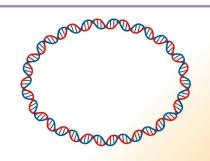


Genome Announcement: Paenibacillus spp.

Accession# SAMEA120513486(NCBI) #INS0022920 (IBDC)



1. Genome Sequencing



BRIC-THSTI

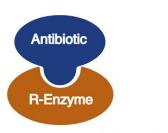
presenting the whole genome sequence of *Paenibacillus* spp. isolated from the pancreatic tissue of the Indian adult.

2. Analysis



The genomic features demonstrates an opportunistic pathogen or a intratumoral microbiota especially in pancreatic diseases warrants a focus on its safety and virulence.

3. Insights



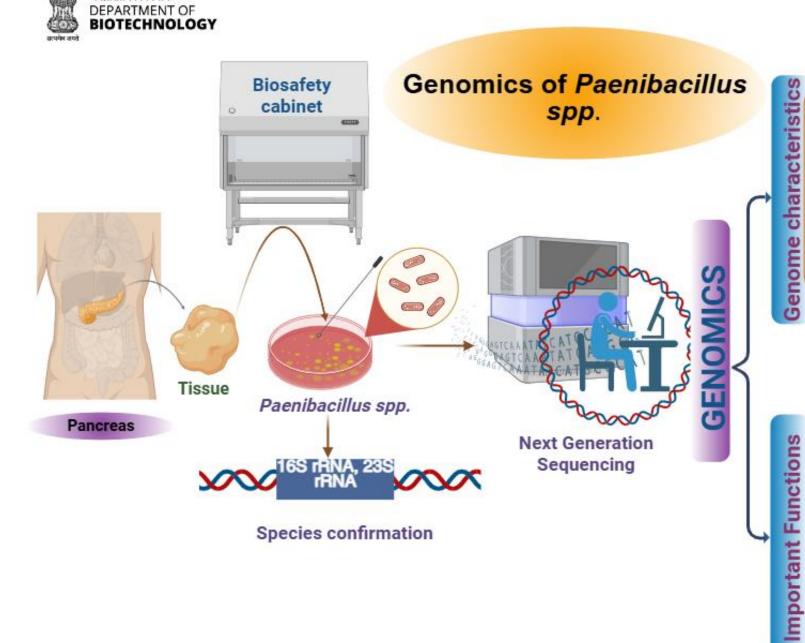
The specific detection of *vatB* against the virginiamycin provides and **CRISPR** based defence system insight into the exact mechanisms of this bacterium uses to protect itself against phages.

4. Translation



Paenibacillus spp.
is exceptionally high,
primarily focusing on
Clinical/Medical and
Public Health
applications, like
understanding its role
in pancreatic health/
disease.





Genome size: 6.22Mb

GC percent: 43.8%

CDS: 5498

Total RNA genes: 61

No. of CRISPRs: 1

AMR genes: vatB