



Government of India
Ministry of Science & Technology
Department of Biotechnology



ONE DAY ONE GENOME

Escherichia coli

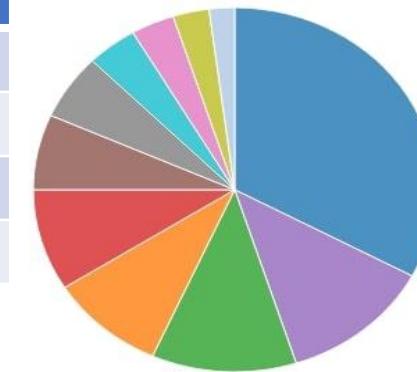
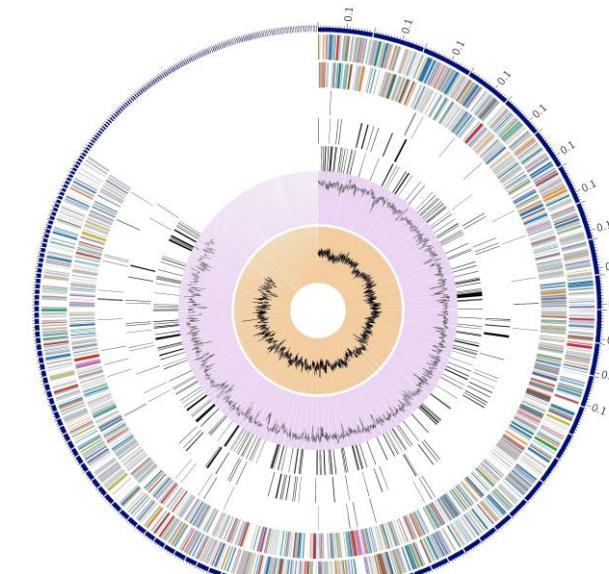


Quality of Genome Assembly and Annotation:Results from indigenously developed **BHARAT** analysis pipeline: (**Bacterial Hybrid genome Assembly and Rapid Annotation Toolset**)**Table 1: Assembly Details**

| | |
|---------------|--------------|
| Contigs | 248 |
| GC Content | 50.36 |
| Contig L50 | 22 |
| Genome length | 4,898,023 bp |
| Contig N50 | 59,341 |

Table 2: Annotated Genome Features

| | |
|----------------|-------|
| CDS | 4,972 |
| tRNA | 81 |
| Repeat Regions | 0 |
| rRNA | 4 |

Subsystem Analysis**Genome Assembly**

| AMR Mechanism | Genes |
|--|---|
| Antibiotic inactivation enzyme | BlaEC family, CTX-M family |
| Antibiotic target in susceptible species | Alr, Ddl, dxr, EF-Tu, folA, Dfr, folP, gyrA, gyrB, inhA, fabI, Iso-tRNA, kasA, MurA, rho, rpoB, rpoC, S10p, S12p |
| Efflux pump conferring antibiotic resistance | AcrAB-TolC, AcrAD-TolC, AcrEF-TolC , AcrZ, EmrAB-TolC, EmrD, EmrE, EmrKY-TolC, MacA, MacB, MdfA/Cmr, MdtABC-TolC, MdtEF-TolC, MdtL, MdtM, SugE, TolC/OpmH |
| Antibiotic target modifying enzyme | Erm(B) |
| Protein altering cell wall charge conferring antibiotic resistance | GdpD, PgsA, |
| Regulator modulating expression of antibiotic resistance genes | AcrAB-TolC, EmrAB-TolC, GadE, H-NS, OxyR |