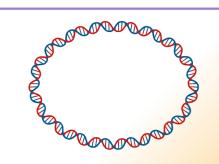


# Genome Announcement: Alcaligenes sp.

Accession#SAMEA116290119 (NCBI) #INS0005067 (IBDC)



#### 1. Genome Sequencing



# **BRIC-THSTI**

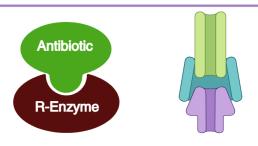
presenting the whole genome sequence of *Alcaligenes sp.* isolated from the Indian wastewater source.

### 2. Analysis



Genomic analysis
uncovers a synergistic
resistance
mechanism: the
AdeABC efflux system
and SMR transporter
work together,
conferring resistance
to industrial
disinfectants.

## 3. Insights



It carries cooccurrence of adeF
and qacG indicates
adaptation to both
clinical (antibiotics)
and industrial
(disinfectants)
pressures suggesting
a potential horizontal
gene transfer

#### 4. Translation



Translational insights propose a new therapeutic strategy involving disinfectant rotation and cross-sector alignment to break resistance transmission cycles.

Alcaligenes sp.: Packs a double punch—adeF and qacG fuel resistance, turning it into a stealthy MDR threat!

e characteristics

Genome size: 3.77Mb

GC percent: 56%

CDS: 3430

rRNA=03, tRNA=54

Important Functions

AMR genes: adeF, qacG

Environmental reservoirs of antimicrobial resistance

