer the fluid samples (instrument channel flush, channel-opening brush fluid) to 50-cc conical tubes
- 4. Concentrate by centrifugation on a benchtop centrifuge equipped for high volume suspensions (range:  $3,500 5,000 \ge g$  for 10 15 min).
- 5. Remove supernatant for a final volume of 1 mL without disrupting the pellet, or re-suspend the pellet to a final volume of 1 mL using PBST
- 6. Transfer the 1 mL sample to TSB (5 mL)
- 7. Incubate at 35°C to 37°C for 48 hrs
- 8. Check and record turbidity at 18 to 24 hrs (overnight) and 48 hrs
- 9. If the sample is turbid, streak broth for isolation onto blood agar and MacConkey II agar plates
- 10. Incubate at 35°C to 37°C; MacConkey II agar for 18-24 hrs (overnight) and blood agar for 48 hrs
- 11. Observe plates for suspect colonies
- 12. Streak suspect colonies for isolation
- 13. Work up pure isolates for characterization of "low- concern" bacteria, which represent flora from skin and the environment, and species identification of "high-concern" bacteria.
  - a. "Low-concern" bacteria include, but are not limited to, coagulase-negative staphylococci, micrococci, diptheroids, *Bacillus* spp. and other gram-positive rods
  - b. "High-concern" bacteria include, but are not limited to, *Staphylococcus aureus*, *Enterococcus* spp., *Streptococcus* sp. viridians group, *Pseudomonas aeruginosa*, *Klebsiella* spp., *Salmonella* spp., *Shigella* spp. and other enteric gram-negative bacilli.

#### **Culture Method B - Quantitative**

Note: Process the irrigation water and PBST negative controls using the same protocol as the samples

- 1. Vortex the sample for 2 minutes in 10 second bursts
- 2. Aseptically, remove the channel-opening brush



- 3. Transfer the fluid samples (instrument channel flush, channel-opening brush fluid) to 50-cc conical tubes
- 4. Concentrate by centrifugation on a benchtop centrifuge equipped for high volume suspensions (range:  $3,500 5,000 \ge g$  for 10 15 min)
- 5. Remove supernatant without disrupting the pellet to a final volume of 1 mL. If needed, add PBST to a final volume of 1 mL and re-suspend.
- 6. Prepare a 1:10 dilution by adding 100  $\mu l$  of sample to 900  $\mu l$  of PBST
- 7. Vortex the sample for 10 sec
- 8. Pipet the following on to blood agar and MacConkey II agar plates in triplicate and spread evenly to allow for counting colonies
  - a. 100  $\mu$ l of the undiluted sample (final dilution 10<sup>-1</sup>)
  - b. 100  $\mu$ l 1:10 dilution (final dilution  $10^{-2}$ )
- 9. Add remainder of sample to TSB (5 mL) for enrichment in order to capture contamination below the detection limit
- 10. Incubate at 35°C to 37°C; MacConkey II agar for 18- 24 hrs (overnight), blood agar for 48 hrs, and TSB for 48 hrs
- 11. For agar plates: check and record growth at 18 to 24 hrs (overnight; MacConkey II and blood agar plates) and approximately 48 hrs (blood agar)
  - a. Count and record number of colonies from plates
  - b. Calculate CFU/sampled duodenoscope, accounting for the dilution of the sample
- 12. For TSB: check and record turbidity at 18 to 24 hrs (overnight) and approximately 48 hrs (two days)
  - a. If the sample is turbid, streak broth for isolation on blood agar and MacConkey II agar plates
  - b. Incubate at 35°C to 37°C; MacConkey agar for 18- 24 hrs (overnight) and blood agar for 48 hrs (two days)
  - c. Observe plates for suspect colonies
- 13. Streak suspect colonies for isolation
- 14. Work up pure isolates for characterization of "low- concern" bacteria, which represent flora from skin and the environment, and species identification of "high-concern" bacteria.
  - a. "Low-concern" bacteria include, but are not limited to, coagulase-negative staphylococci, micrococci, diptheroids, *Bacillus* spp. and other gram-positive rods
  - b. "High-concern" bacteria include, but are not limited to, *Staphylococcus aureus*, *Enterococcus* spp., *Streptococcus* sp. viridians group, *Pseudomonas aeruginosa*, *Klebsiella* spp., *Salmonella* spp., *Shigella* spp. and other enteric gram-negative bacilli.



## Screening colonies for focused identification of "high-concern" bacteria

In this procedure, it is suggested that laboratories focus their efforts on species identification of "high-concern" bacteria to reduce workload. Characterize colonies with morphology consistent with those species using local clinical laboratory procedures. Facilities should consider using a rapid identification system (e.g. MALDI-TOF) for shortening turn-around times of results.

- MacConkey agar: Perform species identification of recovered GNR.
- **Blood Agar:** Characterize by hemolysis and perform preliminary tests (gram-stain, coagulase and other screening biochemicals) to rule out "low-concern" bacteria. Further species identification is required for "high-concern" bacteria.

## **Limitations**

The sensitivity, specificity and limits on quantitation or detection are not established for all organisms with the specified processing methods.

This procedure focuses on the growth of "high-concern" organisms versus overall bioburden. To capture the overall bioburden, facilities may consider requiring lower temperatures of  $30^{\circ}C$  (±2) with an extended incubation time of 5-7 days for samples on additional blood agar plates.