

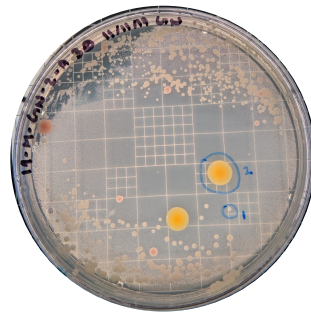
# Isolation and Identification of NTM from the Hawaiian Environment



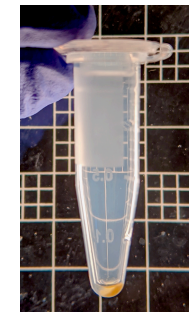
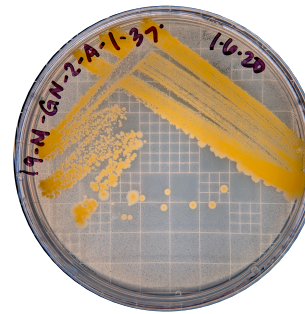
1. Sample Environment



2. Culture Sample



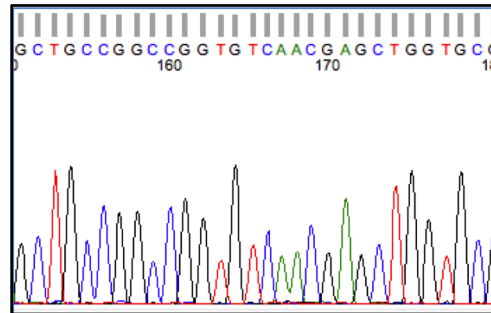
3. Pick Isolate



4. Extract DNA



5. Sequence DNA



6. Identify Isolate

<b>Mycobacterium neoaurum strain ATCC 25795 RNA polymerase beta subunit (rpoB) gene, partial cds</b>				
Sequence ID: <a href="#">JF712875.1</a> Length: 741 Number of Matches: 1				
Range 1: 38 to 741 <a href="#">GenBank</a> <a href="#">Graphics</a> <a href="#">▼ Next Match ▲</a>				
Score	Expect	Identities	Gaps	Strand
1249 bits(676)	0.0	698/708(99%)	4/708(0%)	Plus/Plus

Basic toolkit for identifying environmental NTM isolates.