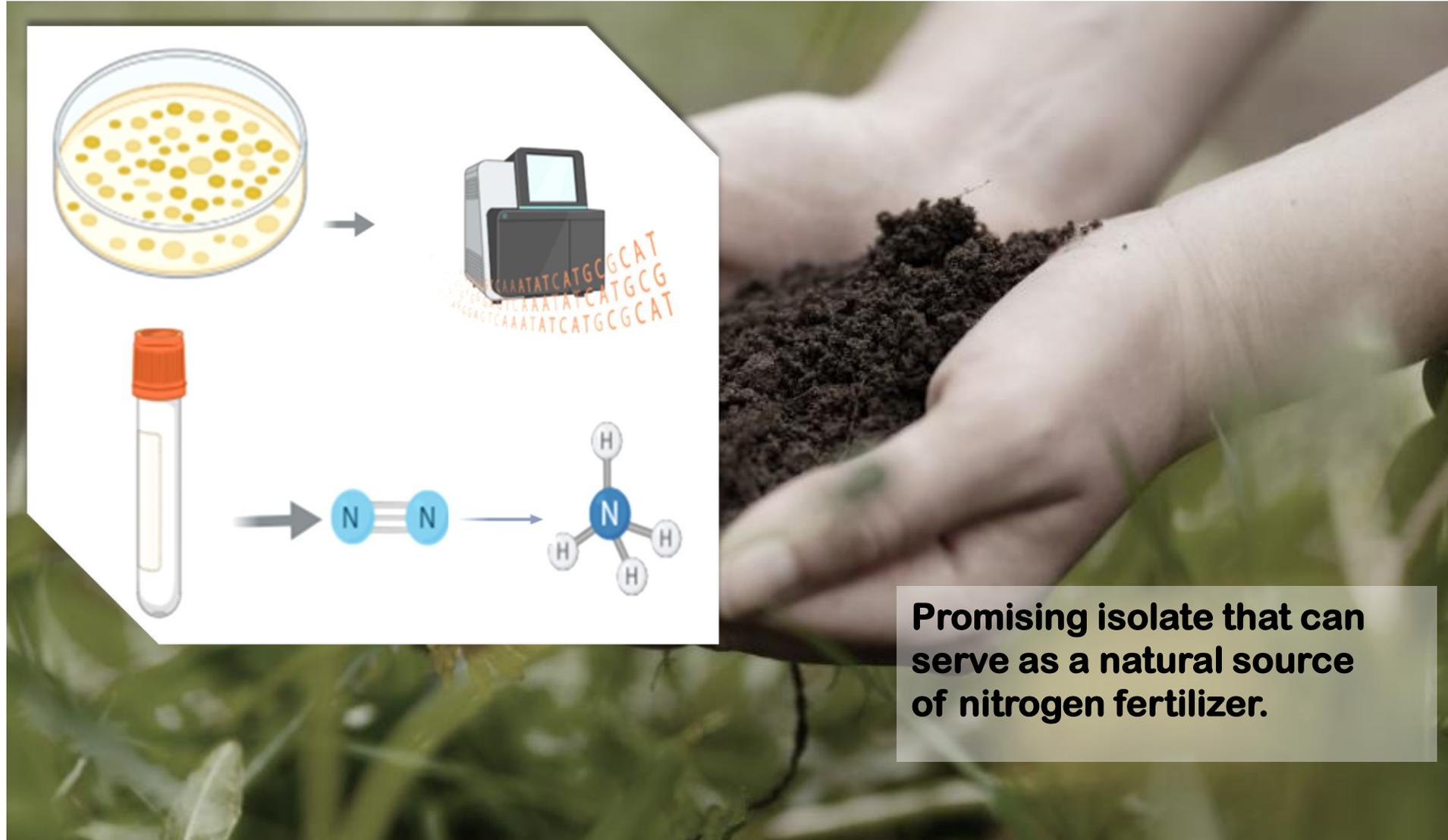




Genome of the Day: *Klebsiella sp.*



Promising isolate that can serve as a natural source of nitrogen fertilizer.

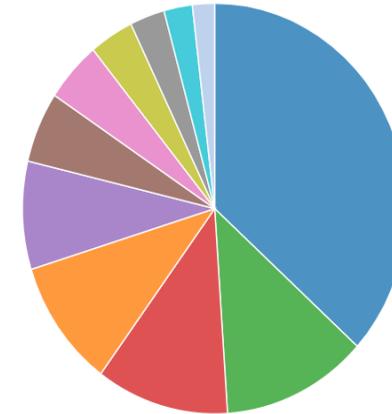
Klebsiella sp.

Genome Accession Number: INRP000250

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



- Subsystem (Subsystems, Genes)
- METABOLISM (138, 1270)
- PROTEIN PROCESSING (46, 267)
- STRESS RESPONSE, DEFENSE, VIRULENCE (42, 212)
- ENERGY (38, 421)
- MEMBRANE TRANSPORT (32, 191)
- DNA PROCESSING (21, 118)
- CELLULAR PROCESSES (18, 94)
- RNA PROCESSING (14, 77)
- CELL ENVELOPE (11, 82)
- MISCELLANEOUS (9, 56)
- REGULATION AND CELL SIGNALING (7, 33)

Table 1: Assembly Details

Table 2: Annotated Genome Features

Table 1: Assembly Details		Table 2: Annotated Genome Features	
Contigs	54	CDS	6.046
GC Content	55.76	tRNA	76
Contig L50	8	rRNA	4
Genome length	6,134,361 bp	Repeat Regions	19
Contig N50	240,986		

Table 3: Antimicrobial Resistance Genes

AMR Mechanism	Genes
Antibiotic activation enzyme	KatG
Antibiotic resistance gene cluster, cassette, or operon	MarA, MarB, MarR
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	BcrC
Efflux pump conferring antibiotic resistance	AcrAB-TolC, AcrAD-TolC, AcrEF-TolC, AcrZ, EmrAB-OMF, EmrAB-TolC, EmrD, MacA, MacB, MdfA/Cmr, MdtABC-TolC, MdtL, MdtM, SugE, TolC/OpmH
Gene conferring resistance via absence	gidB
Protein altering cell wall charge conferring antibiotic resistance	GdpD, PgsA

Genome Assembly

