

Genome size: 4.37mb

GC percent: 42%

CDS: 4034

rRNA=10, tRNA=79

mportant Functions

characteristics

AMR genes: aac(2')-la, aac(6'), ant(2")-la, ant(3")-lia, aph(3'), blaNDM, blaOXA, mphE, cat, arr, tet(B), armA, msrE, dfr, sul

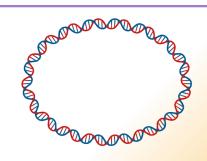


## Genome Announcement: Providencia stuartii

Accession#SAMEA116288275 (NCBI) INS0004956 (IBDC)



#### 1. Genome Sequencing



# **BRIC-THSTI** presenting the whole

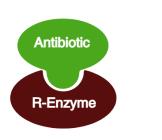
genome sequence of *Providencia* stuartii isolated from the North Indian wastewater source.

#### 2. Analysis



The genomic analysis highlights the AMR dynamics and revealing the presence of diverse resistance genes within the bacterium contributing to its multi drug resistance profile.

### 3. Insights



Genomic insights explore the presence of versatile ARGs including the carbapenemase gene *blaNDM-1*, *blaOXA*, *dfr* and their association with several MGEs.

#### 4. Translation



Providencia stuartii
genomic feature
provide an early
warning system for
potential outbreaks
and inform clinical
guidelines for empiric
treatment and
infection control
measures