

# Laboratory Diagnostics – Whole Genome Sequencing Related to Drug Resistance and Epidemiology

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slide adapted from N. Wengenack, PhD

# Accreditation Statement



## Accreditation Statement

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### AMA

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**UAN Number:** JA0000238-0000-26-035-L99-P

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IPCE CREDIT™

This activity was planned by and for the healthcare team, and learners will receive 1.0 Interprofessional Continuing Education (IPCE) credit for learning and change.

## Other Healthcare Professionals:

A record of attendance will be provided to all registrants for requesting credits in accordance with state nursing boards, specialty societies or other professional associations.

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## Available Credit

- 1.00 ACPE,
- 1.00 *AMA PRA Category 1 Credit™*
- 1.00 ANCC
- 1.00 Attendance
- 1.00 IPCE

# Disclosures

- No relevant financial disclosures to report and no mention of off-label use of any medications or products

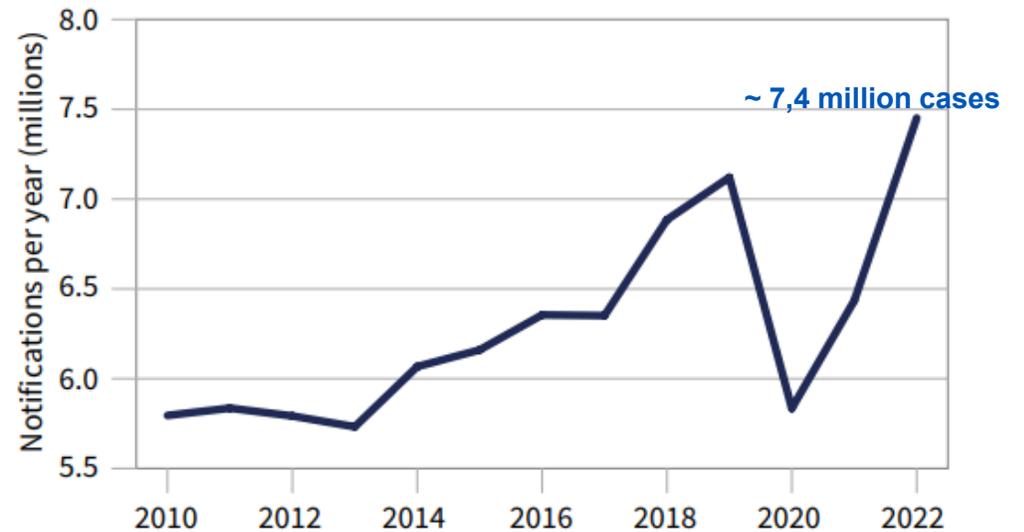
# Learning Objectives

1. Identify essential tools for TB diagnosis
2. Describe molecular testing methods
3. Explain the role of imaging in TB diagnosis

# Global Impact of M. tuberculosis

- Leading cause of death from a single infectious agent
- Each year:
  - 10.6 million cases of active TB worldwide
  - 1.3 million deaths
- >400,000 cases of MDR TB and RR-TB (3.6% of new cases)
- ~ 1.7 billion people have been exposed, don't know extent of latent TB vs cleared infections (1/4 of world population)

**Global trend in case notifications of people newly diagnosed with TB, 2010–2022**



6.2 million people diagnosed with pulmonary TB worldwide in 2022  
~ 80% of Tb case

# WHO: END TB Targets

## A WORLD FREE OF TB

ZERO deaths, disease, and suffering due to TB

## END THE GLOBAL TB EPIDEMIC

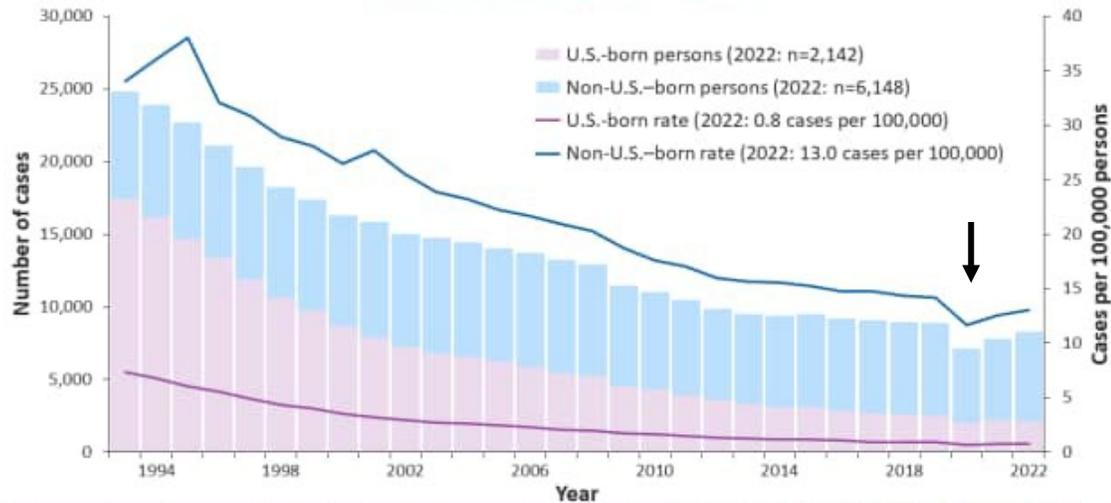
	TARGETS			
	MILESTONES		SDG*	END TB
	2020	2025	2030	2035
<b>Reduction in number of TB deaths</b> compared with 2015 (%)	35%	75%	<b>90%</b>	<b>95%</b>
<b>Reduction in TB incidence rate</b> compared with 2015 (%)	20%	50%	<b>80%</b>	<b>90%</b>

## Diagnosis:

- Access to rapid & accurate diagnostics
  - Target >90% of cases
  - Used in only 47% of cases in 2022
- Universal access to DST
  - Target 100%
  - 2022 WHO Report – RIF, 73%

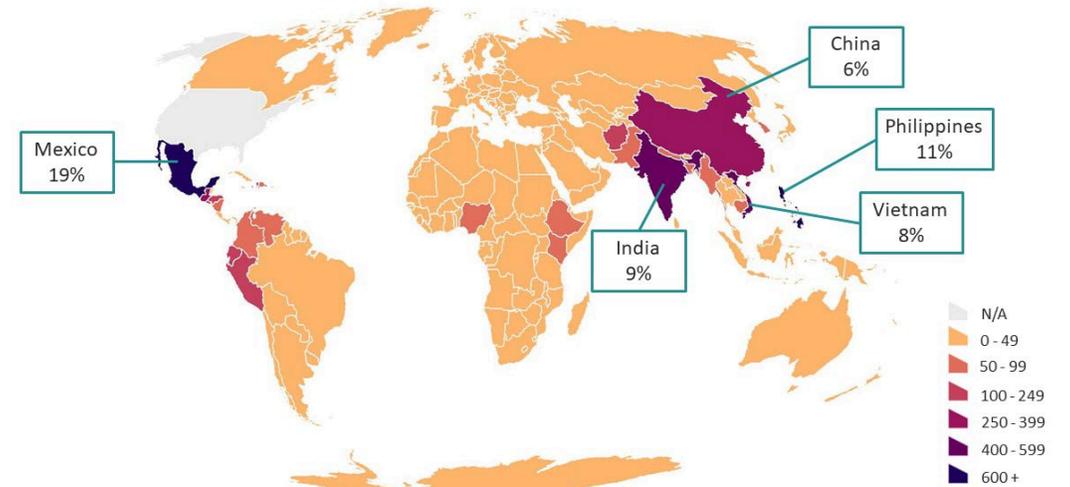
# TB in the US

TB Cases and Incidence Rates by Origin of Birth,\*  
United States, 1993–2022



\*Persons born in the United States, certain U.S. territories, or elsewhere to at least one U.S. citizen parent are categorized as U.S.-born. All other persons are categorized as non-U.S.-born.

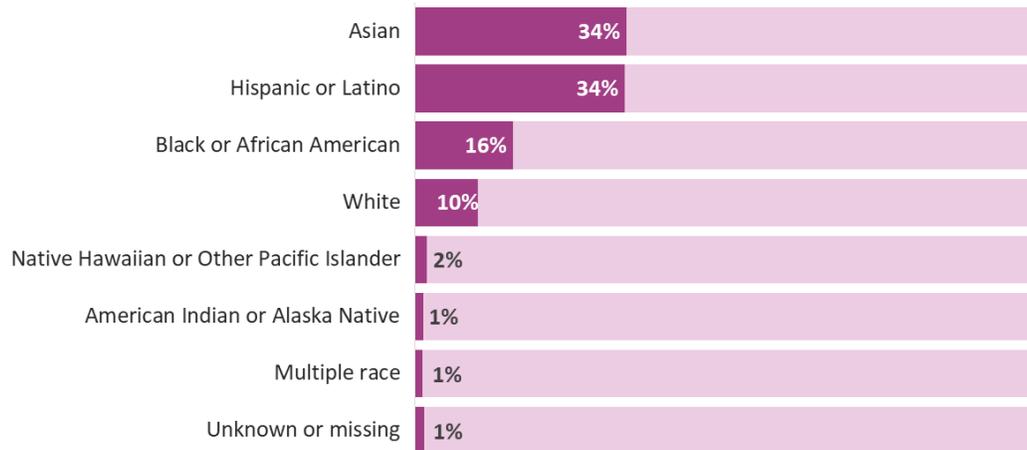
TB Cases by Countries of Birth Among Non-U.S.-Born\* Persons with  
TB, United States, 2022 (N=6,148)



\*Persons born in the United States, certain U.S. territories, or elsewhere to at least one U.S. citizen parent are categorized as U.S.-born. All other persons are categorized as non-U.S.-born.

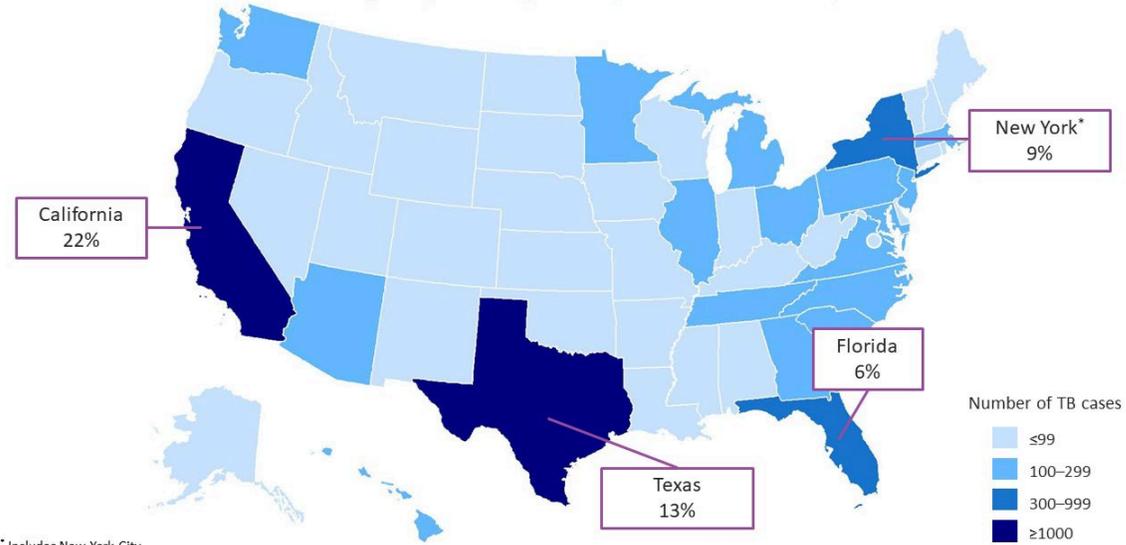
**In 2022, 74% of new cases in the US were non-US born**

### Percentage of TB Cases by Race/Ethnicity,\* United States, 2022 (N=8,331)



\*Persons who identified as Hispanic or Latino were categorized as "Hispanic," regardless of self-reported race. Persons who did not identify as Hispanic or Latino were categorized by self-reported race; if more than one race was reported, the person was categorized as "Multiple race."

### TB Cases by Reporting Area, United States, 2022



\* Includes New York City

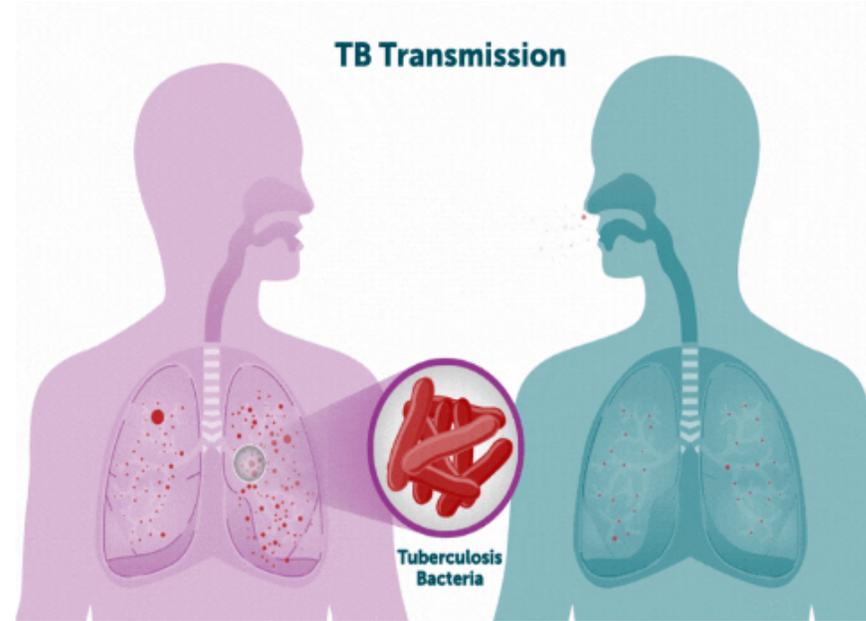
# TB Transmission

TB is spread through the air from person to person. Tiny particles containing *M. tuberculosis* (TB bacteria) may be expelled into the air when a person with infectious TB of the lungs, airway, or larynx:

- Coughs
- Speaks
- Sings

People nearby may breathe in these TB bacteria (illustrated by the person on the right) and become infected.

These particles, called airborne **droplet nuclei**, can remain in the air for several hours, depending on the environment.



*The dots in the air represent airborne droplet nuclei spreading through the air.*

TB is **NOT** spread by:



Sharing toothbrushes



Saliva from kissing



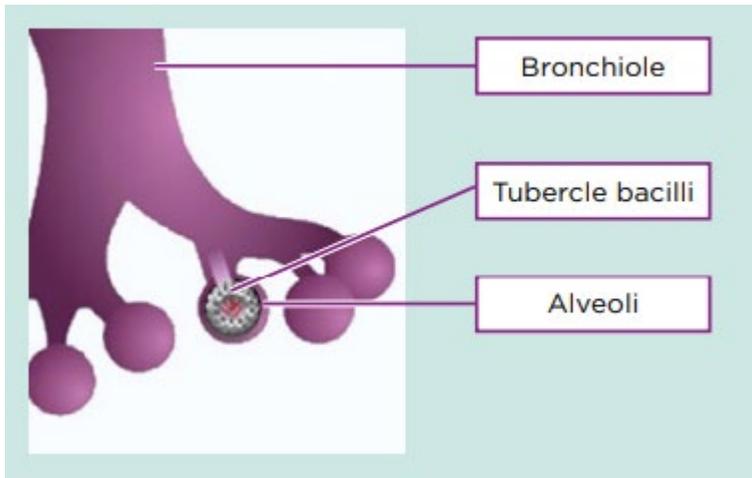
Shaking someone's hand



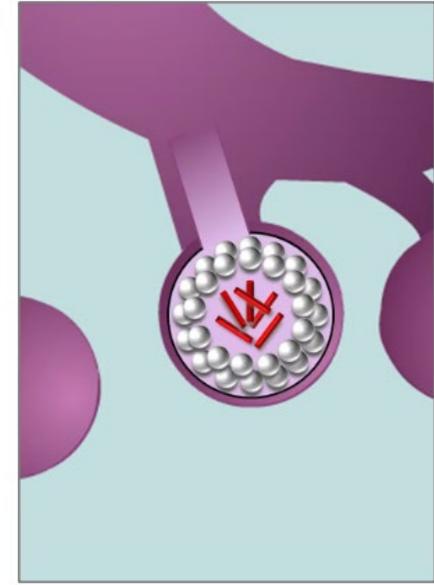
Touching bed linens or toilets



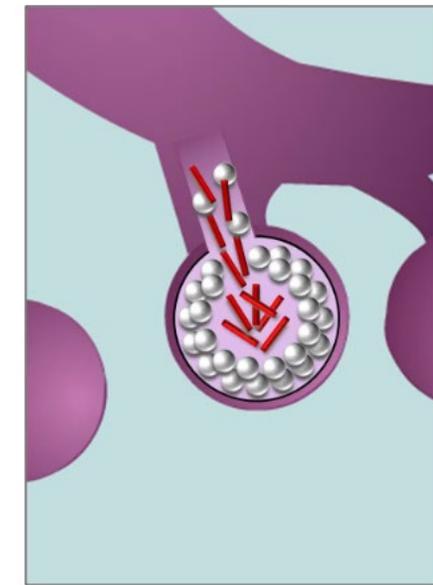
Sharing food, drink, or utensils



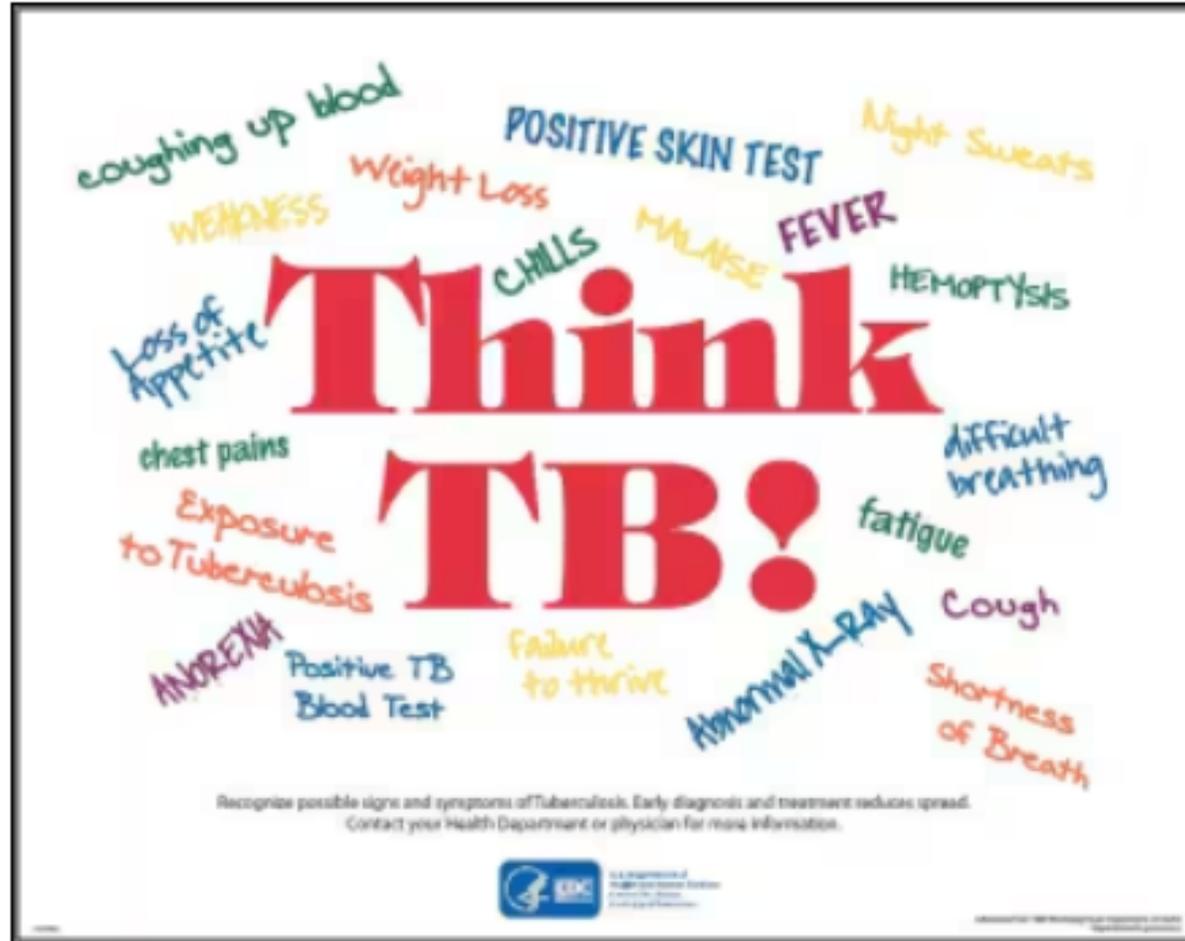
Latent TB



Active TB



Risk Factor	Risk of Developing TB Disease	Description
TB infection and no risk factors	About 10% over a lifetime 	For people with TB infection, <b>no risk factors</b> , and no treatment, the risk is about 5% in the first 2 years after infection and about 10% over a lifetime.
TB infection and diabetes	About 30% over a lifetime 	For people with TB infection, <b>diabetes</b> , and no LTBI treatment, the risk is about 30% over a lifetime (3 times as high as those with no risk factors).
TB infection and HIV infection	About 7% to 10% PER YEAR 	For people with TB infection, <b>untreated HIV infection</b> and with no LTBI treatment, the risk is about 7% to 10% PER YEAR, a very high risk over a lifetime.



*The key to diagnosing TB is for clinicians to “think TB” when they see a patient with signs and symptoms of TB disease.*

# Case Study

IGRA

54-year-old man with DM, HTN and CKD5 admitted for worsening hyperkalemia with plans to start HD.

As part of his pre-dialysis evaluation, he undergoes routine screening, including a QuantiFERON-TB Gold test. Results shown below

<b>QuantiFERON-TB Gold Plus Result</b>	<b>Indeterminate</b>
<b>TB1 Ag minus Nil Result</b>	<b>0.00</b>
<b>TB2 Ag minus Nil Result</b>	<b>0.00</b>
<b>Mitogen minus Nil Result</b>	<b>0.10</b>
<b>Nil Result</b>	<b>0.02</b>

# Which of the following is true regarding this result?

<b>QuantiFERON-TB Gold Plus Result</b>	<b>Indeterminate</b>
<b>TB1 Ag minus Nil Result</b>	<b>0.00</b>
<b>TB2 Ag minus Nil Result</b>	<b>0.00</b>
<b>Mitogen minus Nil Result</b>	<b>0.10</b>
<b>Nil Result</b>	<b>0.02</b>

- A. Patient is likely immunosuppressed
- B. This is due to overstimulated T cells
- C. Indeterminate test means weakly positive

# Which of the following is true regarding this result?

QuantiFERON-TB Gold Plus Result	Indeterminate
TB1 Ag minus Nil Result	0.00
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Mitogen minus Nil Result	0.10
Nil Result	0.02

- A. Patient is likely immunosuppressed
- B. This is due to overstimulated T cells
- C. Indeterminate test means weakly positive

# Interferon gamma release assays (IGRA)



**QuantiFERON® TB-  
Gold Plus (QFT-Plus;  
Qiagen)**



**T.SPOT ®-TB  
(Oxford Immunotec)**

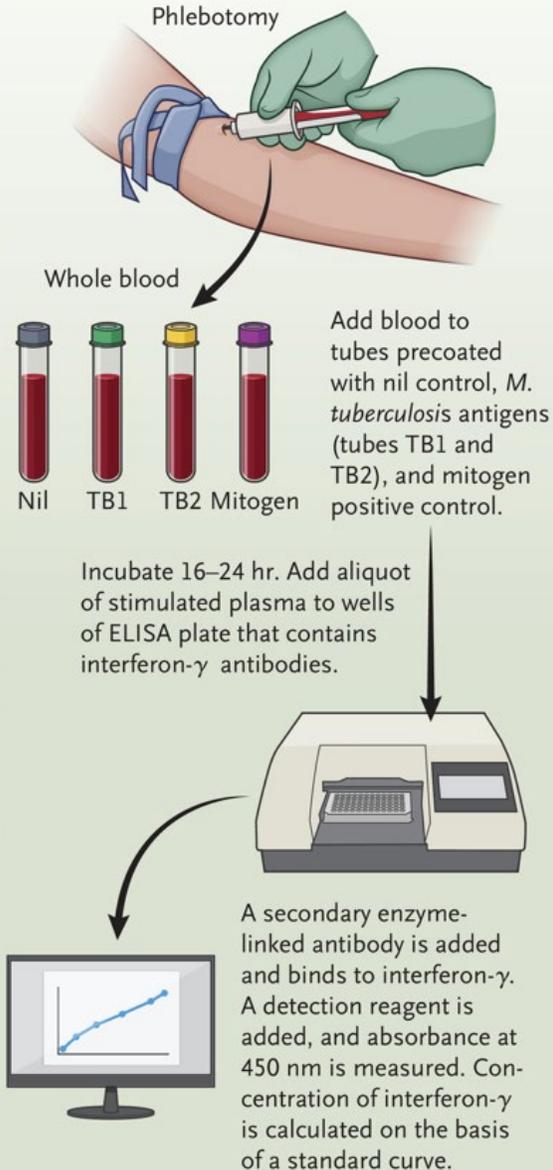
**IGRAs** are **indirect** tests for infection with MTB

Patients infected with MTb have primed T-cells

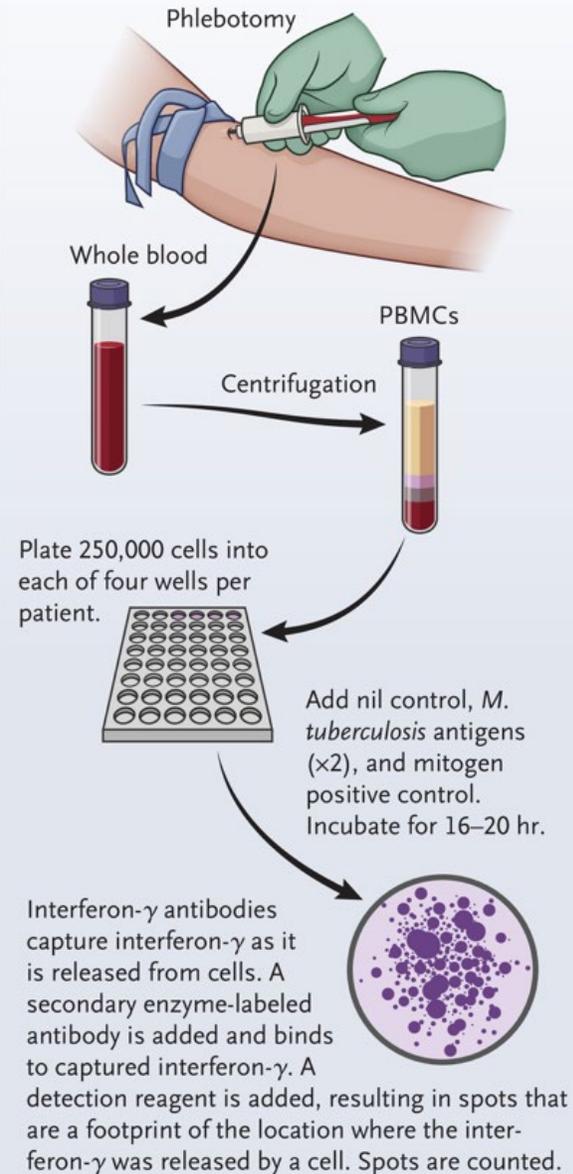
Exposure of primed T-cells to MTb antigens induces  
IFN- $\gamma$  production

IFN- $\gamma$  detected by EIA/CIA or ELISPOT methods

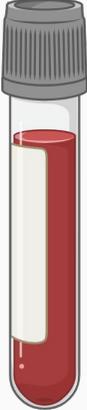
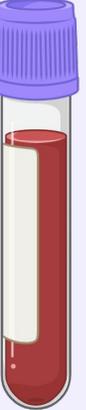
## QuantiFERON-TB Gold Plus IGRA



## T-SPOT.TB IGRA

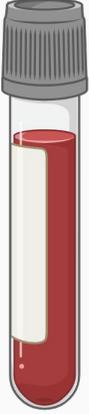
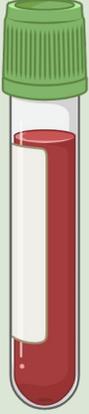
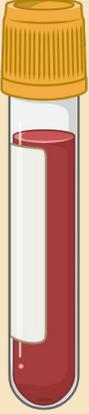
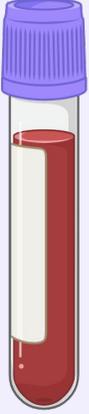


# QuantiFeron-Tb Gold Plus

	Nil Control	Tb 1 Antigen	Tb 2 Antigen	Mitogen Control
				
<b>Antigens</b>	None			PHA phythaemagglutinin-P
<b>T cell stimulated</b>	N/A			Universal T-cell stimulant
<b>Purpose</b>	Adjusts for background IFN- $\gamma$ levels			Low response may indicate inability to generate IFN-

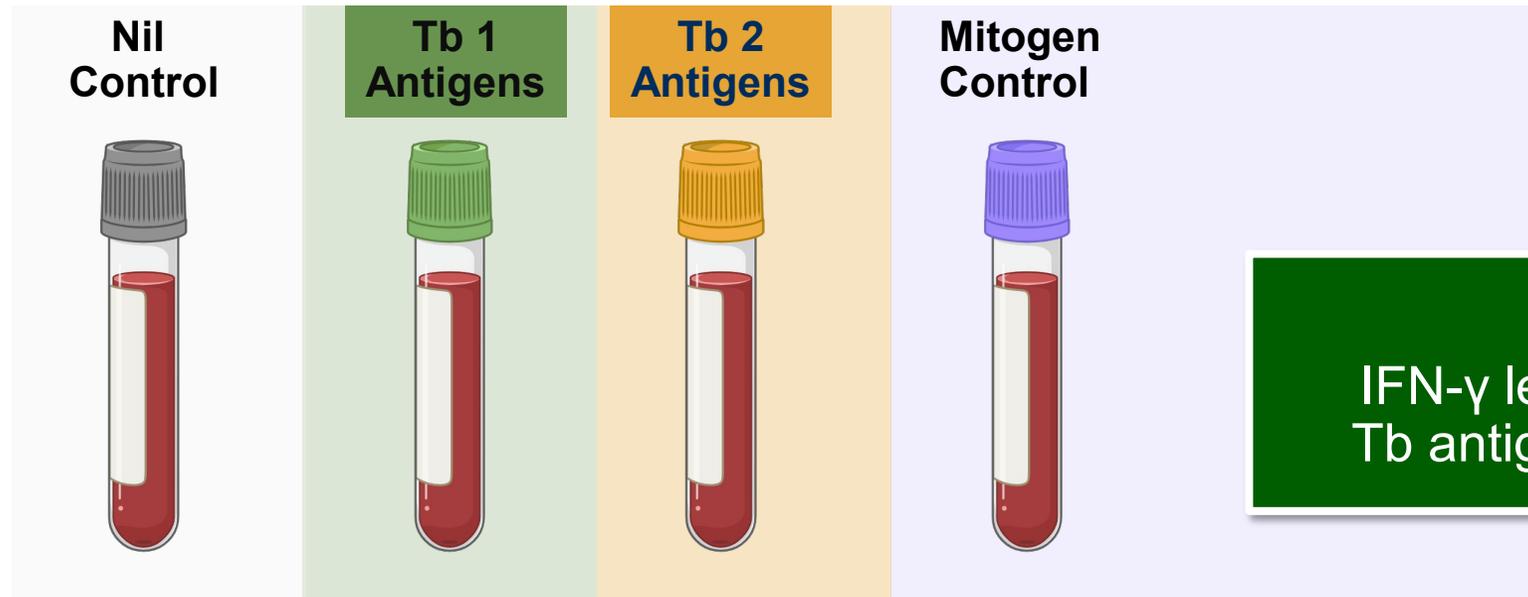
slide courtesy of M. Fida, MD

# QuantiFeron-Tb Gold Plus

	<b>Nil Control</b> 	<b>Tb 1 Antigen</b> 	<b>Tb 2 Antigen</b> 	<b>Mitogen Control</b> 
<b>Antigens</b>	None	<b>ESAT-6</b> <b>CFP-10</b>	<b>ESAT-6</b> <b>CFP-10</b>	<b>PHA</b> phythaemagglutinin-P
<b>T cell stimulated</b>	N/A	<b>CD4+</b>	<b>CD4+/ CD8+</b>	Universal T-cell stimulant
<b>Purpose</b>	Adjusts for background IFN- $\gamma$ levels			Low response may indicate inability to generate IFN-

slide courtesy of M. Fida, MD

# QuantiFeron-Tb Gold Plus



**Negative Result:**  
 IFN- $\gamma$  levels  $<0.35$  IU/mL in *both* Tb antigen tubes and  $<25\%$  of Nil

<b>Antigens</b>	None	ESAT-6 CFP-10	ESAT-6 CFP-10	PHA phythaemagglutinin-P
<b>T cell stimulated</b>	N/A	CD4+	CD4+/ CD8+	Universal T-cell
<b>Purpose</b>	Adjusts for background IFN- $\gamma$ levels			Low response may indicate inability to generate IFN-

**Positive Result:**  
 IFN- $\gamma$  levels  $\geq 0.35$  IU/mL in *either one or both* Tb antigen tubes

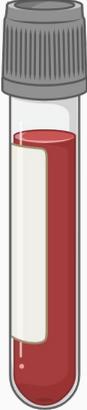
slide courtesy of M. Fida, MD

# QuantiFERON-Tb Gold Plus

**Indeterminate results due to poor mitogen response**

Immunosuppressed patients  
Technical issues

**What to do?**  
Consider getting T spot test

	Nil Control	Tb 1 Antigen	Tb 2 Antigen	Mitogen Control
				
<b>Antigens</b>	None	ESAT-6 CFP-10	ESAT-6 CFP-10	PHA phythaema [g] [un] [r]?
<b>T cell stimulated</b>	N/A	CD4+	CD4+/ CD8+	Universal T-cell stimulant
<b>Purpose</b>	Adjusts for background IFN- $\gamma$ levels			Low response may indicate inability to generate IFN-

Mitogen not reacting

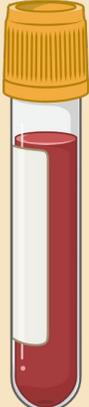
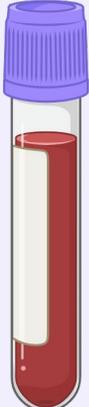
# QuantiFERON-Tb Gold Plus

**Indeterminate results due to Nil reacting**

Elevated baseline IFN- $\gamma$  levels, such as certain infections, autoimmune diseases, or other inflammatory conditions

## What to do?

- Consider repeating QuantiFERON
- Get T-spot test

	Nil Control	Tb 1 Antigen	Tb 2 Antigen	Mitogen Control
				
<b>Antigens</b>	None <b>reacting</b>	<b>ESAT-6</b> <b>CFP-10</b>	<b>ESAT-6</b> <b>CFP-10</b>	PHA phythaemagglutinin-P
<b>T cell stimulated</b>	N/A	<b>CD4+</b>	<b>CD4+/ CD8+</b>	Universal T-cell stimulant
<b>Purpose</b>	Adjusts for background IFN- $\gamma$ levels			Low response may indicate inability to generate IFN-

slide courtesy of M. Fida, MD

# 2021 Meta-Analysis on Performance of QFT-Plus vs. Alternative LTBI Assays

- 24 studies included
  - High- and low-income countries
  - QFT-Plus vs. QFT-Gold vs. T-Spot

		No. of Studies	Pooled Values
Sensitivity (vs. culture)	QFT-Plus	7	92.6%
	QFT-Gold	7	91.8%
	T.Spot	2	90.2%
Specificity	QFT-Plus	2	97.8%
	QFT-Gold	2	98.7%
	T.Spot	1	98.1%

# Variability for the QFT Assays

Assay cut-off for positive:  $\geq 0.35$  IU/mL

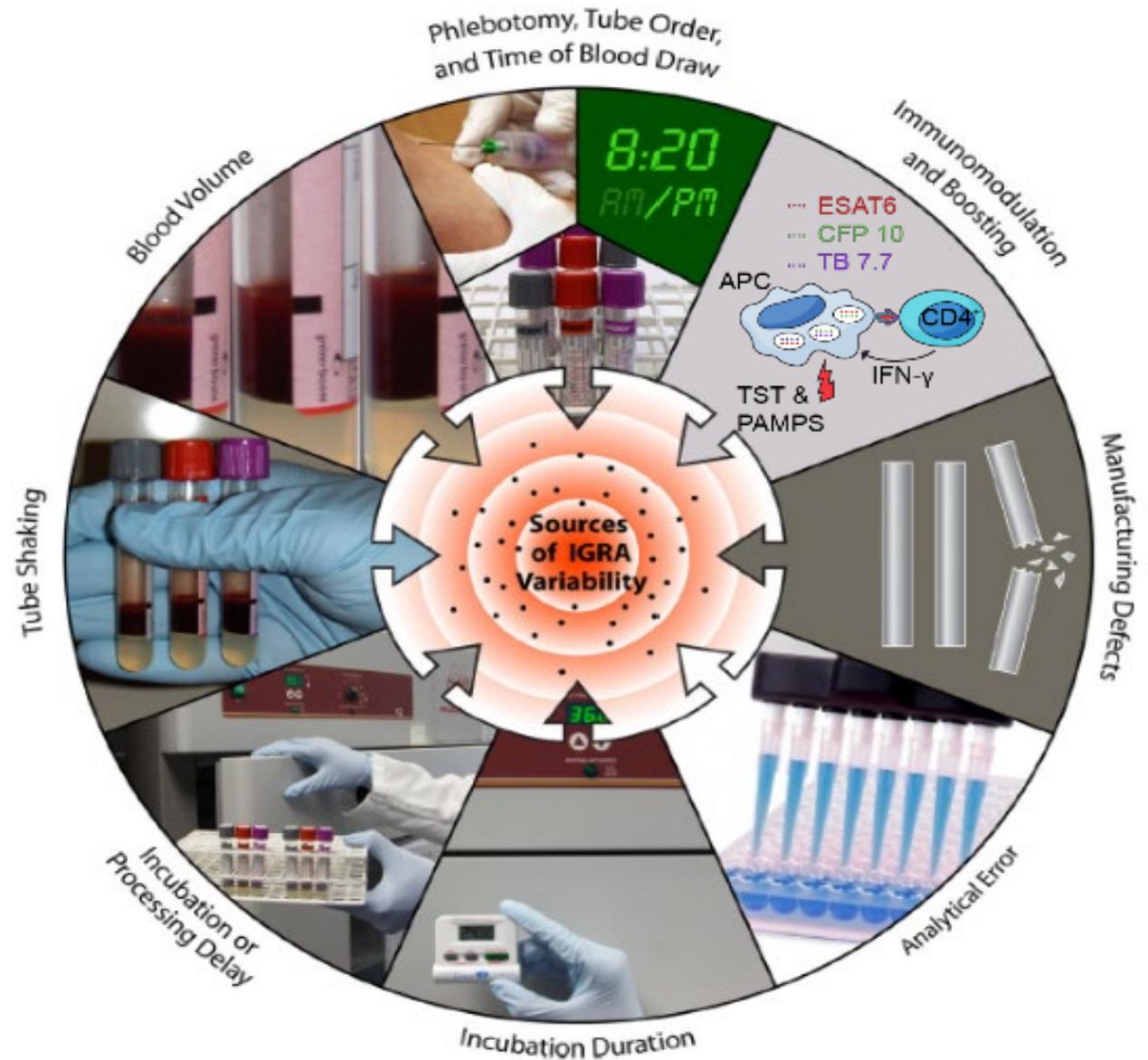
With-in subject variability:  $\pm 0.60$  IU/mL

**Interpret positive QFT values between 0.35 - 0.95 IU/mL with caution!**

High rates of spontaneous conversion/reversion:

QFT-Plus: 22/196 (11%)

QFT-Gold: 16/188 (8.5%)



## Sources of variability for the QFT Assays

Pai M. et. al. *Clin Micro Rev.* 2014;27(1):3-39, Oh CE, et al. *Clin Infect Dis.* 2021;73:e116-e1125

Metcalfe JZ et. al. *Am J Respir Crit Care Med.* 2013;187(2): 206-211

# Detection of Latent TB Infection via IGRAs

- **False Positives:**
  - *M. marinum, M. kansasii, M. szulgai, M. flavescens*
  - Pre-analytic processing errors
- **False Negatives:**
  - Immunosuppression
  - Pre-analytic processing errors
- **Can IGRAs be used to monitor response to therapy?**
  - “...monitoring IGRA changes over time seems to have a speculative value only.” (Chiappini, Clin. Therapy 2012)

# Bacteriological examination

**Patient:** 27-year-old woman, born in Somalia, immigrated to U.S. 8 years ago, visited Kenya recently

**Current symptoms:** Dry cough, shortness of breath, fatigue x 3 months and intermittent night sweats

**CT imaging:** Clustered nodularity in the medial right apex measuring up to 6 x 4 mm, solid 17 x 14 mm suprahilar right upper lobe nodule/lymphadenopathy, separate right hilar lymphadenopathy, and multistation mediastinal lymphadenopathy.

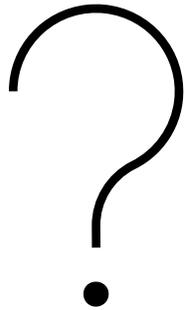
No previous QuantiFERON, Current QuantiFERON is positive

**What is the best next step?**



**Clinical case**

# Polling Question



Given the previous history of TB with incomplete therapy, underlying suspicion for TB was high.

Which of the following will be the best next step?

**A**

3 x sputum samples obtained  $\geq 8$  hours apart for AFB smear and mycobacterial cultures

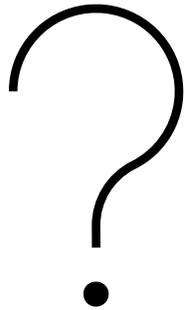
**B**

3 x sputum samples obtained  $\geq 8$  hours apart for AFB smear with mycobacterial cultures and nucleic acid amplification test (NAAT)

**C**

1 x sputum samples for AFB smear and mycobacterial cultures and MTB PCR

# Polling Question



Given the previous history of TB with incomplete therapy, underlying suspicion for TB was high.

Which of the following will be the best next step?

**A**

3 x sputum samples obtained  $\geq 8$  hours apart for AFB smear and mycobacterial cultures

**B**

3 x sputum samples obtained  $\geq 8$  hours apart for AFB smear with mycobacterial cultures and nucleic acid amplification test (NAAT)

**C**

1 x sputum samples for AFB smear and mycobacterial cultures and MTB PCR

- The specimens should be examined and cultured in a laboratory that specializes in testing for *M. tuberculosis*.
- Optimal bacteriologic examination has **5** parts:
  1. Specimen collection, transport, and processing
  2. AFB smear classification
  3. Direct detection is in clinical specimens using nucleic acid amplification tests (NAAT) and, as applicable, molecular detection of resistance
  4. Specimen culture and identification of *M. tuberculosis*
  5. Drug susceptibility testing using growth-based and molecular methods (phenotypic and genotypic)



*Patient coughing up sputum*

# Laboratory Safety for Mycobacteria

*M. tuberculosis* / *M. bovis* are BSL III pathogens when growing in culture



- respiratory protection
- work performed in BSC
- negative pressure room, directional HEPA-filtered exhaust
- solid-front gowns, gloves, etc
- tuberculocidal disinfectants

- At least **3** consecutive sputum specimens are needed, each collected in 8- to 24-hour intervals, with at least one being an early morning specimen.
- If possible, specimens should be obtained in an airborne infection isolation room
- Other acceptable specimens: induced sputum, bronchoalveolar lavage, gastric aspirate



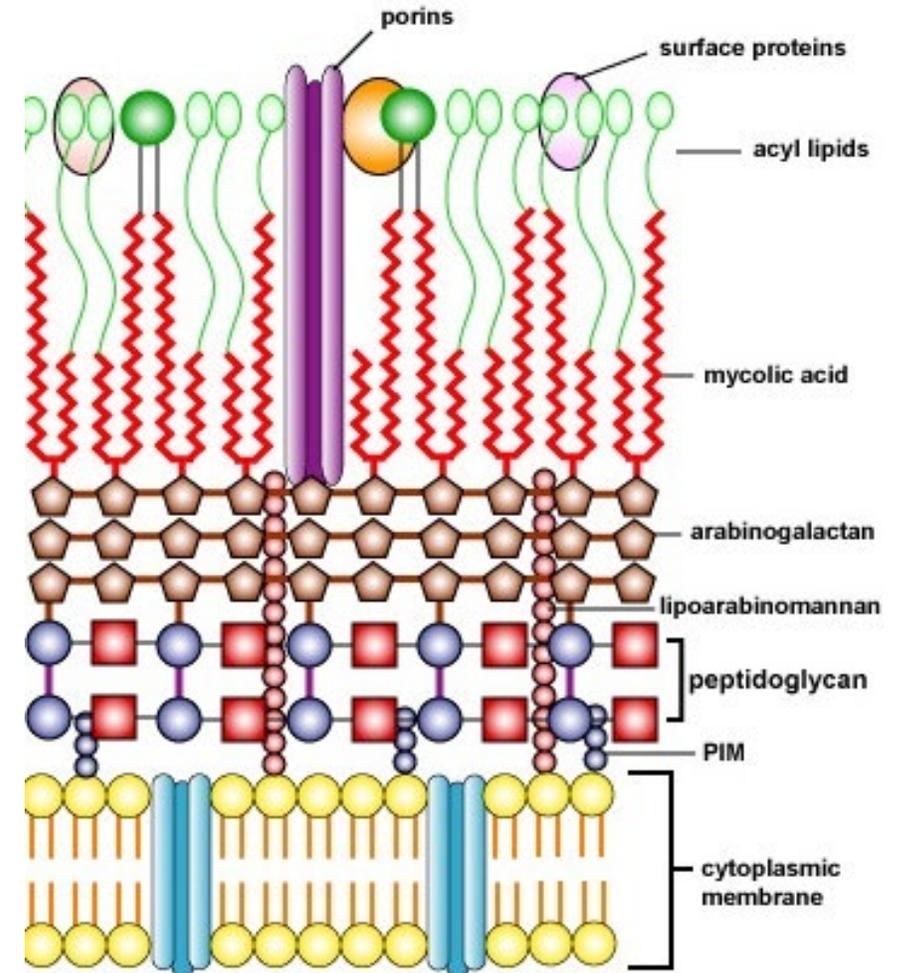
*Patient in a sputum collection booth*

# *Mycobacterium tuberculosis* complex

## General Characteristics

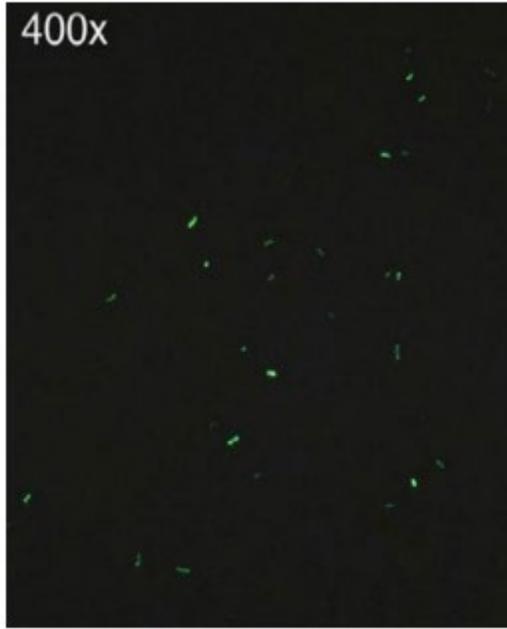
Mycobacteria, including *M. tuberculosis*, are acid-fast bacilli (AFB)

- complex cell wall with very high lipid content that functions as a permeability barrier
- very resistant to drying, desiccation, & many disinfectants and antibiotics
- use special "acid-fast" stains for mycobacteria
  - Auramine-rhodamine (fluorescent)
  - Ziehl-Neelsen
  - Kinyoun
  - Fite (Pathology)

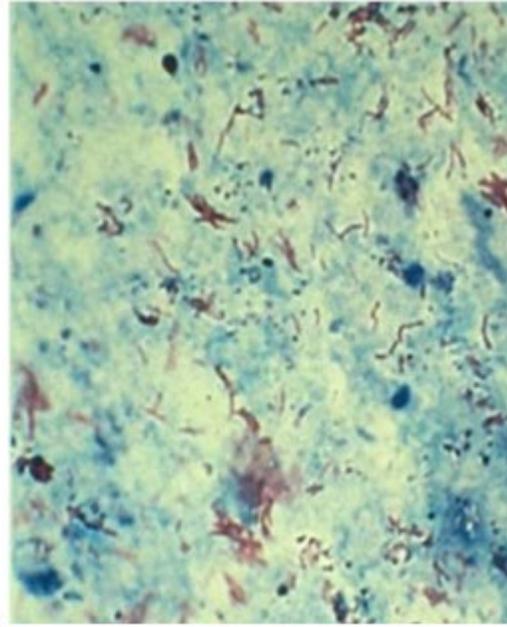


<https://bio.libretexts.org/Bookshelves/Microbiology/>

slide courtesy of N. Wengenack, PhD



Fluorochrome method



Ziehl-Neelsen method with tubercle bacilli shown in red

- AFB stain is relatively inexpensive but not very sensitive (need ~1,000 - 10,000 AFB/mL sputum)
- Fluorescent stains are 10% more sensitive
- Minimum sputum sample is 3 mL; optimal 5-10 mL
- Sensitivity of smear on:
  - Single sample: 53%
  - Two samples: 65%
  - Three samples 70%
- Cannot distinguish *M. tuberculosis* from non-tuberculous mycobacteria

### Fluorescent Microscopy Scale (CDC Scale)

What you see (250X)	What you see (450X)	What to report
0 AFB/ smear	0 AFB/ smear	No AFB seen
1-2/ 30 fields	1-2/ 70 fields	Report exact count; order repeat specimen
1-9/ 10 fields	2-18/ 50 fields	1+
1-9/ field	4-36/ 10 fields	2+
10-90/ field	4-36/ field	3+
>90/ field	>36/ field	4+

Table adapted from Kent, P.T. and G.P. Kubica. *Public Health Mycobacteriology; A Guide for the Level III Laboratory*. U.S. Dept. of HHS/ PHS/ CDC. 1985

# Nucleic Acid Amplification tests (NAAT) or rapid molecular tests

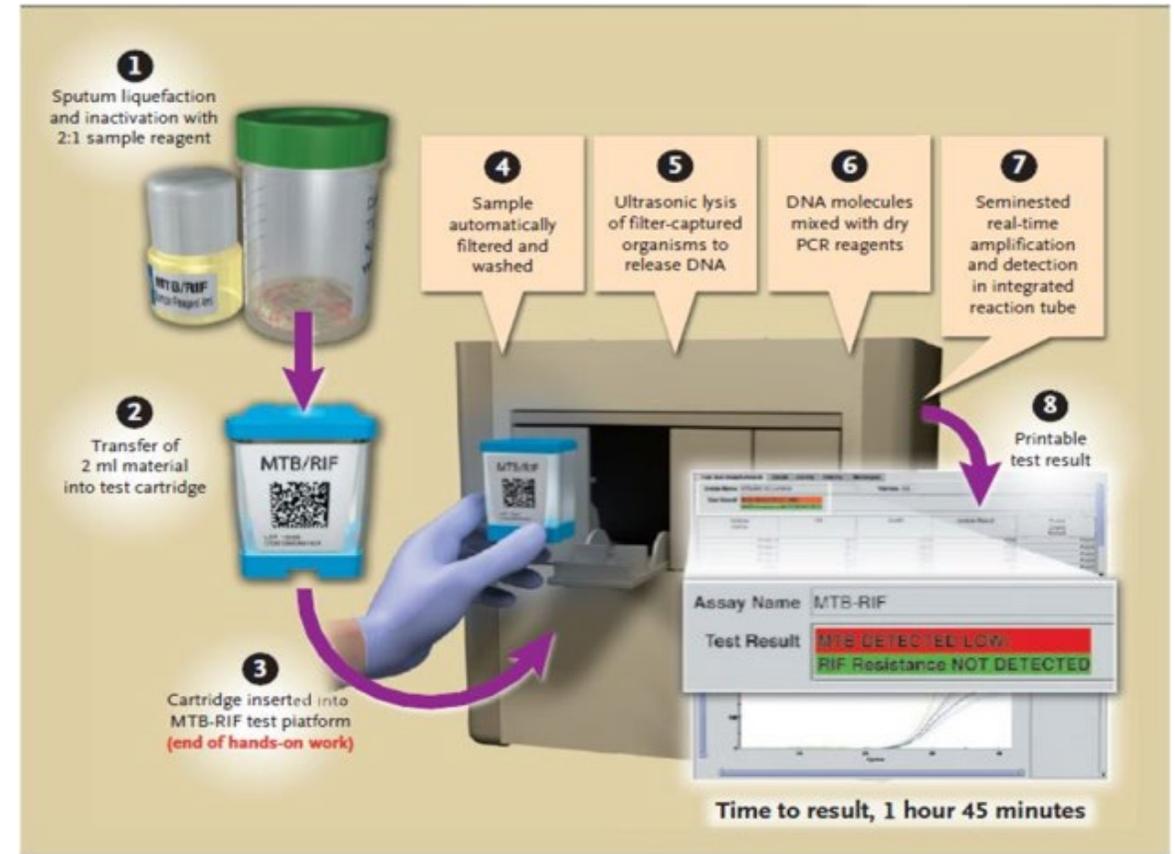
- NAAT: amplify DNA and RNA segments to rapidly identify the microorganisms in a specimen.
- NAAT: detect *M. tuberculosis* DNA in specimens in just hours, compared to a week or more for detection in culture
- CDC recommends that NAA testing be performed on at least one respiratory specimen

	GeneXpert® MTB/RIF	GenoType® MTBDRplus	HAIN GenoType® MTBDRsl
Company	Cepheid	HAIN Lifescience/ Bruker	HAIN Lifescience/ Bruker
Genetic loci	<i>rpoB</i> (for RIF)	<i>rpoB</i> (RIF), <i>katG</i> (INH), and <i>inhA</i> (INH)	<i>gyrA</i> (FQ), <i>rrs</i> (injectables), and <i>embB</i> (EMB)
Format	Semi-automated real-time PCR	Line probe assay	Line probe assay
FDA approved	Yes (market-authorized)	No	No

# Cepheid Xpert® MTB/RIF Test

Direct detection and identification of *M. tuberculosis* from respiratory specimens

- Cepheid GeneXpert platform
- WHO-endorsed
- FDA-approved for respiratory specimens
- off-label use reported for CSF and BAL specimens
- detects *M. tuberculosis* complex and provides information about RIF resistance
- results in about 2 hrs; minimal hands-on needed
- Sensitivity 99% for AFB smear positive and 67% for smear negative
- additional tests outside the U.S.,
  - Xpert MTB/RIF Ultra – enhanced sensitivity
  - Xpert MTB/XDR – additional AST cartridge for 9 target genes (INH, fluoroquinolones, injectibles (amikacin, kanamycin, capreomycin) and ETH)



Source: [www.finddiagnostics.org](http://www.finddiagnostics.org)

## **Benefits of using NAAT Direct Detection of *M. tuberculosis* in Clinical Specimens include:**

- Earlier laboratory confirmation of TB disease.
- Earlier treatment initiation.
- Improved patient outcomes.
- Interruption of transmission by early diagnosis, respiratory isolation, and appropriate treatment.
- Earlier, more efficient use of respiratory isolation (i.e., informed decisions about starting and stopping respiratory isolation).
- Earlier initiation of contact investigation.
- More effective public health interventions.
- Earlier detection of drug resistance when certain NAAs are used (e.g., Gene Xpert)

# Mycobacterial Culture

Sensitivity of culture is better than smear (**only need 10-100 AFB/mL of specimen**)

## Culture Media

- solid medium - Lowenstein-Jensen (LJ) egg-based or Middlebrook enriched agar
- broth/liquid medium - FDA-cleared systems for non-blood specimens are the BACTEC MGIT (BD) and VersaTREK (TF)
- mycobacteria grow faster in broth but there are some strains that prefer solid medium
- utilize PANTA (antibacterials/antifungal) to suppress other bacteria and fungi and OADC enrichment to promote growth

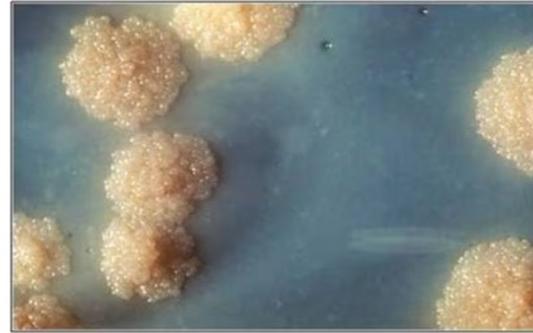
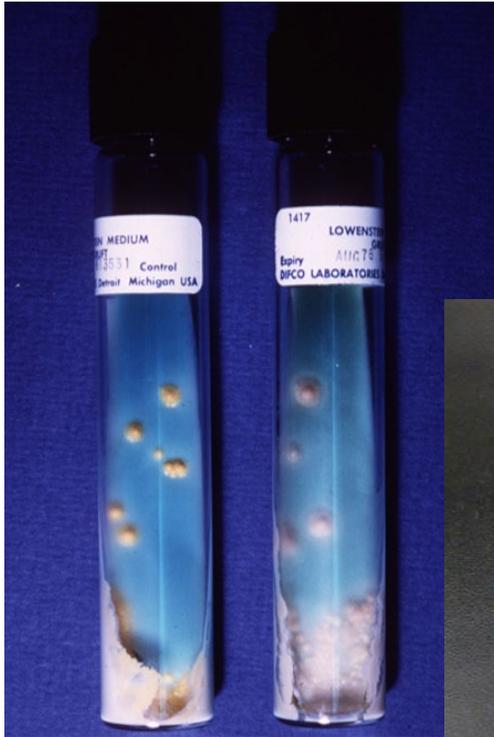
MGIT = Mycobacterial Growth Indicator Tube w/fluorescent detection



VersaTREK – headspace pressure differential indicates growth

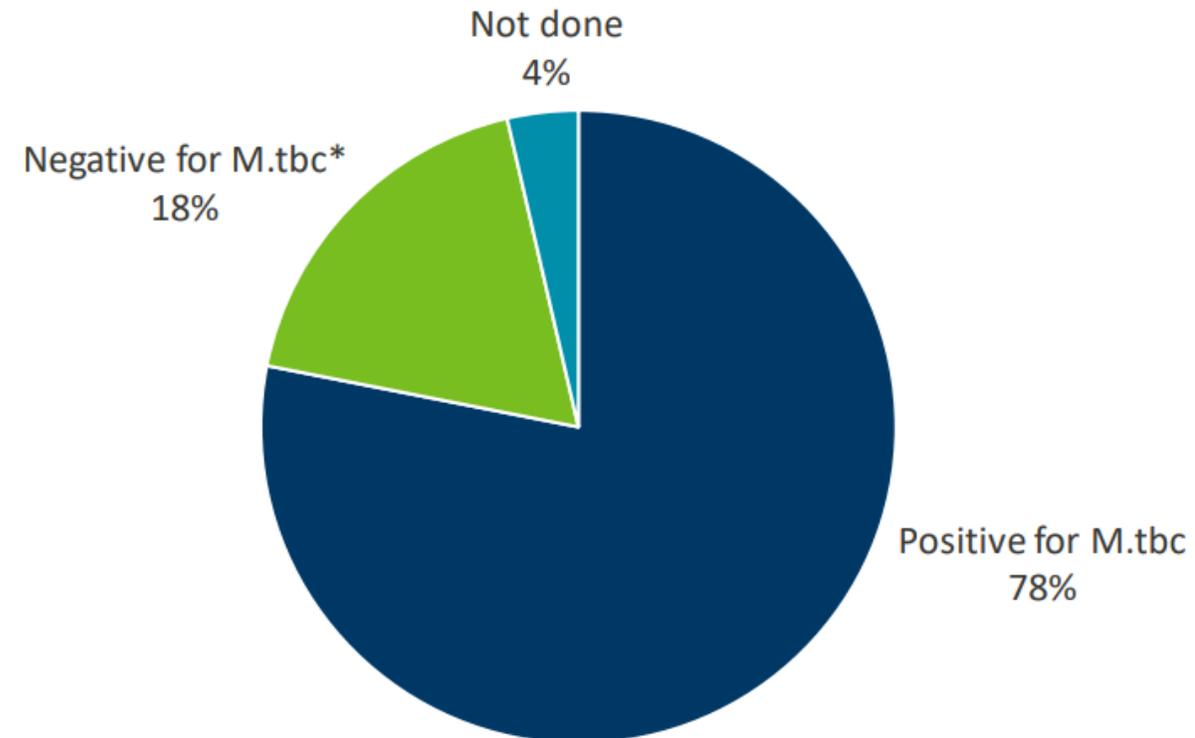


# M. tuberculosis Culture



Solid agar Mtb colony “rough and buff” morphology

## Culture result for Mtb cases in MN, 2017-2021



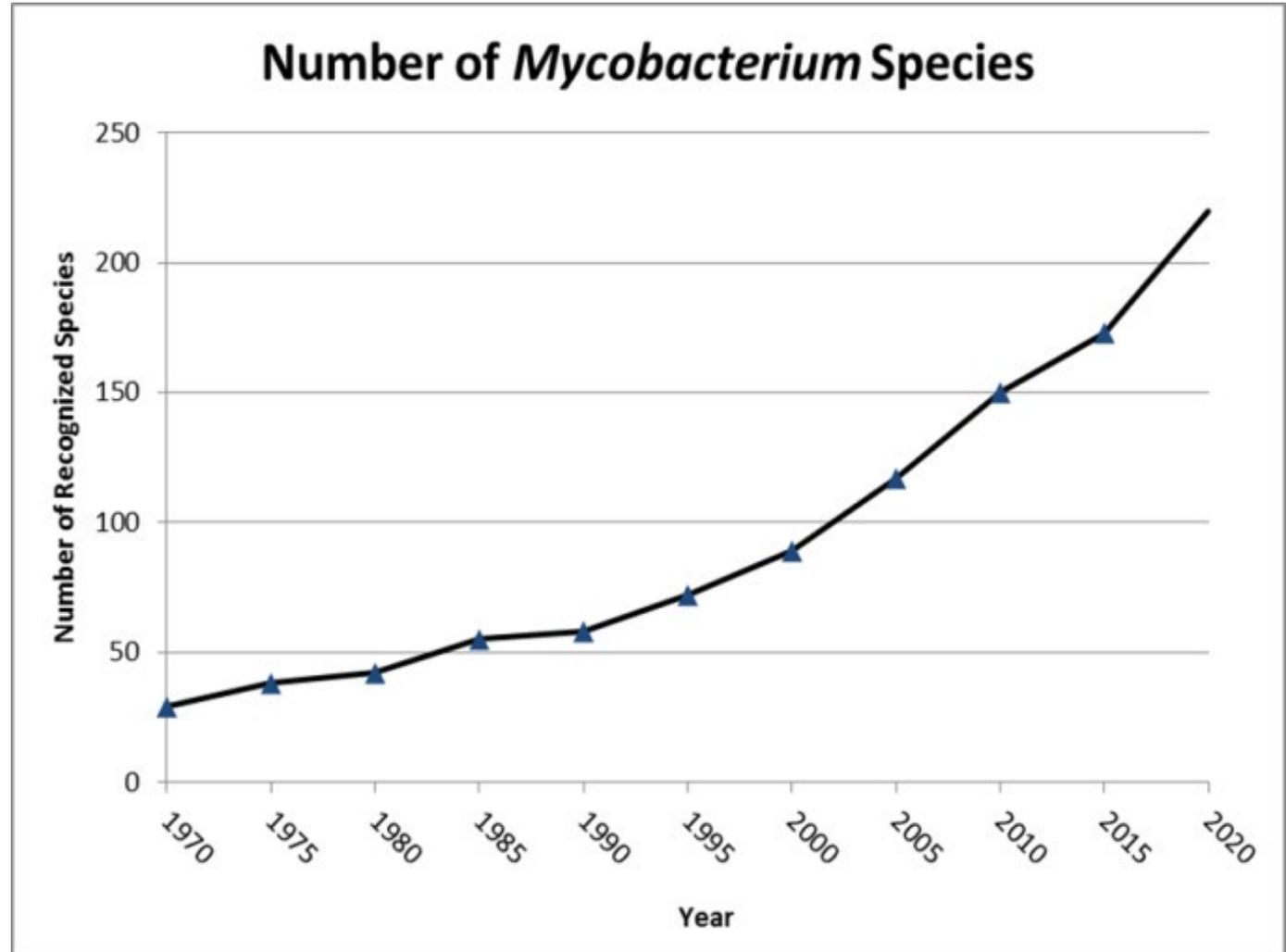
<https://www.health.state.mn.us/diseases/tb/stats/tbepislides.pdf>

slide adapted from N. Wengenack, PhD

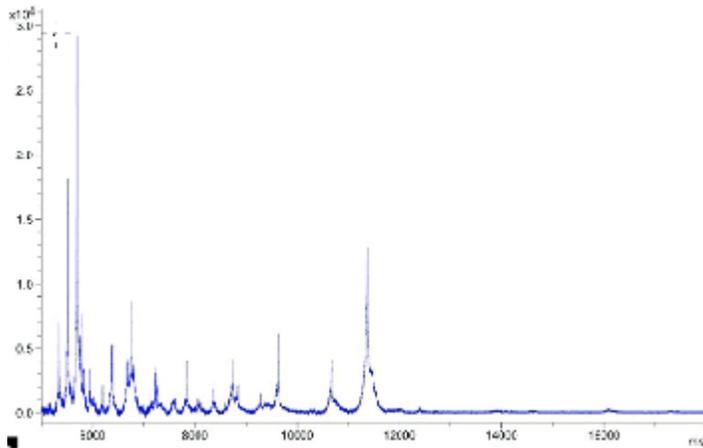
# Mycobacteria (2022)

214 recognized species  
22 subspecies

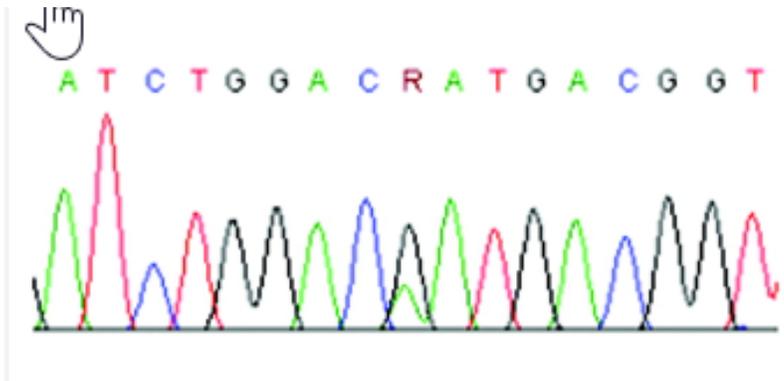
- *M. tuberculosis* complex
  - includes:
    - *M. tuberculosis*
    - *M. bovis* (cattle, unpasteurized milk)
    - *M. bovis* BCG (vaccine strain)
    - *M. africanum*
    - *M. microti* (vole)
    - *M. canetti*
    - *M. caprae*
    - *M. pinnepedi* (seal)
    - *M. mungi* (mongoose)



# Identification of *M. tuberculosis* after growth in culture



- MALDI-TOF MS



Sanger sequencing

## Current laboratory methods for culture identification:

- MALDI-TOF Mass Spectrometry
  - highly accurate; requires decent amount of growth so it may be a little slower than “probes”
- Sanger DNA sequencing
  - good targets are 16S rRNA gene, *rpoB*, *hsp65*
  - technically complex; takes about 1 day after culture growth
- Line Probe Assays
  - Hain Lifesciences (Bruker) and Inno-LIPA (Fujirebio)
  - not approved for diagnostic use in U.S. (clinical labs often don't have them; public health labs may)
- Targeted PCR assays
  - Cepheid GeneXpert MTB/RIF, LDT PCRs – fast, accurate, only for Mtb (not NTMs yet)

# Differentiation of species within the MTB complex

*M. tuberculosis* complex species differentiation PCR available from some labs

- often uses differences in patterns with the Region of Difference repetitive sequence element (RD) in *M. tuberculosis* complex
- performed from pure culture isolate; sometimes from positive MGIT broth
- can differentiate:
  - *M. tuberculosis*, *M. bovis*, *M. bovis BCG*, other species

Expected RD signature patterns	RD1	RD4	RD9	RD12	RD9-2
<i>M tuberculosis</i>	+	+	+	+	+
<i>M bovis</i>	+	-	-	-	+
<i>M bovis BCG</i>	-	-	-	-	+
<i>M africanum</i>	+	+	-	+	+
<i>M canettii</i>	+	+	+	-	+
<i>M caprae</i>	+	+	-	-	+
<i>M microti</i>	-	+	-	+	+
<i>M mungi</i>	+	+	-	+	+
<i>M pinnepedii</i>	+	+	-	-	+

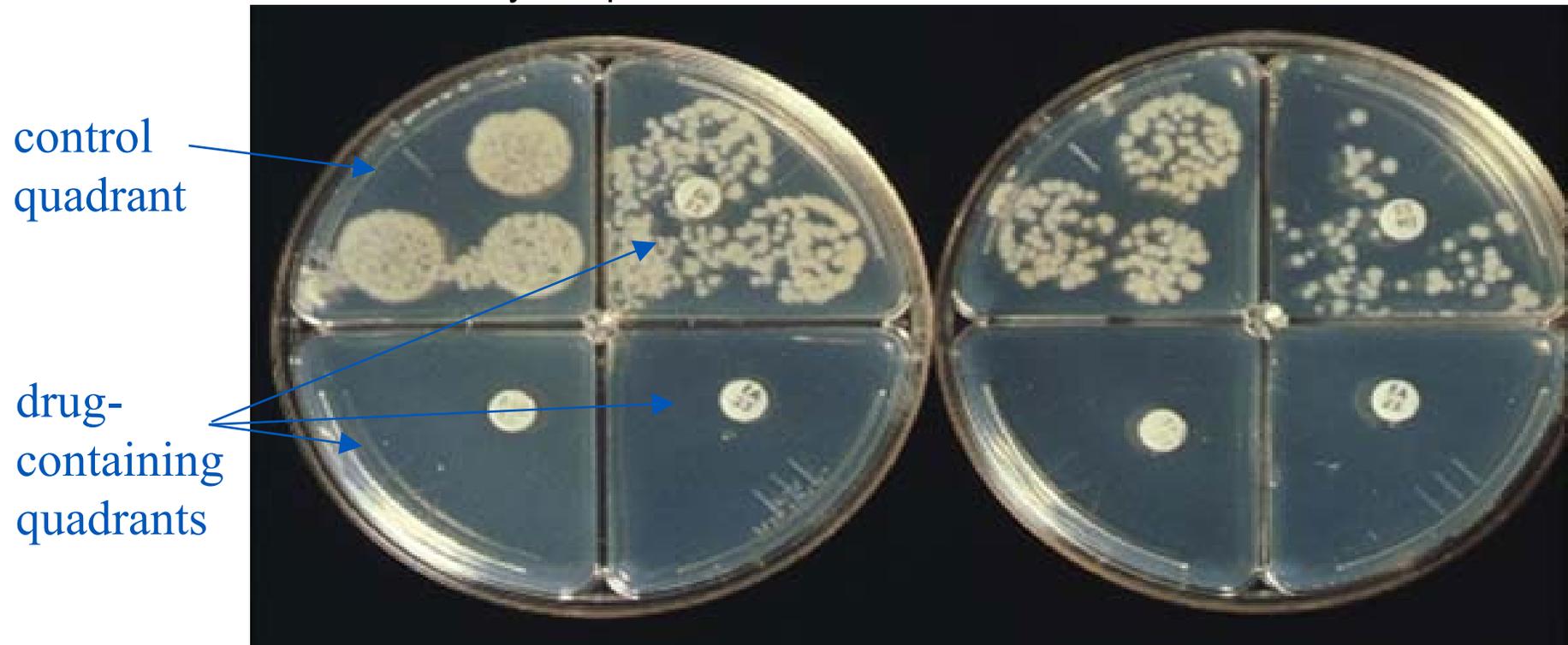
Method adapted from Halse TA, Escuyer VE, Musser KA. Evaluation of a single-tube multiplex real-time PCR for differentiation of members of the Mycobacterium tuberculosis complex in clinical specimens. J Clin Microbiol. 2011 Jul;49(7):2562-7. doi: 10.1128/JCM.00467-11. Epub 2011 May 18. PMID: 21593269; PMCID: PMC3147876.

slide courtesy of N. Wengenack, PhD

# Phenotypic - 1% indirect agar proportion method for *M. tuberculosis* complex susceptibility testing

Gold standard for all drugs except pyrazinamide

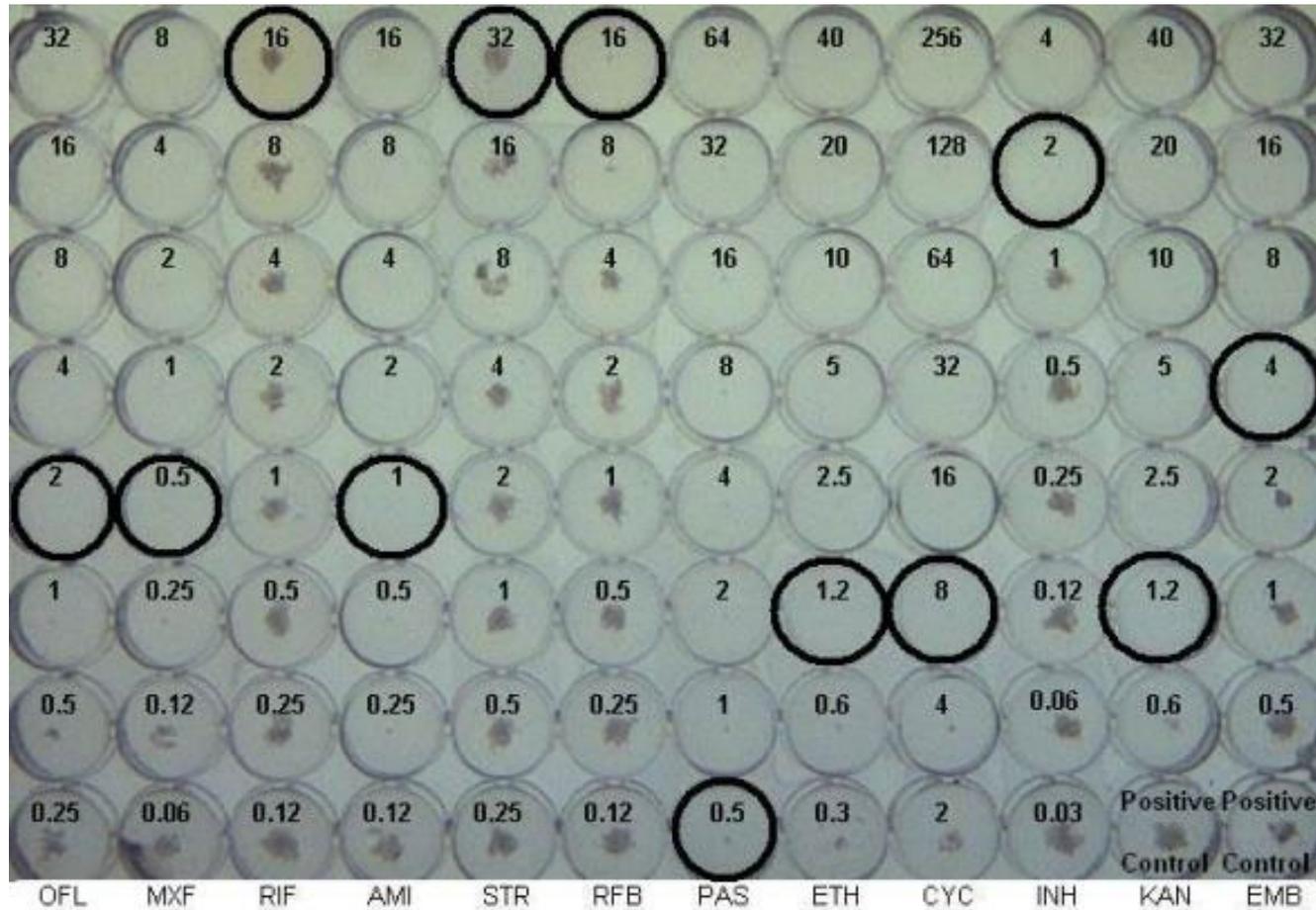
- use a single “critical concentration” of each anti-tuberculous drug
- not rapid (14-21 days)
- labor-intensive, technically complex



Organism is resistant to drug A in the upper right compartment (>1% of inoculum shown by upper left control quadrant is growing in presence of drug). Organism is susceptible to drugs B & C in the lower compartments. Control quadrant in upper left contains no drugs.

slide adapted from N. Wengenack, PhD

# Phenotypic - Broth MIC testing for *Mtb* using the MYCOTB MIC Plate LDT



- Broth microdilution method
- Not rapid (14 days)
- contains INH, RIF, EMB and 9 second-line drugs
- Test 1<sup>st</sup> and 2<sup>nd</sup> line drugs simultaneously with same inoculum
- Provides MIC endpoint
- Requires skilled staff to setup and read plates

Hall L, Jude KP, Clark SL et al., Evaluation of the Sensititre MYCOTB MIC plate for the susceptibility testing of *Mycobacterium tuberculosis* complex against first and second line agents. *J Clin Microbiol.* 2012; 50:3732-4.

# Phenotypic - “Rapid” Broth Susceptibility Testing for *M. tuberculosis* complex

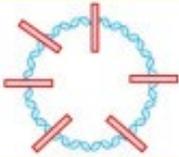
FDA-cleared, semi-automated with MGIT or VersaTREK systems



- compare growth of a standard inoculum in tube +/- critical concentration of the drug
- growth at critical concentration = R
- Turn around time ~ 2 weeks



# Most common Genotypic methods for *M. tuberculosis* complex AST

	Multiplex or specific PCR	Targeted NGS	WGS
			
Principle	<ul style="list-style-type: none"> <li>• Amplification by PCR of specific loci</li> <li>• Detection of mutation(s) by melting curve analysis or Sanger sequencing</li> </ul>	<ul style="list-style-type: none"> <li>• Amplification by PCR of specific loci</li> <li>• Detection of mutation(s) by NGS</li> </ul>	<ul style="list-style-type: none"> <li>• Sequencing of the entire genome by NGS</li> </ul>
Advantages	<ul style="list-style-type: none"> <li>• \$</li> <li>• Rapidity</li> <li>• Sensitivity and specificity</li> </ul>	<ul style="list-style-type: none"> <li>• Detection of numerous mutations for a large panel of antibiotics</li> <li>• Detection of heteroresistance</li> </ul>	<ul style="list-style-type: none"> <li>• Detection of all putative mutations for all antibiotics</li> <li>• Detection of heteroresistance?</li> </ul>
Limits	<ul style="list-style-type: none"> <li>• Analysis limited to well known mutations for few antibiotics</li> <li>• Lack of detection of heteroresistance</li> </ul>	<ul style="list-style-type: none"> <li>• \$\$</li> <li>• Time-consuming</li> <li>• Need of bioinformatic skills +</li> </ul>	<ul style="list-style-type: none"> <li>• \$\$\$</li> <li>• Time-consuming</li> <li>• Need of bioinformatic skills ++</li> </ul>

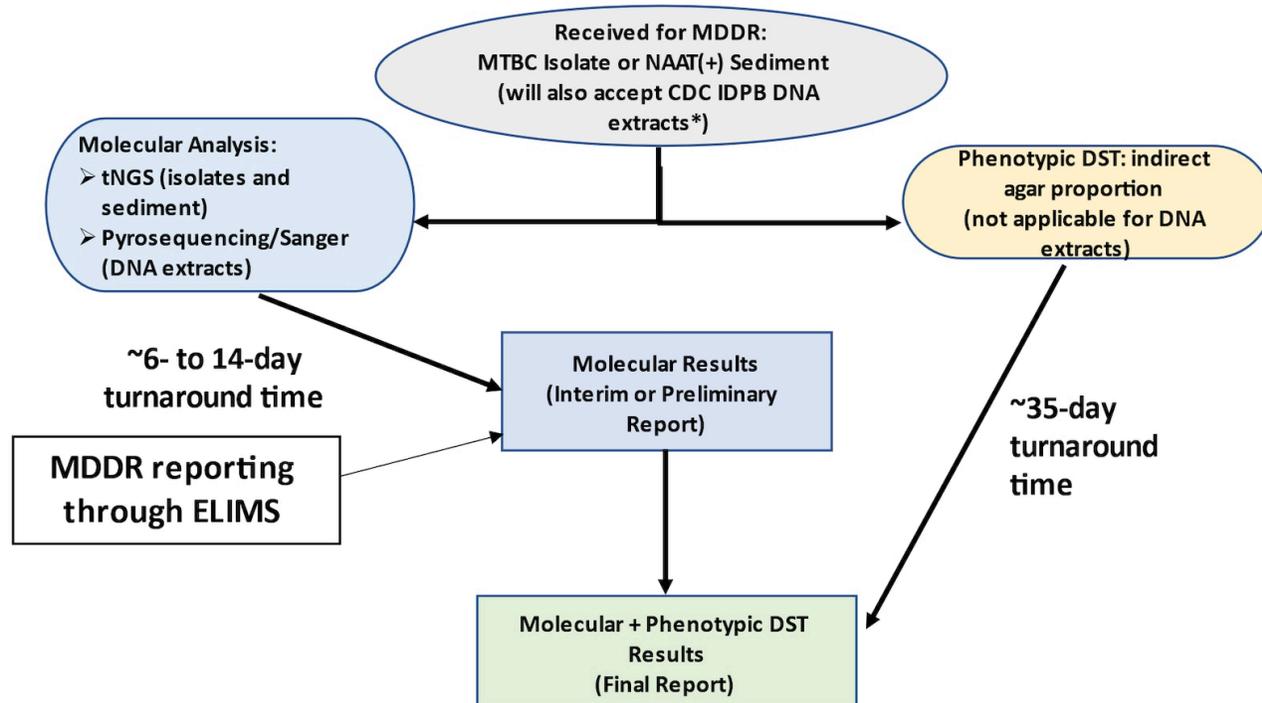
## Multiplex or specific PCR:

- Available in U.S.
  - Cepheid Xpert MTB/RIF – for RIF only
  - LDT PCRs – e.g., MTBRP for INH only (Mayo Clinic)
  - LDT Sanger sequencing – e.g., *pncA* for PZA
- Available mainly outside of the U.S.
  - Cepheid Xpert MTB/RIF Ultra – for RIF only
  - Cepheid Xpert MTB/XDR – to be used in addition to MTB/RIF test; adds several gene targets for additional drugs (INH, ethionamide, fluoroquinolones, plus amikacin, kanamycin, capreomycin)

## Targeted next generation sequencing assay (tNGS):

- Used by the MDDR service at the CDC since February 2023 (and other reference labs)
- Molecular data confirmed with phenotypic testing for majority of drugs.

## MDDR Algorithm



Examines 24 amplicons across 16 genes providing information on more than 12 antituberculosis drugs:

- First line: RIF, INH, EMB, PZA
- Second line: FQ, AMK, CAP, KAN, BDQ, CLF, LZD.

\*DNA extracts only accepted from CDC IDPB and will be tested by conventional sequencing methods (not tNGS)

MTBC *Mycobacterium tuberculosis* complex, IDPB: Infectious Diseases Pathology Branch, ELIMS: enterprise laboratory information management system

## GENE TARGETS FOR GENOSCREEN DEEPLEX MYC-TB ASSAY

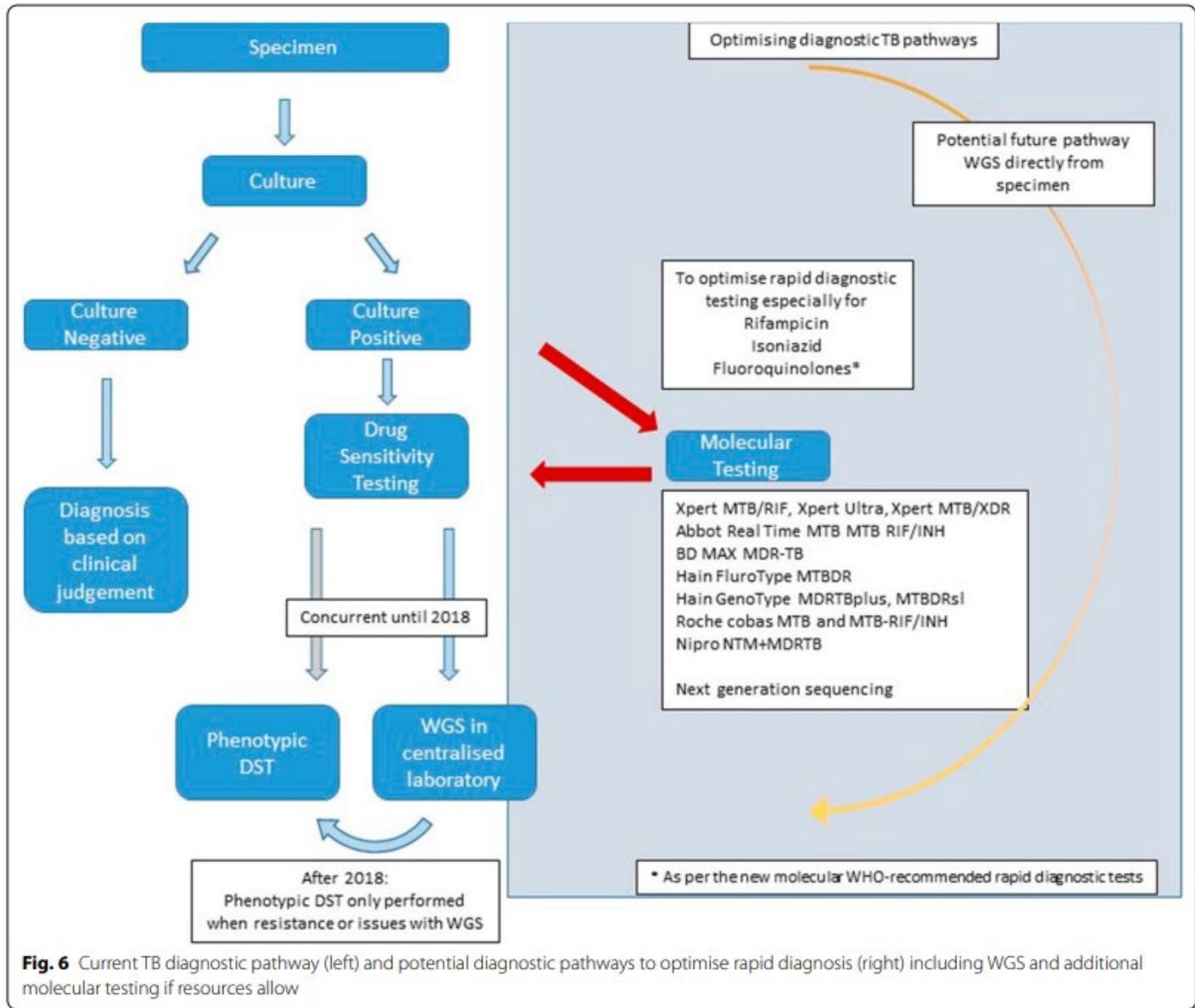
Identification of the species is performed by sequence analysis of the *hsp65* gene in combination with select other targets for closely related species



## 18 drug resistance markers

Genomic target	Drug
<i>rpoB</i>	rifampicin
<i>inhA</i>	isoniazid, ethionamide
<i>fabG1</i>	isoniazid, ethionamide
<i>katG</i>	isoniazid
<i>ahpC</i>	isoniazid
<i>pncA</i> *	pyrazinamide
<i>embB</i>	ethambutol
<i>gidB</i> *	streptomycin
<i>rpsL</i> *	streptomycin
<i>rrs</i> *	streptomycin, amikacin, kanamycin, capreomycin
<i>eis</i>	kanamycin
<i>tlyA</i> *	capreomycin
<i>gyrA</i>	fluoroquinolones
<i>gyrB</i>	fluoroquinolones
<i>ethA</i> *	ethionamide
<i>rrl</i>	linezolid
<i>rplC</i>	linezolid
<i>rv0678</i> *	bedaquiline, clofazimine



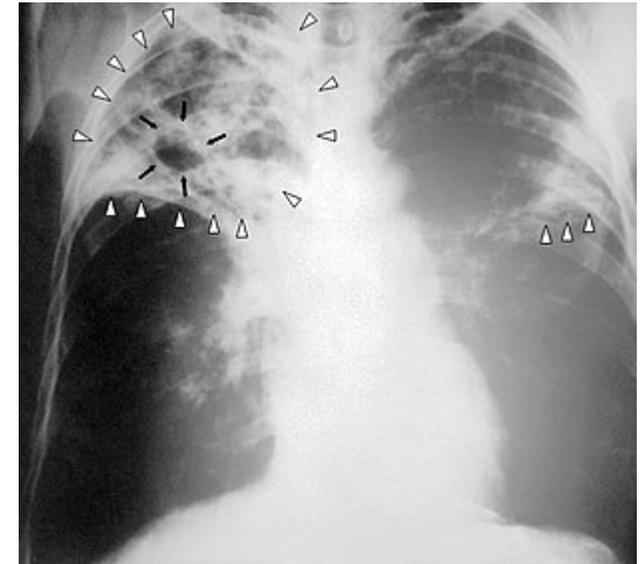
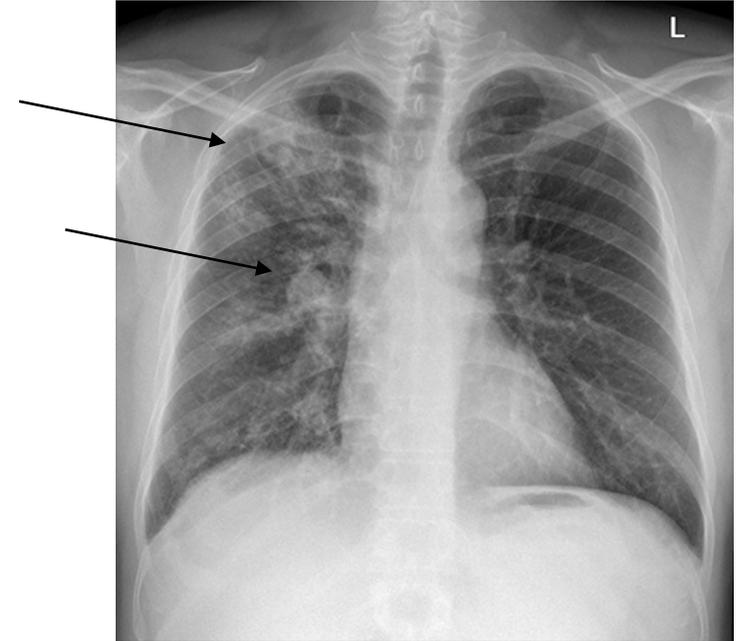


**Fig. 6** Current TB diagnostic pathway (left) and potential diagnostic pathways to optimise rapid diagnosis (right) including WGS and additional molecular testing if resources allow



# Imaging

- A posterior-anterior view is the standard (apical and posterior segments of the upper lobe or in the superior segments of the lower lobe)
- A lateral view may be helpful, especially in children.
- Children: abnormalities tend to be minimal (mostly lymphadenopathy).
- Atypical lesions especially in HIV-infected and immunosuppressed (Lesions anywhere in the lungs and may differ in size, shape, density, and cavitation).
- Cavitory lesions usually in patients with higher CD4 counts (cavitation is thought to occur as a result of the immune response).



In patients with symptoms and signs of TB disease, a  
negative chest radiograph result  
**DOES NOT**  
exclude TB disease

# Case

- 17 yo male in rural MN with chest mass evaluation and intermittent night sweats, otherwise healthy
  - Immigrated to U.S. with family in 2012 from Thailand. Never left U.S. since. Lives with family, attends school.
  - No known TB exposures, positive TB test or prior treatment. BCG vaccine status unknown.
- 
- QuantiFERON Negative
  - Lung nodules noted on chest CT, cervical and bilateral axillary lymph nodes.
  - Sputum samples were AFB smear negative, culture negative, GeneXpert MTB/RIF negative
  - Ultrasound guided FNA of chest mass: Negative for malignancy. AFB stain negative, culture pending
  - No PCR was ordered on fresh chest mass tissue.
  - LDT Mtb PCR performed on FFPE Pathology Block tissue – negative;

## What would you do next?

1. Start RIPE
2. No treatment, wait for cultures
3. Do more tests



Thank you