

Final Identification of Nontuberculous Mycobacteria (NTM) Matters

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COI-I have no disclosures



- Toolbox 1 – specimen
- Toolbox 2 – positive culture
- Good news – bad news
- Pulmonary NTM
- National Jewish Health algorithm
- National Jewish Health laboratory data
- Beyond routine testing
- Patient centered approach

✓ Specimen – sputum, CSF, formalin-fixed tissue

- AFB microscopy
- Solid & broth-based media
- **NAAT-D** (TB complex)
- **NAAT-R** (RIF, INH and more)
- **Direct AST**

Molecular TB Testing 7 Days a Week

- ✓ **AFB positive culture (broth-, solid-based media)**
 - **TB Yes/No** (final identification within TB complex)
 - **NAAT-R**
 - Broth-based AST
 - Agar-based AST
 - Minimal Inhibitory Concentration (MIC)

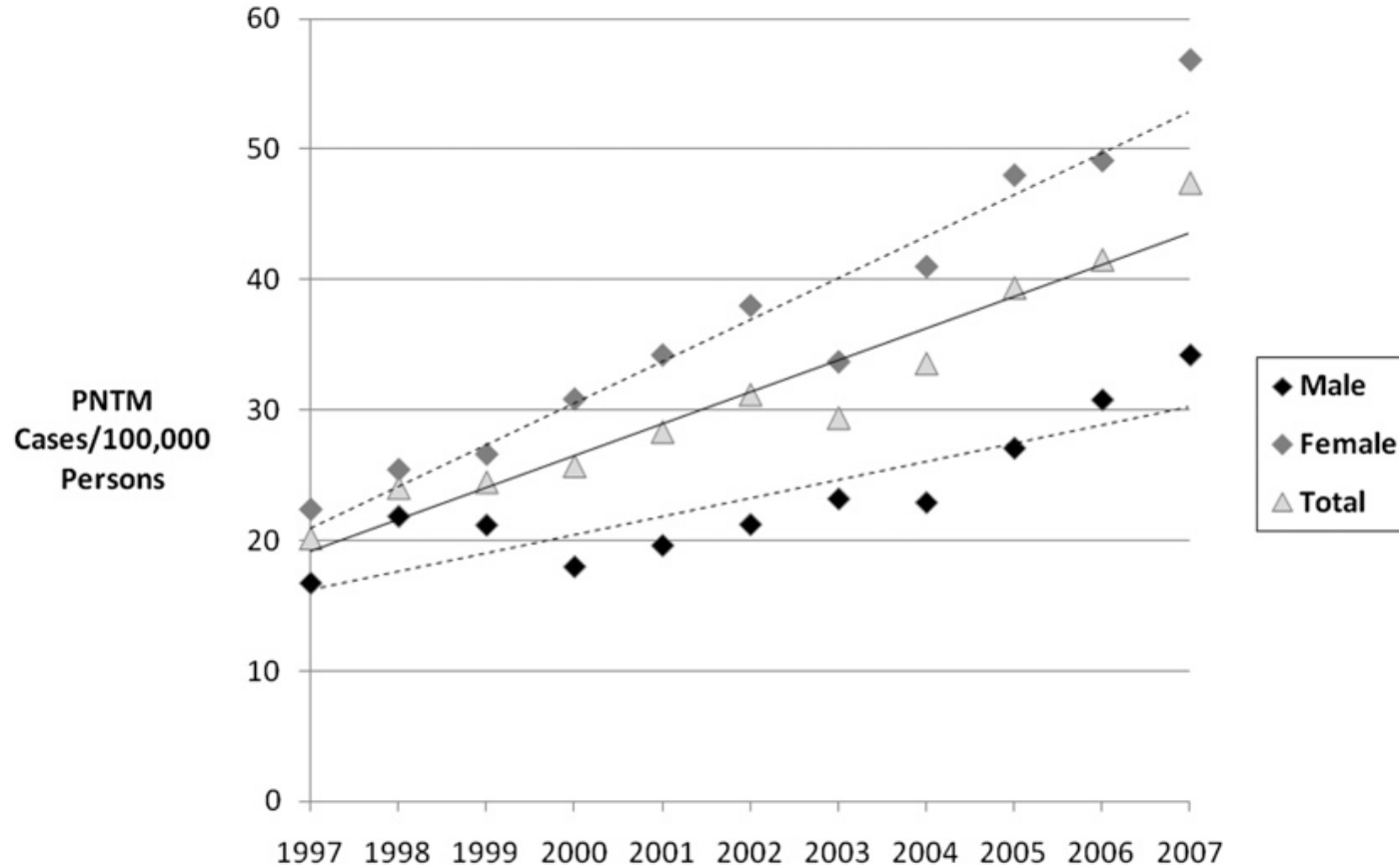
NTM Detected...

Good news for TB Control – No contact investigation, etc.

... and clinically significant...

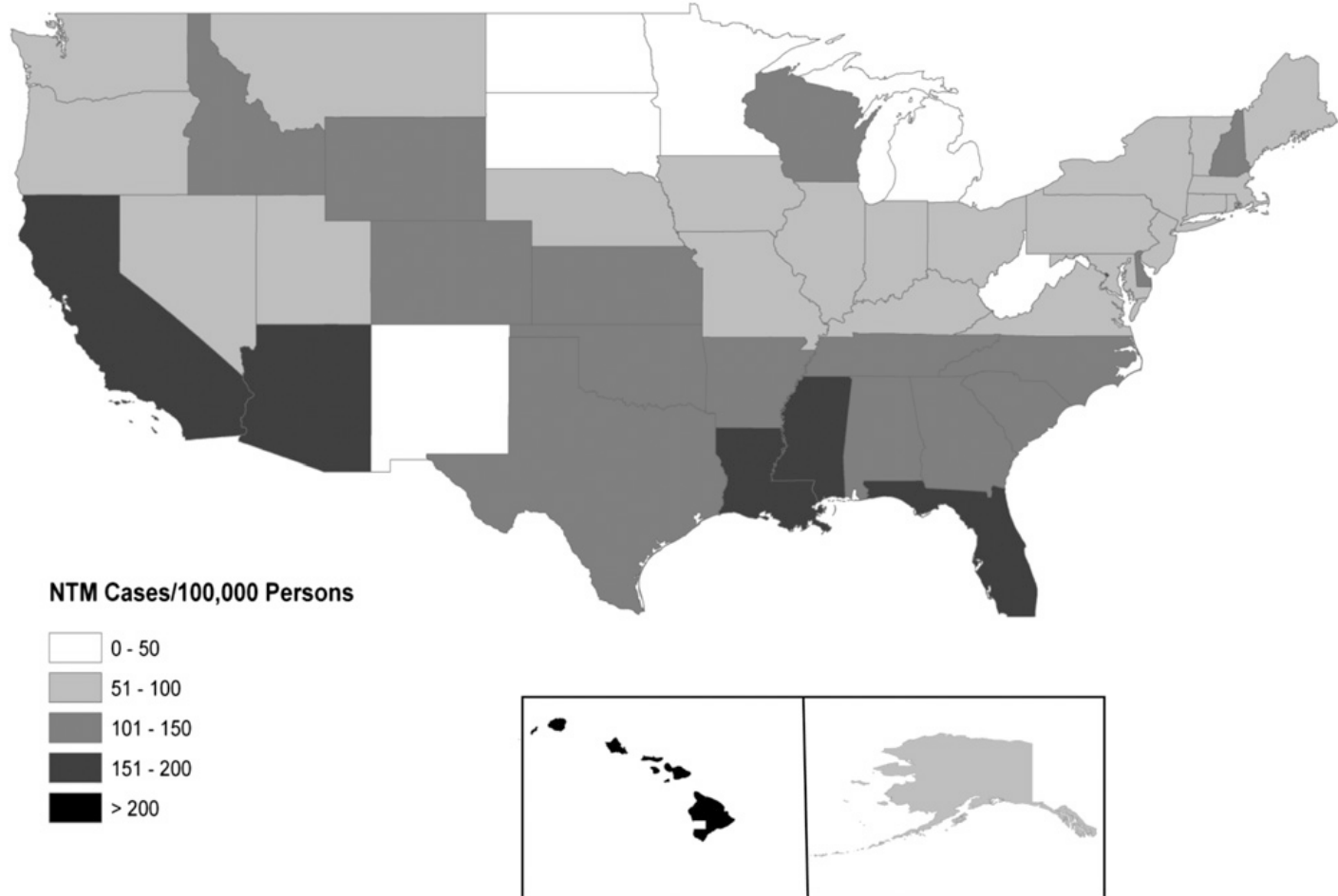
Bad news for the patient – cure rates for *M. avium* complex or *M. abscessus* range from 50-75% but relapses are common and exogenous reinfection is a major challenge

PNTM Medicare Part B



Adjemian et al AJRCCM 185:881-886(2012)
PNTM = pulmonary nontuberculous mycobacteria

PNTM Medicare Part B

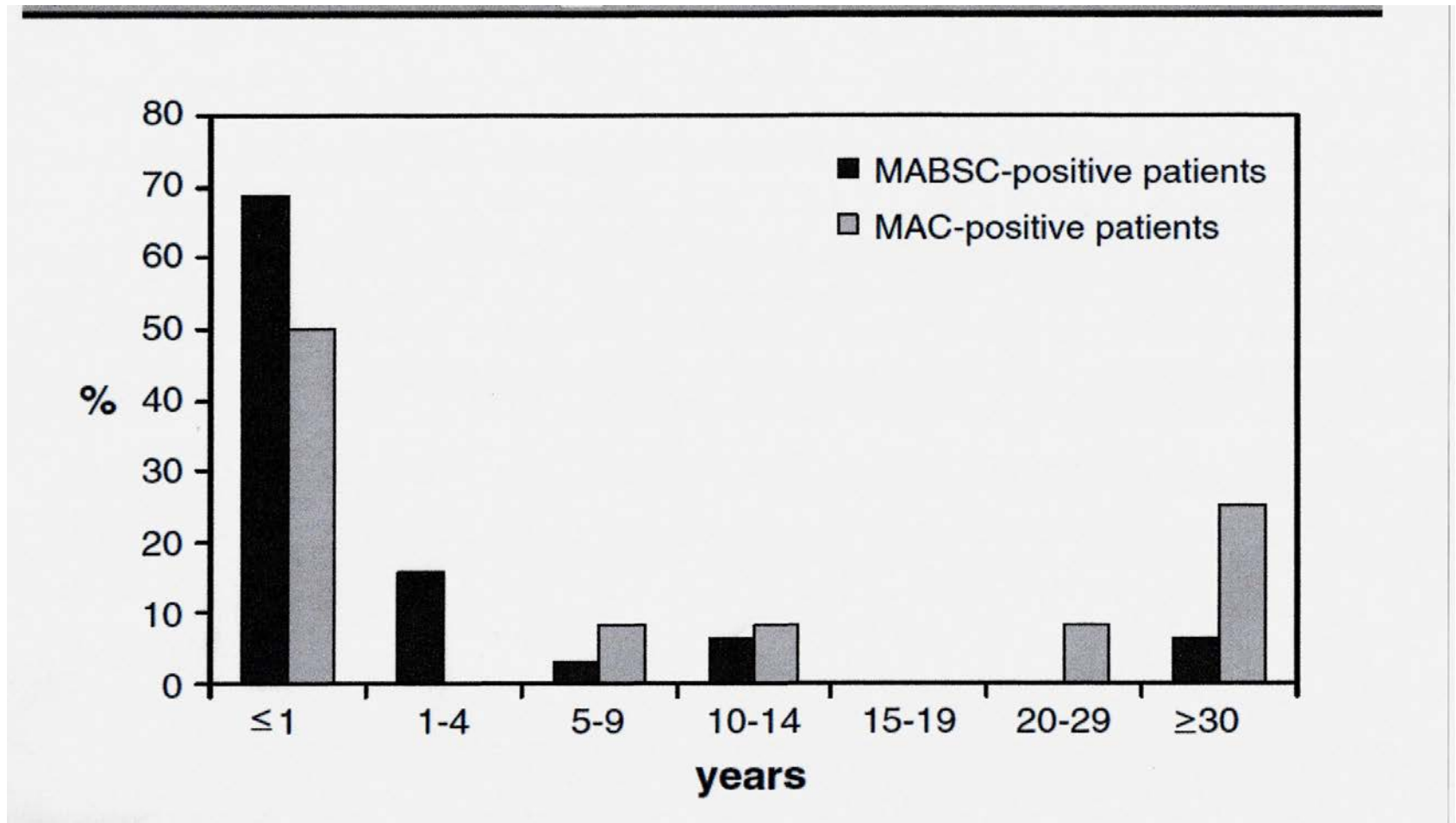


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PNTM Disease, Ontario

Five-year prevalence of pulmonary nontuberculous mycobacterial disease, Ontario, Canada, 1998–2002 and 2006–2010*

Isolate	1998–2002	2006–2010	p value
<i>M. avium complex</i>	18.0	26.5	<0.0001
<i>M. xenopi</i>	7.4	9.5	<0.0001
<i>M. fortuitum</i>	0.63	1.2	0.01
<i>M. abscessus</i>	0.63	1.2	<0.0001
Other nontuberculous <i>Mycobacterium spp.</i>	1.8	3.0	<0.0001
All nontuberculous <i>Mycobacterium spp.</i>	29.3	41.3	<0.0001



- ❑ AFB positive
- ❑ Rule out *M. tuberculosis* complex
- ❑ Sequence *rpoB* gene
 - if *M. abscessus* group, then
 - *erm(41)* and *hsp65*

> 8,800 isolates were analyzed using *rpoB* gene sequencing

Seven *Mycobacterium* species accounted for ~80% of all isolates tested

24.4%	<i>M. abscessus</i> group	}	42.3%
19.9%	<i>M. avium</i>		
16.4%	<i>M. intracellulare</i>		
6.0%	<i>M. chimaera</i>		
5.1%	<i>M. fortuitum</i>		
3.8%	<i>M. gordonae</i>		
3.7%	<i>M. chelonae</i>		



***M. massiliense* is positive for the *erm(41)* gene but contains a 273-bp deletion within the gene rendering the gene nonfunctional.**

<i>Species</i>	Number identified	Percentage
<i>M. abscessus</i>	1,470	71.7%
<i>M. massiliense</i>	420	20.5%
<i>M. bolletii</i>	110	5.4%
<i>other</i>	58	2.4%

Treatment response rates to combination antibiotic therapy including clarithromycin were much higher in patients with *M. massiliense* lung disease than those with *M. abscessus* lung disease.

Koh et al Am J Respir Crit Care Med 183:405-410 (2011)

Outcomes	<i>M. abscessus</i> (n=24)	<i>M. massiliense</i> (n=33)	P value
Symptoms			
Improved	18 (75%)	32 (97%)	0.04
Unchanged	4 (17%)	1 (3%)	
Worsened	2 (8%)	-	
HRCT Findings			
Improved	10 (42%)	27 (82%)	0.003
Unchanged	7 (29%)	5 (15%)	
Worsened	7 (29%)	1 (3%)	
Sputum conversion			
Converted	6 (25%)	29 (88%)	<0.001
Relapsed	4 (17%)	3 (9%)	
Failed	14 (58%)	1 (3%)	

Koh WJ et al. AJRCCM. 2011;183:405-410

Signs and Symptoms:

Cough (chronic); Fatigue; Weight Loss; Hemoptysis; Dyspnea

Radiology

Fibrocavitary – *M. avium* complex

Nodular & interstitial nodular infiltrates – *M. avium* complex

Fibrocavitary – *M. kansasii*

Multi-lobar, reticulonodular or mixed reticulonodular-alveolar opacities – *M. abscessus* group

The NTM species is identified by the laboratory!

- ✓ **Combination MIC**
- ✓ **Extended MIC concentrations for inhaled Amikacin**
- ✓ **Surgery**

“Surgical evaluation in nontuberculous mycobacterial (NTM) infections plays an essential role as part of multidisciplinary management of this complex pulmonary process. Resection of damaged lung parenchyma combined with appropriate antimicrobial therapy may interrupt a cycle of disease progression and relapse in select patients.”

Thorac Surg Clin. 2012 Aug;22(3):277-85

Patient Centered Approach

- ✓ **Pulmonary/ID Expert**
- ✓ **Expert Laboratory**
- ✓ **Experienced Surgeon**



**Mycobacteriology &
Pharmacokinetics staff
(5 not present)**

Thank you!

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