



ONE DAY ONE GENOME

***Moraxella tetraodonis*
sp. nov**

The novel bacterium provides
new insights into antioxidant
roles and microbial functions in
aquatic environments

Quality of Genome Assembly and Annotation:

Results from indigenously developed **BHARAT** analysis pipeline: (**Bacterial Hybrid genome Assembly and Rapid Annotation Toolset**)

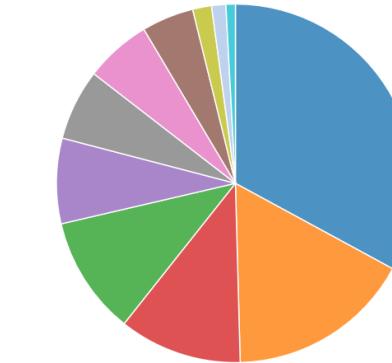
Table 1: Assembly Details

Contigs	7
GC Content	43.38 %
Contig L50	1
Genome length	2,724,669 bp
Contig N50	2,439,788

Table 2: Annotated Genome Features

CDS	2,534
tRNA	48
Repeat Regions	149
rRNA	12

Subsystem Analysis



Subsystem (Subsystems, Genes)
METABOLISM (77, 485)
PROTEIN PROCESSING (39, 216)
STRESS RESPONSE, DEFENSE, VIRULENCE (26, 96)
ENERGY (25, 181)
DNA PROCESSING (18, 77)
MEMBRANE TRANSPORT (15, 65)
RNA PROCESSING (14, 65)
CELLULAR PROCESSES (11, 66)
CELL ENVELOPE (4, 26)
MISCELLANEOUS (3, 8)
REGULATION AND CELL SIGNALING (2, 11)

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	EmrAB-TolC, MacA, MacB, TolC/OpmH
Regulator modulating expression of antibiotic resistance genes	OxyR
Protein altering cell wall charge conferring antibiotic resistance	GdpD, PgsA
Antibiotic target replacement protein	FabG, HtdX

Genome Assembly

