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APPENDICES

UNIVERSITY HEALTH NETWORK/MOUNT SINAI HOSPITAL, DEPARTMENT OF MICROBIOLOGY

NOTE: This document is Uncontrolled When Printed.
Any documents appearing in paper form that are not stamped in red "MASTER COPY" are not controlled and should be checked against the document (titled as above) on the server prior to use.
<table>
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</tr>
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</tr>
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</tr>
<tr>
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<td></td>
</tr>
</tbody>
</table>
METHICILLIN-RESISTANT *Staphylococcus aureus* (MRSA)

I. **Introduction**

These specimens are submitted to identify carriers of methicillin-resistant *S. aureus* (MRSA). Swabs may be submitted from any body site, but the most common are nasal, rectal and wound, or the combined nasal/axilla/groin/perineum (NAGP).

II. **Specimen Collection and Transport**

See Pre-analytical Procedure - Specimen Collection QPCMI02001

III. **Reagents/ Material/ Media**

The OXOID Denim Blue Agar (DBLUE) contains a species-specific chromogen that turns *Staphylococcus aureus* colonies blue. As this chromogen is light sensitive, plates must be stored in their shipping boxes to protect them from unnecessary light exposure until use. After streaking, place directly into plastic bins inside the incubator shielded from light. No more than 4h light exposure by the final read is acceptable.

See Analytical Process - Bacteriology Reagents_Materials_Media List QPCMI10001

IV. **Procedure**

A. Specimen Processing:
   a) Direct Examination: Not indicated
   b) Culture:

<table>
<thead>
<tr>
<th>Media</th>
<th>Incubation</th>
</tr>
</thead>
<tbody>
<tr>
<td>OXOID Denim Blue Agar (DBLUE)*</td>
<td>O₂, 37°C x 24 h -in the dark</td>
</tr>
</tbody>
</table>

*If multiple swabs from a single patient are received individually, then process as separate specimens. If multiple swabs from a single patient are received as a "bundle" with a single label and order number, then process all swabs in the bundle on a single “DBLUE” plate.

Specimens are planted by WASP and incubated within the WAPlab system.
B. Workflow and Culture Interpretations

Images are taken by the WASPlab for a preliminary screening of blue colonies at 18hrs and a final read at 24hrs.

For small amount of plates requiring offline incubation, keep plates in covered container in the walk-in incubator (O₂, 37°C) and screen plates at the beginning of your shift and at time of final incubation (end of your shift or earlier as applicable). When significant downtime occurs, separate plates by shift planted into larger buckets for screening as above until final reads.

Screen remaining MRSA plates through Infection Control phenomatrix software in the WaspApp and offline incubation bucket for denim blue colonies (NOT blue hazes or dark blue pinpoint colonies)

**No denim blue colonies:**

For plates within the Wasplab system, workup will be done through the Waspapp phenomatrix software or IC screening as a backup.

- In MRSA phenomatrix application, 18hr and 24hr images will appear:
  - plates with no blue colonies will be reported automatically through segregation at 18hrs for re-incubation with a prelim status.
  - plates with no blue colonies will be reported automatically through segregation a 24hrs as Negative (see reporting section) and sent to Trash.
- In IC screening
  - At 18hrs: select 18hr-Reinc and send results to re-incubate and status to the MRSA test.
  - At 24hrs: select NO MRSA. Plate will be automatically reported as Negative (see reporting section) and sent to Trash.

For Screening offline document each reading within the LIS.

- For preliminary reads with no blue colonies document DBLUE reading as No Blue and status MRSA test as prelim.
- At the end of your shift or earlier as applicable, perform the 24hr final read. For negative plates at 24hrs, document in DBLUE media 24hr: No blue and report MRSA test as Negative (see reporting section). Trash plates.
Denim Blue colonies:

For each plate with blue colonies, plate with be sent for ID with an isolate quantitation (1-5, scant, light, moderate, heavy)

For plates within the Wasplab system, Reading and Pick plates to send for ID or a subculture. For Screening offline document each reading within the LIS.

i. Send plate for colony identification:
   o If non-sufficient colonies for ID, document and make a subculture plate for next shift.

ii. For ID’s achieved other than S.aureus, suppress the isolate and send out a negative MRSA report (See reporting section)

iii. For ID’s of S.aureus check each patient’s MRSA and VRE history.

Previous VRE history:
   o Regardless of MRSA history (new or prev), if patient has had VRE history *within last 3 months* add “VANCS” before sending an interim result.

Previous Positive MRSA history: (within 3 months)
   o If Vitek MS identified as S. aureus, or Pastorex Staph–Plus is positive, check patient VRE history. If patient has had any VRE, (within the last 3 months) and there is sufficient growth of blue colonies, set up VANCS. If no positive VRE history, report as “MRSA with quantitation”, assign “Interim” status for review.

   o If Vitek MS identified as NOT S. aureus, suppress ID and finalize as “Negative - No methicillin-resistant Staphylococcus aureus (MRSA) isolated”.

iii) If SUBBA grows **an organism other than staphylococcus**, document organism and supplementary tests performed and finalize as “Negative - No methicillin-resistant Staphylococcus aureus (MRSA) isolated”.

New MRSA: follow NEW MRSA work up below
Check “New” MRSA worklist for outstanding specimens from the previous day and ask for replant if any are not accounted for.

Complete leftover old work from the previous day. Check “Old” MRSA worklist for outstanding workup needing completion.

i) **For NEW MRSA**

a) If Vitek MS is negative for *S. aureus*, result as “Negative – No methicillin-resistant Staphylococcus aureus (MRSA) isolated” and status as “Final”.

b) If MS identified as *S. aureus*, perform DENKA (Denka Seiken PBP2a agglutination test).

c) If MS identified as *S. aureus* and DENKA+, <CTRL> “P” as “MRSA” and notify IC and ward as per Isolate Notification and Freezing Table QPCMI15003. Set up oxacillin screen (OXA), vancomycin screen (VANCS), Vitek GPAST and KB mupirocin (MUP\textsubscript{200}) disc. When complete, interim for review as “MRSA”. Set up MUP\textsubscript{200} E-test if MUP\textsubscript{200} zone <19mm. Set up fusidic acid E-test if MRSA is resistant to both SXT and TET. Also set up BHIB for PF (MSH), SUBBA for PFGE at PHL (Baycrest), as appropriate and freeze (FRZ).

If VITEK SXT=R SUPPRESS SXT and confirm result by KB BEFORE reporting. A POP-UP will remind you: “Dsxt=R//uncommon susceptibility result. Suppress and verify w/ KB”

d) If MS identified as *S. aureus* but DENKA-negative, CTRL “P” as “MRSA presumptive identification, confirmation to follow” and notify IC/ward as per Isolate Notification and Freezing Table QPCMI16003, set up OXA/VANCS/MUP/VT GP- AST and set up KB (from 0.5 McFarland suspension) with cefoxitin disc.

e) After overnight incubation, record cefoxitin KB result and perform from colonies that grew closest to the cefoxitin disc.
If induced DENKA is positive, notify IC/ward of confirmed “MRSA”. Document other test results and FRZ, setting up BHIB (for PFGE) or SUBBA for PGFE at PHL when appropriate and status the test as “Interim” for review.

If induced DENKA is negative, refer to How to section in the susceptibility manual.

f) Send to NML in batches when requested by IC for CNISP surveillance

V. Reporting

Negative report: “Negative - No methicillin-resistant Staphylococcus aureus (MRSA) isolated”

Positive report: “Methicillin-Resistant Staphylococcus aureus” with quantitation and appropriate susceptibilities and comments for new cases ().

Scant growth (1-5 colonies) - Upon Infection Control request to replant into BHIB (2mL):
- Confirmed by replanting original specimen in broth
  Add ISOLATE Comment:
  “MRSA confirmed by broth enrichment culture.”
  LIS Code: “\MRSc

- NOT confirmed by replanting original specimen in broth:
  1. Change original isolate to an alpha isolate
  2. Add TEST Comment
     “No MRSA isolated by broth enrichment culture. The previous report of “MRSA isolated” was not confirmed by broth enrichment culture suggesting that the previous report reflects contamination or a very low level positive result. Please send another screening swab as clinically indicated.”
     LIS Code: “\MRSC”

VI. References


VANCOMYCIN-RESISTANT ENTEROCOCCI (VRE)

TEMPORARY PROCEDURE CHANGE IN EFFECT:
Effective Immediately until further notified

Procedure modification: Please set up Cepheid VRE PCR on ANY amount of purple colonies from BVRE plate.

Introduction

These specimens are submitted to identify carriers of vancomycin-resistant E. faecium and/or E. faecalis (VRE). Swabs may be submitted from any body site (other than nasal and axilla), but most commonly are collected from the rectum.

Specimen Collection and Transport

See Pre-analytical Procedure - Specimen Collection QPCMI02001

Specimen Rejection Criteria

Nasal and axilla swabs will not be processed for VRE. Refer to Reporting in Section VI for the appropriate reporting comment.

Reagents/ Material/ Media

See Analytical Process - Bacteriology Reagents_Materials_Media List QPCMI10001
Procedure

A. Processing of Specimen:

Refer to Specimen Processing Procedure MI_SM_PROC

a) Direct Examination: Not indicated

b) Culture in non-outbreak setting:

<table>
<thead>
<tr>
<th>Media</th>
<th>Incubation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Brilliance VRE Agar (BVRE)</td>
<td>O₂, 37°C x 36hrs in the dark</td>
</tr>
</tbody>
</table>

B. Culture for **VRE PCR positive samples** in outbreak setting:

<table>
<thead>
<tr>
<th>Media</th>
<th>Incubation time (all O₂ at 37°C)</th>
</tr>
</thead>
<tbody>
<tr>
<td>i) Place 500uL (0.5 mL mark of transfer pipette) of the eSwab transport medium into:</td>
<td>overnight on shaker</td>
</tr>
<tr>
<td>- 2 mL Brain Heart Infusion broth (BHIB)</td>
<td></td>
</tr>
<tr>
<td>Place 30uL (1 drop from transfer pipette) of the eSwab transport medium onto:</td>
<td>36h in the dark</td>
</tr>
<tr>
<td>- Brilliance VRE Agar (BVRE)</td>
<td></td>
</tr>
<tr>
<td>ii) If BVRE is no growth after overnight incubation, subculture 1 drop from BHIB to:</td>
<td></td>
</tr>
<tr>
<td>- Brilliance VRE Agar (BVRE)</td>
<td>36h in the dark</td>
</tr>
</tbody>
</table>
C. Workflow and Interpretation of cultures:

Workflow is described in the Bench Workflow Manual.

Process specimens as per WASPLab Screening, Reading and Picking Manual.

For specimens processed offline:

a) Label new bin for Planting incubator
b) Read BVRE plates planted from the previous day, separating plates growing purple or blue colonies. Read 36 hrs. plates separating plates growing purple or blue colonies.

VRE cultures will be read at 18hrs, 30hrs and a final reading at 36hrs for VRE \textit{faecium} & \textit{faecalis}

<table>
<thead>
<tr>
<th>Colonies on Briliance VRE Agar:</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Isolate:</strong></td>
</tr>
<tr>
<td></td>
</tr>
<tr>
<td>\textit{Enterococcus faecium}</td>
</tr>
<tr>
<td>\textit{Enterococcus faecalis}</td>
</tr>
<tr>
<td>CNST</td>
</tr>
<tr>
<td>Yeast</td>
</tr>
<tr>
<td>\textit{Enterococcus gallinarum}</td>
</tr>
<tr>
<td>Lactobaclli</td>
</tr>
</tbody>
</table>

A. Royal Blue and Purple colonies:

Check history of patient who’s specimen’s are growing purple colonies.

a) If patient is a \textit{“New” positive with} \geq 5 \textit{purple colonies}; perform PCR and Vitek MS
   a) Inoculate a spot on Vitek MS slide for ID
   b) Pick purple colonies and emulsify them in 0.5 mL saline
   c) Using the same swab, inoculate a vial of PCR sample reagent and set up Cepheid PCR
   d) Using the 0.5mL emulsified saline, inoculate a SUBBA and ¼ BVRE (SBVRE).

\textit{Note: For isolates with unsuccessful ID results, report as Vancomycin-Resistant E.\textit{faecium}, confirmation to follow (with vanA/B positive result)}

b) If patient is \textit{“New” positive with} < 5 \textit{colonies}
   
   - Pick colony(ies) and emulsify into 0.5 mL saline
   - Using the same swab, inoculate a SUBBA and ¼ BVRE
c) If patient is a “Previous” positive (≤3 months)
   • Set up Vitek MS, VANCS, PP,
   • Do NOT report out isolate until VANCS results are know. (Refer to Antimicrobial Susceptibility Manual “How to detect VRE” section.

   Follow : Table 1 VRE Workup Guide –PURPLE COLONIES for further work up.

B. Samples growing Denim/Light Blue colonies:

Observe quantity of suspect colony growth.

   a) Scant growth: inoculate colonies into 0.5mL saline and onto ¼ BVRE (SBVRE)
   b) Moderate/Heavy growth:
      • Inoculate a spot on Vitek MS for ID.
      • If enterococcus faecium or faecalis (or gpc chains when Vitek MS fails) emulsify colonies in 0.5 mL saline and use that swab to inoculate a vial of PCR sample reagent and set up Cepheid PCR.
      • Using the 0.5mL emulsified saline, inoculate a SUBBA and ¼ BVRE (SBVRE).

      Note: For isolates with unsuccessful ID results, report as Vancomycin-Resistant E.faecium, confirmation to follow (with vanA/B positive result)

   Follow : Table 2 VRE Workup Guide –BLUE for further work up.

C. No Royal Blue and Purple or Denim/Light Blue colonies:

Re-incubate negative plates for further incubation as needed.

Enter “__hr: No purple or blue” and status as “Prelim”.
Finalize 36 hr culture as negative. (See VRE reporting section)

Read and report old work. Communicate to ward and/or infection control if necessary as per Isolate Notification and Freezing Table QPCMI16003

Continue to scan BVRE plates and process any that are now growing purple or blue colonies.

Check new VRE worklist after all plates are prelimmed for any missing plates. Document if plate is not found and ask for replant.
VRE Identification:
Rule out VRE as below:

**Table 1**  **VRE Workup Guide –PURPLE COLONIES**

<table>
<thead>
<tr>
<th>NEW Purple/Royal Blue Colonies</th>
<th>PREVIOUS + (&lt;3months)</th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td>BVRE &gt;5cols 24/48 hours</td>
<td>BVRE – scant, &lt;5 cols</td>
<td>BVRE (any amount)</td>
</tr>
<tr>
<td>1. Set up <strong>vanA/vanB Cepheid PCR</strong> and MS and SBVRE and SUBBA</td>
<td>1. Set up SBVRE and SUBBA</td>
<td>1. Set up MS and VANCS</td>
</tr>
<tr>
<td>2. <strong>Cepheid – Positive</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>* Report according to ID as Entfar or Entfer with comment</td>
<td>* Report as No VRE</td>
<td></td>
</tr>
<tr>
<td>* VanA gene positive OR vanB gene positive</td>
<td>3. <strong>SBVRE - Any growth</strong></td>
<td></td>
</tr>
<tr>
<td>* If Cepheid vanB Positive, Roche PCR must be done for MSH patients only (within 24 hours)</td>
<td>* Set up Cepheid PCR &amp; MS</td>
<td></td>
</tr>
<tr>
<td>* Notify ICP/ward</td>
<td>* Cepheid Positive: proceed as #2 (BVRE &gt;5 cols).</td>
<td>2. <strong>VANCS – NG</strong></td>
</tr>
<tr>
<td>* Set up Etest, VANCS</td>
<td>* Cepheid Negative: proceed as #3 BVRE &gt;5 cols. (growth on SBVRE)</td>
<td></td>
</tr>
<tr>
<td>* Vanco Etest ≥8ug/mL,</td>
<td></td>
<td></td>
</tr>
<tr>
<td>* VANCS-growth,</td>
<td></td>
<td></td>
</tr>
<tr>
<td>* SBVRE-growth,</td>
<td></td>
<td></td>
</tr>
<tr>
<td>o Report Entfar or Entfer with phenotype comment</td>
<td>* If “Previous” entfar or entfer: report as NO VRE</td>
<td></td>
</tr>
<tr>
<td>* Vanco Etest ≤4ug/mL,</td>
<td></td>
<td></td>
</tr>
<tr>
<td>* VANCS - NG,</td>
<td></td>
<td></td>
</tr>
<tr>
<td>* SBVRE - NG,</td>
<td></td>
<td></td>
</tr>
<tr>
<td>o Perform PCR again from etest plate to confirm presence of vanA</td>
<td>* If “Previous” entvaa, perform Cepheid:</td>
<td></td>
</tr>
<tr>
<td>o Report as vanco sensitive entvaa or entfva</td>
<td>* Cepheid vanA Positive: report as entvaa</td>
<td></td>
</tr>
<tr>
<td>o Report with comment: “Vanco susceptible phenotype”</td>
<td>* Cepheid Negative: report as NO VRE</td>
<td></td>
</tr>
<tr>
<td>* PFGE (MSH only) &amp; FRZ</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

*Set up etests for Vanco and Teico every 3 months from original isolate to confirm phenotype

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Management System\UHN_Mount Sinai Hospital Microbiology\Standard Operating Procedures\Bacteriology Procedures\
### NEW Purple/Royal Blue Colonies

<table>
<thead>
<tr>
<th>BVRE &gt;5cols 24/48 hours</th>
<th>BVRE – scant, &lt;5 cols</th>
<th>BVRE (any amount)</th>
</tr>
</thead>
</table>

3. Cepheid – Negative

- SBVRE - NG
  - Report as No VRE

- SBVRE - GROWTH
  - Set up Vanco/Teico Etests, VANCS
  - Vanco Etest ≥8ug/mL, VANCS - growth,
    - Add comment non vanA/B to isolate
    - Send for van genotyping & FRZ
    - Notify ICP
Table 2  **VRE Workup Guide – BLUE COLONIES**

<table>
<thead>
<tr>
<th>Blue Colonies (SCANT /LIGHT growth)</th>
<th>Blue Colonies (HEAVY)</th>
</tr>
</thead>
</table>
| NG on SBVRE | Scant Growth on SBVRE | Mod-Heavy Growth on SBVRE | Vitek MS ID of  
| | | | Enterococcus faecium or Enterococcus faecalis. |
| Report – No VRE | Set up VANCS ‘PP’ | Set up Cepheid PCR& MS | Set up Cepheid PCR, SBVRE and SUBBA |
| – Report No VRE | – Follow NEW Purple >5 Cepheid positive workflow. | – Follow NEW Purple >5 Cepheid positive workflow. | |
| 2. VANCS - Growth | 2. Cepheid – Negative | 2. Cepheid – Negative | |
| – Set up MS | – Set up Etests &VANCS | – Set up Etests &VANCS | |
| – Perform Cepheid PCR, Etests. | – If Vanco Etest ≥8ug/mL, VANCS - growth, add comment: ‘non vanA/B’ to isolate | – If Vanco Etest ≥8ug/mL, VANCS - growth, add comment: ‘non vanA/B’ to isolate | |
| – Proceed as Mod-Heavy Growth | – Send to NML for van genotyping & FRZ | – Send to NML for van genotyping & FRZ | |
| – Notify ICP/ward | – Notify ICP | – Notify ICP | |
Table 3 VRE Workup Guide – Cepheid PCR + from E- swab directly

1. Phone/email ward and ICP as per Isolate Notification and Freezing table QPCMI15003.

2. For new or previous VRE patients where NO isolate has been isolated yet proceed as below:

<table>
<thead>
<tr>
<th>Subculture to BHIB broth and BVRE and incubate overnight</th>
<th>If BVRE is No growth, Subculture BHIB to BVRE</th>
</tr>
</thead>
<tbody>
<tr>
<td>NG</td>
<td>Scant Growth (purple or blue colonies)</td>
</tr>
<tr>
<td>Report as “No VRE isolated after broth enhancement”</td>
<td>1. Sub to SUBBA and SBVRE</td>
</tr>
<tr>
<td></td>
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<td></td>
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</tbody>
</table>
VRE PFGE:

If patient is from MSH only, subculture isolate to Brain Heart Infusion Broth for pulsed-field (PFGE).

VI. Reporting

Negative Report: “Negative - No Vancomycin-Resistant Enterococci (VRE) isolated”

Positive Report:

New Positive VRE Patients
Day 1
- **PCR direct from BVRE plate - with isolate**
  ISOLATE: “*Enterococcus (faecium or faecalis)*-vancomycin resistant” “isolated”
  ISOLATE COMMENT:
  “This organism is positive for the vanA or B gene as tested by the Cepheid vanA/B GenXpert Assay (for research only). ~Phenotypic confirmation to follow.” Isolate Comment Code `vaAg or vaBg`

- **PCR direct from BVRE plate - no isolate, from sweep**
  ADD ISOLATE COMMENT:
  “PCR from a sweep of growth on the plate is positive for the vanA gene by the Cepheid vanA/B GenXpert Assay (for research use only) but a distinct vancomycin-resistant or vancomycin-susceptible Enterococcus species that is vanA positive cannot be found.”
  Isolate Comment Code `vaAp`

Day 2
- Vancomycin=R, Teicoplanin=R:
  “*Enterococcus faecium* (or faecalis) -vancomycin resistant” “isolated”
  ISOLATE COMMENT (Code `vaA`):
  “This organism has a vanA phenotype.”

- Vancomycin=R, Teicoplanin=S:
  “*Enterococcus faecium* (or faecalis) -vancomycin resistant”
  ISOLATE COMMENT (Code `vaB`):
“This organism has a vanB phenotype.”

**Previous VRE Positive Patients:**

*Enterococcus (faecium or faecalis)*—vancomycin-resistant isolated.

**ISOLATE COMMENT (Code: \vapr):**

“The Cepheid vanA/B GenXpert Assay was not completed as this patient has had VRE isolated within the past 3 months that has had molecular characterization.”

**Vancomycin=S, vanA gene-positive VRE**

- **Isolate from IC VRE Culture Screen**

  1) Change previous isolate code of entfar to **entvaa** - “*Enterococcus faecium* - vanA gene positive” “isolated”

  **ISOLATE COMMENT (Code: \vaAi)** – “This organism is positive for vanA gene by the Cepheid vanA/B GenXpert Assay (for research use only) but has a vancomycin susceptible phenotype. The effectiveness of vancomycin in this setting is uncertain and is not recommended. Please contact Infectious Diseases or Medical Microbiology for treatment advice.”

  Remove previous duplicated ISOLATE COMMENT.

  2) Change previous isolate code of entfer to **entfva** - “*Enterococcus faecalis* - vanA gene positive” “isolated”

**Vancomycin MIC =>8 by macro Etest, vanA/B-negative by PCR**

“*Enterococcus faecium or faecalis*” “isolated”

**ISOLATE COMMENT (Code: \vanI):**

“This organism has reduced susceptibility to vancomycin but is negative for vanA and vanB genes as tested by the Cepheid vanA/B GenXpert Assay (for research use only). ~This organism has been sent to the National Microbiology ~Laboratory for further testing and results will be ~reported when available.”

**Confirmation from NML:**
Negative – Add the following statement as an ‘Updated Report’: “The previously reported organism has no vancomycin resistance genes as tested by the National Microbiology Laboratory, …. Winnipeg, Specimen No. xxxx”

Positive – Enterococcus faecalis or faecium - vancomycin-resistant “isolated”
ISOLATE COMMENT (Code: vanE):
“This organism is positive for the vanE gene as reported by the National Microbiology Laboratory… NML Specimen No. xxx”
VII. References


2. National Committee for Clinical Laboratory Standards 2003 Performance Standards for Antimicrobial Susceptibility Testing; 13th Informational Supplement Document M100-S13 (M2) for use with M2-A8 – Disk Diffusion NCCLS, Wayne, PA


6. QMP-LS Committee Comments, Survey B-0109 Patterns of Practice with VRE Surveillance Specimens *QMP-LS Bacteriology*; 3, Section 2.2: 663-669


RESISTANT GRAM NEGATIVE BACILLI

I. Introduction

These specimens may be submitted to identify carriage of drug-resistant Gram negative bacilli, to determine cross-transmission between patients or to identify an environmental source of patient infection.

II. Specimen Collection and Transport

Any specimen may be submitted, although rectal swabs and environmental are the most common. Swabs should be transported in an Eswab or Amies transport medium. If a delay in transport or processing is anticipated, the swabs should be kept at 4°C.

III. Reagents/Materials/Media

See Analytical Process - Bacteriology Reagents_Materials_Media List QPCMI10001

IV. Procedure

A. Processing of Specimen:

a) Direct Examination: Not indicated

b) Culture:

<table>
<thead>
<tr>
<th>Media</th>
<th>Incubation</th>
</tr>
</thead>
<tbody>
<tr>
<td>MacConkey Agar (Mac) – no antibiotic</td>
<td>O₂, 35°C x 18 h</td>
</tr>
</tbody>
</table>

For Enterobacteriaceae with fluoroquinolone and/or aminoglycoside resistance but susceptibility to cefpodoxime:

For Serratia marcescens outbreaks,

| CTCZ – with colistin    | O₂, 35°C x 18 h |

B. Interpretation of cultures:

1. Read cultures plates after 18 to 24 hours of incubation.
2. Workup requested organism as per Bacteria Workup Manual
3. Set up susceptibility as per
4. Communicate with requesting Infection Control Practitioner or Microbiologist as appropriate and freeze all positive isolates unless otherwise directed. PFGE will only be performed on request from Infection Control.

For Serratia Screen:
1. Read culture plates after 18 to 24 hours of incubation.
2. For Serratia marcescens, work-up NLF, LLF or orange-red pigmented colonies only. Perform Vitek MS.
   * Phone ward and email ICP if Serratia marcescens is isolated.
3. Set up susceptibility as per Susceptibility Manual.
4. Previously positive Serratia marcescens specimens only require a meropenem screen to be set up.
5. If Serratia is isolated, freeze and set up BHIB for PFGE.

N.B. Susceptibilities can be referred for 3 months

V. Reporting

Negative report: “No <requested organism> isolated”
Positive report: “<requested organism> isolated”
Report their susceptibility results as per Susceptibility Manual.
Add Isolate comment: “Susceptibility testing results are provided for infection control purposes only.” ICSN

VI. References


ESBL and Carbapenemase SCREEN

I. Introduction

These specimens are submitted to identify *Klebsiella* species, *Escherichia coli* and *Proteus mirabilis* with acquired extended spectrum β-lactamases as well as carbapenemases from any *Enterobacteriaceae*.

ESBL testing is only performed on specimens from pregnant patients, specimens originating from mothers and baby units or upon special request.

II. Specimen Collection and Transport

See [Pre-analytical Procedure - Specimen Collection QPCMI02001](#).

III. Reagents/Materials/Media

See [Analytical Process - Bacteriology Reagents_Materials_Media List QPCMI10001](#).

IV. Procedure

A. Processing of Specimen:

   a) Direct Examination: Not indicated

   b) Culture:

<table>
<thead>
<tr>
<th>Media</th>
<th>Incubation</th>
</tr>
</thead>
<tbody>
<tr>
<td>ESBL Isolation Agar with 2 μg/ml cefpodoxime</td>
<td>O₂, 37°C x 18-24 hours</td>
</tr>
</tbody>
</table>

B. Interpretation of cultures:

1. Examine plates after 18-24 hours of incubation for any growth of an *Enterobacteriaceae*.
2. If no *Enterobacteriaceae* are isolated, report as “Negative - no ESBL or Carbapenemase producing organisms isolated.”
3. For all LF and oxidase negative NLF Enterobacteriaceae colony types, set up Vitek MS for identification.
4. Should an isolate ID as an E.coli, Klebsiella spp., or P.mirabilis, check patient history.
   - For a patient with no prior history or with “Previous” positive (>3months) history of E.coli, Klebsiella spp., or P.mirabilis in an IC sample set up ‘KB IC ESBL ’.
   - If a previous positive ESBL was isolated within the last 3 months, set up Meropenem Screen, only by disk diffusion. Refer to the previous sample’s date that susceptibilities were reported. Report isolate with phrase “Phenotypic screening suggests this organism is ESBL POSITIVE as previously confirmed on “yyyy.mm.dd”. LIS isolate comment code \ESBP

Report with Test Comment:
“Negative Carbapenemase screen - No cabapenemase producing organisms isolate. POSITIVE for ESBL screen”.

5. For all other Enterobacteriaceae set up Meropenem Screen only.
6. For CRE work up, refer to
   
V. Reporting

When ESBL screen is requested, report both ESBL and Carbapenemase comments where applicable.

Negative report for both ESBL and carbapenemase:
“Negative - No extended-spectrum-beta-lactamase producing (ESBL) or carbapenemase-producing organism isolated”

Positive reports:
Positive for both ESBL and Carbapenemase:
At TEST Window:

POSITIVE for ESBL screen
POSITIVE Carbapenemase Screen

At ISOLATE Window:
“Escherichia coli” or “Klebsiella species” or “Proteus mirabilis” isolated with one of the following ISOLATE COMMENT:

“The susceptibility pattern suggests that this organism contains a class A extended spectrum beta-lactamase (ESBL).”

OR

“The susceptibility pattern suggests that this organism contains class A and C extended spectrum beta-lactamases (ESBL).”

OR

“The susceptibility pattern suggests that this organism contains a class C extended spectrum beta-lactamase (ESBL).”

OR

“The susceptibility pattern suggests that this organism contains an inducible class C extended spectrum beta-lactamase (ESBL).”

OR

“The susceptibility pattern suggests that this organism contains an extended spectrum beta-lactamase (ESBL) other than class A or C.”

AND

From keypad: ESBL1: ICSN “Susceptibility testing results are provided for infection control purposes only.”

AND

Final Positive CRE Result by CARB-R PCR:

“______ carbapenemase gene DETECTED by Cepheid Xpert CARBA-R Assay (for research use only). This assay is able to detect NDM, KPC, OXA48, OXA181, OXA232, IMP-1, and VIM carbapenemase genes.” Isolate Comment Code: CPC+

OR

Preliminary CRE Result:

Isolate Comment: CNML

AND

UNIVERSITY HEALTH NETWORK/MOUNT SINAI HOSPITAL, DEPARTMENT OF MICROBIOLOGY

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Management System\UHN_Mount Sinai Hospital Microbiology\Standard Operating Procedures\Bacteriology Procedures\
Send updated, Final Result once NML report is available

Negative report:
  a. **Suppress the isolate**
  b. Add the following comment in the TEST window for NOT CONFIRMED carbapenemase:
     Add TEST COMMENT code `KPCN`
  c. Enter the NML results to the LIS ISOLATE Breakpoint panel `kpcrcon`.
  d. E-mail or call Infection Control Practitioner and ward as per Isolate Notification Table.

Positive report:
  a. “Updated Report”
  b. Add the following isolate comment for CONFIRMED carbapenemase:
  c. Add ISOLATE COMMENT code `KPCP`.
  d. Enter the NML results to the LIS ISOLATE Breakpoint panel `kpcrcon`.
  e. E-mail or call Infection Control Practitioner and ward as per Isolate Notification Table.

**Negative report for carbapenemase but POSITIVE for ESBL:**
At TEST Comment: “Negative Carbapenemase Screen - No carbapenemase-producing organism isolated”
  **POSITIVE ESBL Screen**
At ISOLATE Window: “*Escherichia coli*” or “*Klebsiella species*” or “*Proteus mirabilis*” “isolated” with ISOLATE COMMENT: “The susceptibility pattern suggests that this organism contains a class A extended spectrum beta-lactamase (ESBL).” OR “The susceptibility pattern suggests that this organism contains class A and C extended spectrum beta-lactamases (ESBL).” OR “The susceptibility pattern suggests that this organism contains a class C extended spectrum beta-lactamase (ESBL).” OR “The susceptibility pattern suggests that this organism contains an inducible class C extended spectrum beta-lactamase (ESBL).” OR “The susceptibility pattern
suggests that this organism contains an extended spectrum beta-lactamase (ESBL) other than class A or C.”
Report appropriate sensitivity results as per

**Previous ESBL Positive Patient:**

**Negative report for carbapenemase but POSITIVE for ESBL:**

At TEST Comment: “Negative Carbapenemase Screen - No carbapenemase-producing organism isolated”

POSITIVE ESBL Screen”

At ISOLATE Window:

“*Escherichia coli*” or “*Klebsiella species*” or “*Proteus mirabilis*” “isolated” with ISOLATE COMMENT: “Phenotypic screening suggests this organism is ESBL POSITIVE as previously confirmed on “yyyy.mm.dd”.”

LIS isolate comment code: \ESBP

**Negative report for ESBL but POSITIVE for carbapenemase:**

At TEST Comment: “Negative ESBL Screen- No extended spectrum beta-lactamase - producing organism (ESBL) isolated”

POSITIVE Carbapenemase Screen”

At ISOLATE Window:

Report isolate comment as per

**Previous Carbapenemase Positive Patient:**

At TEST Comment: “Negative ESBL Screen- No extended spectrum beta-lactamase – producing organism (ESBL) isolated POSITIVE Carbapenemase Screen”

OR

“POSITIVE ESBL Screen and POSITIVE Carbapenemase Screen”

At ISOLATE Window: Report isolate along with Isolate Comment:
“Phenotypic testing suggests this organism is carbapenemase POSITIVE as previously confirmed on “yyyymmdd”.”
Isolate Comment code CREP

VI. Reference

1. **Clinical and Laboratory Standards Institute** 2016 Performance Standards for Antimicrobial Susceptibility Testing; Documents M100-S26, M2-A12, M7-A10 CLSI, Wayne, PA.
**Carbapenemase (CRE) SCREEN (without ESBL Screen)**

I. **Introduction**

These specimens are submitted to identify carbapenemases from any *Enterobacteriaceae*.

II. **Specimen Collection and Transport**

See [Pre-analytical Procedure - Specimen Collection QPCMI02001](#).

III. **Reagents/Materials/Media**

See [Analytical Process - Bacteriology Reagents_Materials_Media List QPCMI10001](#).

IV. **Procedure**

A. Processing of Specimen:

   a) Direct Examination: Not indicated

   b) Culture:

<table>
<thead>
<tr>
<th>Media</th>
<th>Incubation</th>
</tr>
</thead>
<tbody>
<tr>
<td>ESBL Isolation Agar with 2 μg/ml cefpodoxime</td>
<td>O₂, 37°C x 18-24 hours</td>
</tr>
</tbody>
</table>

B. Interpretation of cultures:

See [IC Carbapenemase Testing Flowchart](#).

1. Examine plate after 18-24 hours of incubation for any growth of an *Enterobacteriaceae*.
2. If no *Enterobacteriaceae* are isolated, report as negative for CRE.
3. For all *Enterobacteriaceae* colony types, set up a meropenem screen disk diffusion test.
4. If isolates >25mm (susceptible) by “MEMS” disk diffusion, report as negative for CRE.
5. For all Meropenem Screen R (<25mm) by disk diffusion, Set up Vitek MS
   - If the isolate is not identified as *Enterobacteriaceae*, report as negative for CRE.
• If the isolate is identified as *Enterobacteriaceae*, suppress the isolate and set up βCARBA (BCARB)

a) **If βCARBA (BCARB) is negative:**
   i. Set up Rosco with Temocillin (breakpoint panel kpcros)
      • If **Temocillin = S** and Rosco disks show no potentiation, send out report as NO CRE.
      • If **Temocillin = R** OR Rosco shows potentiation to MRBO or MRDP,
         o Report isolate with the following ISOLATE Comment: \CNML
         o Phone or e-mail IC and ward as per Isolate Notification Table. Send isolates to NML ASAP (Cannot send on Friday)
         o Order the LIS ISOLATE Breakpoint panel kpcron.
         o Freeze isolate

b) **If βCARBA (BCARB) is positive:**
   – Previous CRE positive (≤ 6 months)
     i. At TEST Comment: “POSITIVE Carbapenemase Screen”
        At ISOLATE Window: Report (un-suppress) isolate along with Isolate Comment: \CREP
        • Phone or e-mail IC and ward as per Isolate Notification Table.

   – New ?positive CRE
     Report (un-suppress) isolate. with comment \PCRB
     Notify ICP
     i. Set up Cepheid CARBA-R PCR (CARBR)
        • If Cepheid CARBA-R PCR (CARBR) is negative
           o Report isolate with the following ISOLATE Comment: \pCRB
           o Phone or e-mail IC and ward as per Isolate Notification Table.
o Send isolates of to NML ASAP (Cannot send on Friday)
o Order the LIS ISOLATE Breakpoint panel kpcrcon
o Set up Rosco with Temocillin (panel kpcros). For epidemiology purposes only. Record and suppress kpcros results.

- If Cepheid CARBA-R PCR (CARBR) is positive
  o Report gene identified by Cepheid using ISOLATE Comment:
    \CPC+
  o Phone or e-mail IC and ward as per Isolate Notification Table.
  o Send to NML and PHOL in batches when requested for
    and .
  o Freeze isolate

V. **Reporting**

See [Carbapenemase Testing Reporting](#).

VI. **Reference**

1. **Clinical and Laboratory Standards Institute** 2016 Performance Standards for Antimicrobial Susceptibility Testing; Documents M100-S26, M2-A12, M7-A10 CLSI, Wayne, PA.

2. **QMP-LS Bacteriology Consensus Practice Recommendations** – Antimicrobial Susceptibility Reporting Toronto ON: QMP-LS QView. c2011
RESISTANT *Pseudomonas aeruginosa* SCREEN

I. **Introduction**

Specimens are submitted for the screening of multi-drug resistant *Pseudomonas aeruginosa*.

II. **Specimen Collection and Transport**

See [Pre-analytical Procedure - Specimen Collection](#)

III. **Reagents/Materials/Media**

See [Analytical Process - Bacteriology Reagents, Materials, Media List](#)

IV. **Procedure**

1. **Processing of Specimen:**

<table>
<thead>
<tr>
<th>Specimen</th>
<th>Processing</th>
<th>Media</th>
<th>Incubation</th>
</tr>
</thead>
<tbody>
<tr>
<td>Water</td>
<td>Centrifuge the entire sample at 3500 rpm for 20 minutes. Pour off all supernatant. Transfer the contents of a 2 mL tube of BHI broth into the falcon tube containing the sediment</td>
<td>BHI Broth</td>
<td>$O_2$ at 35°C overnight</td>
</tr>
<tr>
<td></td>
<td>Subculture BHI broth after overnight incubation to MCPOD by the IC Bench technologist</td>
<td>MCPOD</td>
<td>$O_2$ at 35°C for 24 hours</td>
</tr>
<tr>
<td>Environmental swabs</td>
<td>Incubate the BHI Broth Subculture BHI broth after overnight incubation to MCPOD by the IC Bench tech using a new sterile swab</td>
<td>MCPOD</td>
<td>$O_2$ at 35°C overnight</td>
</tr>
<tr>
<td>Patient pharmaceutical</td>
<td>≤1 mL Subculture BHI broth after overnight incubation to MCPOD by the IC Bench tech using a new sterile swab</td>
<td>TH14 SD14</td>
<td>$O_2$ at 35°C for 14 days $O_2$ at RT° for 4 days</td>
</tr>
</tbody>
</table>
### Specimen Processing

<table>
<thead>
<tr>
<th>Specimen</th>
<th>Processing</th>
<th>Media</th>
<th>Incubation</th>
</tr>
</thead>
<tbody>
<tr>
<td>infusates/injectables (QC bench)</td>
<td>&gt;1 mL</td>
<td>ETH14</td>
<td>O₂ at 35°C for 14 days</td>
</tr>
<tr>
<td></td>
<td></td>
<td>ESD14</td>
<td>O₂ at RT° for 4 days</td>
</tr>
<tr>
<td>Swabs from patients</td>
<td>Directly inoculate MCPOD plate with specimen</td>
<td>MCPOD</td>
<td>O₂ at 35°C for 24 hours</td>
</tr>
</tbody>
</table>

2. Interpretation of Cultures:

For **water, environmental swabs, patient swabs**:
Work up these cultures on the IC Bench.
Work up oxidase-positive gram negative bacilli ONLY.
Set up Vitek MS
When identified as *P. aeruginosa* set up Vitek susceptibility card.
For patient samples, if resistant to all antimicrobials from the vitek card, set up colistin etest.
Freeze resistant strains of *Pseudomonas aeruginosa* into IGR boxes.

For **Patient pharmaceutical infusates/injectables**:
Work up these cultures on the QC/Sterility Bench.
Work up any growth as per Sterility Manual.

### V. Reporting

For **water, environmental swabs, patient swabs**:

Negative Report: No resistant *Pseudomonas aeruginosa* isolated.

Positive (Resistant strains only) Report: *Pseudomonas aeruginosa* with susceptibility result.
Add Isolate comment: “Susceptibility testing results are provided for infection control purposes only.” ICSN

Email / Call ICP.

For **Patient pharmaceutical infusates/injectables**:

Positive: Report *Pseudomonas aeruginosa* with susceptibility result. Call ICP
Add Isolate comment: “Susceptibility testing results are provided for infection control purposes only.” ICSN

### VI. References

**Clinical and Laboratory Standards Institute** 2016 Performance Standards for Antimicrobial Susceptibility Testing; Documents M100-S26, M2-A12, M7-A10 CLSI, Wayne, PA.
GROUP A STREPTOCOCCUS SCREEN

I. **Introduction**

Throat, rectal or wound swabs are the most common that are submitted for the diagnosis of Group A streptococcal infection, to determine cross-transmission between patients or to identify an environmental source of patient infection.

II. **Specimen Collection and Transport**

See Pre-analytical Procedure - Specimen Collection QPCMI02001

III. **Reagents / Materials/ Media**

See Analytical Process - Bacteriology Reagents_Materials_Media List QPCMI10001

IV. **Procedure**

A. Processing of Specimens

See Specimen Processing Procedure

a) Direct Examination: Not routinely performed.

b) Culture:

<table>
<thead>
<tr>
<th>Media</th>
<th>Incubation</th>
</tr>
</thead>
<tbody>
<tr>
<td>CNA (rectal/wound)</td>
<td>AnO₂, 35°C x 18-24 hours</td>
</tr>
<tr>
<td>Carrot Broth</td>
<td>O₂, 35°C x 18-24 hours</td>
</tr>
<tr>
<td>BA (for throat)</td>
<td>AnO₂, 35°C x 18-24 hours</td>
</tr>
</tbody>
</table>

If original plates are negative; Subculture the Carrot Broth to a second CNA plate and incubate overnight in AnO₂, 35°C x 18-24 hours

B. Interpretation of Cultures:

a) Examine the CNA/ BA plate after 18-24 hours incubation and identify all morphologically distinct beta haemolytic colonies by performing:
   i) Catalase test
   ii) Strep grouping

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b) For all specimens processed after 1600 hours, re-incubate CNA/BA anaerobically for a further 24 hours.

c) Subculture the Carrot broth to a second CNA/BA plate and incubate overnight in anaerobic conditions

d) Examine the subculture CNA/BA plate after overnight incubation for distinct beta haemolytic colonies.

e) Perform catalase and strep grouping if any beta haemolytic colonies appear.

f) Freeze all isolates and prepare for PFGE (whether in house or to be sent to PHL)

h) E-mail or call Infection Control Practitioner and ward as per Isolate Notification Table.

V. Reporting

A. Culture:

Negative report: “No Group A streptococcus isolated”.

Positive report: Report as isolate - “Group A streptococcus” with LIS ISOLATE COMMENT: “isolated”

E-mail or call all positive Group A streptococci isolates to ward / Infection Control Practitioners as per Isolate Notification Table.

VI. References

**Clinical and Laboratory Standards Institute** 2016 Performance Standards for Antimicrobial Susceptibility Testing; Documents M100-S26, M2-A12, M7-A10 CLSI, Wayne, PA
KLEBSIELLA OXYTOCA OR KLEBSIELLA PNEUMONIAE SCREEN

I. Introduction

These specimens may be submitted to identify carriage of drug resistant ESBL producing *Klebsiella oxytoca* or *Klebsiella pneumoniae*, to determine cross-transmission between patients or to identify an environmental source of patient infection.

II. Specimen Collection and Transport

See Pre-analytical Procedure - Specimen Collection QPCMI02001

III. Reagents/Materials/Media

See Analytical Process - Bacteriology Reagents_Materials_Media List QPCMI10001

IV. Procedure

A. Processing of Specimen:

   a) Direct Examination: Not indicated

   b) Culture:

      For Cefpodoxime resistant *Klebsiella oxytoca* or *Klebsiella pneumonia*

      | Media                                      | Incubation          |
      |-------------------------------------------|---------------------|
      | ESBL Isolation agar with 2ug/mL cefpodoxime | O2, 350°C x 18 h    |

   B. Interpretation of cultures:

      1. Read cultures plates after 18 to 24 hours of incubation.
      2. Perform id on lactose fermenting gram negative bacilli using Vitek MS.
      3. Set up Double Disk Kirby Bauer (KBESBLR) on those isolates identified as *Klebsiella oxytoca* or *Klebsiella pneumonia* to rule out ESBL producers.
      4. Record susceptibility results and freeze organism
5. Communicate with requesting Infection Control Practitioner or Microbiologist as appropriate and freeze all positive isolates. PFGE will only be performed on request from Infection Control.

N.B. Susceptibilities can be referred for 3 months

V. **Reporting**

Negative report: “No <requested organism> isolated”

Positive report: “<requested organism> isolated”

Report their susceptibility results as per Susceptibility Manual with one of the following ISOLATE COMMENT if applicable:

“The susceptibility pattern suggests that this organism contains a class A extended spectrum beta-lactamase (ESBL).” OR

“The susceptibility pattern suggests that this organism contains class A and C extended spectrum beta-lactamases (ESBL).” OR

“The susceptibility pattern suggests that this organism contains a class C extended spectrum beta-lactamase (ESBL).” OR

“The susceptibility pattern suggests that this organism contains an inducible class C extended spectrum beta-lactamase (ESBL).” OR

“The susceptibility pattern suggests that this organism contains an extended spectrum beta-lactamase (ESBL) other than class A or C."

AND

“Susceptibility testing results are provided for infection control purposes only.”

VI. **References**

**Clinical and Laboratory Standards Institute** 2016 Performance Standards for Antimicrobial Susceptibility Testing; Documents M100-S26, M2-A12, M7-A10 CLSI, Wayne, PA
APPENDIX II
HOW TO SET UP AND INTERPRET A MIC PANEL

I. Materials

MIC panel
Panel inoculator set
Sterile distilled water
Sterile transfer pipettes
Blood agar plate
Sealable bag

II. Procedure

1. Remove the desired MIC panel from the –70°C freezer. Place a cover over the panel and place into the O₂ incubator to thaw.

2. When thawed, label the panel and a blood agar plate with the LIS order number.

3. Make a suspension of the organism in saline to match a 0.5 McFarland standard.

4. Place 1.5 mL of organism into a 50mL tube. Add sterile distilled water to reach 40mL on same falcon tube (~38.5mL). Pour into the inoculator base. Gently mix by agitating slowly

5. Place the inoculator into the base making sure there are no bubbles and that all prongs are in contact with the bacterial suspension.

6. Align the left side (lettered) of the panel with the left side (lettered) of the inoculator.

7. Lift the inoculator straight up and place it, prong side down, into the wells of the MIC panel.

8. Using a transfer pipette, transfer 1 drop of suspension from within the inoculator base to a blood agar plate and streak for isolated colonies.

9. Pour the suspension into a sharps container containing hypochloride and discard the inoculator into a sharps disposal box.
10. Place a lid onto the panel and place into a sealable bag. Seal the bag and incubate the panel in the appropriate atmosphere and temperature (See below).

<table>
<thead>
<tr>
<th>Panel</th>
<th>Temp.</th>
<th>Atmosphere</th>
<th>Incubation</th>
</tr>
</thead>
<tbody>
<tr>
<td>VRE (vancomycin)</td>
<td>35°C</td>
<td>O₂</td>
<td>24 h</td>
</tr>
<tr>
<td>MRSA (oxacillin)</td>
<td>35°C</td>
<td>O₂</td>
<td>24 h</td>
</tr>
<tr>
<td>GNB</td>
<td>35°C</td>
<td>O₂</td>
<td>24 h</td>
</tr>
</tbody>
</table>

III. Interpretation

Use a coordinating MIC panel sheet to record wells with any growth. Each panel contains a positive growth control well (no antibiotic) and a negative growth control well (no inoculum). The MIC for each drug is the lowest dilution showing no growth. Record results in the LIS.

Interpretation of MIC results is performed in accordance with NCCLS breakpoint criteria found in the Performance Standards for Antimicrobial Susceptibility Testing Informational supplement M100-S**. This informational supplement is updated annually and breakpoint criteria for all antibiotics used should be checked yearly for changes.

MIC breakpoints for antimicrobial agents tested in MIC panels that do not have NCCLS criteria available should be obtained from the literature (see references for agents such as mupirocin and fusidic acid). When breakpoints are not available in the literature, no interpretation of MIC should be reported.

IV. References


3. Fuchs P. C., R. N. Jones, A. L. Barry  
Interpretive Criteria for Disk Diffusion Susceptibility Testing of Mupirocin, a Topical Antibiotic  
*J Clin Microbiol* 1990; 28: 608-609

4. Skov R., N., Frimodt-Moller, F. Espersen  
Correlation of MIC methods and tentative interpretive criteria for disk diffusion susceptibility testing using NCCLS methodology for fusidic acid  
*Diag Microbiol Infect Dis* 2001; 40: 111-116
### Record of Edited Revisions

**Manual Section Name: Infection Control Manual**

<table>
<thead>
<tr>
<th>Page Number / Item</th>
<th>Date of Revision</th>
<th>Signature of Approval</th>
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<tr>
<td>Annual Review</td>
<td>March 13, 2002</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td>October 25, 2003</td>
<td>Dr. T. Mazzulli</td>
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<td>Annual Review</td>
<td>May 26, 2004</td>
<td>Dr. T. Mazzulli</td>
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<td>May 12, 2005</td>
<td>Dr. T. Mazzulli</td>
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<tr>
<td>Annual Review</td>
<td>July 23, 2006</td>
<td>Dr. T. Mazzulli</td>
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<tr>
<td>Link to <a href="#">IC\Infection Control Pulsed-field Gel Electrophoresis.doc</a> added</td>
<td>January 30, 2007</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Link to <a href="#">IC\VRE PCR Procedure.doc</a> added</td>
<td>January 30, 2007</td>
<td>Dr. T. Mazzulli</td>
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<td>Enter the no. of pink colonies grown on MRSA-Select if &lt;5</td>
<td>January 30, 2007</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Added quantitation for MRSA</td>
<td>February 28, 2007</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Change to Denim Blue plates for MRSA Screen</td>
<td>March 13, 2007</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Change negative resulting phrases for MRSA, VRE and ESBL screen</td>
<td>March 13, 2007</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Included P. mirabilis for ESBL screen</td>
<td>March 13, 2007</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td>March 13, 2007</td>
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</tr>
<tr>
<td>Revised VRE Identification Procedure</td>
<td>March 22, 2008</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>VRE – VANCS resistant <em>E. faecium</em> or <em>E. faecalis</em> report to MSH ICP if it is MSH patient; change to report as Presumptive VRE to all ICP</td>
<td>September 20, 2008</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Pseudo screen, patient swabs – change incubation period from 48 hours to 24 hours</td>
<td>September 20, 2008</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td>September 20, 2008</td>
<td>Dr. T. Mazzulli</td>
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<tr>
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<td>September 20, 2009</td>
<td>Dr. T. Mazzulli</td>
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<tr>
<td>Annual Review</td>
<td>September 20, 2010</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>ESBL screen updated to include KPC and NDM screen</td>
<td>November 10, 2010</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Removed send by taxi for carbapenemase PCR send out for Monday, Wednesday and Thursday</td>
<td>January 20, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Modified carbapenemase screening procedure to match Susceptibility manual</td>
<td>April 04, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Change VRE screening to Brilliance VRE Agar</td>
<td>April 04, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Page Number / Item</td>
<td>Date of Revision</td>
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<td>----------------------------------------------------------------------------------</td>
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<tr>
<td>Removed VRE Table 3; added link to Susceptibility manual</td>
<td>May 11, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>VRE Screen, added VANCS back to heavy growth from BVRE or SBVRE</td>
<td>May 31, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>VRE Screen – modified, finalized all day 2 reading in the morning of day 2</td>
<td>October 17, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td>October 17, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Modified Serratia screen</td>
<td>November 25, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Modified VRE resulting phrases</td>
<td>December 13, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Added CRE only screen</td>
<td>December 13, 2011</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Modified VRE reporting for vanA gene positive but phenotype vancomycin=S strains</td>
<td>February 1, 2012</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Added link to VRE PCR by Cephied</td>
<td>July 16, 2012</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Modified planting volume into BHI broth for VRE/MRSA</td>
<td>August 28, 2012</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td>August 28, 2012</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td>May 31, 2013</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Manual updates in each section (Maldi procedure review)</td>
<td>October 10, 2013</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Changed CRE screen from ERTA to MERO discs</td>
<td>October 10, 2013</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Update VRE Identification</td>
<td>April 19, 2014</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td>April 19, 2014</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>CRE reporting changes (Mero screen I/R)</td>
<td>June 27, 2014</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>VITEK SXT=R SUPPRESS SXT confirm result by KB BEFORE reporting.</td>
<td>June 27, 2014</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>vanB gene detected by Cepheid Xpert vanA/vanB Assay reporting</td>
<td>August 6, 2014</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Insert proper headers/footers, UHN/MSH Logo</td>
<td>August 12, 2014</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Fix broken link to appendix 2</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Added Group A Strep Screen and Klebsiella Screen</td>
<td>September 30, 2014</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Reviewed and updated procedure steps in all sections</td>
<td>September 30, 2014</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Added MRSA scant growth repeat broth culture comments</td>
<td>December 10, 2014</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td></td>
<td></td>
</tr>
<tr>
<td>MRSA Removed MACRO use</td>
<td>May 19, 2015</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>p.6 add “Scant growth (1-5 colonies) Upon Infection Control request to replant into BHIB (2mL):”</td>
<td></td>
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<tr>
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<tr>
<td>p.19 changed serratia media to new ctcz media’</td>
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</tr>
<tr>
<td>p.22 ESBL testing is only performed on specimens from pregnant patients, specimens originating from mothers and baby units or upon special request.</td>
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<tr>
<td>p.35 BA (for throat) added to GAS screen procedure culture media</td>
<td></td>
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<tr>
<td>Changed Mac with cefpodoxime to ESBL isolation agar with cefpodoxime</td>
<td></td>
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<tr>
<td>p.39 added ESBL comments to reporting comments</td>
<td></td>
<td></td>
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<tr>
<td>p.24: added suscep. comment: \MRSS</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Specimen collected and transport for each section transferred to Specimen collection manual QPCMI02001 And replaced with link to specimen collection manual</td>
<td>May 26, 2015</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>VRE outbreak: Temporary Procedure change in effect: VRE PCR on any amount of purple colonies</td>
<td>June 11, 2015</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>VRE outbreak: Temporary Procedure change ended. Section removed.</td>
<td>July 15, 2015</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Prev + ESBL and Prev + CRE new statements</td>
<td>August 20, 2015</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>previously ESBL reporting phrased changed from “Susceptibility not done, please refer to sample collected on <strong>Date</strong>” to “Phenotypic screening suggests this organism is ESBL POSITIVE as previously confirmed on “yyyy.mm.dd”.” LIS isolate comment code: \ESBP</td>
<td>December 2, 2015</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Updated ESBL+CRE and CRE sections with new reporting phrases</td>
<td>December 21, 2015</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Updated CRE section with new BCARB/CARB-R/ROSCO procedure</td>
<td></td>
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</tr>
<tr>
<td>MRSA reporting section: added link for susceptibility comments to MRSA reporting phrases in susceptibility manual.</td>
<td>April 4, 2016</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Resistant GNB reporting section &amp; Pseudomonas screen section added for Positive reports: Add comment: “Susceptibility testing results are provided for infection control purposes only.” \ICSN</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Added links in TOC to CNISP Surveillance Study for MRSA, VRE, and CRE and PHOL CRE Surveillance Study as well as in the CRE procedure section.</td>
<td>April 12, 2016</td>
<td>Dr. T. Mazzulli</td>
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<td>VRE outbreak: Temporary Procedure change in effect: VRE PCR on any amount of purple colonies</td>
<td>June 13, 2016</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>MRSA for new MRSA added step “Send to NML in batches when requested by IC for CNISP surveillance” VRE - added HEAVY growth workup</td>
<td>July 29, 2016</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>VRE: Updated commntnb \vaAi to include: “The effectiveness of vancomycin in this setting is uncertain and is not recommended. Please contact Infectious Diseases or Medical Microbiology for treatment advice.”</td>
<td>December 1, 2016</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td>May 20, 2017</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Updated Direct VRE PCR results with instructions to phone/email as per Isolate Notification and freezing table.</td>
<td>February 2, 2018</td>
<td>Dr. T. Mazzulli</td>
</tr>
<tr>
<td>Annual Review</td>
<td>May 22nd, 2018</td>
<td>Dr. T. Mazzulli</td>
</tr>
</tbody>
</table>
| Temporary Procedure change in effect: VRE PCR on any amount of purple colonies. PFGE for all new VRE from MSH NOT UHN. Instructions to release ID once PCR is done of \textit{E. faecium} or \textit{faecalis} for suspect colonies if Vitek MS fails, confirmation of ID to follow. WASPLAB screening/incubation time changes:  
- MRSA 12hr & 24hrs modified to one 24hr read on Wednesday May 18\textsuperscript{th}, 2018 evening.  
- MRSA 24 hr read changed to 18 & 24 hr read to Monday May 21, 2018 evening.  
- VRE changed from 12 & 36 to 18, 30, 36 on Monday May 21, 2018 evening. | | |
| Minor format change | September 14, 2018 | Dr. T. Mazzulli |
| Addition of Wasplab changes to MRSA section. | December 11, 2018 | Dr. T. Mazzulli |
| Update of MUP to MUP\textsuperscript{200} | November 11, 2019 | Dr. T. Mazzulli |
| Updated CPO flowchart. Removed reporting and notification for BCARB negative enterobacteriaceae (\textit{NCRB}), and icp notification of enterobacteriaceae that are BCARB \& Rosco negative reported as No CPO. | December 19, 2019 | Dr. T. Mazzulli |