



Government of India
Ministry of Science & Technology
Department of Biotechnology



BRIC
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राष्ट्रीय जैववैज्ञानिक
जीनोमिक्स संस्थान
National Institute of
Biomedical Genomics

One Day One Genome

Stutzerimonas stutzeri



Competitive interactions with other
microbes to maintain skin homeostasis

Nutrient cycling

Stutzerimonas stutzeri (NIBMG_HMR_CI17)

Genome Accession Number: INS0016145

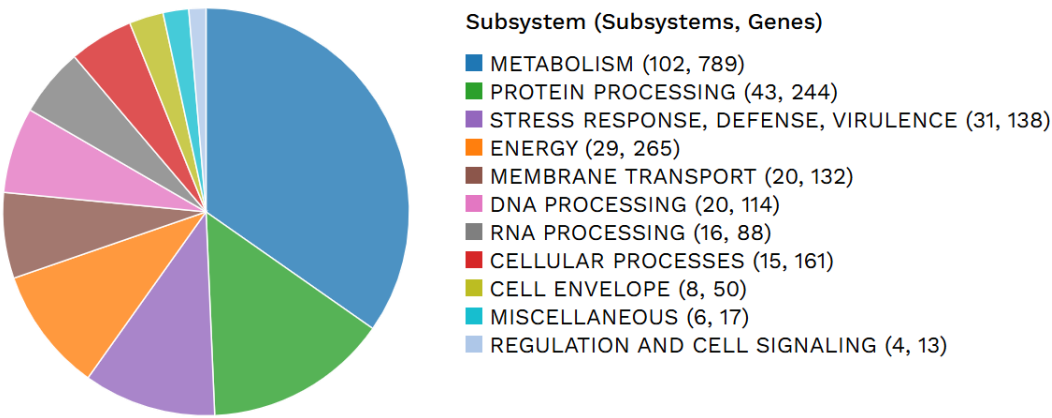
Quality of Genome Assembly and Annotation:

Results from indigenously developed **BHARAT** analysis pipeline: (**B**acterial **H**ybrid genome **A**ssembly and **R**apid **A**nnotation **T**oolset)

Table 1: Assembly Details		Table 2: Annotated Genome Features	
Contigs	51	CDS	4,270
GC Content	64.10 %	tRNA	58
Contig L50	7	Repeat Regions	0
Genome length	4,499,401 bp	rRNA	3
Contig N50	286,117		

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	MdtABC-OMF, MdtABC-TolC, MexAB-OprM, MexEF-OprN, MexEF-OprN system, MexJK-OprM/OpmH, MexVW-OprM, TolC/OpmH
Protein modulating permeability to antibiotic	OccD4/OpdT, OccD6/OprQ, OccK5/OpdH, OprD family, OprF
Regulator modulating expression of antibiotic resistance genes	OxyR
Protein altering cell wall charge conferring antibiotic resistance	GdpD, PgsA
Antibiotic inactivation enzyme	LCR/NPS family

Subsystem Analysis



Genome Assembly

