

SCIENCE UPDATE – RESEARCH RELEVANT TO BIOSECURITY IN THE TOP OF THE SOUTH REGION

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SABELLA SPALLANZANII

IMPACTS, BIOLOGY AND NOVEL DETECTION TOOLS

1. Experiments to determine ecological impact of fanworms in soft sediments
2. Reproductive biology of *Sabella*
3. Optimisation of eDNA based surveillance tools



1. How does *Sabella spallanzanii* modify the functioning of soft sediment communities?

- 2016 – 18, Rangitoto Channel
- Transplants – worms and mimics
- Density gradients (0-50 per m²)
- 6 months

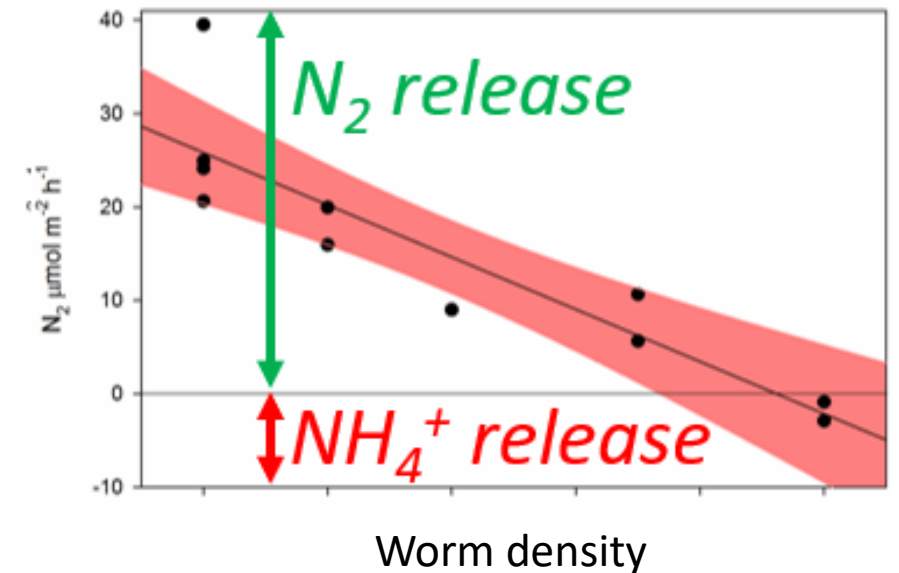
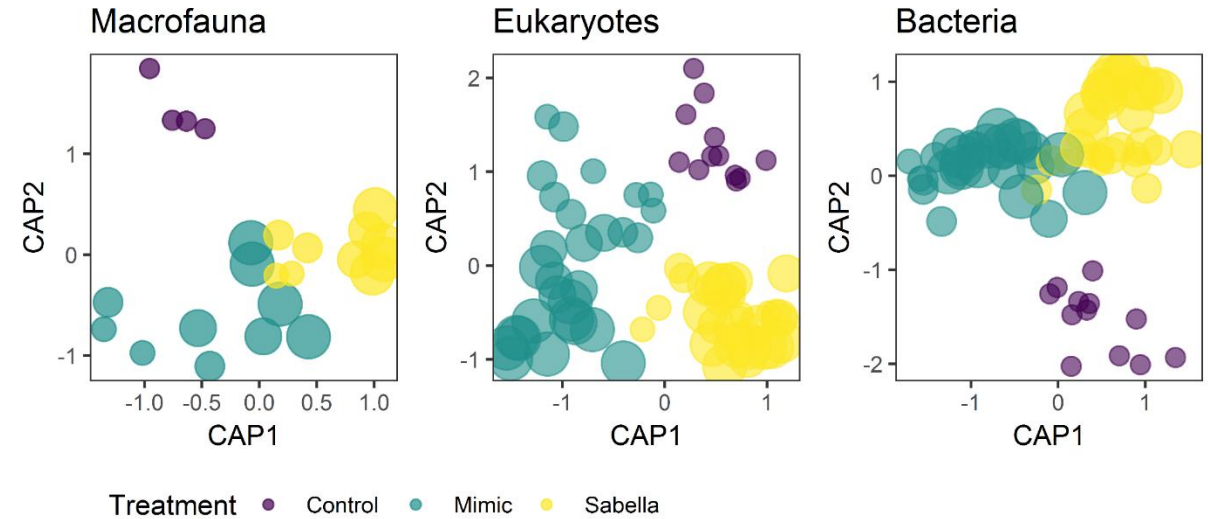
Examined impacts on:

- Soft-sediment fauna (macro/micro)
- Community respiration
- Denitrification
- Primary productivity

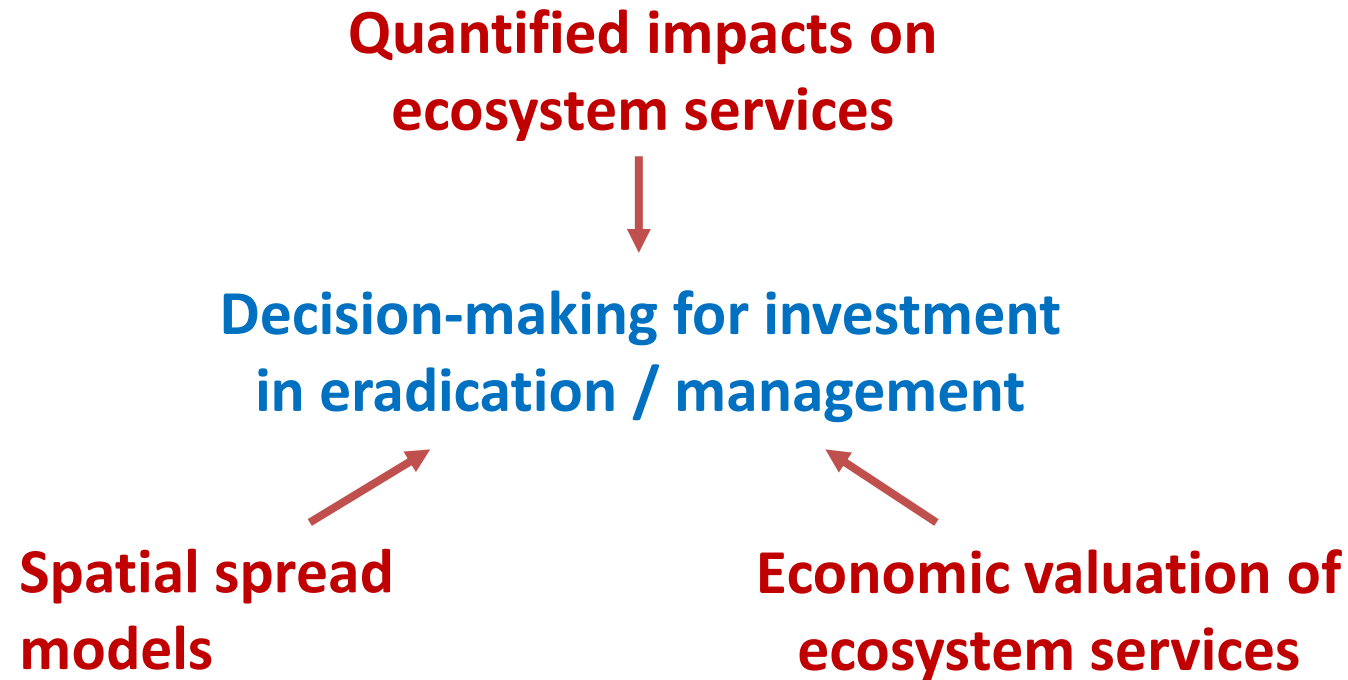


1. How does *Sabella spallanzanii* modify the functioning of soft sediment communities?

- No impacts on sediment biodiversity
- But shifts in the composition of sediment communities
- Increased community metabolism and reduced denitrification
- Worms change structure of above-sediment physical habitat



1. How does *Sabella spallanzanii* modify the functioning of soft sediment communities?



Impact research leads
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2. PhD project: *Sabella spallanzanii* in the context of mussel farms in the Coromandel

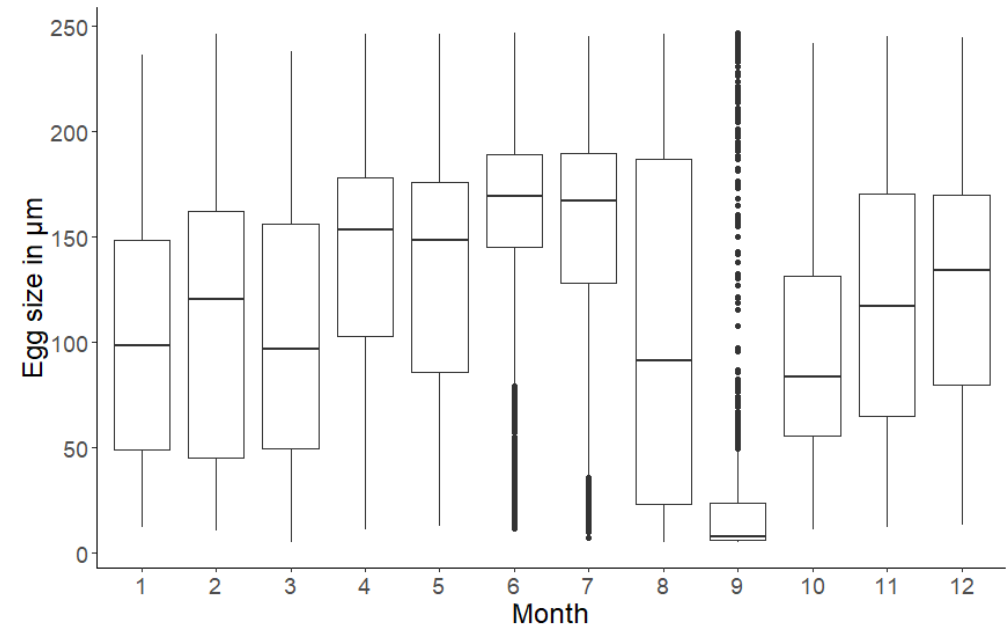
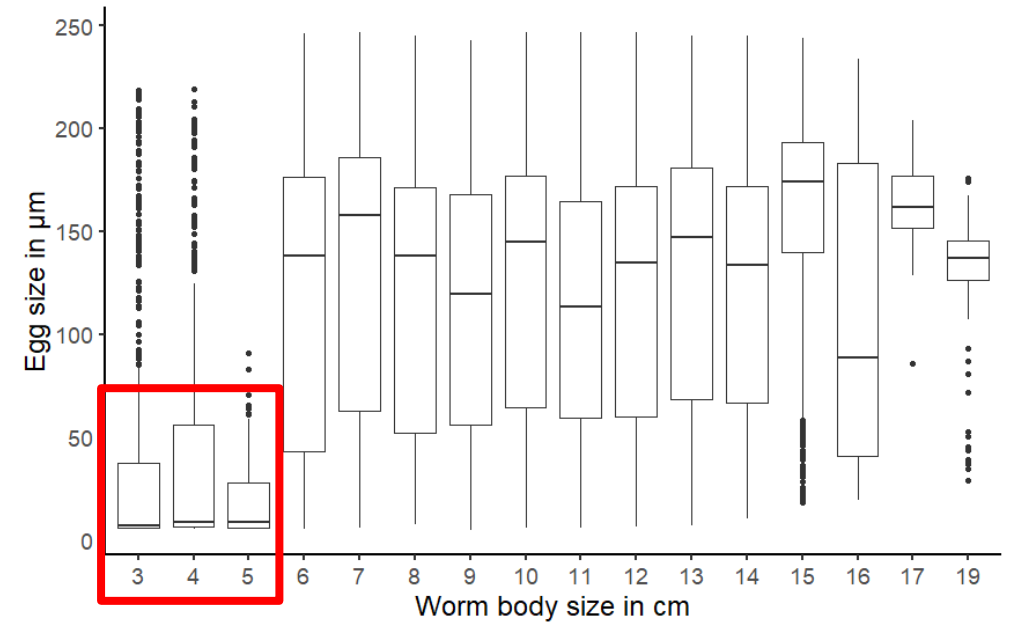
Project aims

- Reproductive cycle and output
- Seasonal presence of larvae around mussel farms
- Larval development and settlement preferences
- Potential for regeneration after fragmentation

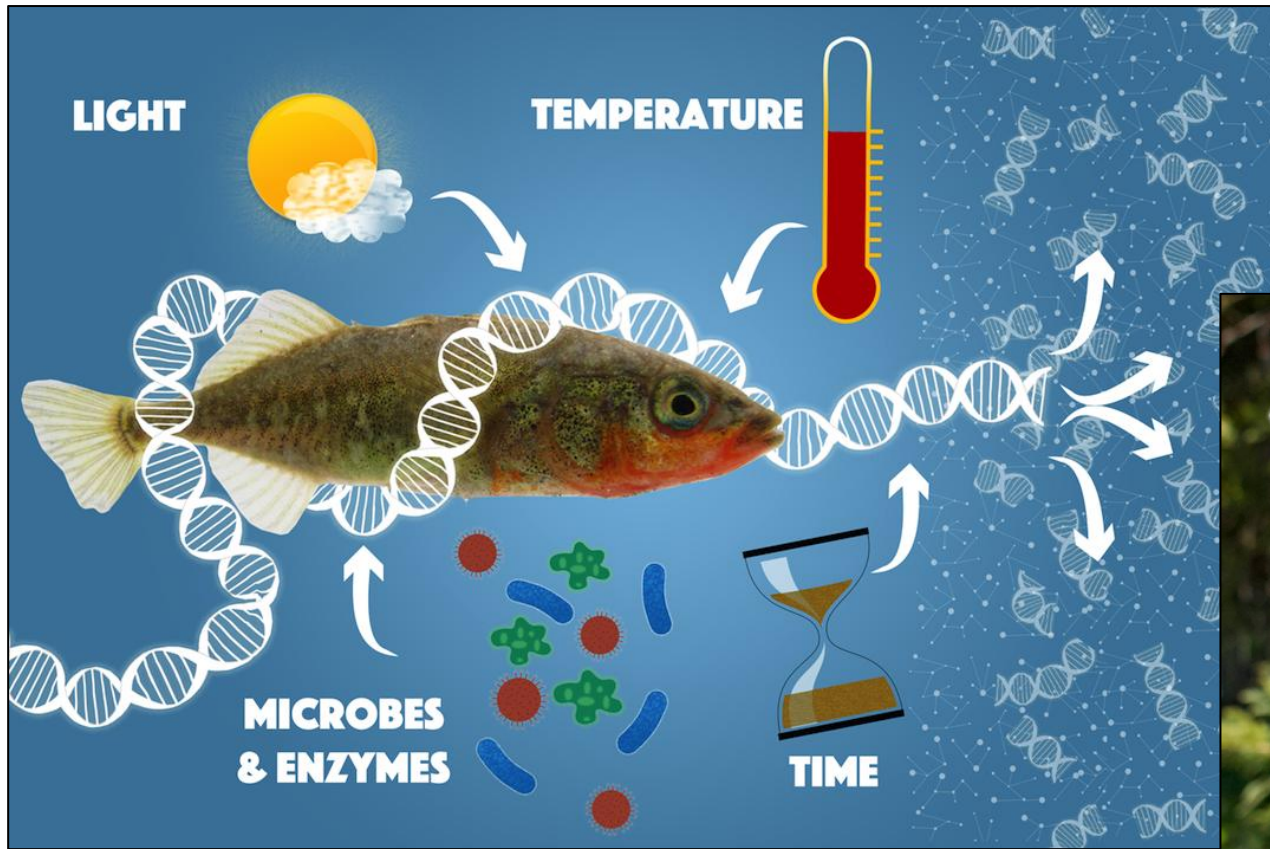


Reproductive cycle and output

- 1,200 worms sampled over 1y
- Worms <5cm (<10cm tube length) generally not reproductively mature
- Sex ratio of 1:1 not confirmed
- Wide range of egg sizes produced each month
- Continuous egg production and gametogenesis, *potentially* enabling spawning events year-round
- Evidence for a spawning event in Aug/Sept



3. Environmental DNA approaches to finding Sabella (and others)



eDNA sample

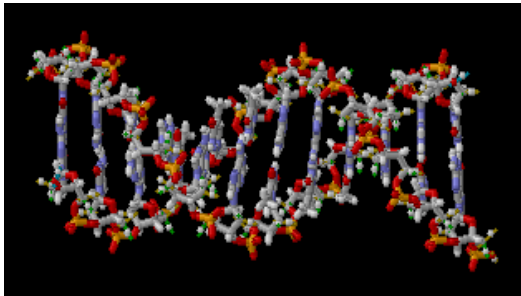


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Marine biosecurity applications

Target species detection

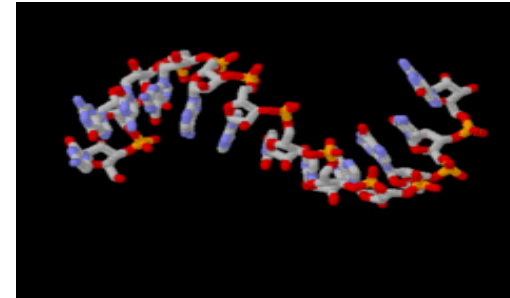
- 🧬 Species-specific assays
- 🧬 qPCR, ddPCR
- 🧬 One-to-few known target taxa
- 🧬 High specificity



DNA (Deoxyribonucleic acid)

Biodiversity screening

- 🧬 Community-wide inventories
- 🧬 HTS metabarcoding
- 🧬 Multiple taxa, incidental detections
- 🧬 Validation of positive detections is recommended



RNA (Ribonucleic acid)

Targeted detection surveys to:

- 🧬 Verify the results of traditional surveillance over various spatial scales
- 🧬 Assess large-scale pest occupancy (distribution)
- 🧬 Optimise surveillance designs and management programmes
- 🧬 Verify eradication success over a constrained area (in prog.)



Image: J. Atalah

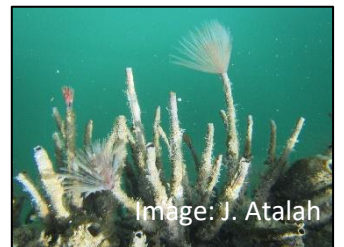
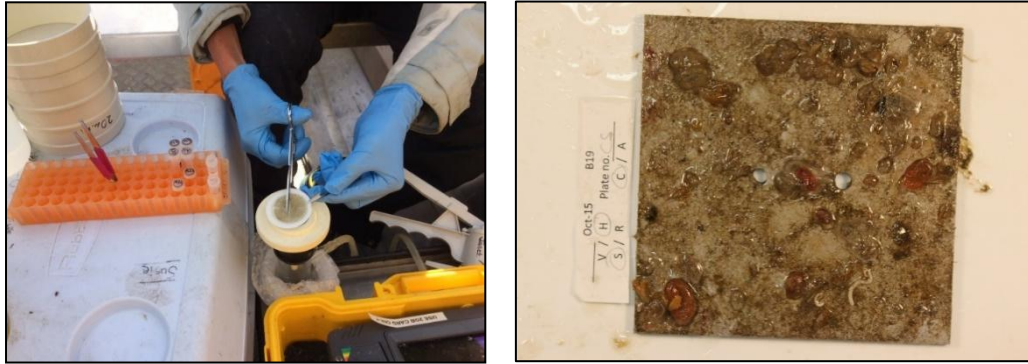


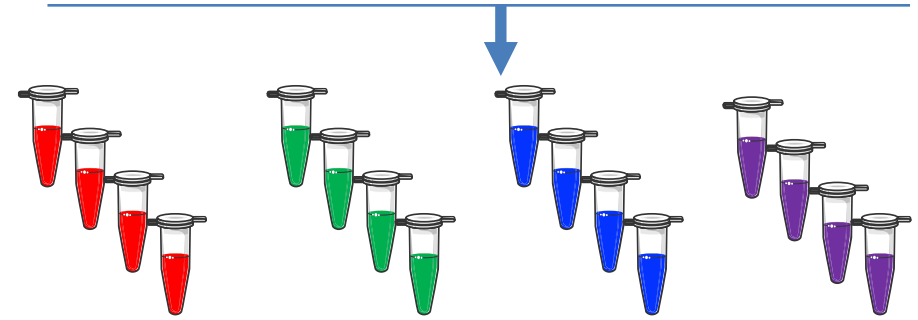
Image: J. Atalah

Method development/optimisation

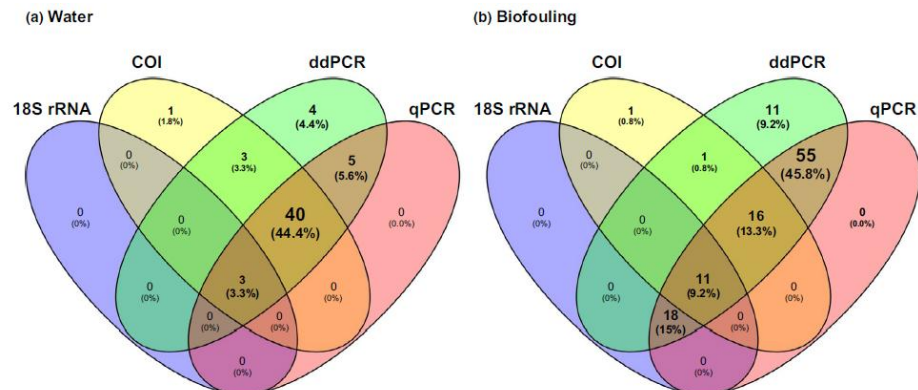
Sampling methods/matrices: filtered water samples vs settlement plates vs sediment



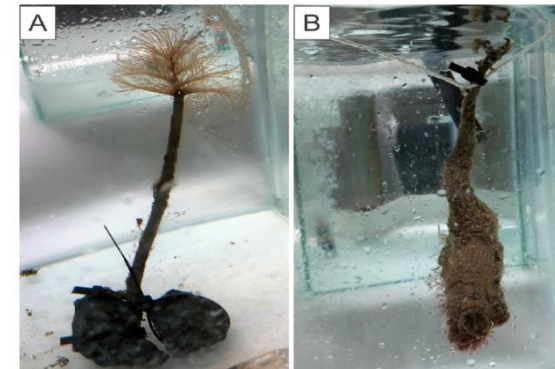
International validation and standardization of pipelines



Analytical platforms: qPCR vs ddPCR vs metabarcoding



eDNA and eRNA fate in water: Experimental study of shedding and decay rates

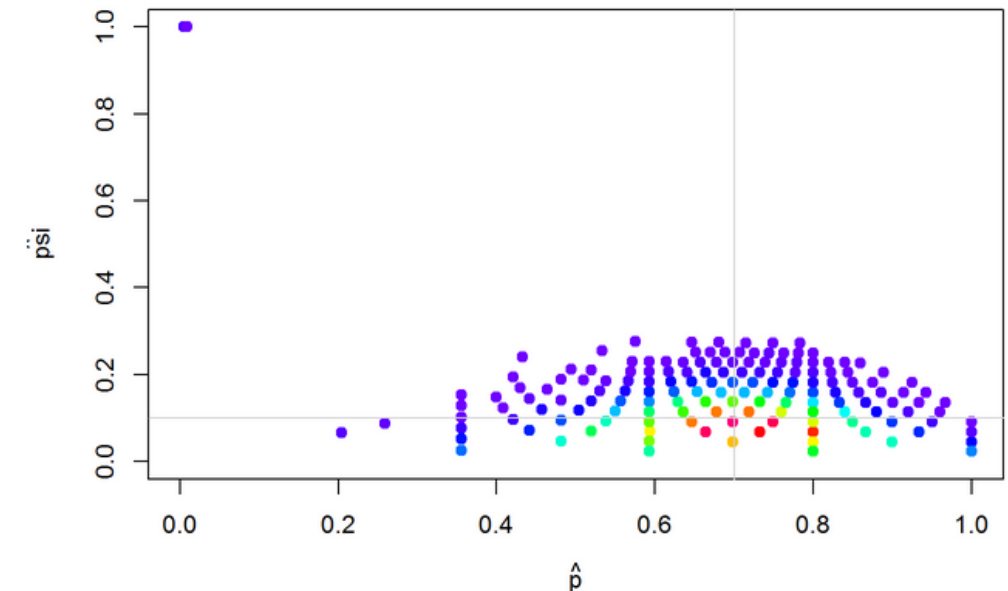


Coupling eDNA signals with probabilistic models to:

- 🧬 Estimate occupancy of the species (accounting for potentially imperfect detection)
- 🧬 Estimate probability of detection for particular method/ sampling approach
- 🧬 Design fit-for-purpose surveys - optimised detection probabilities for a given sampling effort
- 🧬 Informed interpretation of non-detections

In development: coupling eDNA with hydrodynamic models and novel sampling methods

	qPCR	ddPCR	COI	18S rRNA
Water				
ψ (SE)			0.74 (0.1)	
θ (SE)			0.91 (0.02)	
p (SE)	0.93 (0.02)	1.0	0.57 (0.05)	0.27 (0.05)
Biofouling				
ψ (SE)			1.0	
θ (SE)			0.91 (0.02)	
p (SE)	0.87 (0.03)	1.0	0.40 (0.04)	0.16 (0.03)



FOR MORE INFORMATION....

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