

## Diagnosis of TB Disease: Laboratory

Benjamin Alpers, BA September 13, 2023

TB Intensive September 13 – 15, 2023 Richmond, TX

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## **Benjamin Alpers, BA** has the following disclosures to make:

- No conflict of interests
- No relevant financial relationships with any commercial companies pertaining to this educational activity





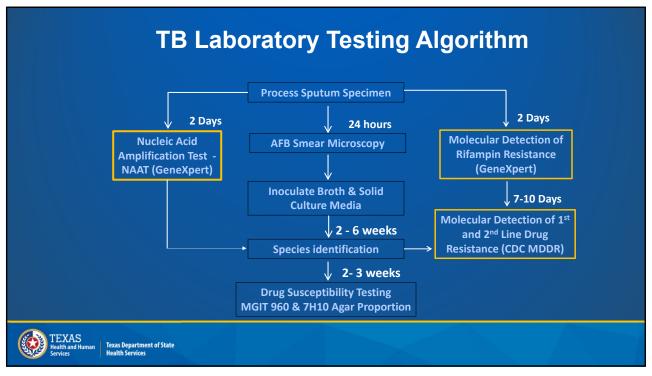
## Laboratory Diagnosis of Tuberculosis

or

**Desperately Seeking Tuberculosis** 

**Benjamin Alpers** 

Applications Scientist/TB Reference Team Lead DSHS Austin Laboratory

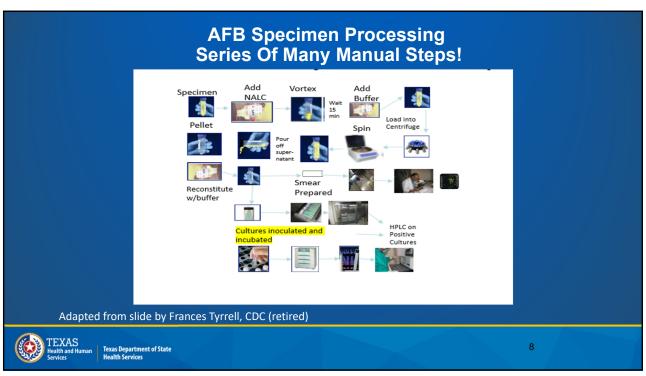


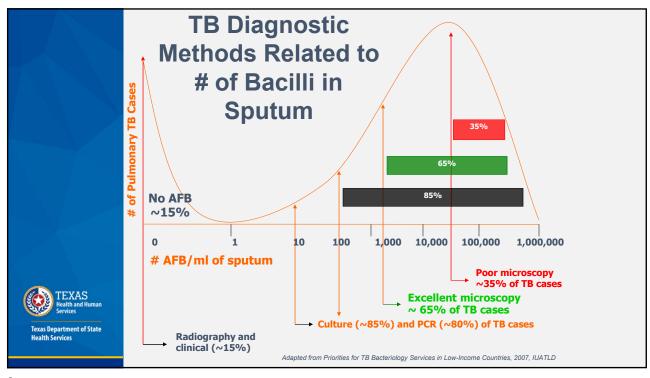
## **Specimen Quality**

- Accurate laboratory results are directly related to the quality of the specimen
- Sputum
  - Recently discharged material from the bronchial tree, with minimal amounts of upper respiratory tract secretions
    - Well coached patient, collect at least 3ml
    - Label tube, form, and indicate test:
      - · initial Dx: Smear, NAAT, & Culture
      - Follow-up: Smear and Culture
      - Release from respiratory isolation? Order Smear only
- Transport to lab cool and quickly

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### **AFB Culture**

- · More sensitive than smear
  - 5,000 to 10,000 AFB/ml for smear
  - ~10 viable AFB/ml for culture
- Positive for only ~85% of Pulmonary TB
  - Requires a quality specimen
  - May be invalid due to contamination
- Used to monitor patient response to treatment (like smear)
- Required for conventional drug susceptibilities & genotype
- Lengthy
  - 1-6 weeks by liquid media
  - 2-8 weeks by solid media

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## **Becton Dickinson BACTEC<sup>TM</sup> MGIT<sup>TM</sup>** 960

- A fluorescent compound is embedded in silicone on the bottom of tubes.
- Tube initially contains dissolved oxygen which quenches emissions from the compound and little fluorescence can be detected.
- Later, actively respiring microorganisms consume the oxygen and allow the fluorescence to be observed (instrument takes a reading once an hour).



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## Becton Dickinson BACTEC<sup>TM</sup> MGIT<sup>TM</sup> 960, cont.

- Fluorescence can also be manually observed using a transilluminator.
- Particulates are often observed at bottom of tube.
- Modified 7H9 broth that requires supplement for each process (growth and antibiotic).
- The same MGIT<sup>TM</sup> tube used for IIRE drug susceptibility testing (tube with different pH used for PZA).



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# Solid Media for MTBC Detection • Middlebrook 7H11 agar • Primary media • Morphology can be viewed microscopically • Middlebrook 7H10 agar • Used as secondary media (less selective) • Lowenstein Jenson slant • Long life span for storage

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# Methods for Diagnosis Used in Conjunction with Culture • Acid Fast Bacilli Microscopy (AFB Smear) • Nucleic Acid Amplification Test (NAAT) • Cepheid GeneXpert ® • High Performance Liquid Chromatography (HPLC) • MALDI-TOF Mass Spectrometry • Real-time Polymerase Chain Reaction (PCR) • Used in M. tb cx. speciation

## TE:

## **Acid Fast Bacilli Microscopy (AFB Smear)**

- Has many qualities of an ideal diagnostic test
  - Rapid & universally available
  - Detects the most infectious cases
  - Used to support diagnosis and identify need to isolate
  - Helps monitor response to therapy
  - Identifies priority cases for nucleic acid amplification (NAA)
- Problems
  - Not sensitive misses ~50% of TB
  - Not specific in low TB prevalence areas (e.g. Texas)
    - Positive smear may be NTM (~20% at DSHS-Austin)
- Highly specific where TB is highly prevalent

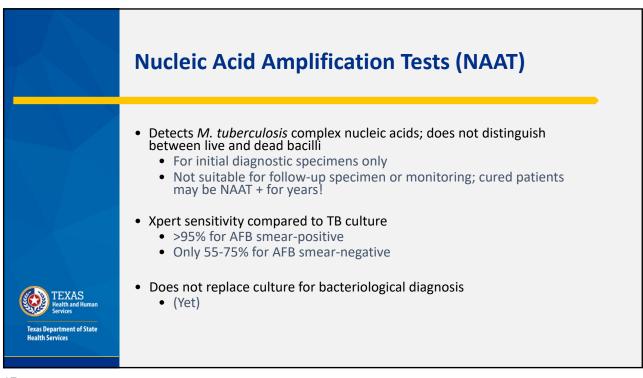
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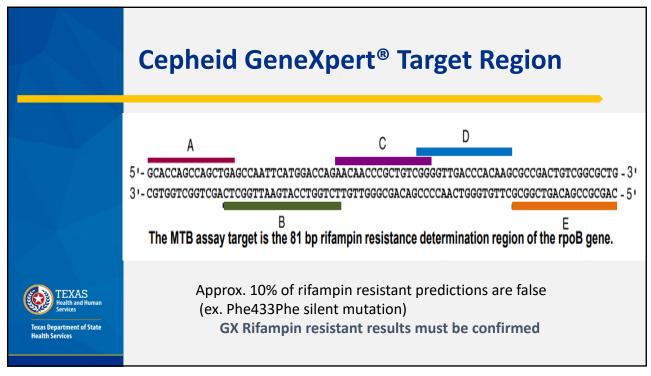
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## **Nucleic Acid Amplification Tests (NAAT)**

- Tiny amounts of DNA/RNA are amplified (copied) until there is enough for easy detection
- GeneXpert examines DNA for:
  - Identification
  - Detection of Rifampin Resistance
- Test turnaround time measured in hours







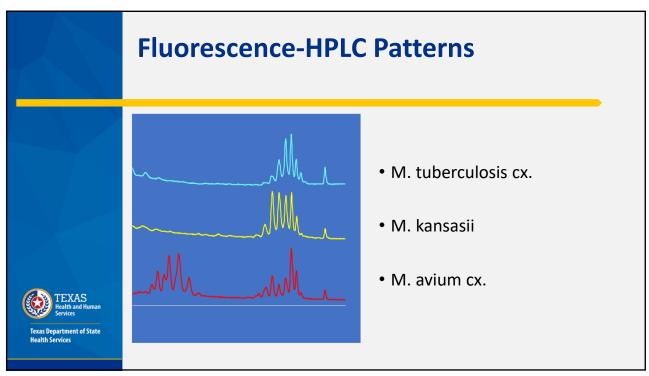
## High-Performance Liquid Chromatography (HPLC)

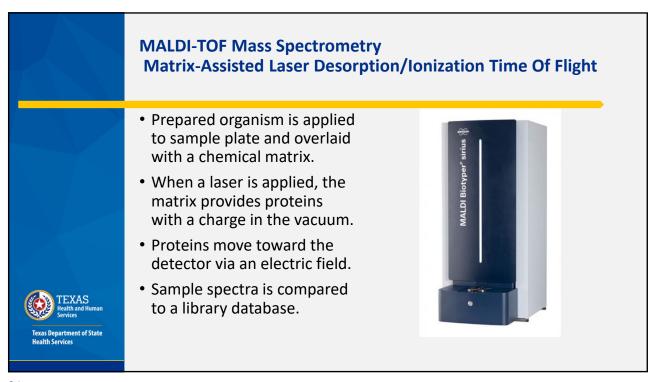
- Identifies mycobacteria by analysis of mycolic acids in the thick cell wall.
- Mycolic acids are chemically cleaved, separated, and extracted from cell wall.
- Mycolic acid esters are then separated and detected with reverse-phase HPLC with fluorescent detection (FL-HPLC) to produce a chromatogenic pattern with diagnostic peaks.

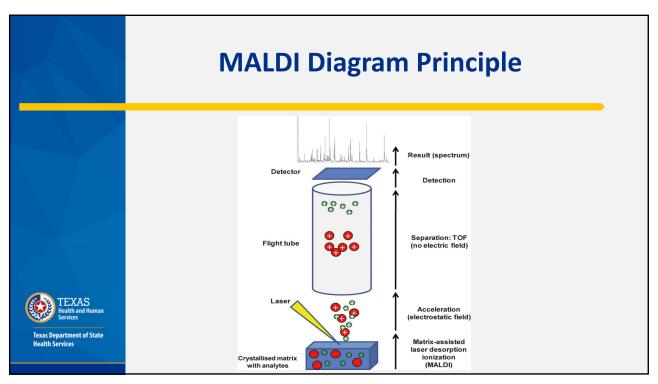
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## MALDI vs. HPLC

- MALDI able to identify many more species than HPLC, also break some groups and complexes up into distinct organisms
- MALDI is less sensitive than HPLC, especially for MTB.
   Liquid media may need to incubate an additional couple days for valid ID.
- MALDI is currently validated filamentous mold and Candida yeast; working on AFB validation

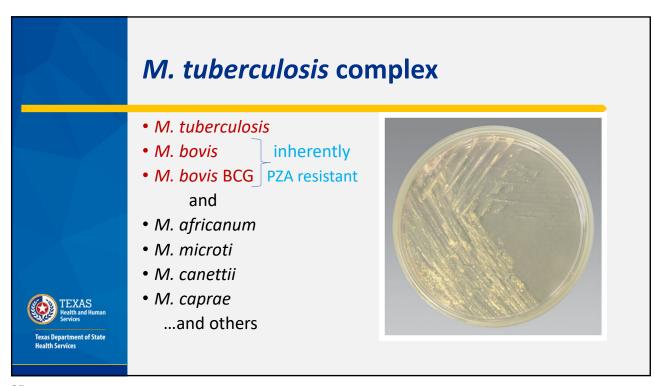
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## M. tb cx. Speciation through Real-Time Polymerase Chain Reaction (PCR)

- New laboratory developed test
- Currently send to CDC when bovis or bovis BCG suspected with an average of 4 week turnaround
- Once implemented should produce a result in <4 days
- Can differentiate between species within the complex using 5 sets of probes and primers targeting known regions of difference (RD)

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	Species	RD 1	RD 4	RD 9	RD 12	Ext-RD9
	M. tuberculosis	+	+	+	+	+
	M. bovis	+	-	-	-	+
	M. bovis BCG	-	-	-	-	+
	M. africanum	+	+	-	+	+
	M. microti	-	+	-	+	+
TEVAC	M. canettii	+	+	+	-	+
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## Conventional growth-based method performed at DSHS Molecular Detection of Drug Resistance (MDDR) performed at the CDC TEXAS Realth and Ruman Services Texas Department of State Health Services

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# Conventional Drug Susceptibility Testing (DST) of M. tuberculosis Complex Current Laboratory Protocol Initial isolate should be tested against first-line drugs (FLD) Isoniazid, rifampin, ethambutol, pyrazinamide plus a fluoroquinolone (ofloxacin) Repeat test if patient cult+ after 3 mo. Rx For isolates resistant to Rifampin or to any 2 FLDs: test second-line drug panel! Rifabutin, ethionamide, & injectable (capreomycin, kanamycin)

## **Important Considerations**

- A rapid report of "INH & RMP susceptible" is the single most common & important DST report issued by the TB lab.
  - ~90% of U.S. cases are susceptible to the primary drugs
  - "INH & RMP susceptible" predicts primary drug panel efficacy
  - "INH & RMP susceptible" allows discontinuation of relatively toxic antibiotics in treatment continuation phase

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## **Additional Considerations**

- CDC recommends RMP DST results be reported within 17 days after M. tuberculosis culture identification
- Most U.S. laboratories use a rapid commercial system for DST (MGIT 960)
- Commercial DST methods miss some clinically significant RMP resistance that can be detected by agar proportion
- Agar Proportion (AP) is the "gold standard" method for conventional DST...however AP is not a rapid method; conventional AP method takes 21 days (3 weeks) for full results
- DSHS is able to regularly report INH & RMP susceptibility within 17 days of culture identification

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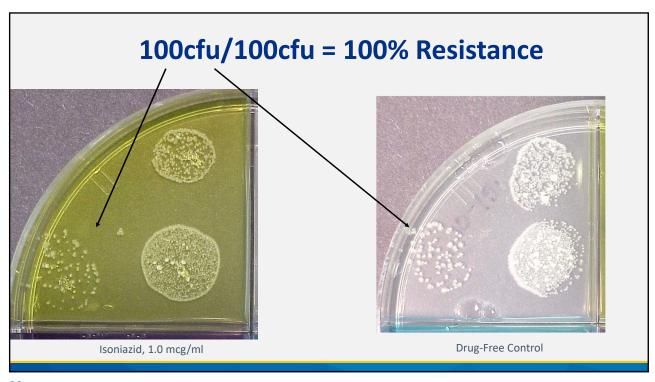
## Agar Proportion DST Method Principle

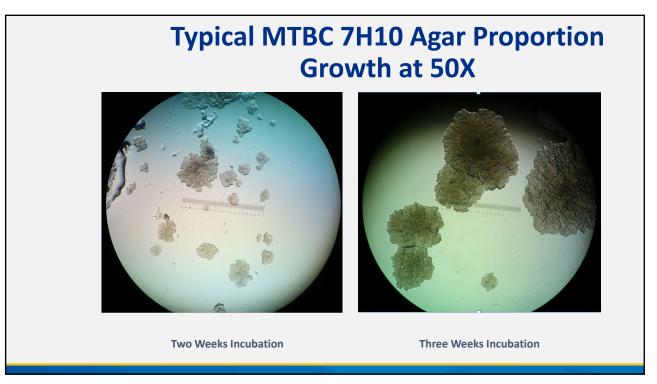
- A standardized suspension of M. tuberculosis is inoculated to quadrant plates of drug-containing Middlebrook 7H10 agar and a drug-free control.
- If growth of M. tuberculosis on the drug quadrant is 1% or greater than the growth on the control, the drug can no longer be counted on as being effective for treatment.



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## Molecular Detection of Drug Resistance (MDDR)

- Can provide rapid detection of drug resistance
- Both NAAT positive and culture positive specimens are candidates
- Particularly useful for high-risk patients, RMP positive Xpert sediment, contaminated specimens, or those specimens that do not grow well or are non-viable in standard TB media
- Examines 24 amplicons across 16 genes providing information on more than 12 antituberculosis drugs

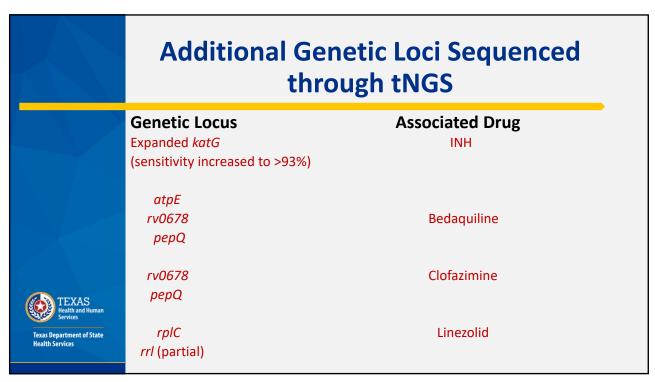
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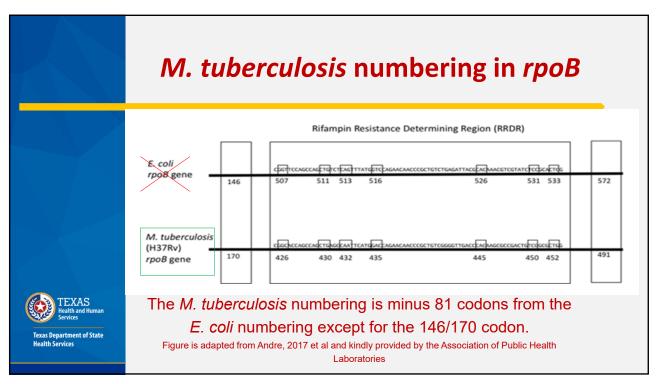
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### **Genetic Loci Sequenced through MDDR Associated Drug Genetic Locus** RRDR within the rpoB gene with Rifampin (RMP) the addition of two codons outside of the RRDR inhA, katG, fabG1 Isoniazid (INH) Ethambutol (EMB) embB Pyrazinamide (PZA) pncA Fluoroquinolones gyrA, gyrB Amikacin Capreomycin rrs Kanamycin eis Kanamycin

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# Changes in CDC's MDDR Assay New sequencing method using Targeted Next Generation Sequencing (tNGS) More regions sequenced including genes associated with bedaquiline, clofazimine, and linezolid Improved limit of detection for heteroresistance More TB DNA needed for successful amplification than pyrosequencing (1-10/field smear result or greater) Turnaround time now 7-10 days





## CDC Molecular Detection of Drug Resistance (MDDR) vs. Agar Proportion (AP) Method

- MDDR provides 7-10 day DNA sequence analysis for drug resistance prediction, AP 11-22 day growth-based susceptibility
  - 1st and 2nd line drugs tested
- MDDR complements conventional DST
  - Used alone, MDDR and conventional DST are imperfect
  - Used together, accuracy of drug resistance or susceptibility detection can be improved.
- Conventional DST results are still needed, or at least desirable, to confirm susceptibility to individual drugs.

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## Whole Genome Sequencing (WGS)

- Current genotyping method
- CDC began performing this in 2012, regularly in 2014, and exclusively in 2022
- Primarily used to determine relatedness between strains infecting individuals for epidemiological purposes
- Can be helpful in false-positive investigations
- Potential to establish reactivation vs. reinfection
- Reflexively alerts to any significant rpoB mutation

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## WGS (con't)

- 2,690 genetic loci examined and compared
- Those that are 99.7% similar clustered by wgMLSType
- This translates as <8 SNPs difference to at least one isolate in cluster
- Phylogenic trees can be created within clusters
- Not indicative of drug resistance pattern!

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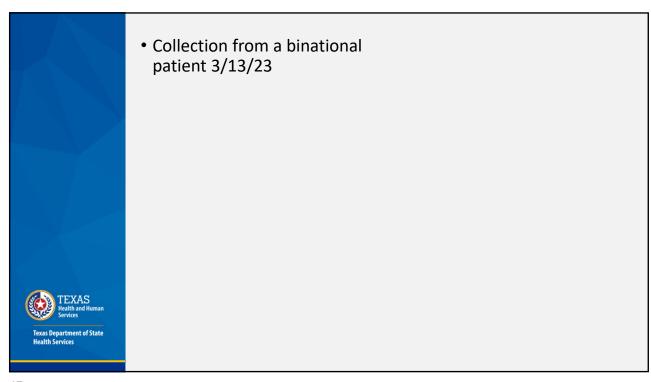
MTBC000025 in Texas Analysis updated with Results received 2/10/2021 20RF5776 and 20RF5462 (isolates from the same patient) 19RF2847, MRCA 19RF4405, 20RF2369 19RF4755 19RF2304 19RF4223 19RF1160 18RF1535 **Excluded isolates:** 19RF3857 - contaminated 18RF6976 - contaminated \*Isolates denoted with an asterisk are from the 18RF6084 – contaminated same patient. Isolate 20RF5462 was collected from a sputum specimen and 20RF5776 was 18RF4713 - contaminated  $collected \ from \ a \ urine \ sample.$ 18RF3636 - low sequence coverage

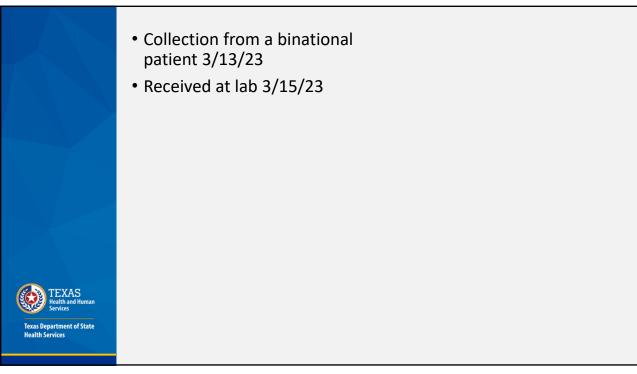
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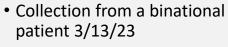
## WGS (con't) DSHS plans to incorporate this method into our algorithm for predictive DST Can only be performed on isolate One isolate per patient per year in most instances Will report mutations detected from genes associated with primary drug resistance (INH, RMP, EMB, FQN, and PZA) MDDR will not be replaced by this; used as an alternative unless MDDR indicated initially or reflexively

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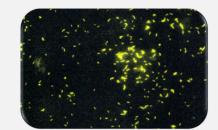
## Diagnostics for an MDR Patient A Case Study for the Way Things Should Work TEXAS Services Texas Department of State Realth Services Realth Services



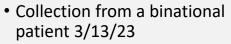




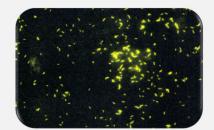
- Received at lab 3/15/23
- Smear reported 3/15/23
- >10/field



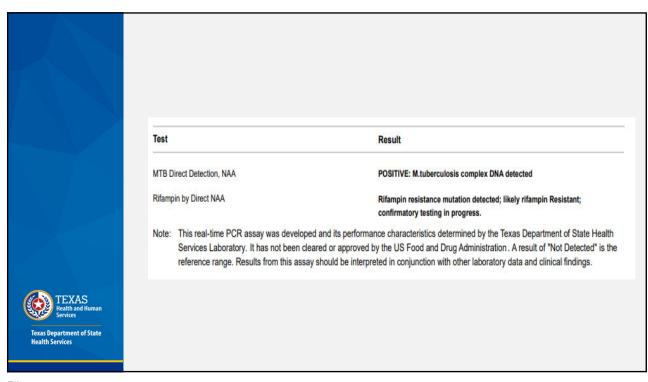


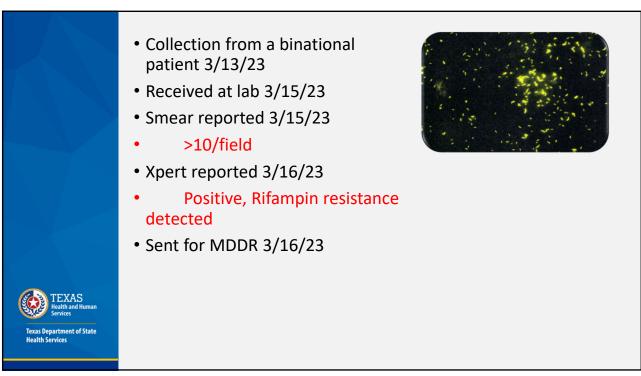


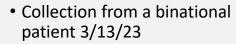
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- Xpert reported 3/16/23
- Positive, Rifampin resistance detected



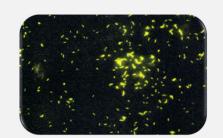






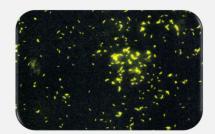


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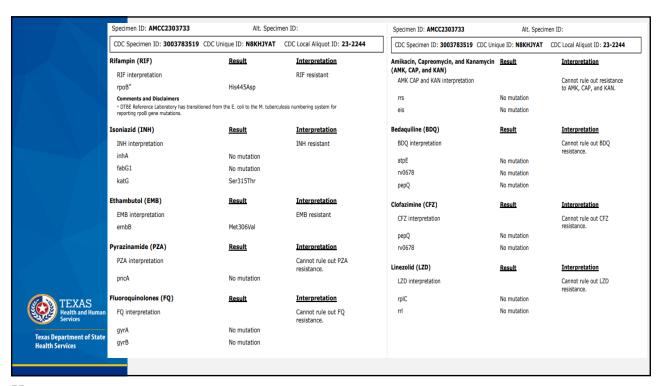


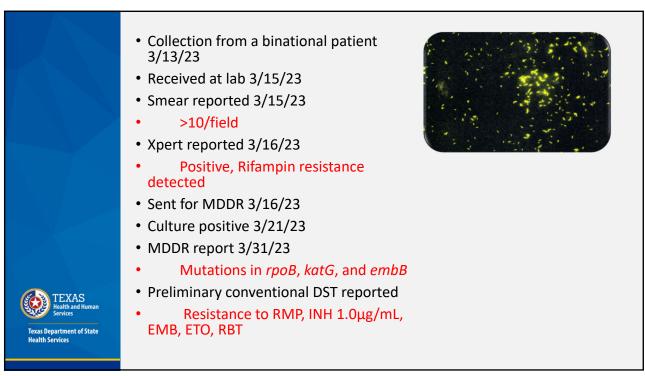


- Collection from a binational patient 3/13/23
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- MDDR report 3/31/23
- Mutations in rpoB, katG, and embB



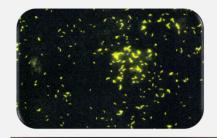








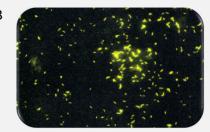
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- Preliminary conventional DST reported
- Resistance to RMP, INH 1.0 $\mu$ g/mL, EMB, ETO, RBT
- Final report 4/12/23
- Resistance to SM





	Isoniazid 0.2 mcg/ml by Agar Proportion	Resistant
	Note: MEDICAL EMERGENCY: This patient is res filing this report.	sistant to one or more drugs. Notify responsible physician and Infection Control prior to
	Rifampin 1.0 mcg/ml by Agar Proportion	Resistant
	Pyrazinamide 100 mcg/ml by MGIT 960	Susceptible
	Ethambutol 5.0 mcg/ml by Agar Proportion	Resistant
	Isoniazid 1.0 mcg/ml by Agar Proportion	Resistant
N. /	Ethionamide 5.0 mcg/ml by Agar Proportion	Resistant
	Streptomycin 2.0 mcg/ml by Agar Proportion	Resistant
	Ofloxacin 2.0 mcg/ml by Agar Proportion	Susceptible
	Rifabutin 2.0 mcg/ml by Agar Proportion	Resistant
TEXAS	Kanamycin 5.0 mcg/ml by Agar Proportion	Susceptible
Health and Human Services	Capreomycin 10.0 mcg/ml by Agar Proportion	Susceptible
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- CDC final 4/27/23

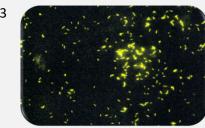




Specimen ID: AMCC2303733	Alt. Specir	nen ib.
CDC Specimen ID: 3003783519 CDC Ur	nique ID: N8KHJYAT	CDC Local Aliquot ID: 23-2244
MTBC Agar Proportion Susceptibility*	% Resistant	<u>Interpretation</u>
Isoniazid 0.2 μg/mL	100 %	Resistant
Isoniazid 1.0 µg/mL	100 %	Resistant
Isoniazid 5.0 μg/mL	0 %	Susceptible
Rifampin 1.0 μg/mL	100 %	Resistant
Ethambutol 5.0 µg/mL	33 %	Resistant
Streptomycin 2.0 µg/mL	67 %	Resistant
Streptomycin 10.0 µg/mL	0 %	Susceptible
Rifabutin 2.0 µg/mL	67 %	Resistant
Ciprofloxacin 2.0 µg/mL	0 %	Susceptible
Kanamycin 5.0 μg/mL	0 %	Susceptible
Ethionamide 10.0 µg/mL	33 %	Resistant
Capreomycin 10.0 μg/mL	0 %	Susceptible
PAS 2.0 μg/mL	0 %	Susceptible
Ofloxacin 2.0 µg/mL	0 %	Susceptible
Amikacin 4.0 µg/mL	0 %	Susceptible
Comments and Disclaimers  - Susceptibility testing method: Indirect agar prop drug-containing medium compared to drug-free m This test has not been cleared or approved by the the DTBE Reference Laboratory.	edium).	
MTBC Pyrazinamide Susceptibility*	Result	
Pyrazinamide 100 μg/mL	Susceptible	
Comments and Disclaimers  * Susceptibility testing method: Mycobacteria Grov	vth Indicator Tube (MGIT)	



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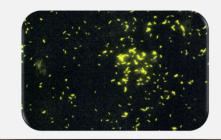








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- Resistance to SM
- CDC final 4/27/23
- Agrees with DSHS
- Repeat susceptibilities after 3 months of culture positive 6/27/23
- No acquired additional resistance





## **Genotype Investigation**

DST	Relation	Originating Lab		Submitter Number		wgMLSType	GENType	SpoligoType	MIRU	MIRU2
		GRUPO SIN FRONTERAS								
MDR	(current pt)	BINATIONAL PROJ	03/13/2023	AMCC2303733	03/29/2023	MTBC002441				

### **Genotype Investigation** Originating Date of Submitter **Date** GRUPO SIN FRONTERAS BINATIONAL PROJ 03/13/2023 AMCC2303733 03/29/2023 MTBC002441 MDR (current pt) SOUTH TEXAS LABORATORY 09/24/2012 AMRC1202584 10/31/2012 777776777760771 224325153314 333334213338 MDR (father) G11225 GRUPO SIN FRONTERAS MDR (brother-in-law) BINATIONAL PROJ 08/06/2018 AMRC1802226 10/03/2018 MTBC002441 G40790 777776777760771 224325133314 333334213338 **GRUPO SIN** pre-XDR (b-in-law's sister) BINATIONAL PROJ 07/09/2022 AMCC2209108 08/04/2022 MTBC002441

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