



#OneDayOneGenomeInitiative

Staphylococcus aureus

Spa type t021, ST243, CC30

Causes **mastitis** i.e. inflammation (pain, redness, and swelling) in the udder of cows, also contaminating the milk they produce.

Isolated from



Milk of affected cows



Genomic Analysis



Revealed genetic markers associated with efflux pumps and genes conferring resistance to quinolones and metallothiols - drugs commonly used against this disease.

Insights



Staph. aureus can
escape cow's
immune system using
evasion proteins. It
also has secretion
systems, & toxins
including
enterotoxins and
hemolysins!

Application



Part of the first study
to present a
comprehensive
whole-genome
analysis of bovine
mastitis-associated S.
aureus isolates from
India!

Staphylococcus aureus GCA_018995185.1

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	39
GC Content	32.73
Contig L50	6
Genome Length	2,732,334 bp
Contig N50	170,521

Table 2: Annotated Genome Features		
CDS	2,593	
tRNA	55	
rRNA	4	

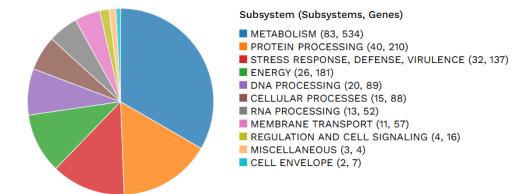
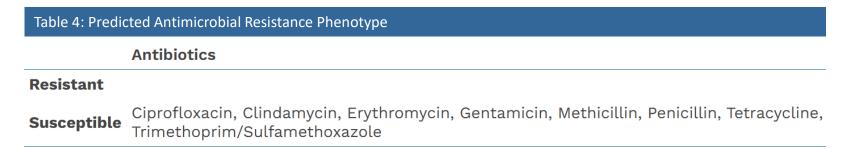


Table 3: Antimicrobial Resistance Genes		
AMR Mechanism	Genes	
Antibiotic inactivation enzyme	FosB	
Antibiotic resistance gene cluster,cassette,or operon	TcaA, TcaB, TcaB2, TcaR	
Antibiotic target in susceptible species	Alr, Ddl, EF-G, EF-Tu, folA, Dfr, folP, gyrA, gyrB, inhA, fabI, Iso-tRNA, kasA, MurA, rho, rpoB, rpoC, S10p, S12p	
Efflux pump conferring antibiotic resistance	BceA, BceB, NorA, Tet(38)	
Gene conferring resistance via absence	gidB	
Protein altering cell wall charge conferring antibiotic resistance	GdpD, MprF, PgsA	
Regulator modulating expression of antibiotic resistance genes	BceR, BceS, LiaF, LiaR, LiaS	



Genome Assembly

