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## MANGROVE MICROBIOME AND ECOSYSTEM HEALTH

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MANGROVE MICROBIOME

Fig 4.(A) Non-metric multidimensional scaling (NMDS) of microbial community structure (Bray-Curtis). (B) Alpha diversity: inverse Simpson index. (C)Relative abundance of top 20 taxa

Q1.) What's the mangrove microbiome **community structure** and diversity?

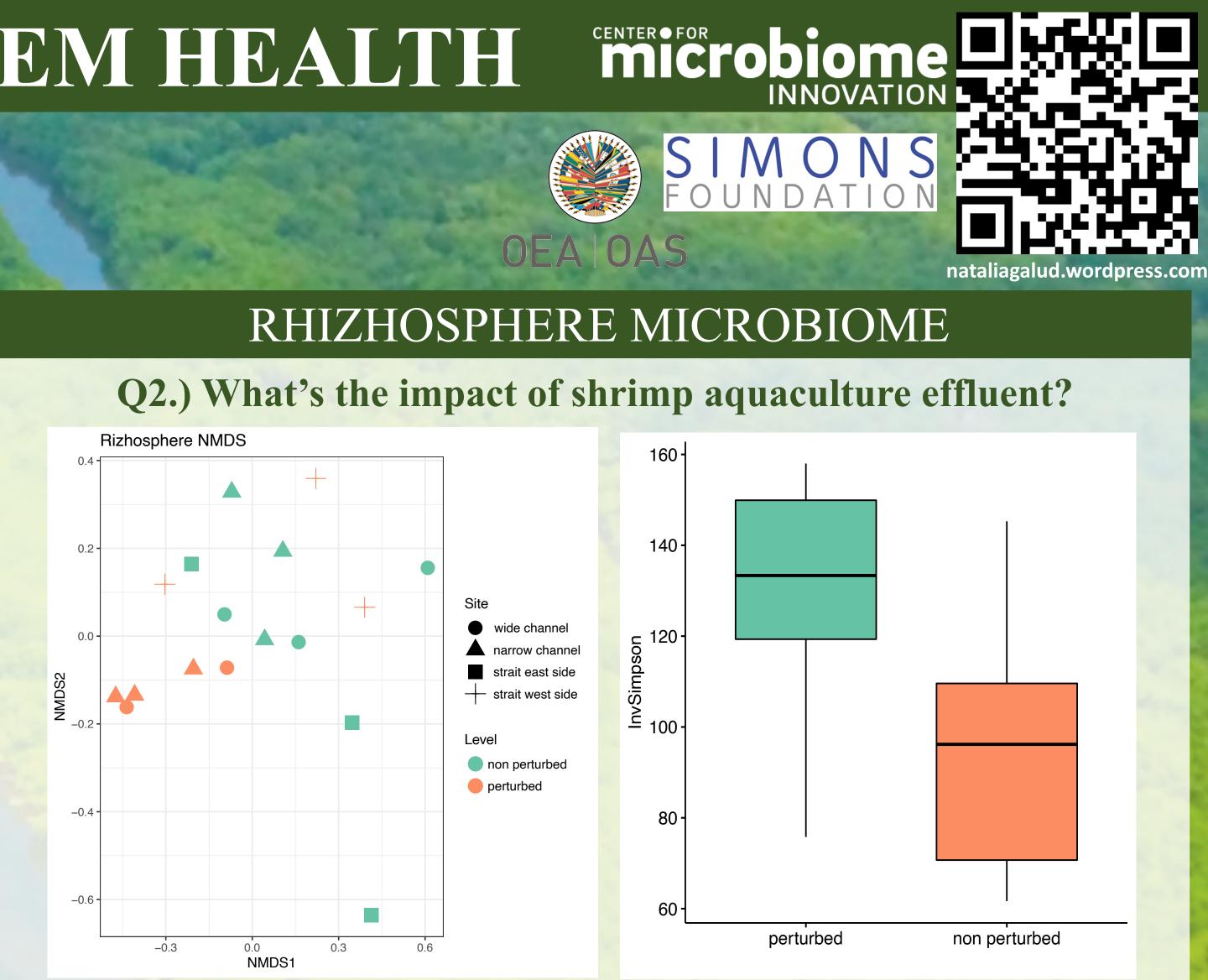
**Distinctive microbial community for** different environments

**High diversity in the** soil: it has more ecological niches than the water

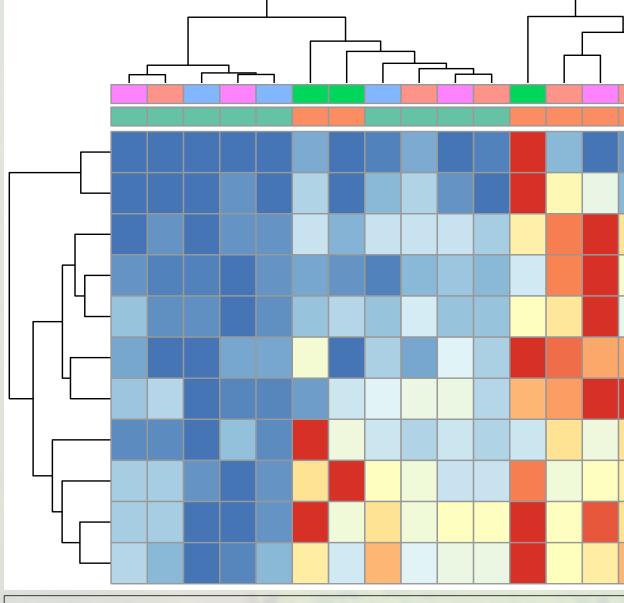
ower diversity for he rhizosphere: pecialized community

Candidatus Dependentiae Candidatus Gracilibacteria bacterium HOT-871 Haliscomenobacter hydrossis DSM 1100 Candidatus Carsonella ruddi. Calothrix sp. NIES-4071 Oscillatoria acuminata PCC 6304 Planctomycetes <phylum> Cystobacterineae Brevefilum fermentans Sulfuriflexus mobilis Thiohalobacter thiocyanaticus Dehalococcoidia Anaerolineaceae Desulfatibacillum aliphaticivorans Vampirococcus sp. LiM Luteitalea pratensis Desulfobulbus oralis Pelolinea submarina Anaerolineaceae Caldithrix abyssi DSM 13497

0.8 mudflat aerial root 0.6 riznosphere sediment 0.4 water 0.2







## **Perturbed:**

**Candidatus Gracilibacteria:** link to necrotic conditions of larvae in bivalves<sup>3</sup> Candidatus Carosnella rudii: symbiont in psyllid plant-pathogen pest

Fig 6. Image of *Rhizhophora mangle* and fisherman collecting bivalves. Differential abundance of bacteria between perturbed vs non perturbed trees. Identified 11 significant taxa p-value <0.05 and potential pathogens.

- to monitor changes in the ecosystem health.
- biogeochemical cycles with severe implications: eutrophication, hypoxia, and reduction of ecosystem services & health.

## Fig 5. NMDS of rhizosphere microbial community and alpha diversity

	L			
		Site	1	Site
		Level		wide channel
		Calothrix sp. NIES–4071	0.8	
		Candidatus Carsonella ruddii	0.6	strait east side strait west side
		Chloracidobacterium thermophilum B	0.4	Level
		Candidatus Xiphinematobacter	0.4	non perturbed perturbed
		Haliangium ochraceum DSM 14365	0.2	perturbed
		Tistrella mobilis	0	
		Cystobacterineae		
		Ignavibacterium album JCM 16511		
		Candidatus Gracilibacteria bacterium HOT-871		
		Brevefilum fermentans		
		Candidatus Dependentiae		

CONCLUSION

Distinctive microbiomes in the water, sediment, and root samples. Presence of pathogenic microbes in perturbed rhizosphere could be used Changes in microbial community could lead to decoupling of