



MYCOBACTERIUM TUBERCULOSIS WHOLE GENOME SEQUENCING REPORT

NOT FOR DIAGNOSTIC USE

| | | | |
|---------------|-------------|-------------|------------|
| Patient Name | GORDON COLE | Patient ID | 23456789 |
| Birth Date | 1862-12-12 | Location | CAMBRIDGE |
| Sample Type | SPUTUM | Sample Date | 1916-12-26 |
| Reporting Lab | OXFORD | Report Date | 1917-01-06 |

Summary

The specimen was positive for **Mycobacterium tuberculosis**. It is predicted to be **resistant to Isoniazid and Rifampin**. It belongs to a cluster, suggesting **recent transmission**

Organism

The specimen was positive for **Mycobacterium tuberculosis**

Drug Susceptibility

Drug susceptibility is predicted by the detection of known *M. tuberculosis* drug resistance conferring genetic mutations

- ☐ No drug resistance predicted
- ☐ Mono-resistance predicted
- ☒ Multi-drug resistance predicted
- ☐ Extensive drug resistance predicted

| Drug class | Prediction | Drug | Resistance Gene (Amino Acid Mutation) |
|-------------|------------|---------------|---------------------------------------|
| First-line | Sensitive | Ethambutol | No resistance mutation detected |
| | | Pyrazinimide | No resistance mutation detected |
| | Resistant | Isoniazid | katG (S315T) |
| | | Rifampin | rpoB (S531L) |
| Second-line | Sensitive | Streptomycin | No resistance mutation detected |
| | | Ciprofloxacin | No resistance mutation detected |
| | | Moxifloacin | No resistance mutation detected |
| | | Amikacin | No resistance mutation detected |
| | | Kanamycin | No resistance mutation detected |
| | | Capreomycin | No resistance mutation detected |



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Cluster Detection

Current specimen was found to be closely clustered with previous specimens suggesting **recent transmission**

| Relatedness | Number of prior matching isolates |
|---------------------------------------|-----------------------------------|
| Likely Related (< 5 SNPS apart) | 2 isolates |
| Possibly Related (6 to 30 SNPS apart) | 2 isolates |

