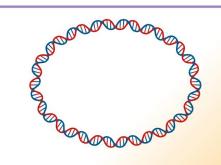


# Genome Announcement: Staphylococcus haemolyticus

Accession #SAMEA120513482 (NCBI) #INS0022916 (IBDC)

### 1. Genome Sequencing



### **BRIC-THSTI**

presenting the whole genome sequence of **Staphylococcus haemolyticus** isolated from the neosepsis blood sample.

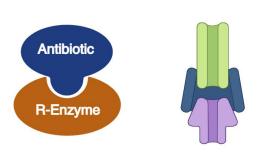
### 2. Analysis



Genomic analysis of a clinically significant bacterium,

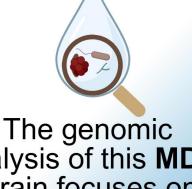
Staphylococcus haemolyticus,
revealing a high level of multidrug resistance and mobile genetic elements.

## 3. Insights



The presence of multiple, distinct ARGs like APH(3')Illa, blaZ, SAT-4
confirms this strain's multidrug-resistant (MDR) nature, which is a hallmark of S. haemolyticus.

#### 4. Translation

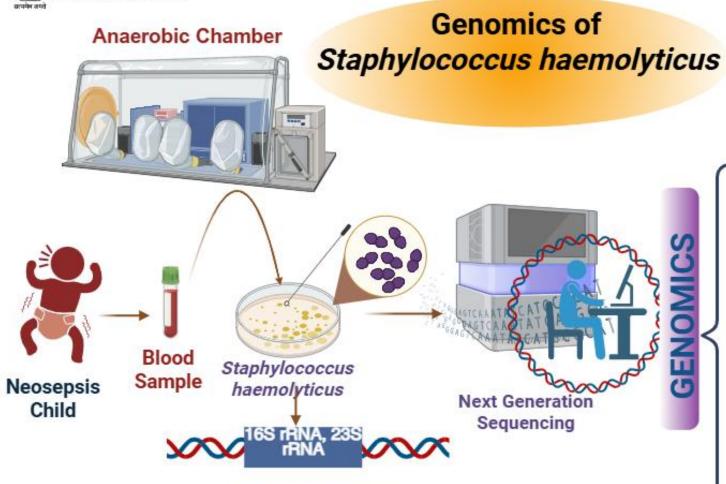


analysis of this MDR
strain focuses on
providing important
insights for
managing public
health risks and
developing new
drugs, including
public health
monitoring

Staphylococcus haemolyticus: The Hospital Superbug is armed with ARGs and a mobile escape route.







Species confirmation

characteristics Genome size: 2.3Mb GC percent: 32.71% CDS: 2254

Total RNA=66

Plasmids: 3

AMR genes: APH(3')-IIIa, PC1\_betalactamase\_(blaZ), SAT-4, mgrA