

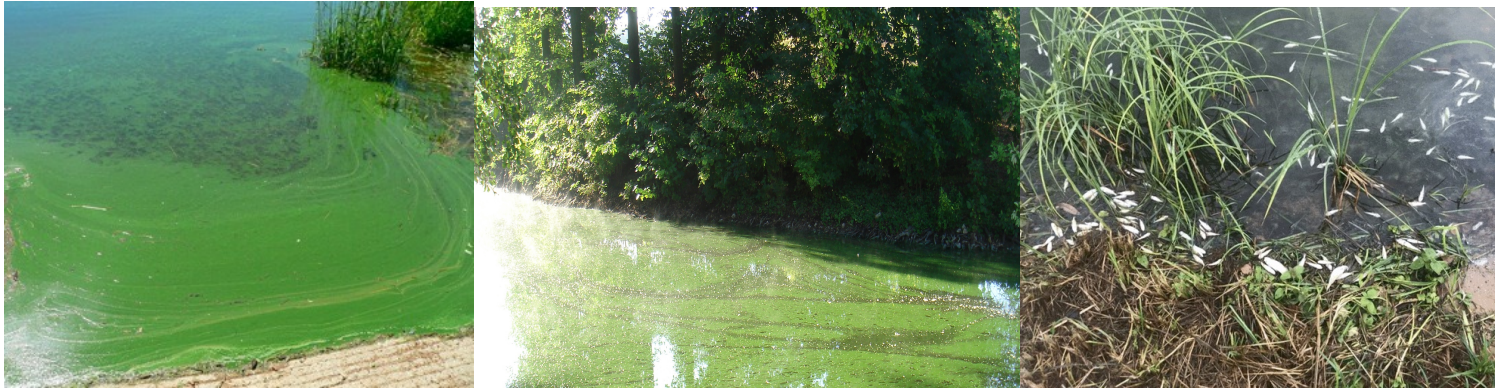
MICROBIAL DYNAMICS DURING A CYANOBACTERIAL BLOOM COLLAPSE

A MESOCOSM APPROACH

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Phytoplankton blooming impact ?

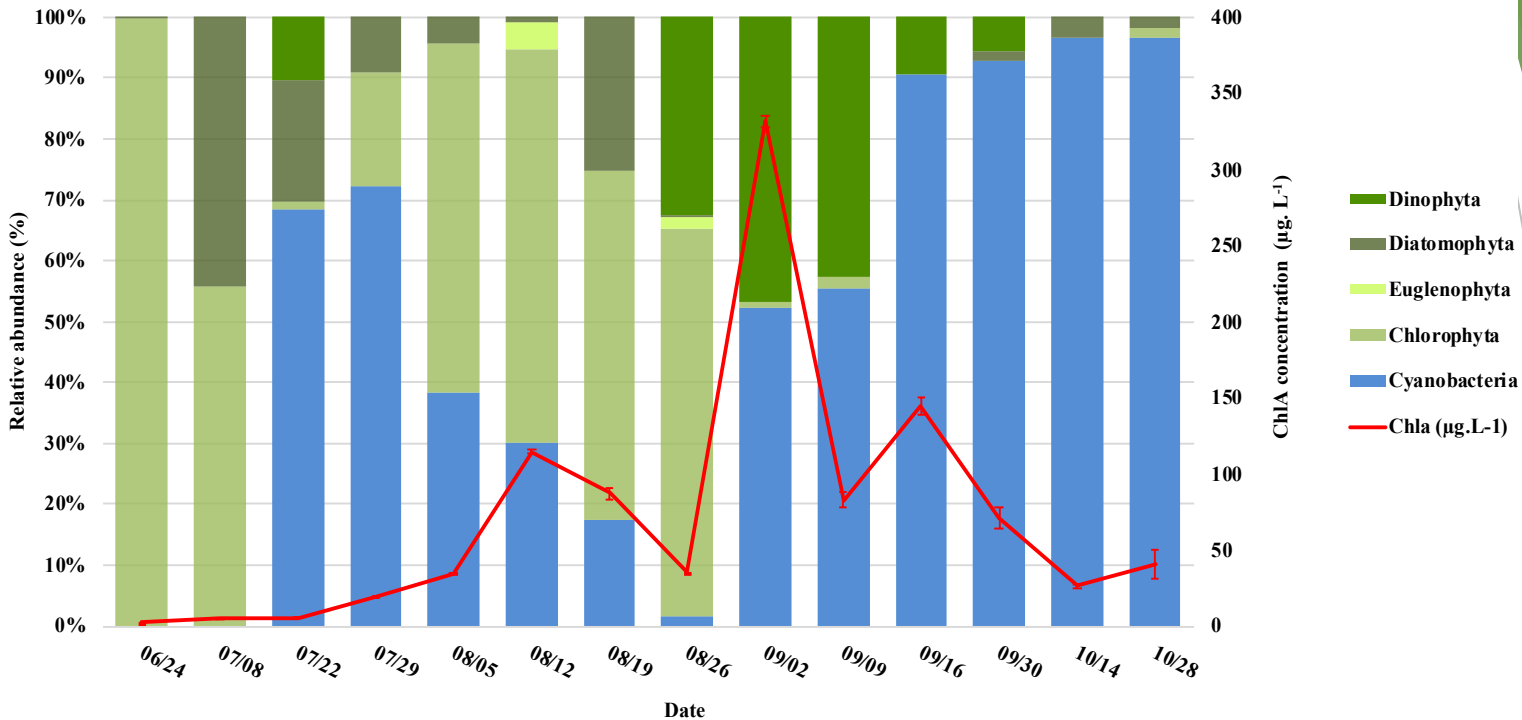
Phytoplankton blooms, mainly cyanobacterial ones, are annual events leading to strong perturbations in eutrophic ecosystems such as the decrease of phytoplanktonic diversity, potential release of toxins, and hypoxia



=> One of the main consequences is the huge release of OM when the bloom collapses

=> A good scenario to disentangle the mechanisms underlying the synergy between OM and microbial heterotrophs

Phytoplankton blooming impact ?

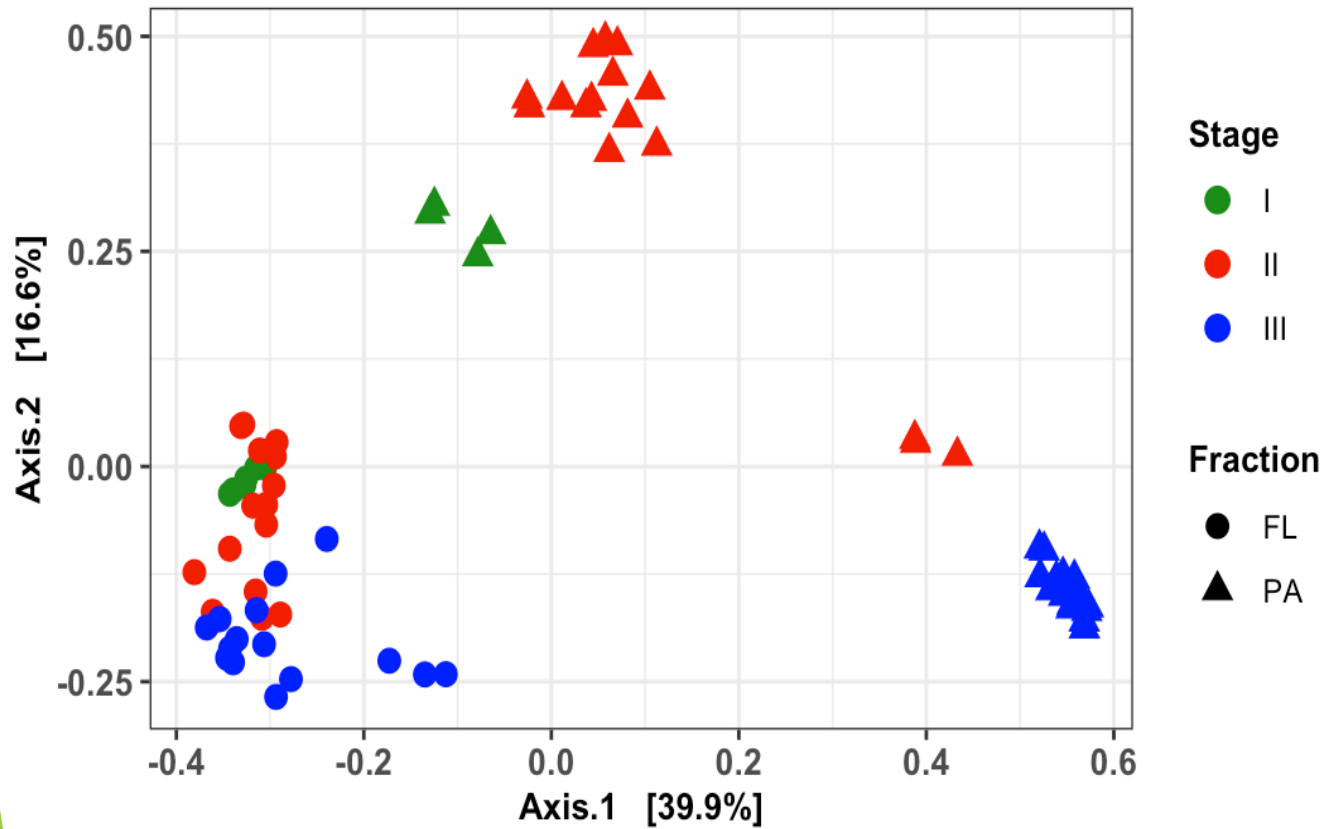


Low biomass
and richness

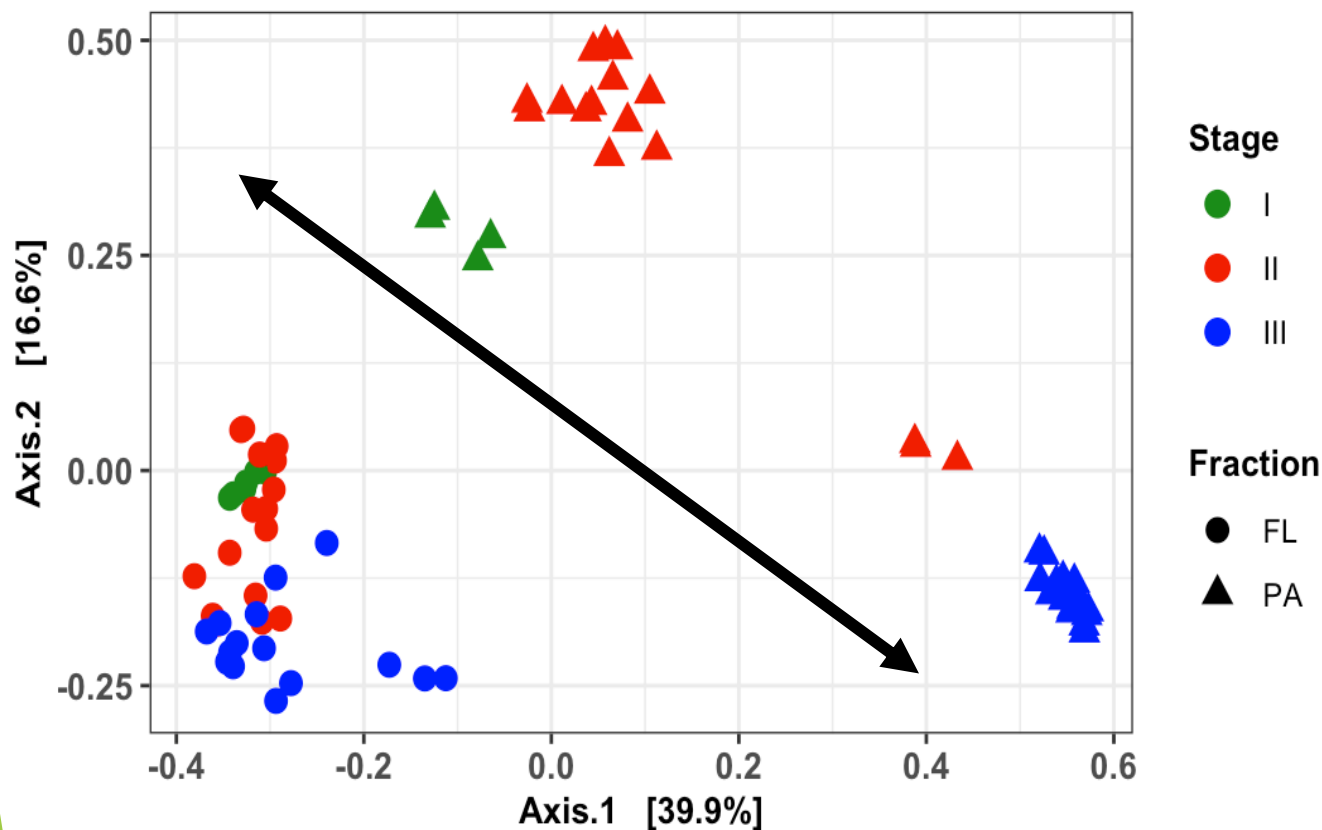
High biomass
and richness

High biomass but
low richness

Phytoplankton selectivity ?



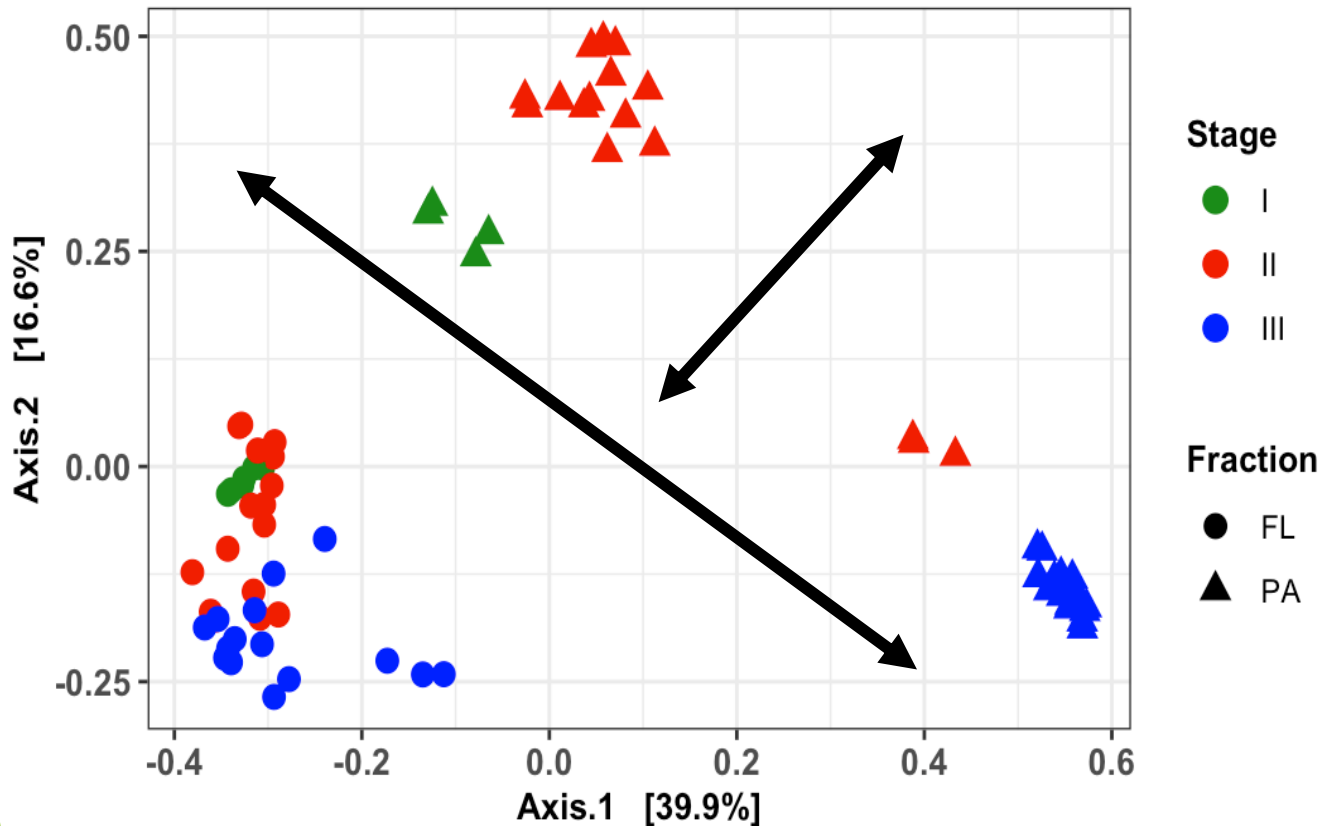
Phytoplankton selectivity ?



=> Strong differentiation according to the living mode

Louati et al., 2023

Phytoplankton selectivity ?



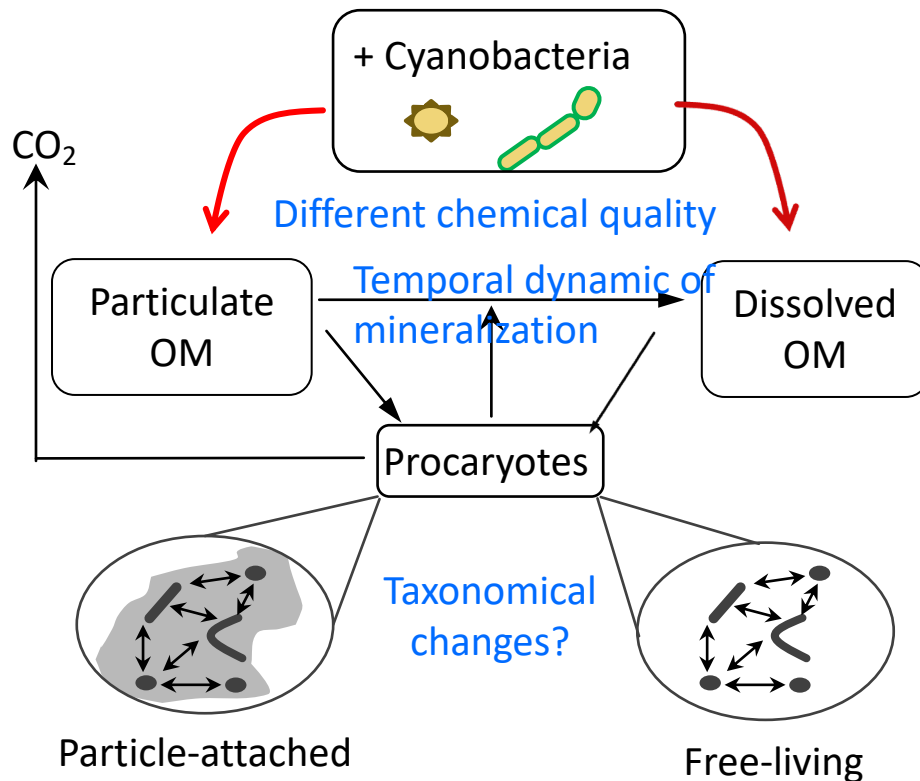
=> Strong differentiation according to the living mode

=> Strong differentiation according to the phytoplankton community

Phytoplankton selectivity ?

=> Synergy established between microbial heterotrophs and organic matter

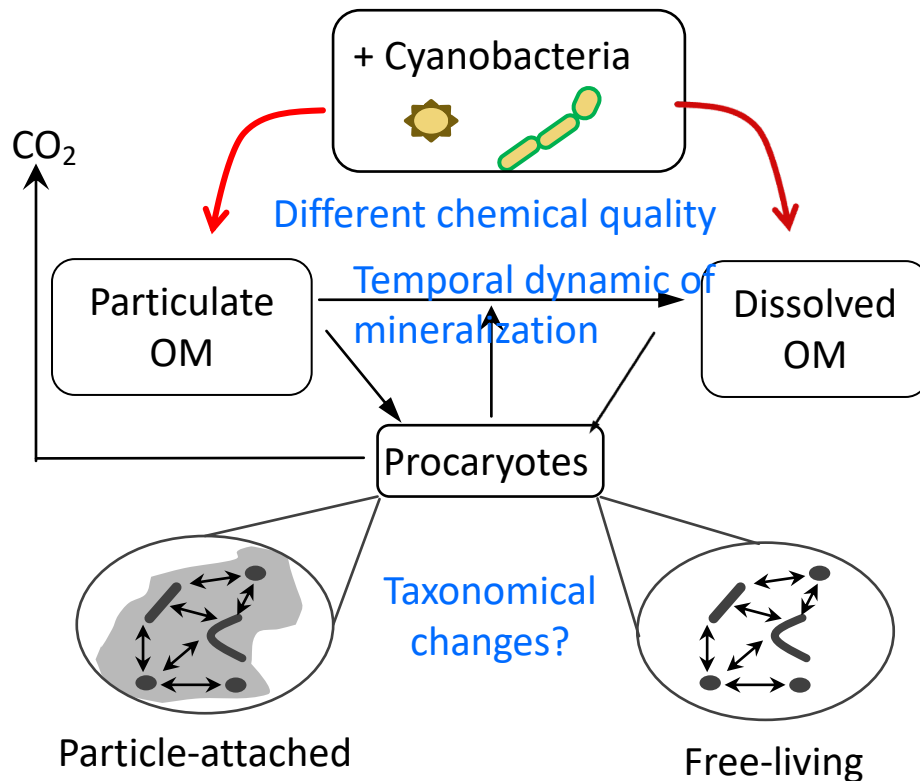
=> Mainly driven by its bioavailability, chemical composition as well as its biotic origin



Phytoplankton selectivity ?

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What is happening at the end of the bloom?

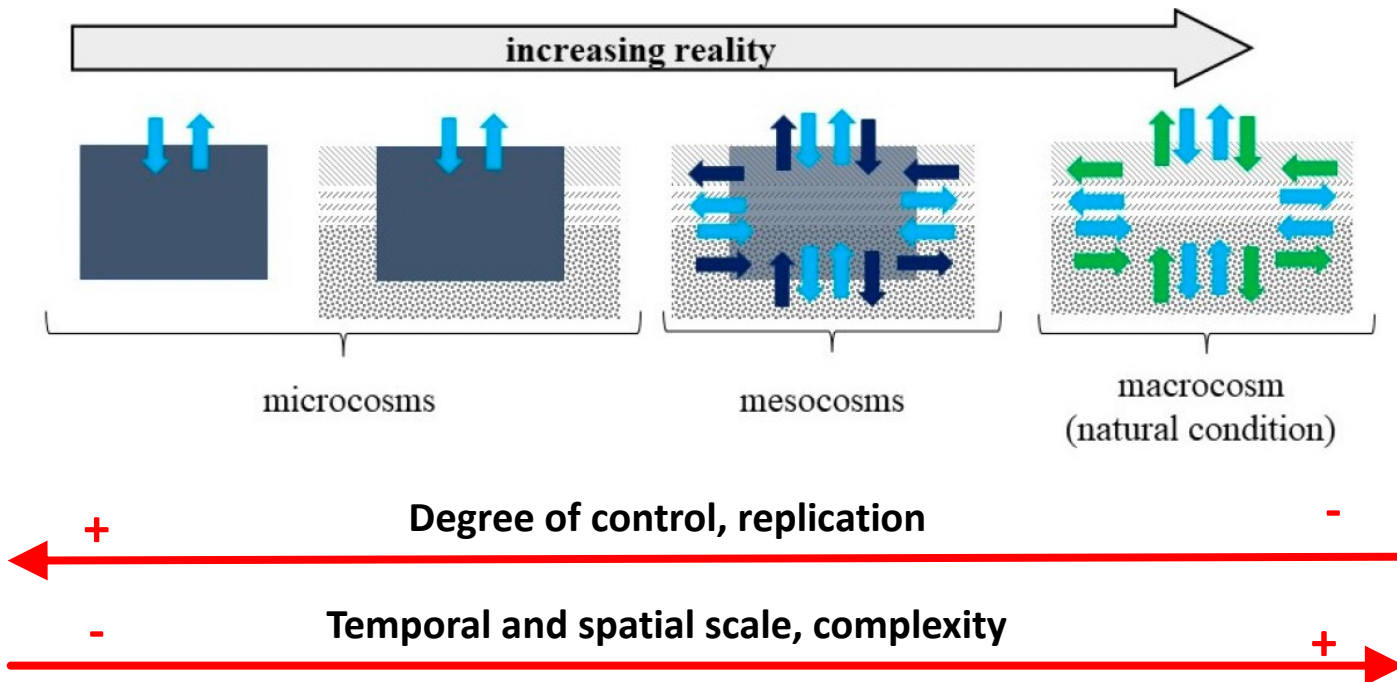


Simulation of a the strong release of particulate and dissolved OMOM



How to do so?

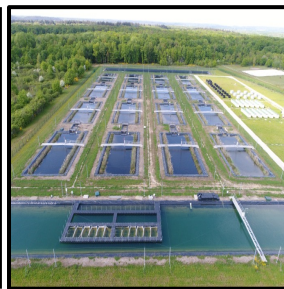
Size matters in aquatic ecology?



Microcosm



Mesocosm



Macrocosm



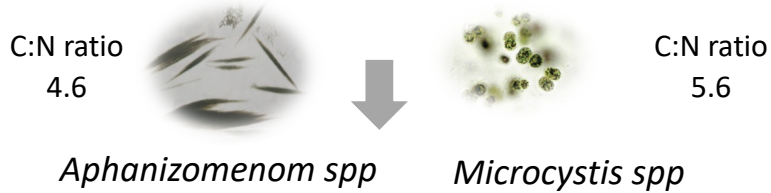
Manipulation of natural ecosystems



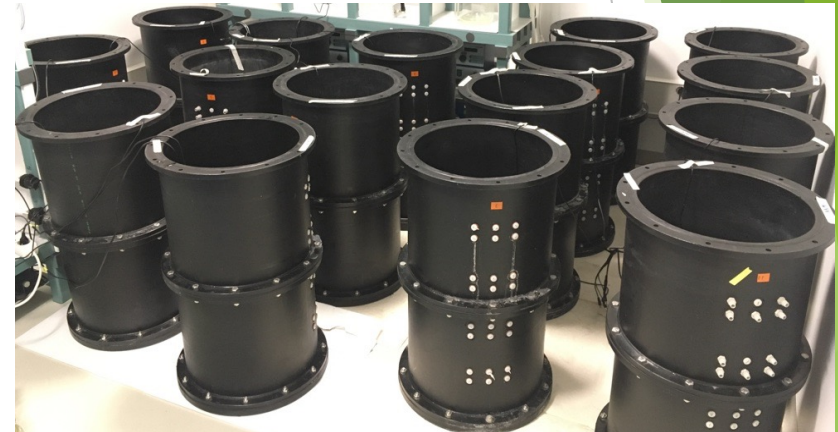
Observation

Mesocosm approach

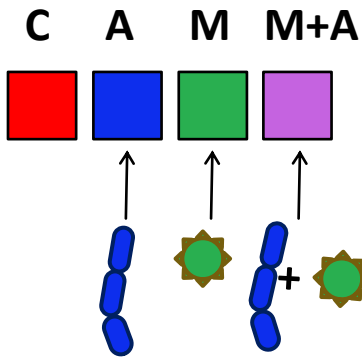
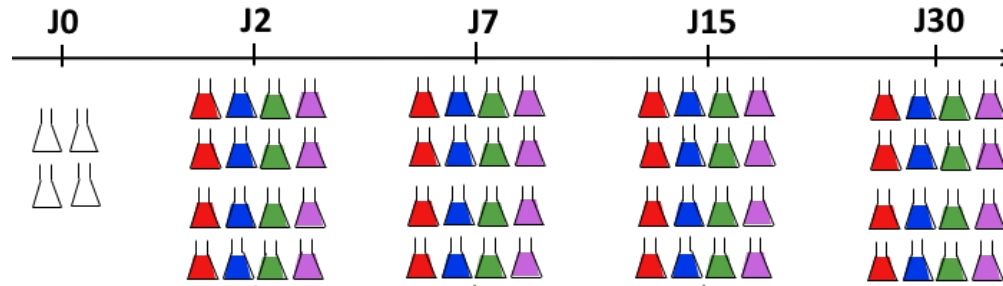
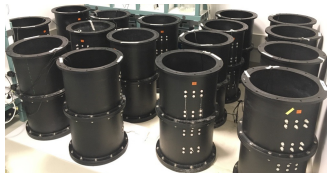
N₂-fixing and non-fixing
bloom-forming cyanobacteria



CEREEP-Ecotron IDF - St-Pierre-lès-Nemours
<https://www.cereep.bio.ens.psl.eu/>



Mesocosm approach

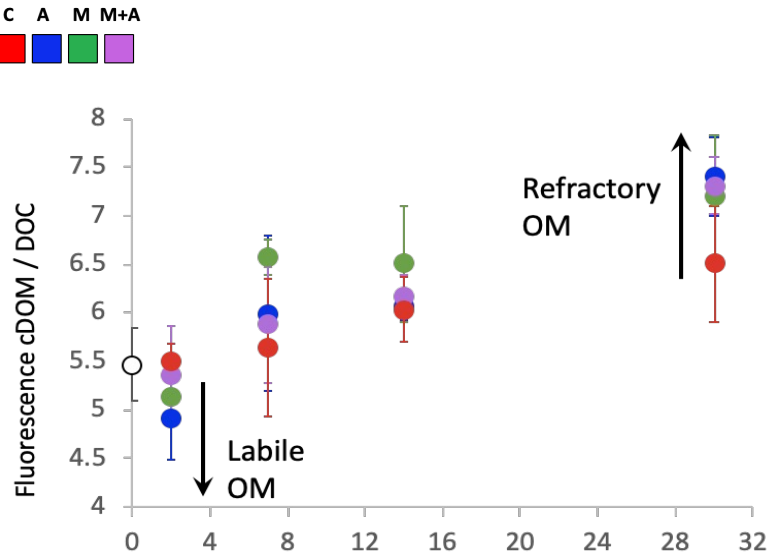


30 days-long incubation in the dark at
23°C in 20L-mesocosms

- Optical characterization of the OM
- Chemical identification of fatty and amino acids
- Nutrient concentrations

- Functional diversity (Biolog®)
- Microbial diversity (16S rRNA metabarcoding)
- Viral and bacterial abundances (cytometry)

How organic matter respond ?



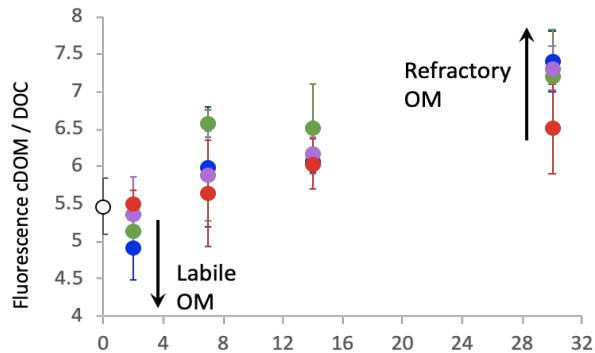
=> Significant mineralization of the labile pool of OM by microbial heterotrophs in a short term

=> No difference of mineralization according to the biotic origin

=> Refractory OM accumulation

How organic matter respond ?

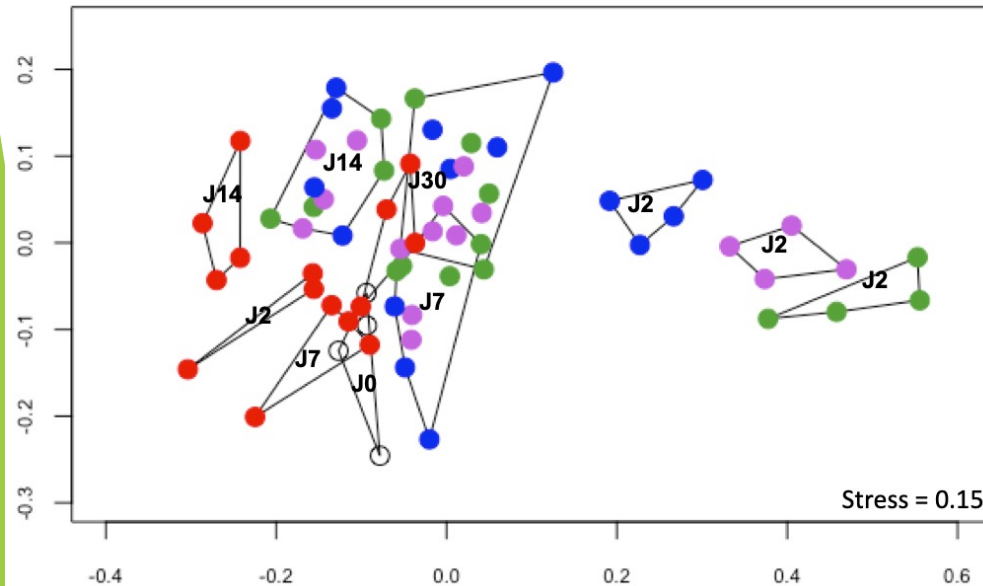
C A M M+A
■ ■ ■ ■



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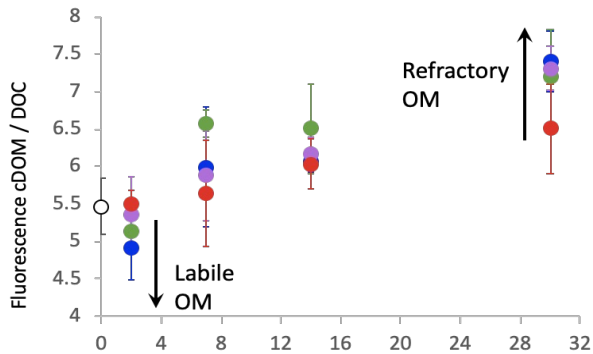


=> Rapid evolution of the OM quality during mineralization

=> OM trajectory according to the biotic origin, with different resilience response

How organic matter respond ?

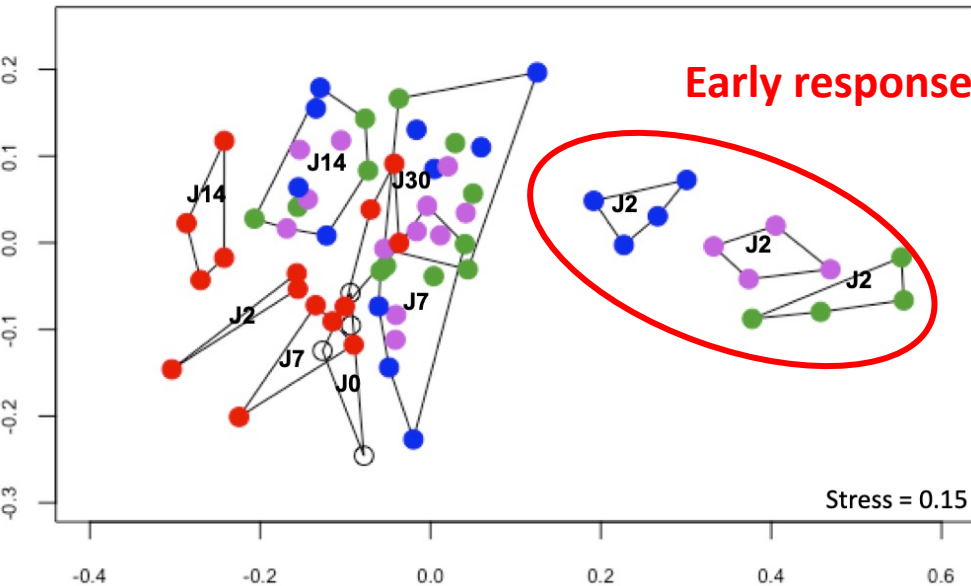
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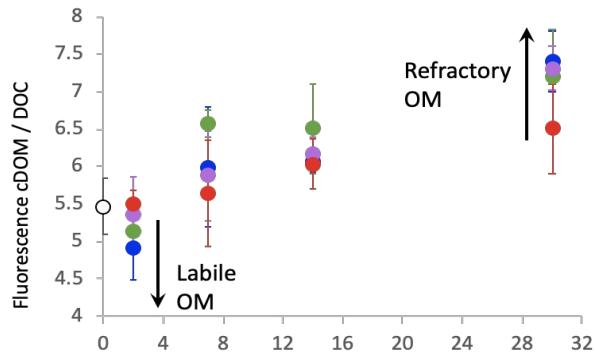


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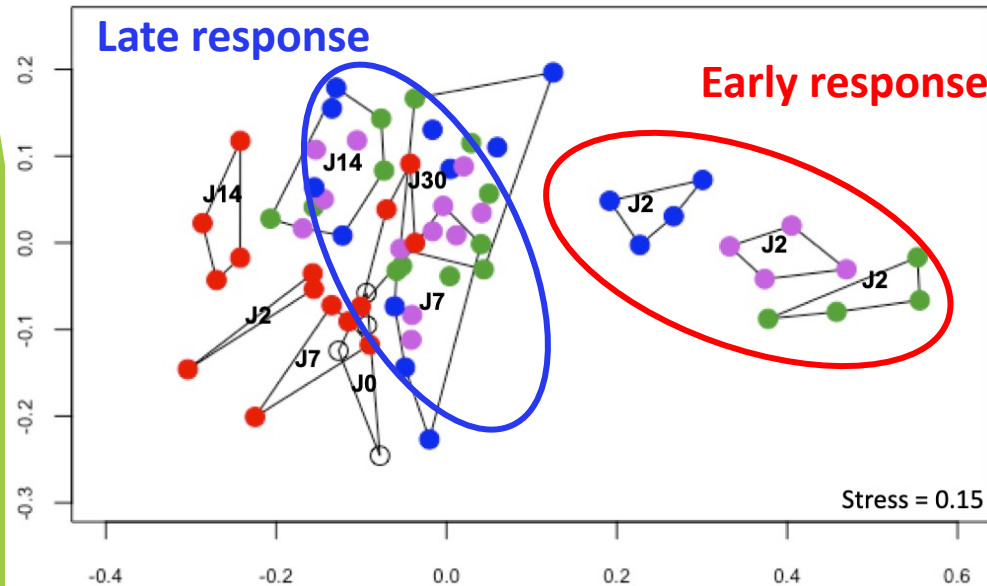
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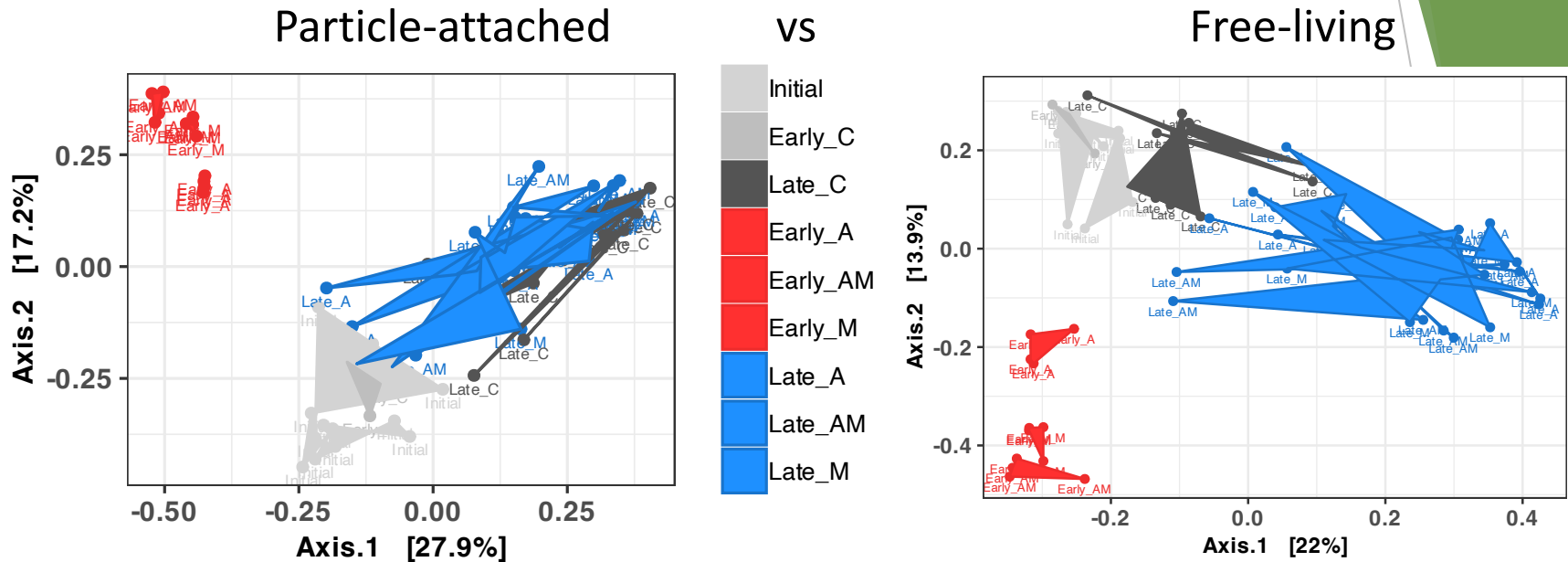
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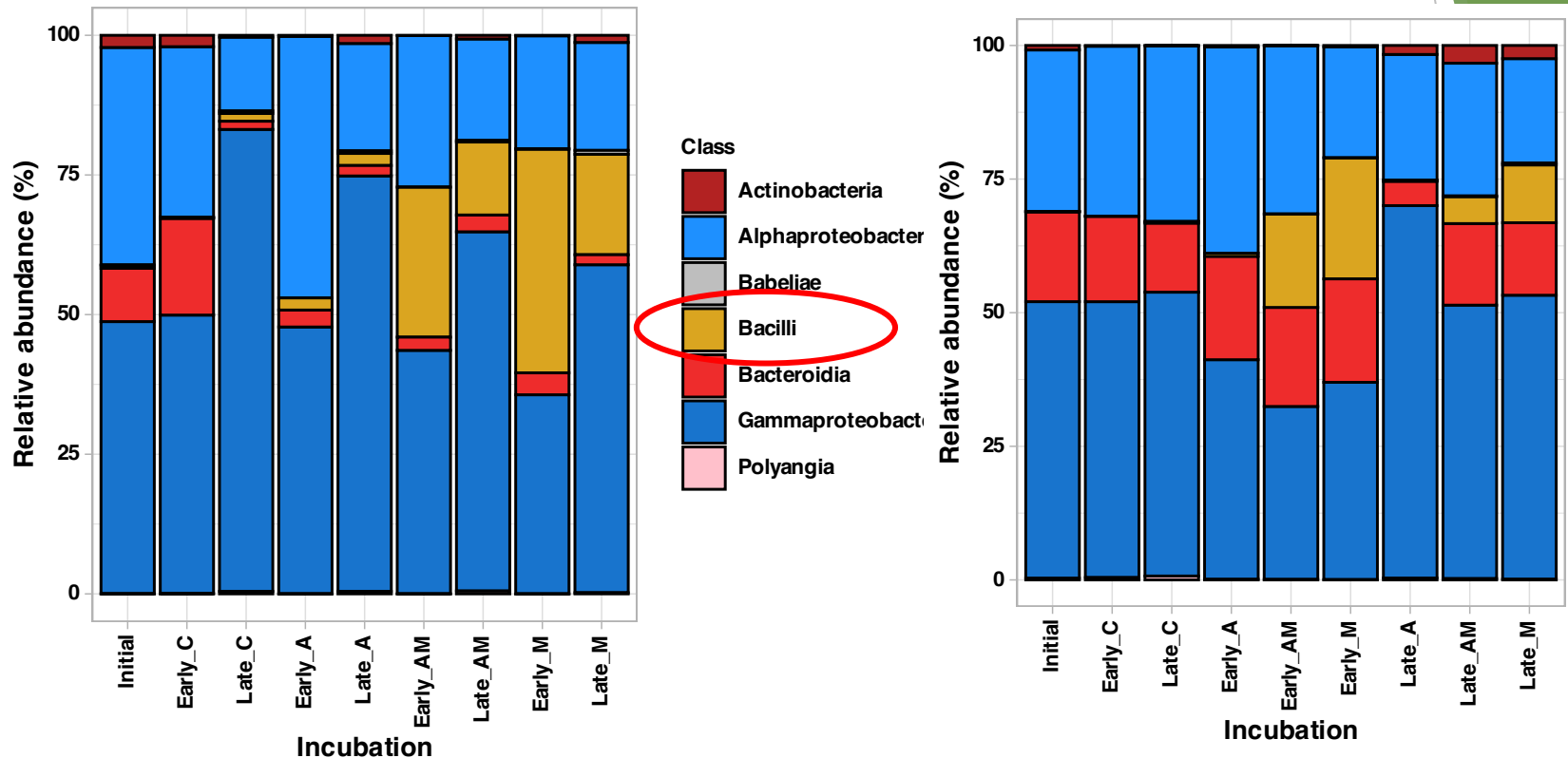
=> OM trajectory according to the biotic origin, with different resilience response

How microbial communities respond ?



- => Strong and rapid impact of the cyanobacterial senescence
- => Highly resilient communities in PA mode
- => Resilient but without return to the initial state

How microbial communities respond ?



- => Emergence of specific microbial groups = Bacilli
- => According to the biotic origin of OM
- => Remain after 30 days

To conclude

Cyanobacterial bloom senescence

- = Strong and impact on the OM pool with a **more refractory quality**
- = Intense and rapid impact on the PA and FL microbial communities
- = **No return to the initial state** of the natural freshwater communities
- = **Appareance of Bacilli** members, specifically *Exiguobacterium* spp followed by *Bacillus* spp