



One Day One Genome

Ligilactobacillus salivarius

A lactic acid bacterium with promising probiotic potential known for its antimicrobial activity and immunomodulatory effects



Isolation

A bacterial isolate from the gut microbiota of laboratory mice (C57BL/6J mice)



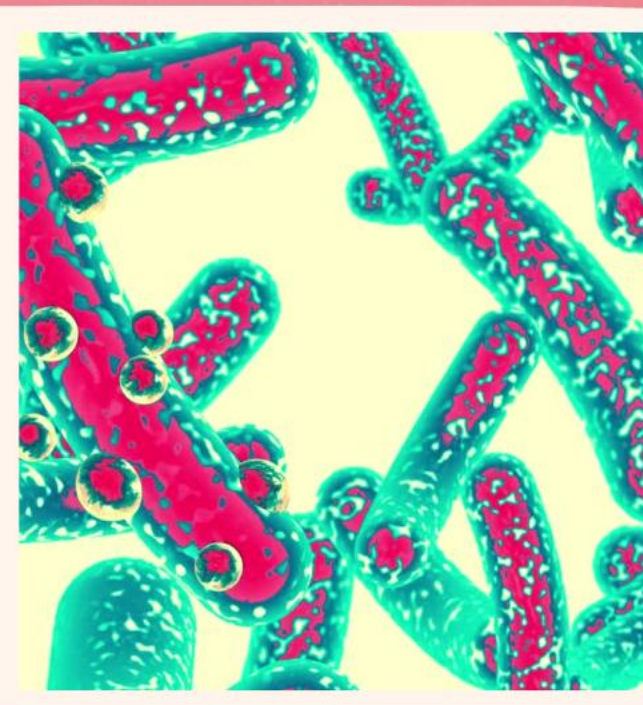
Insights

High-throughput sequencing reveals genes linked to nutrient metabolism and antimicrobial resistance



Application

Insight into the role of gut microbiota in host health especially their therapeutic potential against infections



Ligilactobacillus salivarius (C57BL6J)

Genome Accession Number: INS0010333

Quality of Genome Assembly and Annotation:

Results from indigenously developed **BHARAT** analysis pipeline: (**B**acterial **H**ybrid genome **A**ssembly and **R**apid **A**nnotation **T**oolset)

Table 1: Assembly Details	
Contigs	41
GC Content	32.74
Contig L50	4
Genome length	1,936,431 bp
Contig N50	119,833

Table 2: Annotated Genome Features	
CDS	1,902
tRNA	44
Repeat Regions	21
rRNA	4

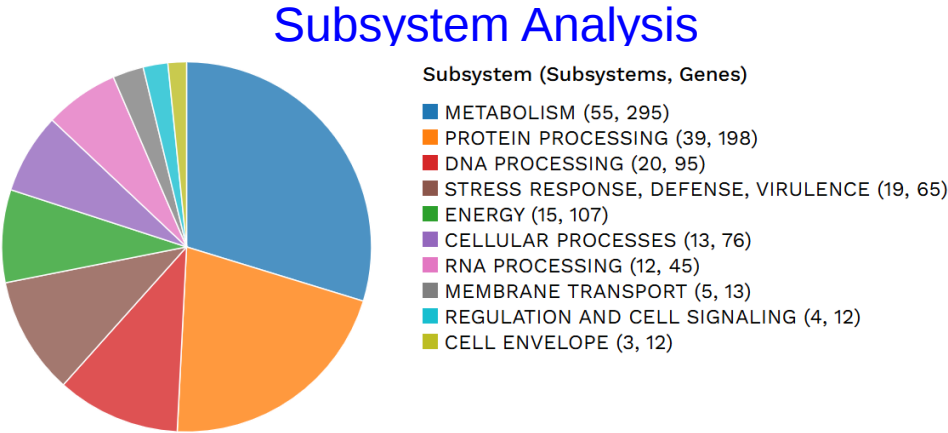


Table 3: Antimicrobial Resistance Genes	
AMR Mechanism	Genes
Antibiotic target in susceptible species	Alr, Ddl, EF-G, EF-Tu, folA, Dfr, gyrA, gyrB, inhA, fabI, Iso-tRNA, kasA, MurA, rho, rpoB, rpoC, S10p, S12p
Antibiotic target modifying enzyme	RlmA(II)
Antibiotic target replacement protein	FabK
Gene conferring resistance via absence	gidB
Protein altering cell wall charge conferring antibiotic resistance	MprF, PgsA

