



## *Staphylococcus epidermidis* (NIBMG\_HMR\_19)

Beneficial Bacteria of Human Skin

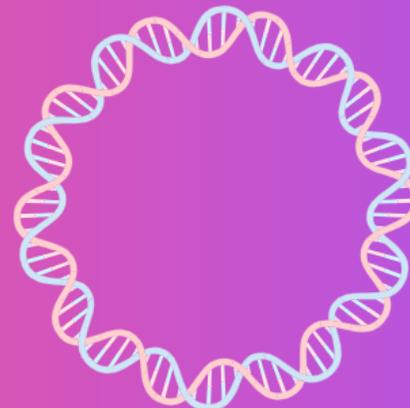
Isolated from Healthy Human Skin

✓ Protects against Pathogenic Bacteria

✓ Produces Antimicrobial Peptides

✓ Modulate Immunity

✓ Helps in Wound Healing



# *Staphylococcus epidermidis* (NIBMG\_HMR\_19)

Genome Accession Number: [INS0009755](https://www.ncbi.nlm.nih.gov/assembly/INS0009755)

## 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	47
GC Content	32.05
Contig L50	6
Genome length	2,465,463 bp
Contig N50	129,718

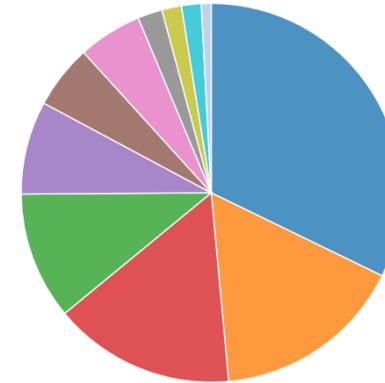
**Table 2: Annotated Genome Features**

CDS	2,379
tRNA	53
rRNA	4
Repeat Regions	0

**Table 3: Antimicrobial Resistance Genes**

AMR Mechanism	Genes
Antibiotic inactivation enzyme	APH(3')-III/APH(3')-IV/APH(3')-VI/APH(3')-VII, BlaZ family, CatA7 family, FosB, Mph(C) family
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, YkkCD
Protein altering cell wall charge conferring antibiotic resistance	GdpD, MprF, PgsA

## Subsystem Analysis



Subsystem (Subsystems, Genes)

- METABOLISM (77, 518)
- PROTEIN PROCESSING (39, 211)
- STRESS RESPONSE, DEFENSE, VIRULENCE (37, 137)
- ENERGY (26, 190)
- DNA PROCESSING (19, 85)
- CELLULAR PROCESSES (13, 79)
- RNA PROCESSING (13, 50)
- MEMBRANE TRANSPORT (5, 32)
- REGULATION AND CELL SIGNALING (4, 16)
- MISCELLANEOUS (4, 7)
- CELL ENVELOPE (2, 7)

## Genome Assembly

