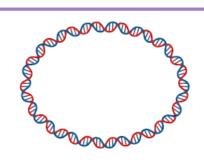


Genome Announcement: Prevotella copri Accession#NMPZ00000000 (NCBI)

INGCA000000188(INDA)



1. Genome Sequencing



BRIC-THSTI

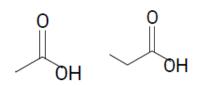
presenting the whole genome sequence of Prevotella copri isolated from the gut of a healthy Indian adult.

2. Analysis



Through this genomic analysis, we gain valuable insights for its ability to degrade a wide range of complex carbohydrate including fibres, polysaccharides and oligosaccharides.

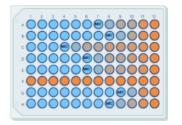
3. Insights



Short Chain Fatty Acids

P. copri plays an important role in immunomodulation, with the potential for both beneficial and detrimental effects depending on host health and diet.

4. Translation



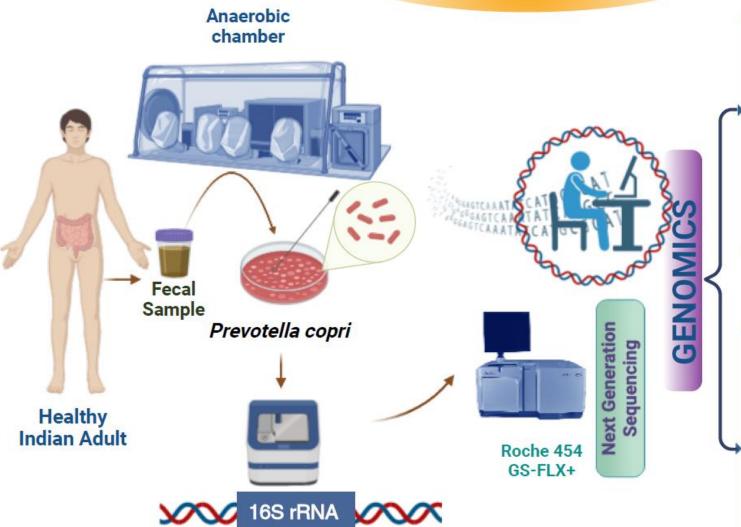
Understanding this genomic feature is important to developed live biotherapeutics based on the active biomass of P. copri.





Genomics of Prevotella copri





Sanger Sequencing confirmation

Genome size: 3.65Mb

GC percent: 45.4%

CDS: 3,112

Genome characteristics

Stable RNAs: 94

Important functions SCFAs, CAZymes, BSH, Alcohol Dehydrogenase