SUPPLEMENTARY INFORMATION FOR

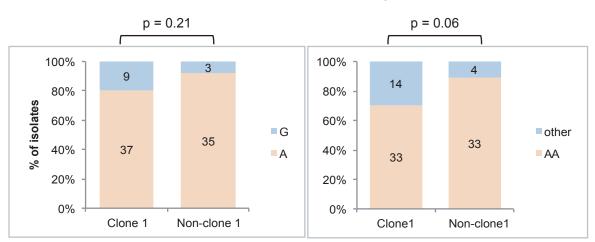
Genomic Characterization of Sporadic Isolates of the Dominant Clone of *Mycobacterium abscessus* subspecies *massiliense*

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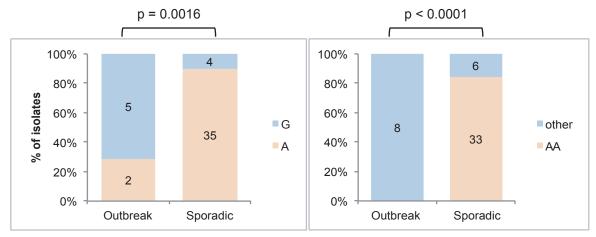
Running Title: Dominant Clone of Mycobacterium massiliense

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Supplemental Figure 1. Proportions of known antimicrobial resistance (AMR) mutations in *M. massiliense* isolates. The 16S rRNA position 1408 for amikacin mutational resistance and 23S rRNA positions 2058 and 2059 for clarithromycin mutational resistance were analyzed in *M. massiliense* genomes using one isolate per patient. Genotypes in orange are the wild type, and genotypes in blue are the AMR genotype. Percent of isolates are shown on y-axes. **A.** *M. massiliense* isolates are classified as Clone 1 vs. Non-Clone 1. **B.** Clone 1 isolates are classified as Outbreak vs. Sporadic.

A. 16S rRNA (position 1408) amikacin mutational resistance

