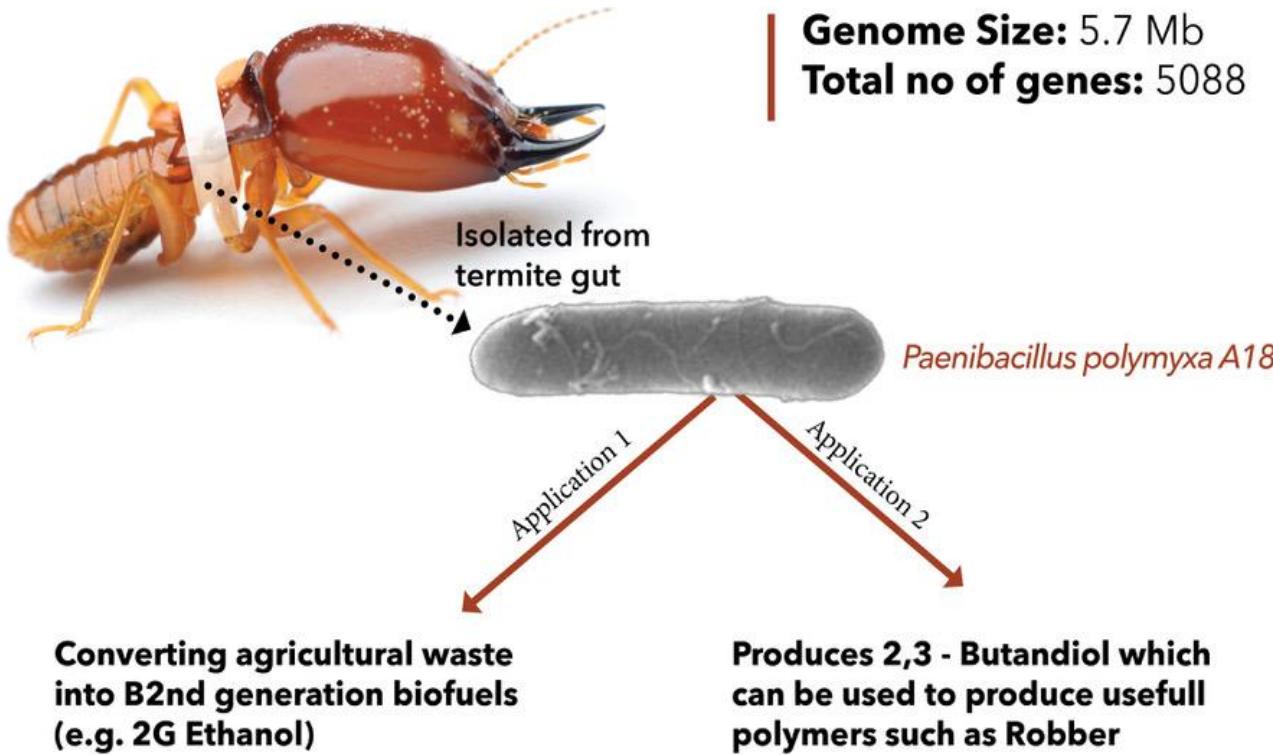




#OneDayOneGenomeInitiative

Today's Genome: *Paenibacillus polymyxa* *A18* bacteria



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

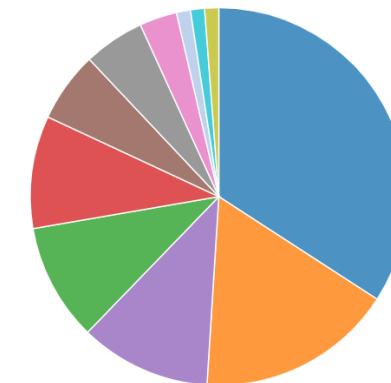
Contigs	28
GC Content	45.31
Contig L50	2
Genome length	5,714,252 bp
Contig N50	1,538,476

Table 2: Annotated Genome Features

CDS	5,273
tRNA	107
Repeat Regions	39
rRNA	2

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
Efflux pump conferring antibiotic resistance	BceA, BceB, YkkCD
Protein altering cell wall charge conferring antibiotic resistance	GdpD, MprF, PgsA
Antibiotic inactivation enzyme	ANT(6)-I, BlaZ family, FosB
Regulator modulating expression of antibiotic resistance genes	BceR, BceS, LiaR, LiaS

Subsystem Analysis

Subsystem (Subsystems, Genes)

- METABOLISM (85, 638)
- PROTEIN PROCESSING (42, 235)
- STRESS RESPONSE, DEFENSE, VIRULENCE (28, 96)
- CELLULAR PROCESSES (25, 171)
- ENERGY (24, 140)
- DNA PROCESSING (15, 68)
- RNA PROCESSING (13, 54)
- MEMBRANE TRANSPORT (8, 55)
- MISCELLANEOUS (3, 7)
- CELL ENVELOPE (3, 9)
- REGULATION AND CELL SIGNALING (3, 11)

Genome Assembly