



Wisconsin State Laboratory of Hygiene



Why are We Concerned with Non-Tuberculous Mycobacteria?

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Outline

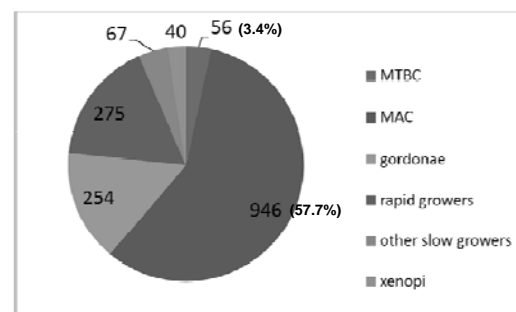
- Incidence of Non-tuberculous mycobacteria (NTM) in Wisconsin
- Clinical significance of NTM
- Identification of NTM
- Susceptibility testing of NTM

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3

2012 Wisconsin State-wide Surveillance Mycobacteria Isolates, n=1638



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4

Clinical Significance of Mycobacteria

- Identification of *Mycobacterium tuberculosis* complex (MTBC) is the most important finding in the laboratory
- Finding of MTBC has serious clinical and public health consequences
 - Isolation almost always signifies disease
 - Exception: A single patient specimen in the absence of clinical indications may be a false positive
 - MTBC is not found in the environment

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5

Clinical Significance of Non-Tuberculous Mycobacteria (NTM)

- NTM are free-living mycobacteria, usually found in association with water or soil habitats.
- Although not components of the microbiota of humans, NTM may be isolated as “bystanders” from the skin, upper respiratory tract, intestinal tract and genital tract in asymptomatic individuals
- Not all NTM isolations are clinically significant as sources of human disease
- Due to their ubiquitous nature, the question of clinical significance is important and often difficult to answer.

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6

Clinical Significance of NTM

- Considerations for clinical relevance
 - Clinical setting and host
 - Organism species and its pathogenic potential
 - Source of the culture isolate & likelihood for contamination/colonization
 - Quantification of organisms detected both in culture and AFB smears
 - Number of positive cultures

Clinical Significance of NTM

- Likely significant if collected surgically or aseptically: sterile body fluids, tissue biopsies
- For respiratory specimens:
 - Multiple culture-positive specimens
 - 2 positive sputa or 1 bronch
 - 1 trans-bronch or lung biopsy
 - Single positive sputum specimen not likely to be significant
- Laboratory clues:
 - Abundant growth on primary culture
 - More than one culture positive
 - Liquid and solid media positive

Clinical Relevance of Mycobacteria Isolated from Respiratory Specimens


	<ul style="list-style-type: none"> • <i>M. tuberculosis</i> • <i>M. kansasii</i> • <i>M. avium-intracellulare</i> • <i>M. abscessus</i> • <i>M. fortuitum</i> • <i>M. xenopi</i> • <i>M. goodii</i> 	<ul style="list-style-type: none"> • Severe immunocompromised • HIV • Immunosuppressive drugs • Bronchiectasis and other underlying lung disease • Older age • Cystic fibrosis • Diabetes, system illness • Otherwise healthy
	Species pathogenicity	Patient risk factors

Figure adapted from Clinical and Laboratory Standards Institute (CLSI). Laboratory Detection and Identification of Mycobacteria, M88-A, 2008

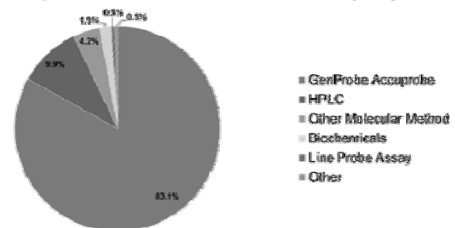
Common NTM Diseases

NTM	Disease
<i>M. abscessus</i>	Skin and soft tissue infections following traumatic injury and surgical or medical procedures, pulmonary disease in CF patients
<i>M. avium</i> complex	Tuberculosis-like pulmonary disease, chronic pulmonary disease, disseminated disease in patients with AIDS, lymphadenitis in children
<i>M. chelonae</i>	Skin and soft tissue infections following traumatic injury and surgical or medical procedures
<i>M. fortuitum</i> group	Skin and soft tissue infections following traumatic injury and surgical or medical procedures, nail salon infections
<i>M. goodii</i>	Rarely a pathogen, contamination from tap water
<i>M. kansasii</i>	Chronic pulmonary disease resembling tuberculosis
<i>M. marinum</i>	Cutaneous infections ("fish tank granuloma")
<i>M. xenopi</i>	Lung infections

Identification of Mycobacteria

National TB Laboratory Services Survey

Primary Method for Identification of MTBC from Culture (n=213)



APHL National TB Laboratory Services Survey Report, 2012

Wisconsin Mycobacteriology Laboratory Network (WMLN)

- 32 labs around the state that provide some level of mycobacteriology testing
 - 30 clinical labs
 - Milwaukee City Health Department Lab
 - Wisconsin State Lab of Hygiene (WSLH)
- 7 labs perform identification of mycobacteria
 - GenProbe Accuprobe
 - Biochemical reactions
 - DNA sequencing
 - MALDI-TOF

Identification of Mycobacteria

CLASSICAL METHODS

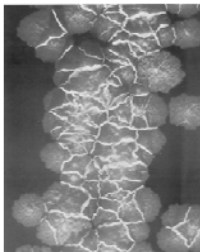
Growth Characteristics

- Growth Rate
 - Can be observed on the primary solid media, but dependent on appropriate incubation temperature and number of organisms in the primary specimen
 - To perform a standardized growth test from subculture
 - Inoculate a defined suspension of mycobacteria on solid media
 - Incubate at 30 C and 35–37 C
 - Observe for growth at 5–7 days and weekly thereafter
 - Rapidly Growing Mycobacteria: form visible colonies within 7 days of incubation (usually 3–4 days)
 - Slowly Growing Mycobacteria: require more than 7 days for visible colonies to form

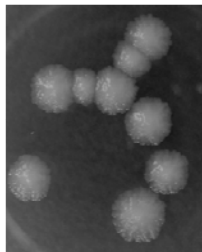
Growth Characteristics (2)

- Colony Morphology
 - Observe on solid media, use stereoscope to visualize young and small colonies
 - Colony texture and consistency
 - Rough, smooth, dry, granular, mucoid, translucent
 - Colony pigment
 - Color range: Non-pigmented, buff, yellow, orange, pink
 - Photochromogen: require light to form pigment
 - Scotochromogen: form pigment in either light or dark
 - Non-photochromogens: no pigment

Growth Characteristics (3)



M. tuberculosis complex



M. goodii

Pictures from Manual of Clinical Microbiology 10th Edition, p 475, 510

Conventional Biochemical Testing

- Classical approach to identification
- Requires sufficient amount of bacterial cells and several weeks of incubation
- New mycobacterial species cannot be reliably identified by biochemical and other phenotypic tests
- Current recommendations are for rapid methods

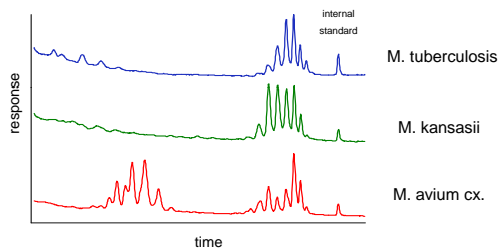
High Performance Liquid Chromatography (HPLC)

- Cell wall mycolic acids are extracted and derivatized to fluorescent or UV-adsorbing esters and then separated by chromatography
- A pattern of peaks (chromatogram) is generated as mycolic acids are detected
- Identification is based on comparison of isolate's pattern to a database or library of chromatograms

Considerations for HPLC

- Advantages
 - Some methods can identify *Mycobacterium tuberculosis* complex (MTBC) and nontuberculous mycobacteria (NTM) from broth culture and directly from clinical specimens
 - Cost of individual sample testing is relatively inexpensive
 - FDA-cleared system commercially available
- Limitations
 - Initial equipment costs are high
 - Some methods require mature solid medium growth
 - Problematic for identification of rapidly-growing mycobacteria; limited ability to resolve some NTM groups/complexes
 - Extraction uses hazardous chemicals; hazardous waste production

Fluorescence-HPLC Patterns



Identification of Mycobacteria MOLECULAR AND EMERGING METHODS

GenProbe Accuprobe

- In-solution hybridization assay for identification of growth on solid or liquid media
 - Nucleic acids are extracted after organisms are lysed and made non-viable
 - Single-stranded labeled DNA probes (in tubes) are allowed to anneal to target RNA
 - If present, RNA:DNA hybrids are detected by chemiluminescence
- Commercially available kits for identification of MTBC, *M. avium* complex, *M. gordonae*, *M. kansasii*

Considerations for Accuprobe

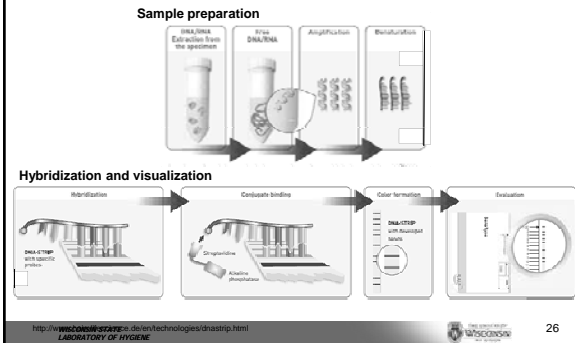


- Advantages
 - Identifies four frequently-isolated mycobacteria; three clinically significant
 - Performed routinely by many laboratories
 - Relatively easy to use
 - FDA-cleared
- Limitations
 - No nucleic acid amplification occurs during this assay; sufficient culture growth is necessary for identification
 - Beware of relative light units (RLU) values that are near the cutoff; "high negative" values could indicate that the target organism is present in low numbers

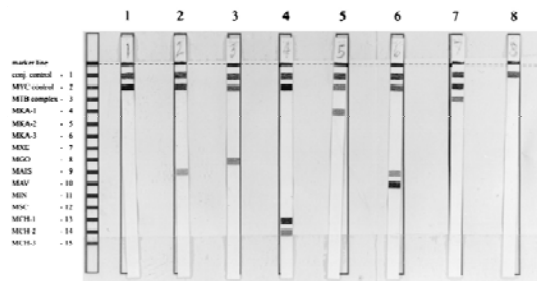
Line Probe Assay

- Platform is nitrocellulose strip used for identification of MTBC and commonly-isolated NTM by reverse hybridization
 - Hybridization of denatured DNA to probes on the membrane strip
- Commercially developed assays
 - HAIN GenoType Mycobacterium CM (MTBC + 24 NTM) and GenoType Mycobacterium AS (19 NTM)
 - Innogenetics INNO-LiPA Mycobacteria v2 (MTBC + 7 NTM)

Line Probe Assay (2)



INNO-LiPA

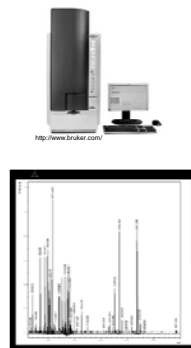


Considerations for Line Probe Assay

- Advantages**
 - Nucleic acid amplification for increase sensitivity
 - Some assays detect mutations associated with MTBC drug resistance
 - Relatively low implementation costs
- Limitations**
 - Not FDA approved
 - Can be difficult to differentiate bands with visual inspection
 - Sometimes difficult to identify species within *Mycobacterium fortuitum* complex, *M. chelonae/abscessus* group

MALDI-TOF

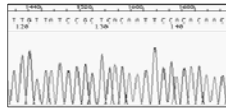
- "Matrix-Assisted Laser Desorption Ionization Time-of-Flight"
- A form of mass spectrometry
- Used to analyze proteins and other macromolecules
- Recent advances have allowed application to the clinical realm
- Charged particles are accelerated by a laser. Time of flight is proportional to the ion's mass. Patterns are compared to a library for identification.



Considerations for MALDI-TOF

- Advantages**
 - Small sample size
 - Rapid identification
 - Can also be used for identification of many bacteria and fungi in the laboratory
- Limitations**
 - Difficult heat inactivation/cell disruption for Mycobacteria
 - Database limitations
 - Initial cost investment high
 - Cannot identify to species within the MTBC

DNA Sequencing



- Determining the precise order of nucleotides
- DNA sequence is compared to a database of sequences from known/characterized organisms
- 16S rRNA, rpoB and hsp65 genes are commonly sequenced for identification

<http://seqware.bmf.med.utoronto.ca/seqmag.html>

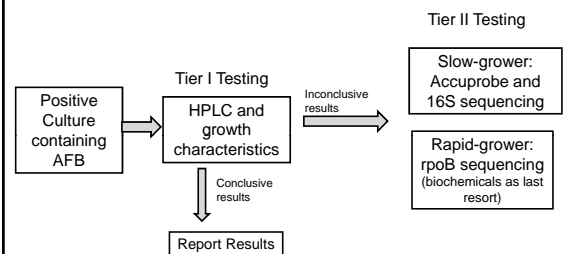
Limitations of 16S sequencing

- Cannot differentiate between
 - Members of the TB complex
 - *M. marinum* vs. *M. ulcerans*
 - *M. chelonae* vs. *M. abscessus*
 - *M. kansasii* vs. *M. gastri*

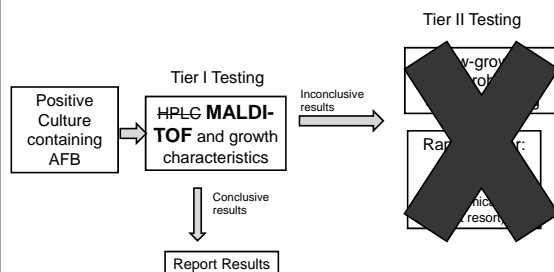
Considerations for Molecular and Emerging Methods

- Advantages
 - Quicker turnaround time (TAT)
 - Uses liquid or solid media cultures, possibly smear-positive clinical specimens
 - Ability to recognize new strains
- Limitations
 - Cost, in particular the initial setup cost
 - Computer and Database management
 - Specialized equipment
 - Requires new expertise and training
 - New methods do not always replace standard methods

WSLH Algorithm for Identification



WSLH Algorithm for Identification



Mycobacteria Taxonomy and Nomenclature

- More than 130 currently validated (fully characterized) species of NTM
- Molecular techniques increase discrimination for better identification to species and appropriate grouping of genetically similar organisms
- Nomenclature and taxonomy are constantly changing.

Slowly Growing Mycobacteria

Group or Taxon	Species within the group or taxon
<i>M. avium</i> complex ²	<i>M. avium</i> subsp. <i>avium</i> , <i>M. avium</i> subsp. <i>silvaticum</i> , <i>M. avium</i> subsp. <i>paratuberculosis</i> , <i>M. avium</i> subsp. <i>hominissuis</i> , <i>M. intracellulare</i> , <i>M. chimera</i> , <i>M. colombiense</i> , <i>M. vulneris</i> , <i>M. marseillense</i> , <i>M. timonense</i> , <i>M. bouchardurhonense</i> .
<i>M. simiae</i> clade ¹	Over 150 species: <i>M. triplex</i> , <i>M. genovense</i> , <i>M. florentinum</i> , <i>M. lentiflavum</i> , <i>M. palustre</i> , <i>M. kubicae</i> , <i>M. parascrofulaceum</i> , <i>M. heidelbergense</i> , <i>M. interjectum</i> , <i>M. simiae</i> , <i>M. longobardum</i>
<i>M. terrae</i> complex	<i>M. nonchromogenicum</i> , <i>M. terrae</i> , <i>M. trivale</i> , <i>M. arupense</i>

(1) Tortoli et al., IJSEM 2012
(2) Manual of Clin. Micro 10th Edition

Major Groups of Rapidly-growing Mycobacteria

Group or Taxon	Species within the group or taxon
M. fortuitum group	<i>M. fortuitum</i> , <i>M. peregrinum</i> , <i>M. senegalense</i> , <i>M. setense</i> , <i>M. septicum</i> , <i>M. porcinum</i> , <i>M. houstonense</i> , <i>M. boenickelii</i> , <i>M. brisbanense</i> , <i>M. neworleansense</i>
<i>M. chelonae</i> / <i>M. abscessus</i> group	<i>M. chelonae</i> , <i>M. immunogenum</i> , <i>M. abscessus</i> subsp. <i>abscessus</i> , <i>M. abscessus</i> subsp. <i>bolletii</i> , <i>M. salmoniphilum</i>
<i>M. mucogenicum</i> group	<i>M. mucogenicum</i> , <i>M. aubagnense</i> , <i>M. phocaicum</i>
Early pigmented rapidly-growing mycobacteria	<i>M. neoaurum</i> , <i>M. canariense</i> , <i>M. cosmeticum</i> , <i>M. monacense</i>
Late Pigmented rapidly-growing mycobacteria	<i>M. smegmatis</i> , <i>M. goodii</i> , <i>M. mageritense</i> / <i>M. wolinskyi</i>

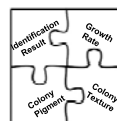
Adapted from Manual of Clin
Micro 10th Edition

Mycobacteria Identification Reporting

- Report identification as TB or not TB as soon as possible (≤ 21 days)
 - Preliminary information (colony morphology) can be helpful to health care providers)
- If sending to a reference laboratory, ensure that this lab is meeting the expected turn-around times.

Mycobacteria Identification

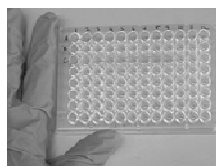
- Identification to species or group is important to help determine clinical significance and treatment
- Use a multi-faceted approach
 - Ensure that identification result matches phenotypic results (e.g., colony morphology and growth rate) before issuing final report
 - Biochemical reactions no longer recommended
 - May need molecular methods for accurate ID
 - Send tricky isolates to reference lab with expertise



Susceptibility testing of Non-tuberculous Mycobacteria

NTM Susceptibility Testing:

Broth microdilution method

[illegible]

***M. avium* complex Susceptibility Testing**

- Which isolates to test?
 - Initial isolates to establish baseline value
 - Isolates from patients on prior macrolide therapy
 - Isolates from patients who develop bacteremia on macrolide prophylaxis
 - Isolates from patients who relapse on macrolides
- Testing should be repeated in 3 months for patients with disseminated disease and 6 months for patients with no improvement of chronic pulmonary disease.

CLSI M24-A2

***M. avium* complex: Clinically active drugs used for therapy**

- macrolide
 - clarithromycin
 - azithromycin
- ethambutol
- rifamycins
 - rifampin
 - rifabutin
- aminoglycosides
 - streptomycin
 - amikacin

***M. avium* complex Susceptibility Testing**

- Routine testing of clarithromycin testing only (broth based)
- Secondary drugs: moxifloxacin and linezolid
- No correlation between in vitro susceptibility results for anti-tuberculous agents (rifampin/rifabutin, ethambutol) with clinical outcome
- No 1st line anti-tuberculous agents should be reported

CLSI M24-A2

***M. avium* complex Susceptibility Testing Available**

- WSLH
 - Broth microdilution, clarithromycin only
- National Jewish, Denver, CO
 - 8-drug panel: amikacin, ciprofloxacin, clofazimine, clarithromycin, ethambutol, rifabutin, rifampin, streptomycin, rifampin + ethambutol synergy
 - 12-drug panel: 8 drug panel plus cycloserine, ethionamide, kanamycin moxifloxacin, rifampin + ethambutol synergy

***M. marinum* Susceptibility Testing**

- Routine susceptibility testing not recommended (all untreated strains have the same drug pattern)
- MICs performed at 3 months if still culture positive
- Clinically active drugs used for therapy: clarithromycin, rifampin, doxycycline/minocycline, trimethoprim-sulfamethoxazole, rifampin + ethambutol

CLSI M24-A2

***M. marinum* Susceptibility Testing Available**

- WSLH
 - Susceptibility testing not performed.
- National Jewish, Denver, CO
 - 10 drug panel plus 3 single drugs:
 - Agar proportion: isoniazid, rifampin, ethambutol, ethionamide, streptomycin, amikacin, kanamycin, capreomycin, cycloserine, PAS
 - radiometric MIC: clarithromycin, ciprofloxacin, rifabutin
 - Susceptibility testing of single drugs

***M. kansasii* Susceptibility Testing**

- Clinically active drugs used for therapy: clarithromycin, ethambutol, rifampin/rifabutin, INH
- Routine testing of rifampin and clarithromycin as primary agents
- Test secondary agents only if rifampin resistant (amikacin, ciprofloxacin, levofloxacin, ethambutol, linezolid, moxifloxacin, rifabutin, trimethoprim-sulfamethoxazole)

CLSI M24-A2

***M. kansasii* Susceptibility Testing Available**

- WSLH
 - Susceptibility testing not performed
- National Jewish, Denver, CO
 - 10 drug panel plus 3 single drugs:
 - Agar proportion: isoniazid, rifampin, ethambutol, ethionamide, streptomycin, amikacin, kanamycin, capreomycin, cycloserine, PAS
 - radiometric MIC: clarithromycin, doxycycline, Bactrim/SXT
 - Susceptibility testing of single requested drugs

RGM: Which Isolates to Perform Susceptibility Testing on?

- Follow ATS criteria for respiratory specimens:
 - Multiple culture-positive specimens
 - 2 positive sputa or 1 bronch
 - 1 trans-bronch or lung biopsy
- Clinically significant isolates from blood, sterile body fluids, skin and soft tissue
- Repeat susceptibility in 6 months if cultures remain positive

Rapidly-Growing Mycobacteria Susceptibility Testing

- Agents that should be tested: amikacin, cefoxitin, ciprofloxacin, clarithromycin, doxycycline (minocycline), imipenem, linezolid, moxifloxacin, trimethoprim/sulfamethoxazole, tobramycin
- No anti-tuberculous agents reported
- Clarithromycin MIC's
 - Read at 3-5 days for mutational resistance
 - Final reading at 14 days to detect inducible resistance due to *erm* gene

CLSI M24-A2

***erm* Gene**

- Clarithromycin resistance in *M. fortuitum* and *M. abscessus*
- Patients with isolates containing the *erm* gene have delayed treatment response and possible failures compared to those patients whose isolates do not contain functional *erm* gene
- *M. abscessus* subsp. *abscessus* (*erm* +)
- *M. abscessus* subsp. *bolletii* (*erm* negative)
- No *erm* gene in *M. chelonae*

Rapid Grower Susceptibility Testing Available

- WSLH
 - 9 drugs by broth microdilution: amikacin, cefoxitin, ciprofloxacin, clarithromycin, imipenem, linezolid, doxycycline, SXT, tobramycin
- National Jewish, Denver, CO
 - 15 drug panel: amikacin, kanamycin, imipenem, ciprofloxacin, tobramycin, trimethoprim/sulfa, linezolid, augmentin, azithromycin, clarithromycin, gentamycin, ceftriaxone, cefepime, cefotaxime, minocycline
 - Susceptibility testing of single requested drugs

NTM Susceptibility Testing Summary

- Not all drugs used for patient therapy have standardized protocols for *in vitro* susceptibility testing or guidelines for interpretation for results.
 - Correlation between *in vitro* susceptibility tests and clinical response has not demonstrated in controlled clinical trials
 - Not adequately studied, breakpoints have not been established
 - Technical difficulties with drug testing
 - Poor drug solubility in assays
 - Poor reproducibility of testing

NTM Susceptibility Testing Summary

- Perform testing on clinically significant isolates
- WSLH performs testing per CLSI recommendations for MAC and RGs
- Further testing can be performed at National Jewish (Denver, CO)

To Request Add-on Susceptibility Testing for Clinically Significant Isolates (WSLH)

Test	Test Codes (old/new)	CPT code	Price
<i>M. avium</i> complex susceptibility (clarithromycin)	652MAC/MM00202	87186 X 1	\$135
Rapid grower susceptibility (9 drugs)	652RG/MM00207	87186 X 9	\$145

To Request Add-on Susceptibility Testing for Clinically Significant Isolates (WSLH)

- Call WSLH Mycobacteriology for help, 608-262-1618.
- WSLH Reference manual and fee schedule: www.slh.wisc.edu
- WSLH requisition Other Tests: "Isolate already at WSLH", indicate desired test codes
- Fax requisition form to 608-890-4891.

To Request Add-on Susceptibility Testing for Clinically Significant Isolates (National Jewish)

Test	Test Code	CPT code	Price
<i>M. avium</i> complex susceptibility (8 drugs)	Panel 3G	87188 X 8	\$305.70
<i>M. avium</i> complex susceptibility (12 drugs)	Panel 3H	87188 X 12	\$427.98
Rapid grower susceptibility (15 drugs)	Panel 3I	87186 X 15	\$265.05
<i>M. Marinum</i> (13 drugs)	Panel 3A + 3 drugs	87190 X 10 87188 X 3	\$234
<i>M. Kansalii</i> (13 drugs)	Panel 3A + 3 drugs	87190 X 10 87188 X 3	\$234

To Request Add-on Susceptibility Testing for Clinically Significant Isolates (National Jewish)

- Call WSLH Mycobacteriology for help, 608-262-1618.
- WSLH requisition Other Test: "Isolate already at WSLH, send to National Jewish for panel ____"
- Fax requisition form to 608-890-4891.
- \$90 shipping and handling fee

Summary

- Non-tuberculous mycobacteria make up the majority of mycobacteria isolations in Wisconsin.
- NTM can cause clinically significant disease, sometimes mimicking tuberculosis
- Identification of mycobacteria requires a multi-faceted approach
- Susceptibility testing for clinically significant NTM is available

For More Information

- WSLH Mycobacteriology Lab: 608-262-1618
- WSLH Customer Service: 1-800-862-1013
- Julie Tans-Kersten
Wisconsin State Lab of Hygiene
(608) 263-5364
Fax: (608) 890-2548
julie.tanskersten@slh.wisc.edu

References

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- Clinical and Laboratory Standards Institute (CLSI), Susceptibility Testing of Mycobacteria, Nocardiae and other Aerobic Actinomycetes; M24-A2, 2011
- www.slh.wisc.edu

WSLH Laboratory Team



Nate



Dave



Youngmi and Ana



Julie B.



Don



Julie TK