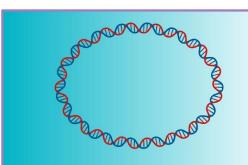


Genome Announcement: Providencia manganoxydans Accession#SAMEA116288277(NCBI) **Comparison** **Comparison**

INS0004959 (IBDC)



1. Genome Sequencing



BRIC-THSTI

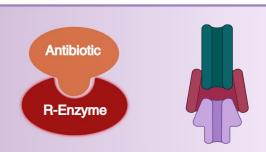
presenting the whole genome sequence of Providencia manganoxydans from the North Indian sewage water source.

2. Analysis



Genomic analysis suggests potential resistance to multiple classes of antibiotics and identifies the presence of an efflux regulator gene crp which may contribute to the resistance mechnism.

3. Insights



The genomic Insights illustrate the presence of multiple resistance genes (vanG, qacJ, FosA8, KpnH, KpnF) including the crp, a global regulator gene of the efflux pump.

4. Translation

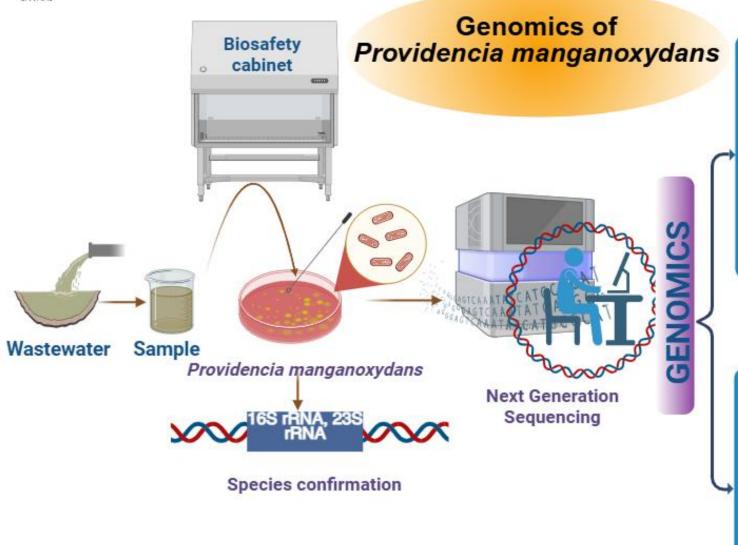


This genomic evaluation calls for a for specific actions, including improved surveillance and infection control for the betterment of the public health.

Providencia manganoxydans: An environmental bacterium with clinical resistance profile







Genome size: 4.86Mb

GC percent: 41.6%

CDS: 3987

Genome characteristics

rRNA= 8, tRNA=83

AMR g AMR g Kp

AMR genes: vanG, qacJ, FosA8, KpnH, KpnF, CRP