





Wastewater Sample Enterococcus raffinosus

Next Generation

Species confirmation

Sequencing

Genome characteristics

Genome size: 4.33mb

GC percent: 40%

CDS: 4232

rRNA=7, tRNA=64

Important Functions

AMR genes: aac(6')-leaph(2")-la, aph(3'), aad(6), erm, sat-4, tetM

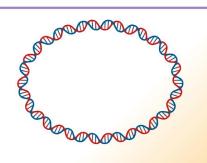


Genome Announcement: *Enterococcus raffinosus*

BRIC thsti

Accession#SAMEA116288592 (NCBI) INS0005074 (IBDC)

1. Genome Sequencing



BRIC-THSTI

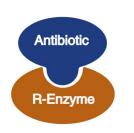
presenting the whole genome sequence of *Enterococcus* raffinosus isolated from the Indian urban wastewater.

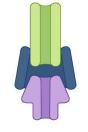
2. Analysis



This genomic analysis carries highrisk resistance genes with mobile genetic potential, posing a clinical and public health threat requiring urgent surveillance and infection control measures.

3. Insights





The insights of the *Enterococcus* raffinosus showing multi drug resistance against 4+ antibiotic classes located on MGEs like plasmids/transposons.

4. Translation



Genomic surveillance of Enterococcus raffinosus in wastewater enables tracking of antibiotic-resistant strains across hospital and community environments.

Enterococcus raffinosus: 'A Genomic Sentinel' for Antibiotic Resistance Surveillance