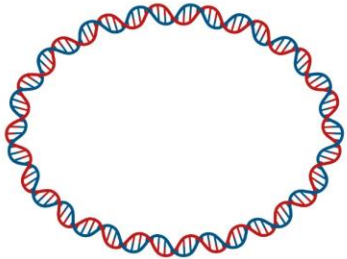


Genome Announcement: XDR *Acinetobacter baumannii*

(Acc#SRR23106361)

1. Genome Sequencing



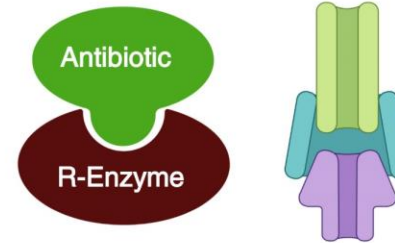
We present the whole genome sequences of a XDR nosocomial pathogen *Acinetobacter baumannii* isolated from the blood of a sepsis patient

2. Analysis



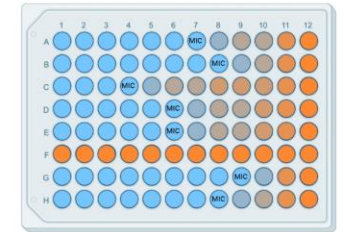
This genomic analysis provides insights into the organism's resistance mechanisms, virulence factors, and potential pathogenicity.

3. Insights



AMR is highly heterogeneous, with various factors reducing the efficacy of commonly used antibiotics against *A. baumannii* infections.

4. Translation



Understanding these genomic features is crucial for developing effective therapies and diagnostics for a better clinical outcomes .

