



ONE DAY ONE GENOME

Erythrobacter sp.



The microbe showcases strong metabolic plasticity, environmental resilience, and potential for degrading aromatic pollutants

Erythrobacter sp.

Genome Accession Number: *JBKFDH01*

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)

Subsystem Analysis

Table 1: Assembly Details		
Contigs	5	
GC Content	64.81	
Contig L50	1	
Genome length	3,597,984 bp	
Contig N50	3,050,932	

Table 2: Annotated Genome Features		
CDS	3,440	
tRNA	48	
Repeat Regions	10	
rRNA	3	

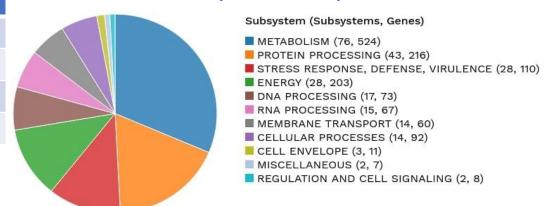


Table 3: Protein Features		
Proteins with functional assignments	2,142	
Proteins with EC number assignments	781	
Proteins with GO assignments	660	
Proteins with Pathway assignments	604	
Proteins with PATRIC genus-specific family (PLfam) assignments	2,470	
Proteins with PATRIC cross-genus family (PGfam) assignments	2,962	

Genome Assembly

