



Government of India
Ministry of Science & Technology
Department of Biotechnology

सत्यमेव जयते



a DBT Organization

Biotechnology & Biomanufacturing



**ONE DAY
ONE GENOME**

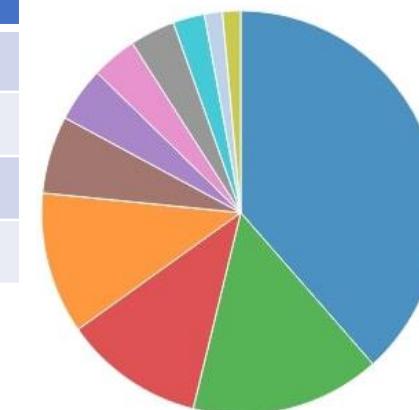
Rhodococcus erythropolis C27

UNDERSTANDING ITS GENOMIC POTENTIAL AIDS IN
DESIGNING ECO-SOLUTIONS FOR DETOXIFYING
PHENOLIC COMPOUNDS FROM INDUSTRIAL AND OIL-
CONTAMINATED SITES

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	1
GC Content	62.53
Contig L50	1
Genome length	6,274,870 bp
Contig N50	6,274,870

Table 2: Annotated Genome Features	
CDS	5,974
tRNA	52
Repeat Regions	17
rRNA	3

**Subsystem Analysis**

Subsystem (Subsystems, Genes)	Count
METABOLISM (105, 1063)	105, 1063
PROTEIN PROCESSING (42, 231)	42, 231
STRESS RESPONSE, DEFENSE, VIRULENCE (31, 167)	31, 167
ENERGY (31, 283)	31, 283
DNA PROCESSING (17, 82)	17, 82
CELLULAR PROCESSES (12, 92)	12, 92
MEMBRANE TRANSPORT (10, 48)	10, 48
RNA PROCESSING (10, 43)	10, 43
CELL ENVELOPE (7, 36)	7, 36
REGULATION AND CELL SIGNALING (4, 16)	4, 16
MISCELLANEOUS (4, 43)	4, 43

Table 3: Antimicrobial Resistance Genes

AMR Mechanism	Genes
Antibiotic target in susceptible species	Alr, Ddl, dxr, EF-G, EF-Tu, folA, Dfr, folP, gyrA, gyrB, inhA, fabI, Iso-tRNA, kasA, MurA, rho, rpoB, rpoC, S10p, S12p
Antibiotic target replacement protein	FabG, HtdX
Gene conferring resistance via absence	gidB
Protein altering cell wall charge conferring antibiotic resistance	GdpD, PgsA,
Regulator modulating expression of antibiotic resistance genes	LpqB, MtrA, MtrB, OxyR

Genome Assembly