

# Frequency of Mixed *Mycobacterium* Species Infections Detected by the INNO-LiPA MYCOBACTERIA v2 Hybridization Line Probe Assay

## January 2015 to October 2016

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### Abstract

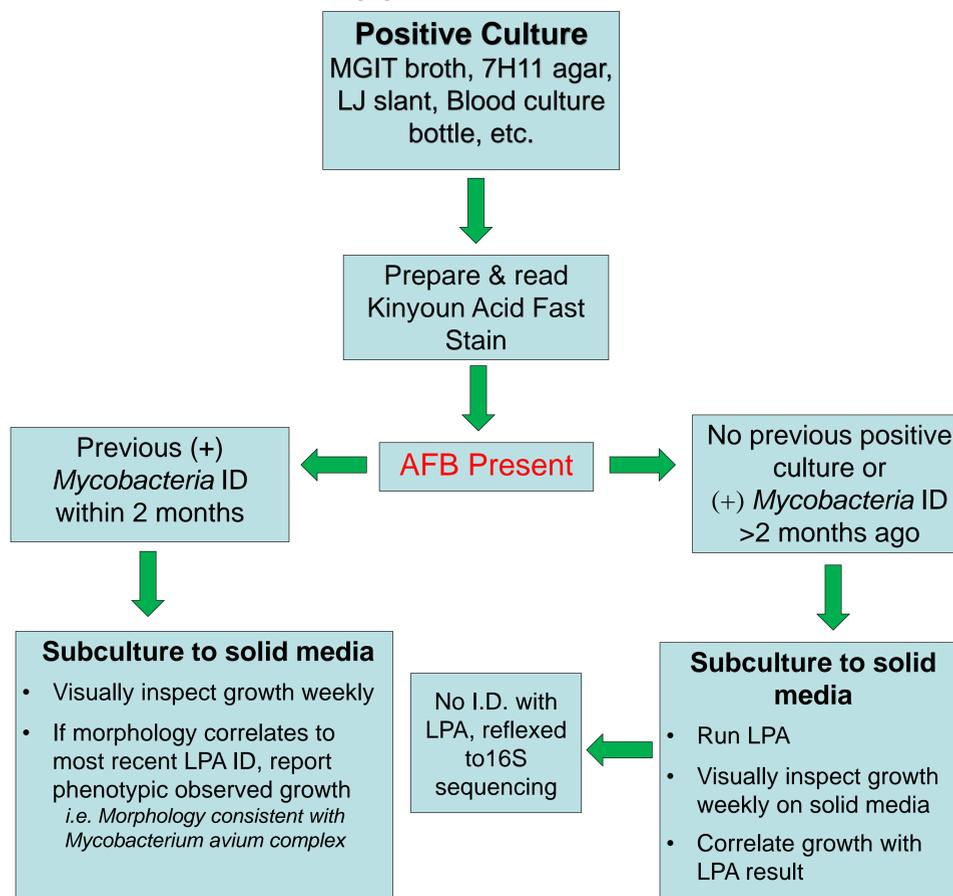
**Objective:** To evaluate the frequency of mixed mycobacteria infections detected from 2015 to 2016 using the Fujirebio (Ghent, Belgium) manufactured INNO-LiPA MYCOBACTERIA v2 line probe hybridization assay (LPA), which identifies 18 clinically relevant *Mycobacterium* species.

**Study Design:** PCR is used to amplify the 16S-23S internal transcribed spacer region of *Mycobacterium* species. This biotinylated amplified material is reverse hybridized with specific oligonucleotide probes immobilized on membrane strips. A final colorimetric step produces characteristic bands that are associated with specific *Mycobacterium* species. More than one *Mycobacterium* species can be detected on a single strip. Twenty-two months of data (January 2015 through October 2016) was assessed for the number of specimens where more than one species of mycobacteria was detected by the LPA, compared to the total number of LPA positive specimens.

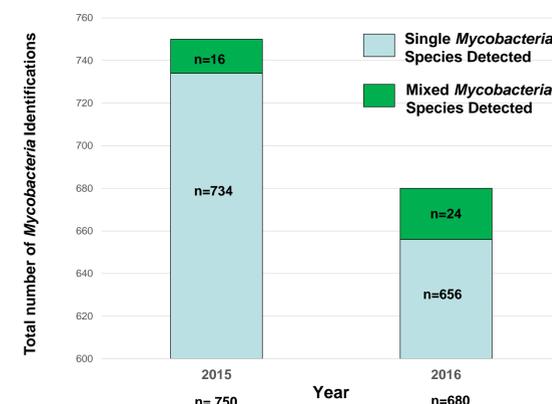
**Results:** During the study period, a total of 1,430 LPA identifications were performed, of which the LPA detected 3% (40) mixed mycobacterial infections. Most of the mixed populations were *M. avium complex* with *M. intracellulare* infections followed by *M. avium complex* with *M. abscessus* infections. Other clinically relevant mixed infections were also detected in this study such as *M. tuberculosis complex* with *M. avium complex* in far lower numbers but nonetheless considered significant for patient care.

**Conclusions:** In Minnesota (MDH TB Lab), mixed mycobacteria infections do occur with low frequency, which can be easily and rapidly detected using the INNO-LiPA MYCOBACTERIA v2 hybridization line probe assay. Accurate mycobacterial identification and detection of mixed infections is essential for not only epidemiological investigations but for correct diagnosis and appropriate patient treatment.

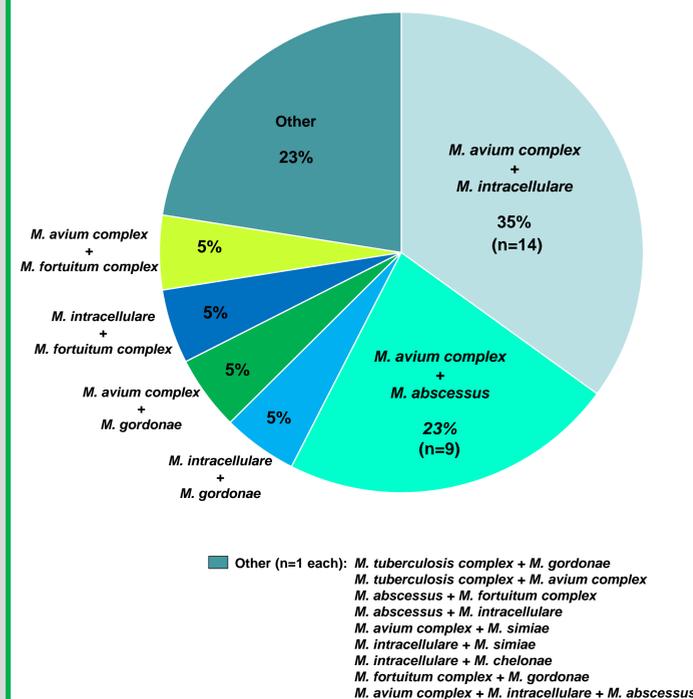
### 2. AFB (+) Culture Workflow



### 4. Total Line Probe Identifications January 2015 - October 2016



### 5. Mixed *Mycobacterium* spp. Detected January 2015 - October 2016



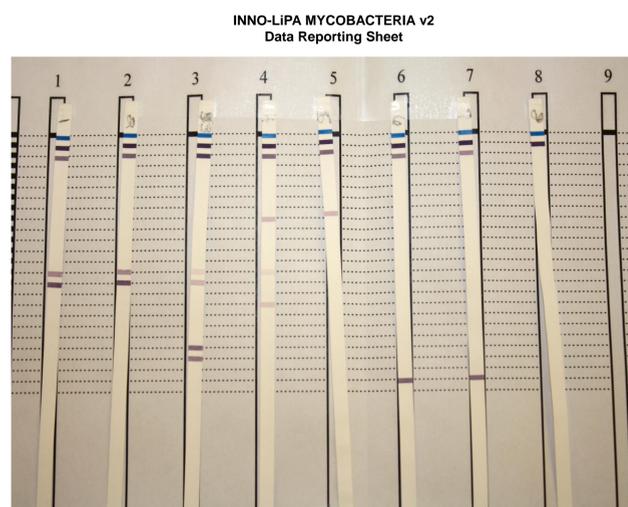
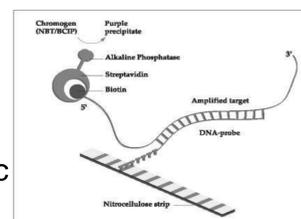
### 1. Interpretation Chart: Banding Patterns on INNO-LiPA Probe Strips

Assay can detect 18 clinically relevant *Mycobacterium* spp.

Line	Probe Names	<i>Mycobacterium</i> sp.	MTBC	<i>M. kansasii</i>	<i>M. kansasii/gastri</i>	<i>M. xenopi</i>	<i>M. goodii</i>	<i>M. genavense</i>	<i>M. simiae</i>	<i>M. marinum/M. neoaurum</i>	MAC	MAC	<i>M. intracellulare</i>	<i>M. intracellulare</i>	<i>M. scrofulaceum</i>	<i>M. malmoense</i>	<i>M. haemophilum</i>	<i>M. chelonae</i>	<i>M. abscessus</i>	<i>M. chelonae</i>	<i>M. fortuitum complex</i>	<i>M. smegmatis</i>
1	Conj	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1	1
2	MYC	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2	2
3	MTB	3																				
4	MAA-1	4																				
5	MAA-2	5																				
6	MAA-3	6																				
7	MXE	7																				
8	MGO	8																				
9	MGV	9																				
10	MSI	10																				
11	MSU	11																				
12	MCE	12																				
13	MAIS	13											13	13	13	13	13	13*				
14	MAV	14																				
15	MIN-1	15																				
16	MIN-2	16																				
17	MSC	17																				
18	MML	18																				
19	MHP	19																				
20	MCH-1	20																	20	20		
21	MCH-2	21																		21		
22	MCH-3	22																			22	
23	MFO	23																			23	23
24	MSM	24																				24

### 3. INNO-LiPA Line Probe Assay Methods

- Step 1:** Amplification of 16S-23S rRNA spacer region
- Step 2:** Hybridization & stringent wash of amplified 16S-23S spacer region with specific probes immobilized on strip
- Step 3:** Color development of membrane strips
- Step 4:** Interpretation of signal pattern



- M. avium complex*
- M. avium complex*
- M. avium complex* + *M. chelonae*
- M. intracellulare* + *M. gordonae*
- M. gordonae*
- M. fortuitum complex*
- M. fortuitum complex*
- Not *Mycobacterium* species

### 6. Conclusions

- The INNO-LiPA MYCOBACTERIA v2 hybridization line probe assay (Fujirebio) serves as our primary means of identification.
  - One strip replaces multiple tests; easy to perform
  - Detection of 18 clinically relevant mycobacterial species
  - Applicable to early liquid culture and solid media
  - Quick visual interpretation
  - Mostly automated using the Auto-LiPA 48
  - Mixed populations easily identified**
- Mixed mycobacteria infections are detected in the MN TB Laboratory at a frequency of 3%, with *M. avium complex* + *M. intracellulare* (35%) and *M. avium complex* + *M. abscessus* (23%) being the two most common mixed populations.
- Accurate and rapid laboratory detection of mixed mycobacterial infections can impact patient management and appropriate public health control measures.

### Acknowledgements:

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