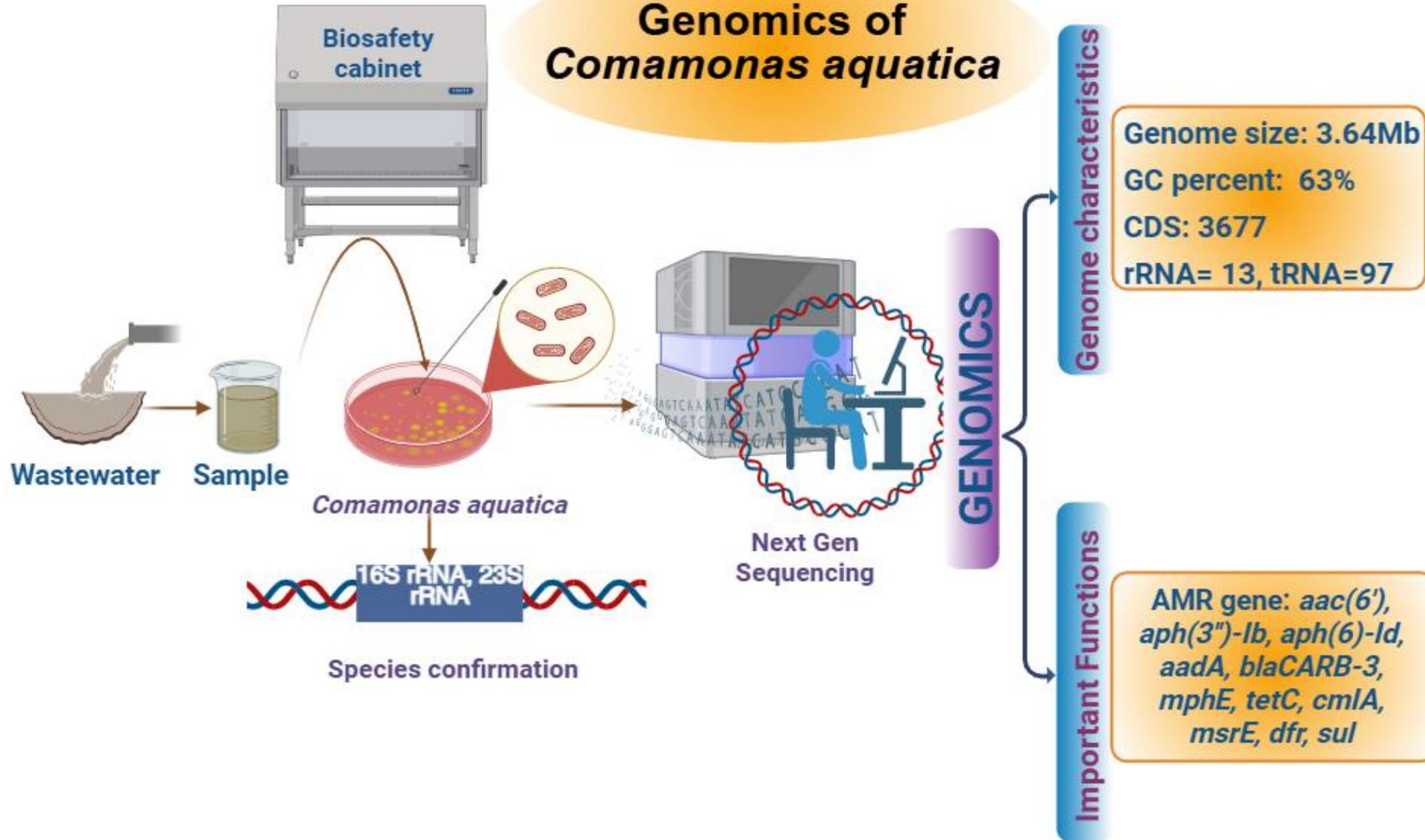




Genomics of *Comamonas aquatica*



Genome characteristics

Genome size: 3.64Mb
GC percent: 63%
CDS: 3677
rRNA= 13, tRNA=97

GENOMICS

Important Functions

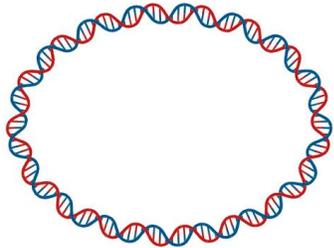
AMR gene: *aac(6')*,
aph(3'')-Ib, *aph(6)-Id*,
aadA, *blaCARB-3*,
mphE, *tetC*, *cmlA*,
msrE, *dfr*, *sul*

Genome Announcement: *Comamonas aquatica*

Accession#SAMEA116293470 (NCBI)

INS0005206 (IBDC)

1. Genome Sequencing



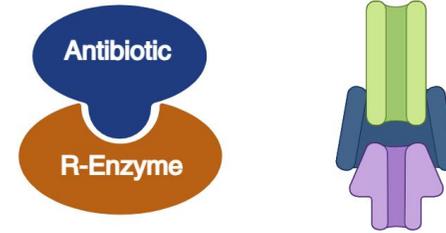
BRIC-THSTI presenting the complete genome sequence of *Comamonas aquatica* isolated from the Indian wastewater source.

2. Analysis



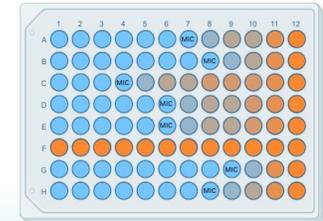
This genomic analysis demonstrates as breeding grounds for MGE-driven resistance dissemination with a potential threat on multi-drug co resistance.

3. Insights



Comamonas aquatica insights into the resistance against the multiple drugs including Aminoglycosides, β -Lactams, Macrolides, Tetracyclines, Chloramphenicol, and Sulfonamides/Trimethoprim

4. Translation



These insights bridge genomics to real-world solutions-arming clinicians with diagnostic tools, guiding policymakers on containment strategies, and fueling R&D for next-generation antimicrobials.

***Comamonas aquatica*: Exposes the Hidden Highway of Antibiotic Resistance Spread**