# CDC Model Performance Evaluation Program (MPEP) for Mycobacterium tuberculosis Drug Susceptibility Testing

Attachment 11
Expected Results Report

### March 2024 TB MPEP

## **Expected Results Report**

Results listed are expected results and may not necessarily be consistent with aggregate data when results from all participating laboratories are analyzed. Please refer to the forthcoming Final Report for aggregate data and detailed information for each isolate.

#### **Expected Growth-based Results**

Isolate	RIF	INH	EMB	PZA	FQ	Second-line Drug Resistances:
2024A	S	R (low-level*)	S	S	S	
2024B	S	S	S	S	R	
2024C	S	R (high-level†)	S	S	S	
2024D	R	S	S	S	S	RBT
2024E	S	S	S	R	S	

Note: RIF=rifampin, INH=isoniazid, EMB=ethambutol, PZA=pyrazinamide, FQ=fluoroquinolones, RBT=rifabutin, S=susceptible, R=resistant. Growth-based DST results were obtained by agar proportion method, except for pyrazinamide, in which MGIT™ was performed.

### **Expected Molecular Results**

Isolate	rpoB*	inhA	katG	gyrA	pncA
2024A		C-15T			
2024B				Ala90Val	
2024C			Ser315Thr		
2024D	Ser450Leu				
2024E					His57Asp

Note: Empty cell=No mutation detected. Molecular results were obtained by whole genome sequencing. High confidence mutations were not detected in these loci: fabG1, embB, ethA, eis, rrs, and tlyA.

<sup>\*</sup>Resistant at 0.2  $\mu$ g/ml by agar proportion.

 $<sup>^{\</sup>dagger}$ Resistant at 0.2 and 1.0  $\mu$ g/ml by agar proportion.

<sup>\*</sup> *M. tuberculosis* numbering system used [1, 2]

<sup>1.</sup> Andre, E., et al., Consensus numbering system for the rifampicin resistance-associated rpoB gene mutations in pathogenic mycobacteria. Clin Microbiol Infect, 2017. 23(3): p. 167-172.

<sup>2.</sup> APHL, Issues in Mycobacterium tuberculosis complex (MTBC) Drug Susceptibility Testing: Rifampin (RIF), in APHL Issues in Brief: Infectious Diseases. 2019, Association of Public Health Laboratories: Washington, D.C.