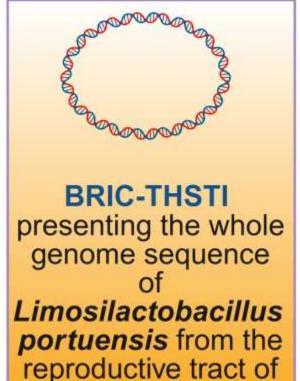
Genome Announcement: Limosilactobacillus portuensis



Accession#SRR28593025 (NCBI) #INS0004703 (IBDC)

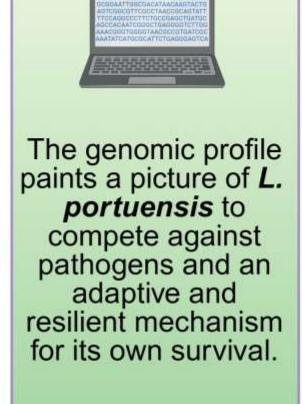


1. Genome Sequencing

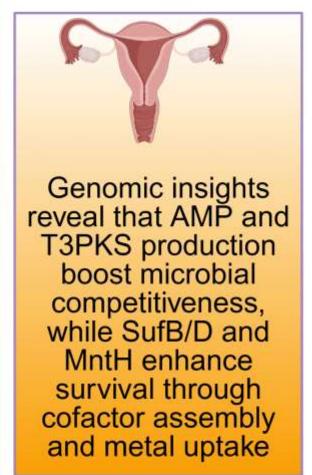


Indian women.

2. Analysis



3. Insights



4. Translation



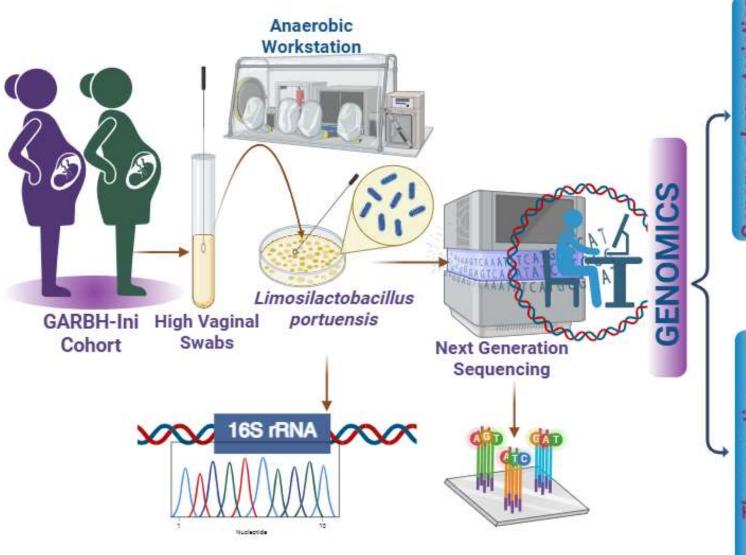
This genomic analysis offers a roadmap to leverage Limosilactobacillus portuensis for drug discovery, health improvement, and industrial innovation.

Limosilactobacillus portuensis: A microbial treasure chest to battle against the pathogens





Whole Genome sequence of Limosilactobacillus portuensis



Genome size: 1.87mb

GC percent: 40.1

RNAs: 70,

Subsystems: 194

Transcription regulators:

Helix turn helix, LysR

Family

Antimicrobial peprtides:
T3PKS (TypeIII

Polyketide synthase)

Secretory signal

peptides: 4

SufB: 1, SufD: 1, MntH: 1