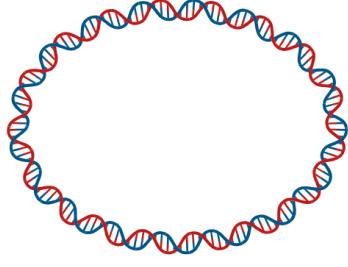


Genome Announcement: *Providencia spp.*

Accession# SAMEA116293622 (NCBI)
INS0005012 (IBDC)

1. Genome Sequencing



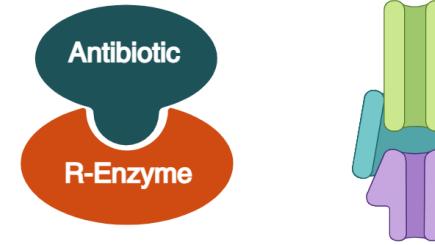
BRIC-THSTI
presenting the
complete genome
sequence of
Providencia spp.
isolated from Indian
urban wastewater.

2. Analysis



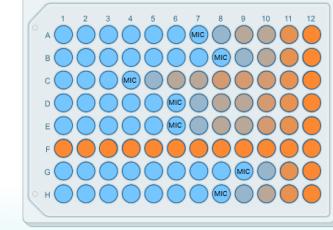
This genomic analysis provides key insights into the specific genes driving antibiotic resistance and virulence, conferring resistance to aminoglycosides, chloramphenicol, and tetracyclines..

3. Insights



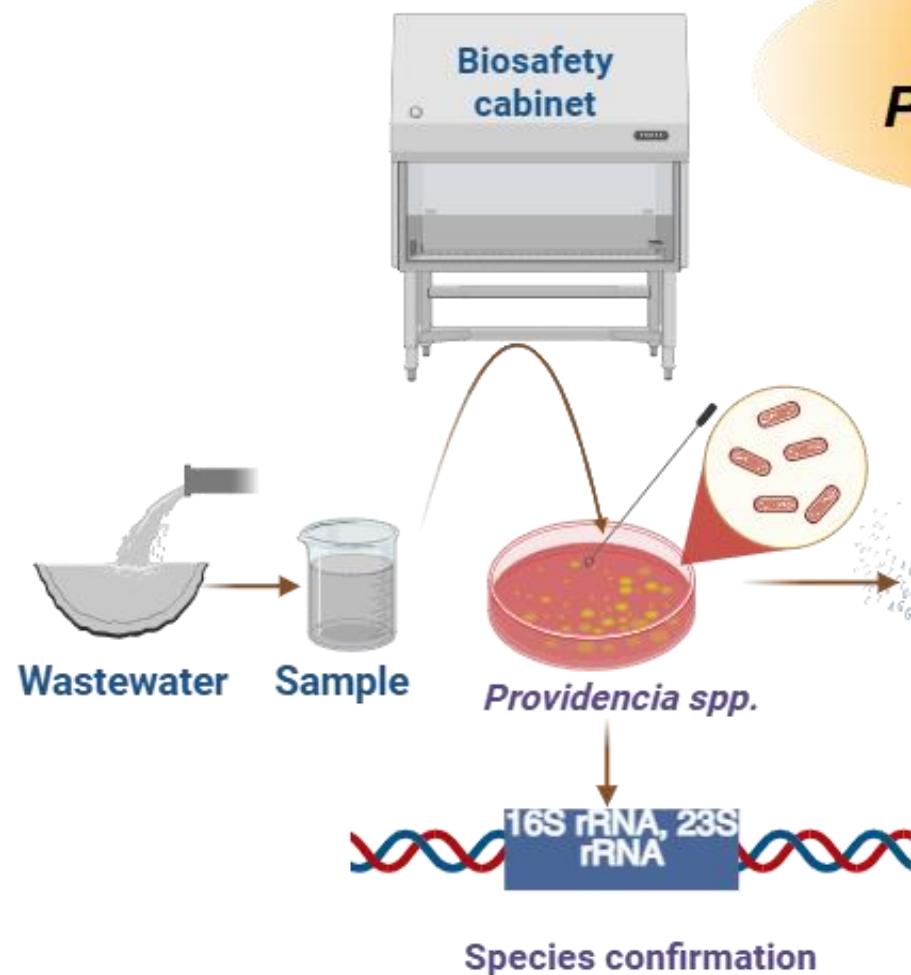
Providencia spp. strain carries resistance genes against aminoglycosides ***aac(2')*-*Ia***, chloramphenicol (***cat***), and tetracyclines (***tet(B)***) resistance genes.

4. Translation



Linking genomic data to clinical action, this translational approach highlights how resistance genes in *Providencia spp.* can inform targeted antibiotic choices, guide diagnostics, and support infection control strategies.

***Providencia spp.*: A resilient opportunist, quietly spreading resistance in tough environments.**



Genomics of *Providencia spp.*

GENOMICS

Genome size: 2.44Mb
GC percent: 45%
CDS: 4026
rRNA=12 , tRNA=80

Important Functions

AMR genes: *aac(2')*-
la, *cat*, *tet(B)*