Bacterial diversity of Bhitarkanika mangrove ecosystem, Odisha, India Surajit Das* and Krishna Palit

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Abstract

Bhitarkanika estuary is a highly diverse tropical mangrove ecosystem situated along the river delta of Brahmani and Baitarani before meeting the Bay of Bengal. Large number of heterotrophic bacteria present within the sediment and water are involved in different ecological functions and are capable of interfering in nutrient cycling thereby biological productivity. Heterotrophic bacterial diversity with respect to seasonal variation of several ecological parameters was studied from the five transects namely Dangmal, Ekakula, Gupti, Habalikathi and Kalibhanjadiha, within this ecosystem. During monsoon, highest heterotrophic bacterial population was recorded at Dangmal (39.10±3.60x10³ cfu/ml and 43.93±5.28x10⁵ cfu/g) and lowest total heterotrophic bacterial population was recorded from Ekakula $(7.95\pm2.05 \times 10^3)$ cfu/ml and $6.8\pm2.17 \times 10^5$ cfu/g) from both the water and sediment samples respectively. During summer, highest heterotrophic bacterial population was recorded at Habalikathi sediment sample $(165.58\pm41.76 \times 10^4 \text{ cfu/g})$ and Kalibhanjadiha water sample $(10.287\pm11.82 \times 10^3 \text{ cfu/ml})$ whereas lowest was recorded from Gupti sediment sample (3.835±3.41x10⁴cfu/g) and Ekakula water sample (1.843±1.52x10³ cfu/ml). Physico-chemical parameters such as pH, Organic carbon content, Organic matter content, Conductivity, CEC, Temperature, TDS, Ammonia content, Phosphate content, Nitrate content, Chloride content, Salinity, Dissolved Oxygen were studied to understand the impact of seasonal variation on heterotrophic biodiversity.

Culture independent bacterial diversity from sediment and water samples were studied by performing illumina sequencing of V3-V4 region of 16S rRNA which showed that Dangmal sediment sample contain the highest number of bacterial species (21,207 species) whereas Ekakula sediment sample contain the lowest (5,077 species). In case of water sample, highest no of bacterial species was recorded in Habalikhati (34,601 species) and lowest was recorded in Gupti (24,846 species). Species richness and chao 1 index was also carried out using alpha-diversity estimation followed by refraction curve and rank abundance plot for the sediment and

water samples. Comparison between microbial communities were done using Jaccard and Bray-Curtis index. This study revealed that unculturable fraction of bacteria varies greatly with respect to the culturable method. Predominant genera are *Acidovorax, Acinetobacter, Bacillus, Bergeriella, Burkholderia, Clostridium, Corynebacterium, Enterobacter, Proteobacteria, Escherichia, Moraxella, Neisseria, Pseudomonas, Serratia, Staphylococcus* and *Yersinia* from monsoon samples and *Clostridium, Acidovorax, Burkholderia, Pseudomonas, Streptococcus, Neisseria* and *Bacillus* from summer samples. The distribution and diversity of the heterotrophic bacterial population has not been well documented for Bhitarkanika mangrove ecosystem. Therefore, this study will form the database for bacterial genera available in Bhitarkanika Mangrove ecosystem.

Keywords: heterotrophic bacteria, Bhitarkanika mangrove, physico-chemical parameters, bacterial diversity.

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Background and Objectives

Background

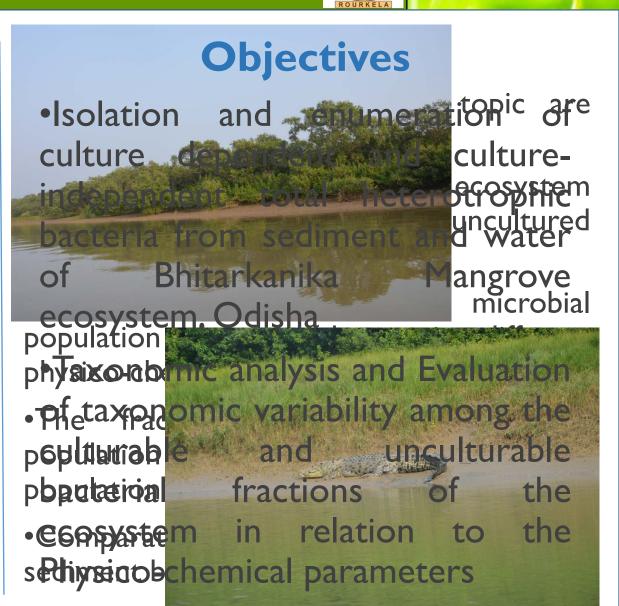
• Mangrove ecosystem of Bhitarkanika is highly rich in biological diversity as well as significant habitat for wildlife.

• Its brackish water environment is highly rich in organic matter due to **microbial enzymatic and metabolic activities**.

• It has been documented that microbial population in Bhitarkanika varies from 10⁵ to 10⁷ CFU/g of soil whereas 10⁴ to 10⁷ CFU/ml incase of water sample and **Gram negative bacteria** are more in comparison to Gram positive bacteria (Dash and Das., 2014; Thatoi et al., 2012)

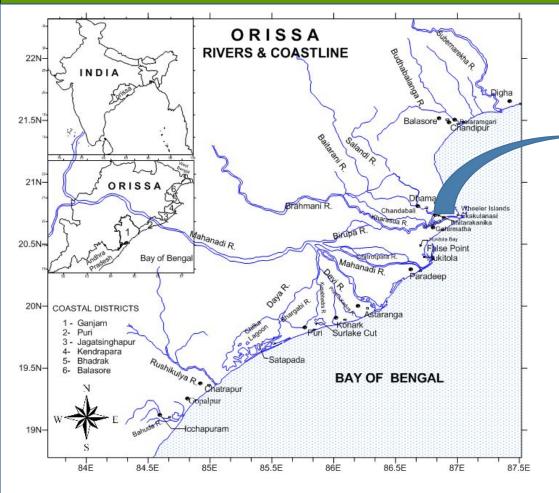
• Population diversity have been found to shift seasonally, however role and function of these microbes in the ecosystem and effect of seasonal variation is not known.

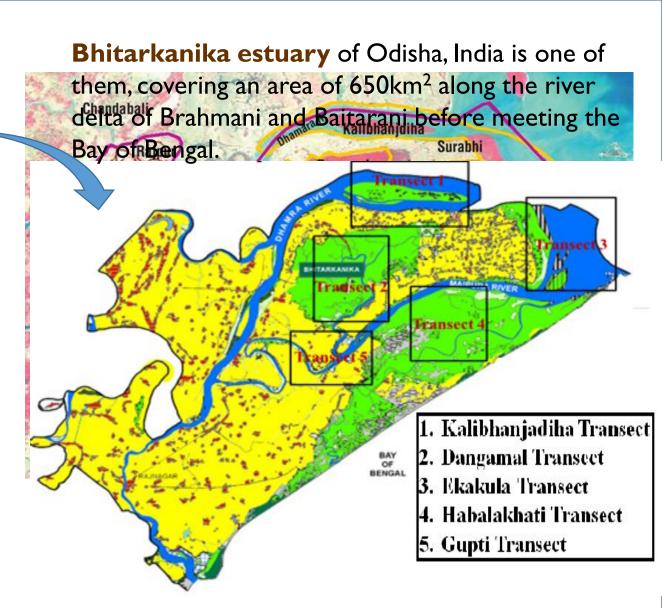
• Vast microbial flora of this ecosystem is still untapped because of un-culturable nature of the microbes.





Study area







Methodology



Study site and collection of sample

• Five transects include Kalibhanjadian, Dangmal, Ekakula, Habalikhati and Gupti. Water (100 ml) and sediment samples (100 g) were collected in duplicates twice a year i.e. Monsoon (August 2016) and Summer (Post-monsoon) (March 2017) and on site analysis of physico-chemical parameters were conducted.

Analysis of physicochemical parameters

- pH, temperature and salinity (On site)
- Dissolved oxygen, Total Dissolved Solid, conductivity, chloride content, nitrate, phosphate, magnesium and calcium content and its total hardness will be tested in the laboratory (APHA, 1992).

Culture-dependent heterotrophic bacterial count

- Sediment and water samples were subjected to serial dilution followed by spread plating on Sea Water Nutrient agar (SWNA) and Zobell's Marine agar (ZMA) plates.
- Incubation at temperature as per the temperature of the samples for 24-48h.
- The total heterotrophic bacterial load was estimated for the respective samples using the standard formula: CFU/ml or g = No. of colonies x inverse of dilution factor/ volume taken.

Culture-independent bacterial population

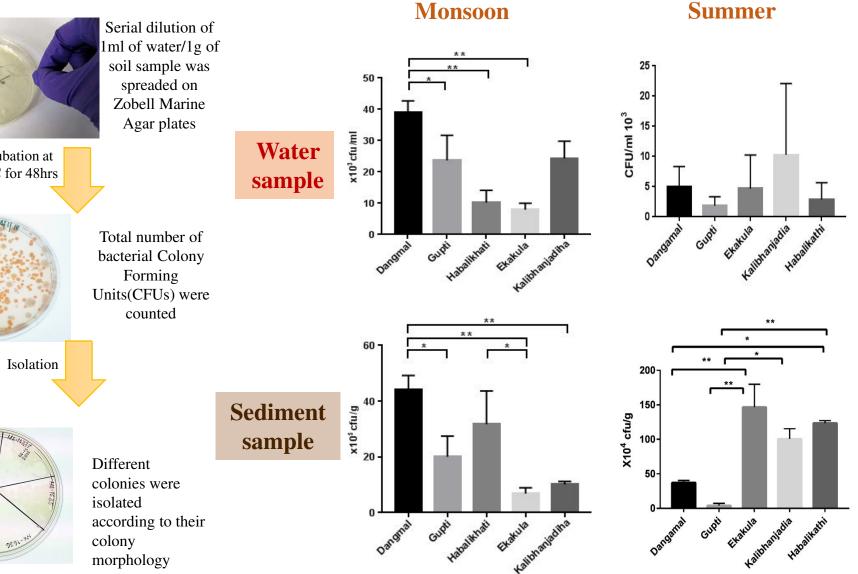
 Culture-independent bacterial population of water and sediment samples were estimated by metagenomic analysis of the V3-V4 (Product size ~459bp) region of 16S rRNA at the Illumina platform.



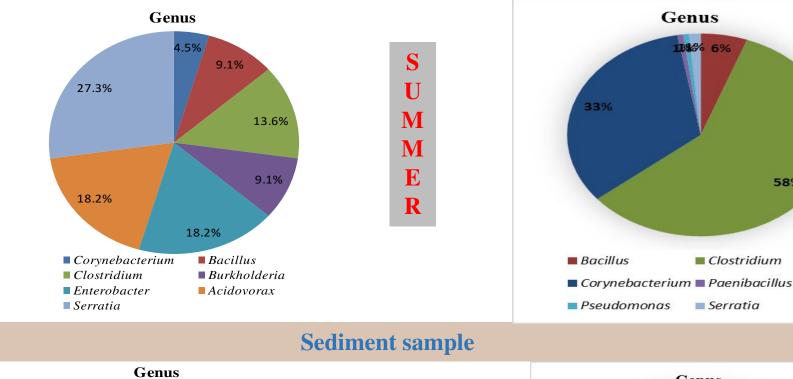
Results



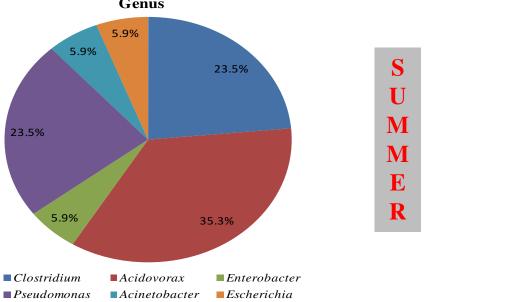


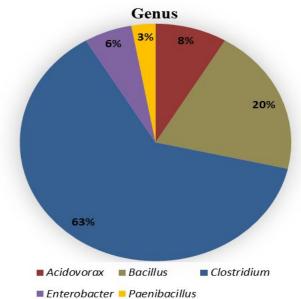


Water sample









58%

Clostridium

🔳 Serratia

(Dangmal

Μ

0

N

S

0

0

N

Μ

0

N

S

0

0

N

23.5%

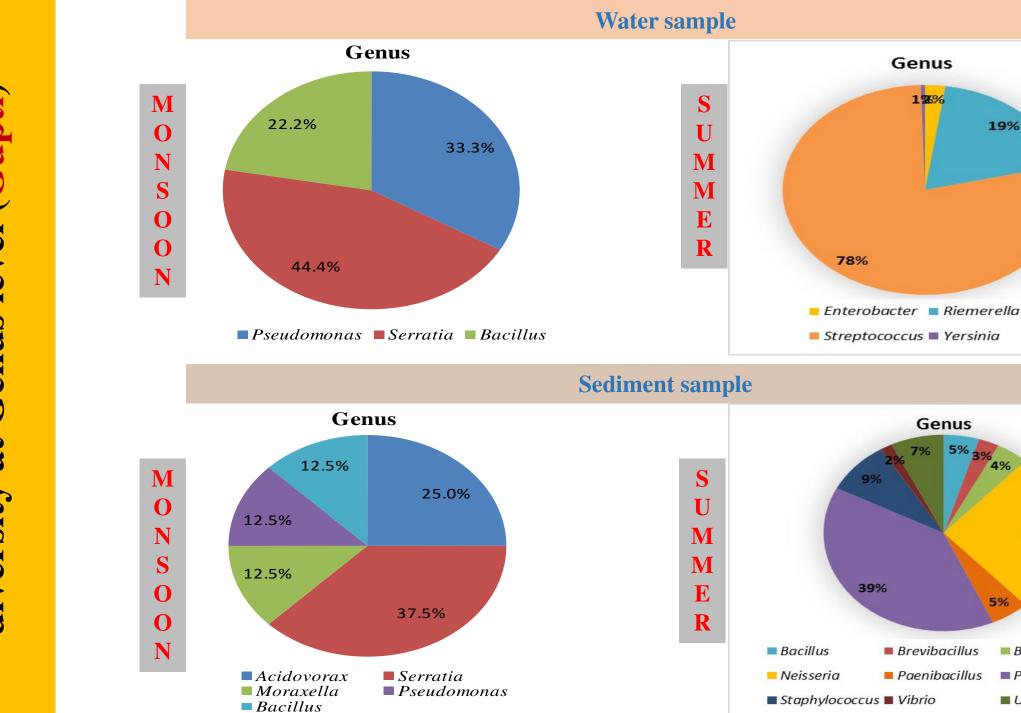
5.9%

Clostridium

bacterial diversit

0f

Relative abundance



19%

26%

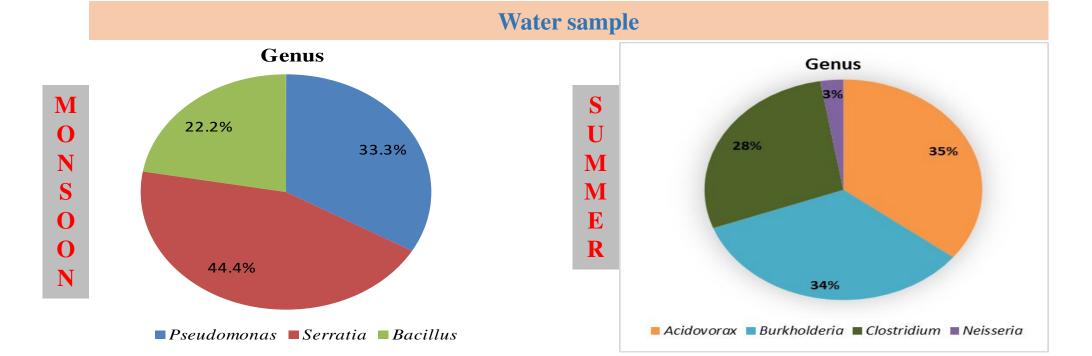
Burkholderia

Pseudomonas

Unidentified

5%

Relative abundance of bacteria Gub level enus **U** at diversity



S

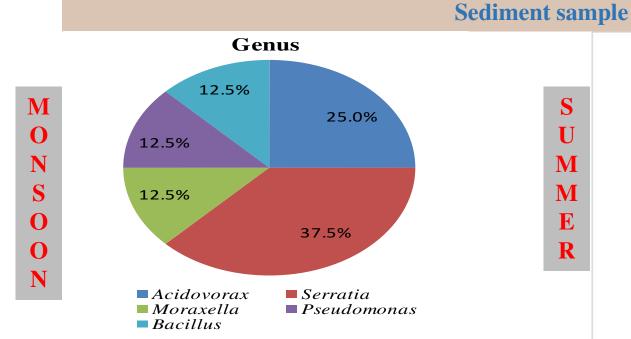
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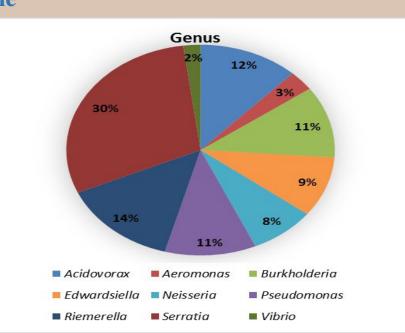
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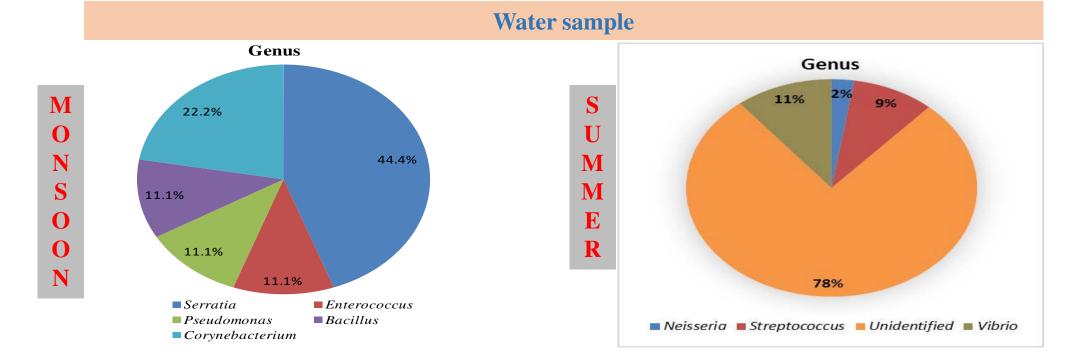
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Sediment sample

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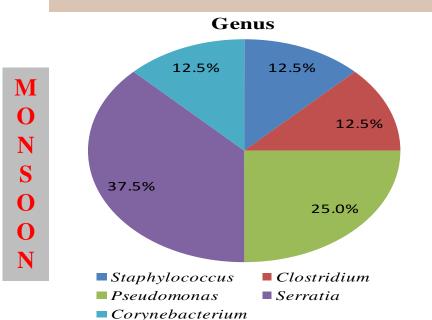
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bacteria

Relative abundance of

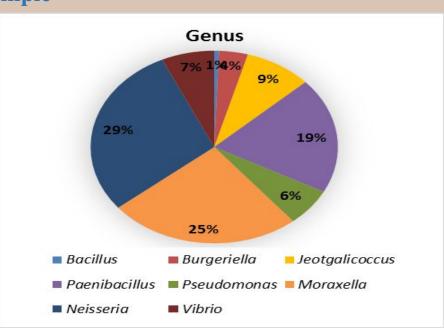
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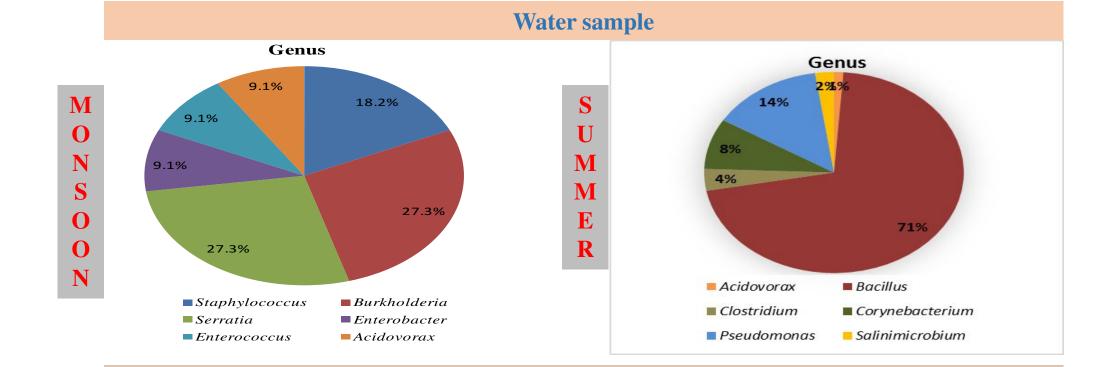
level

Genus

at

diversity

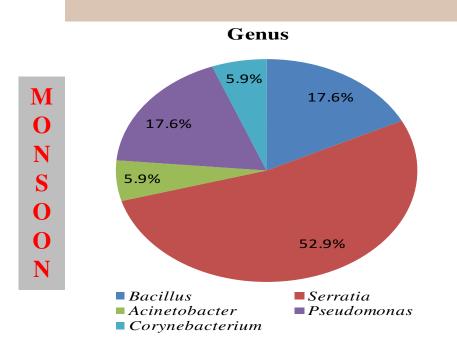


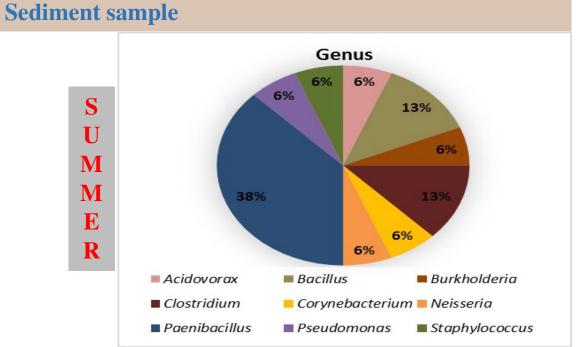


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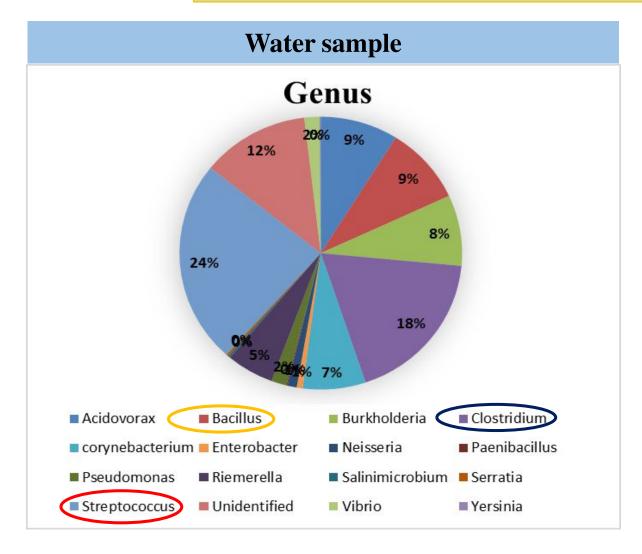
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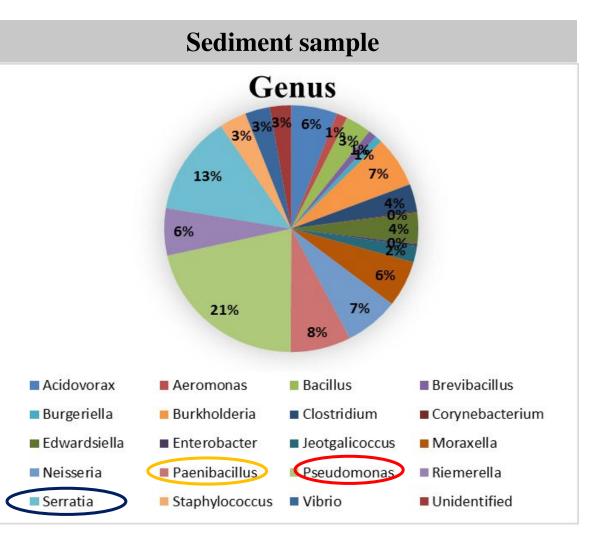
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Overall GENRIC distribution of bacteria in Bhitarkanika Mangrove ecosystem







Physico-chemical parameters



Table I. Correlation matrix for the physico-chemical parameters of water samples

	-	_						_			_			
Paramete	ТНВ	pН	Temp	Ammonia	Phosphate	Nitrate	Ca++	Mg++	Hardness	Chloride	Salinity	Conductivity	TDS	DO
rs	/ \													
ТНВ	1.00													
рН	0.15	1.00												
Temp	0.04	0.87	1.00											
Ammonia	0.06	0.05	0.16	1.00										
Phosphate	0.05	0.08	0.24	0.99***	1.00									
Nitrate	0.62	-0.20	-0.46	0.44	0.36	1.00								
Ca++	-	-0.13	0.14	-0.29	-0.23	-0.90*	1.00							
	0.87*													
Mg++	-0.22	0.41	0.59	-0.61	-0.52	-0.88*	0.60	1.00						
Hardness	-0.54	0.21	0.45	-0.54	-0.45	-0.99***	0.85	0.93*	1.00					
Chloride	-0.77	-0.13	0.11	-0.49	-0.42	-0.93*	0.98**	0.71	0.91*	1.00				
Salinity	-0.58	0.11	0.44	-0.35	-0.25	-0.98**	0.89*	0.85	0.97**	0.92*	1.00			
Conductivi	-0.82	-0.05	0.17	-0.45	-0.39	-0.94*	0.98**	0.70	0.91*	0.99***	0.91*	1.00		
ty														
TDS	-0.83	-0.06	0.14	-0.46	-0.40	-0.93*	0.98**	0.68	0.90*	0.99***	0.89*	1.00	1.00	
DO	-0.08	0.32	-0.15	-0.19	-0.28	0.31	-0.27	-0.27	-0.30	-0.27	-0.49	-0.20	-0.17	1.00

Values in the table indicate Pearson's r value. Level of significance ***P < 0.001; **P < 0.01: *P < 0.05 at n = 3

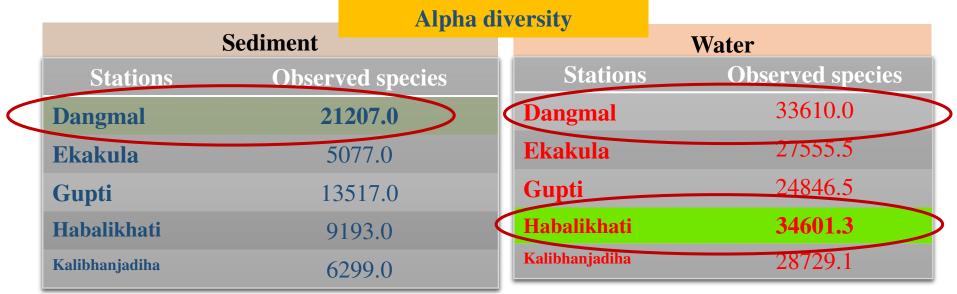
Table 2. Physico-chemical propertiesand the total heterotrophic bacterialpopulation of sediment samples ofBhitarkanika mangrove ecosystem.

Parameters	Dangmal	Gupti	Habalikhati	Ekakula	Kalibhanjadiha
THB (X10 ⁵ cfu/g)	43.96±5.28	20.01±7.45	31.76±11.93	6.8±2.17	7.98±3.56
рН	7.34±0.19	7.48±0.42	7.73±0.41	7.27±0.04	7.92±0.26
Organic Carbon (%	1.06±0.07	0.38±0.09	0.89±0.02	0.17±0.02	0.85±0.08
Organic Matter (%)	1.82±0.12	0.66±0.16	1.53±0.04	0.30±0.04	1.47±0.14
Sediment texture	Clay	Clay	Clay	Sandy Loam	Clay
Conductivity (mS/cm)	0.32±0.08	1.09±0.07	0.98±0.07	0.51±0.05	1.31±0.08

Culture-independent studies

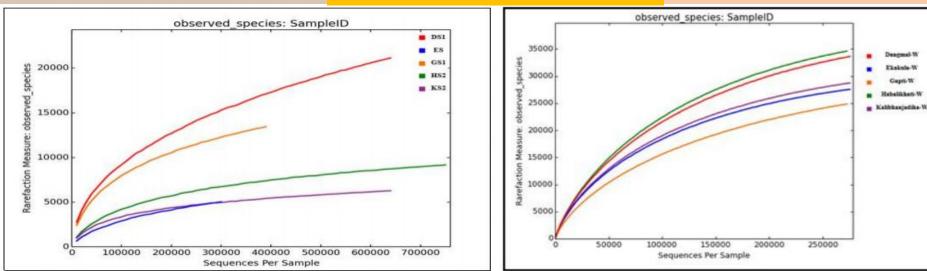


Water





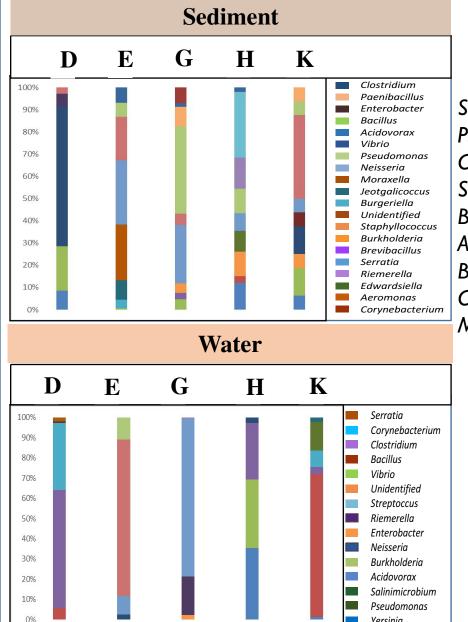






CULTURABLE vs UNCULTURABLE FRACTION





Yersinia

Genera (Cuturedependent) Streptococcus Pseudomonas Clostridium Serratia **Bacillus** Acidovorax Burkholderia Corynebacterium Moraxella

Genera (Culture independent) Marinobacter Alcanivorax Kordiimonas Pseudospirillum Sedimentibacter Limnobacter Streptomyces

0.921% (~1%) of total

microbial population of

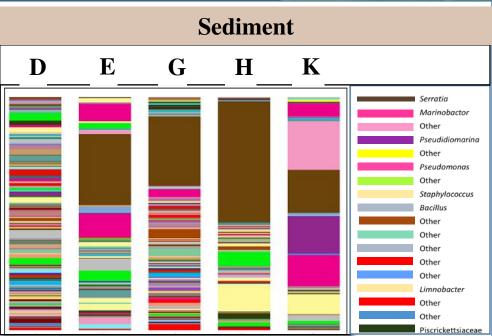
Bhitarkanika Mangrove is

culturable and remaining

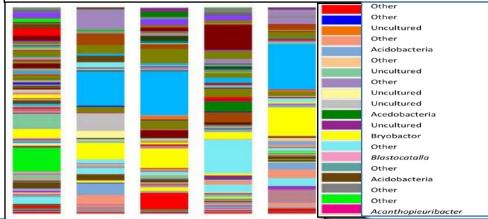
~99% is uncultured.

D

E



Water G Η K





Highlights

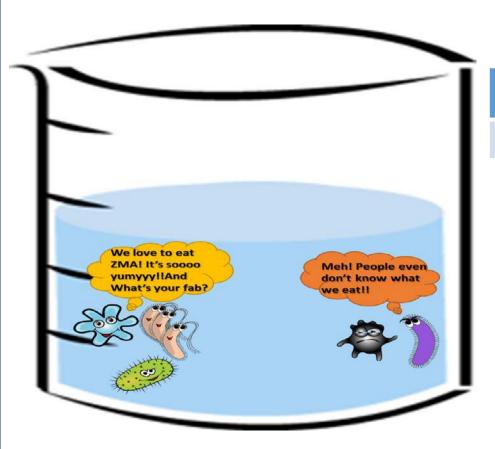


- Occurrence of pathogenic bacterial genera such as **Streptococcus, Serratia, Staphylococcus, Neisseria, Clostridium** showed that the sites such as Dangmal, Gupti and Kalibhanjadiha water and sediment are influenced by the anthropogenic activities.
- Mangrove ecosystem accounts for 25% of the earth's coastline and 75% of the tropical coastline. Bacteria, having role in mangroves is vital for biogeochemical cycles and transformations of most nutrients.
- However, the question still exists: what are all they doing?
- Bhitarkanika mangrove ecosystem is lagging behind to elucidate the question. A complete document on the microbial diversity is way far for this ecosystem.
- So a detailed analysis and further study is ongoing to elucidate the role of the microbial community in nutrient cycling and productivity in the mangrove ecosystem.



Conclusions





• As many as 21000 species from sediments and 30000 species from water have been recorded.

	Total no of cultured bacteria (Sediment)		Total no of cultured bacteria (Water)
40249	663	113269	751

• Cultivable fraction is 1.64% in sediment sample and 0.663% in water sample (of total microbial population).

• Overall, 0.921% of total microbial population of Bhitarkanika Mangrove is culturable and remaining ~99% is uncultured.

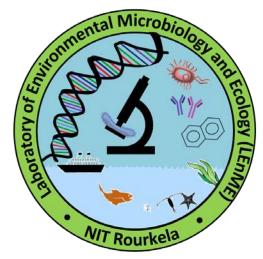
• Seasonal variation has been studied and found that during monsoon bacterial population and species diversity is more.

• Major physico-chemical factors regulating the bacterial population are nitrate and calcium in water and organic carbon in sediment.









Acknowledgement

MoEFCC, Gol through AICOPTAX project for financial support THANK YOU