



Government of India
Ministry of Science & Technology
Department of Biotechnology



ONE DAY ONE GENOME

Natronococcus pandeyae sp. nov.

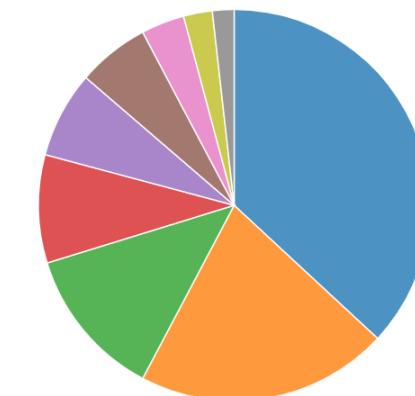


Quality of Genome Assembly and Annotation:Results from indigenously developed **BHARAT** analysis pipeline: (**Bacterial Hybrid genome Assembly and Rapid Annotation Toolset**)**Table 1: Assembly Details**

Contigs	50
GC Content	63.19 %
Contig L50	6
Genome length	5,316,806 bp
Contig N50	288,393

Table 2: Annotated Genome Features

CDS	5,893
tRNA	52
Repeat Regions	0
rRNA	4

Subsystem Analysis

Subsystem (Subsystems, Genes)

METABOLISM	(62, 429)
PROTEIN PROCESSING	(35, 151)
ENERGY	(21, 133)
RNA PROCESSING	(15, 66)
STRESS RESPONSE, DEFENSE, VIRULENCE	(12, 33)
DNA PROCESSING	(10, 28)
MEMBRANE TRANSPORT	(6, 27)
MISCELLANEOUS	(4, 6)
CELLULAR PROCESSES	(3, 8)

Table 3: Antimicrobial Resistance Genes

AMR Mechanism	Genes
Antibiotic target in susceptible species	folA, Dfr, folP, gyrA, gyrB, Iso-tRNA
Antibiotic activation enzyme	KatG
Protein altering cell wall charge conferring antibiotic resistance	GdpD

Genome Assembly