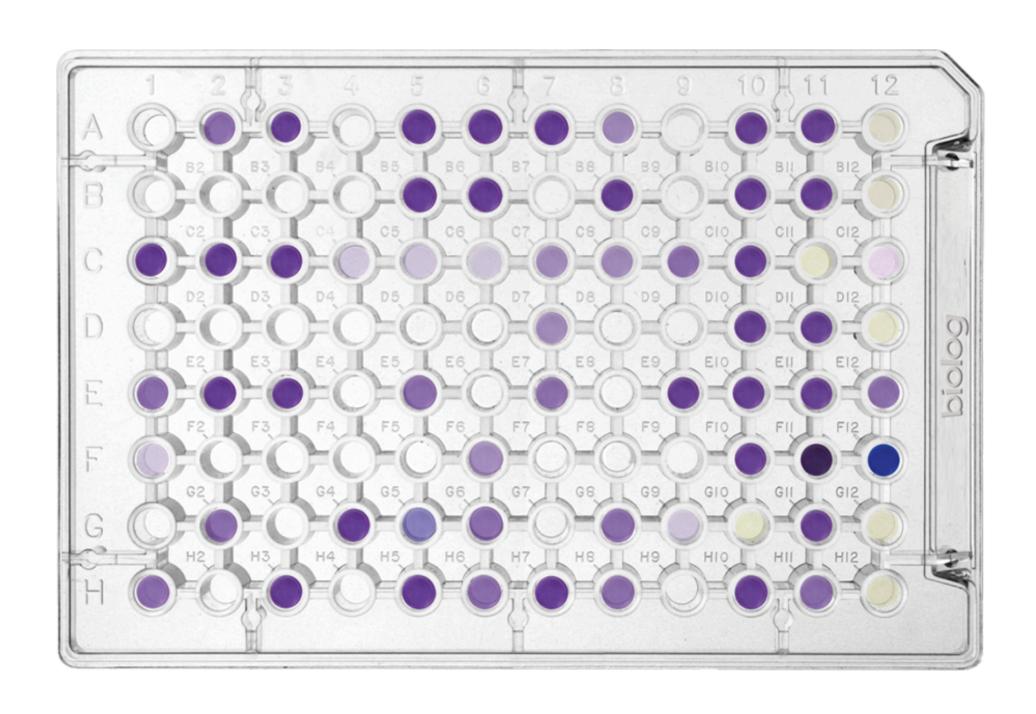
EPFL

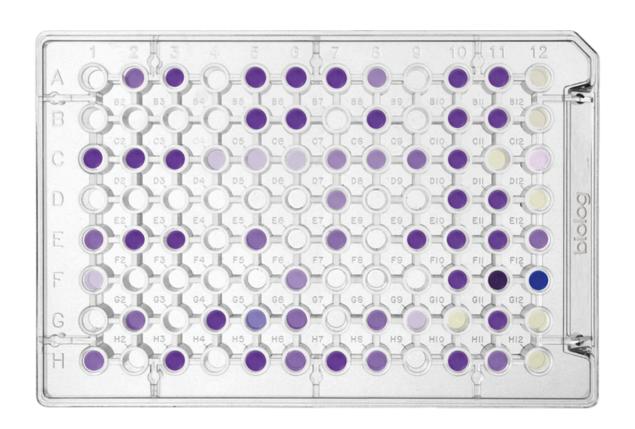


Biolog EcoPlates

Microbial Ecology

Fanny Bitoun & Miyuka Laurenson

General Presentation:

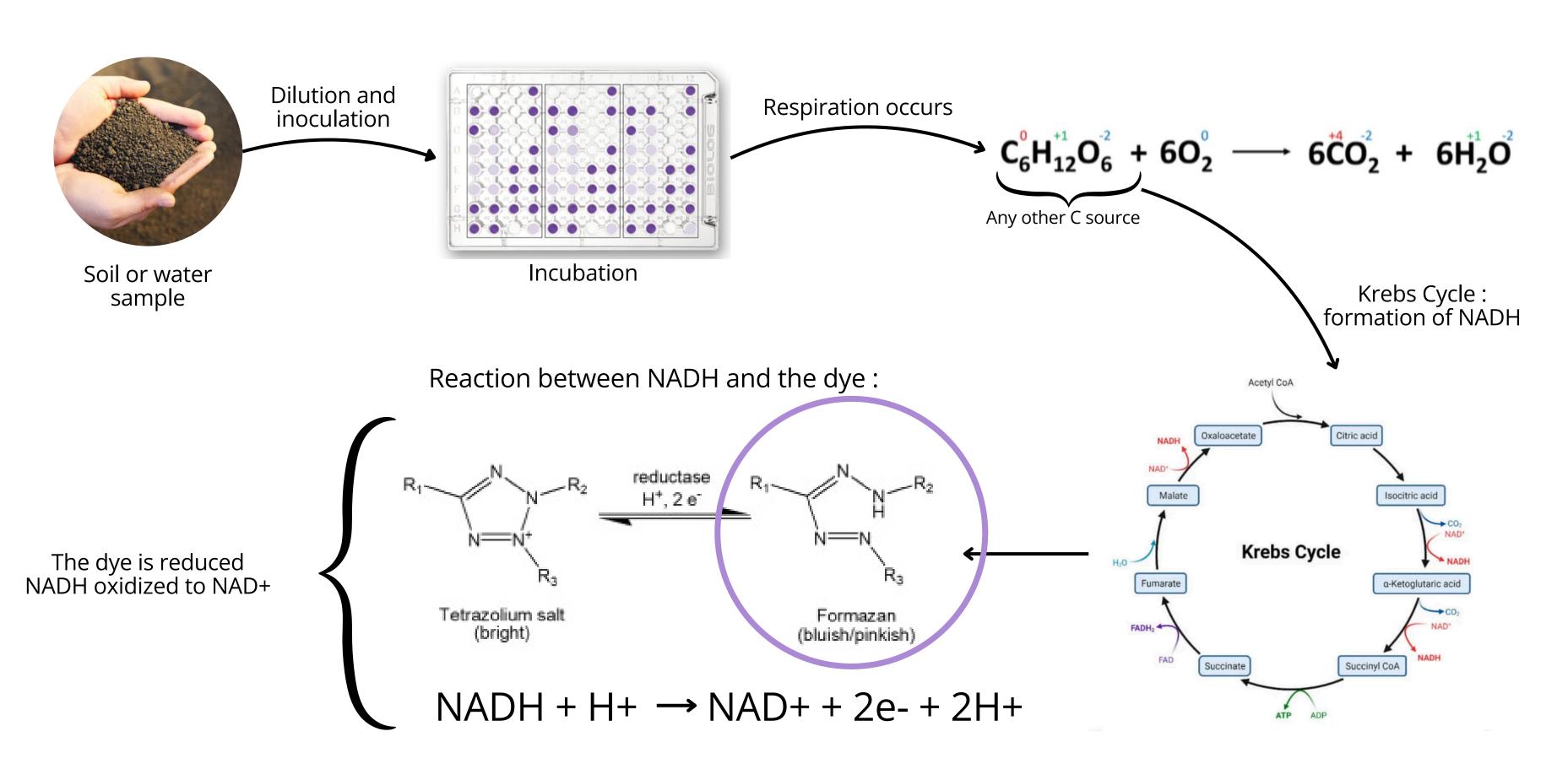


- Assessment of bacterial substrate use and metabolic potential in the environment
- 31 environmentally relevant organic carbon substrates + 1 control
- 3 sets of 32 wells : data replicates
- Dye present in each well

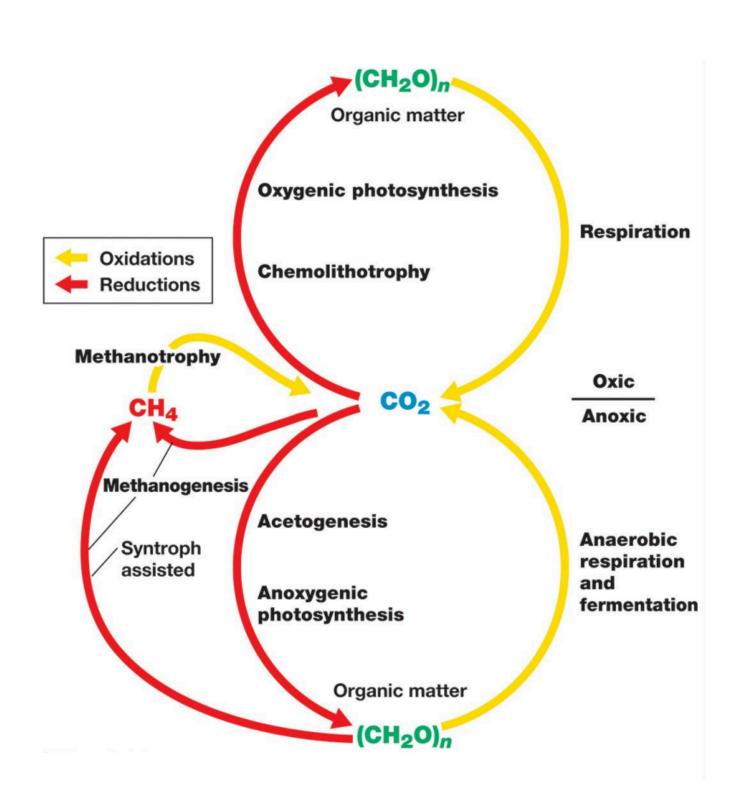
- Carbon sources classed in 6 categories
- 9/31 are reported plant exudates

Polymers Carbohydrates Carboxylic Acids Amino-acids Amines Phenolic compounds

Science behind the method

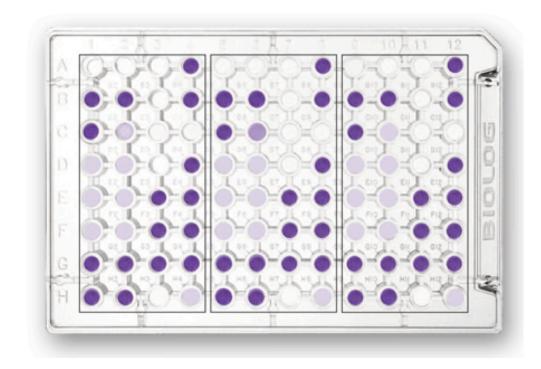


Reminder: Carbon cycling



- Carbon: essential growth & metabolic component
- Ecoplates target specific growth strategies :
 - Organic C source : Heterotrophy
 - Organic e- and H+ donors : Chemo organotrophy
 - Mostly used in aerobic conditions

Key Parameters to Analyze:



- Pattern development : **similarity** between replicates
- Rate of color change in each well: microbial activity
- Richness of well response : diversity

In class:

- Isolated microorganism
- No replicates
- Activity and diversity qualitatively assessed

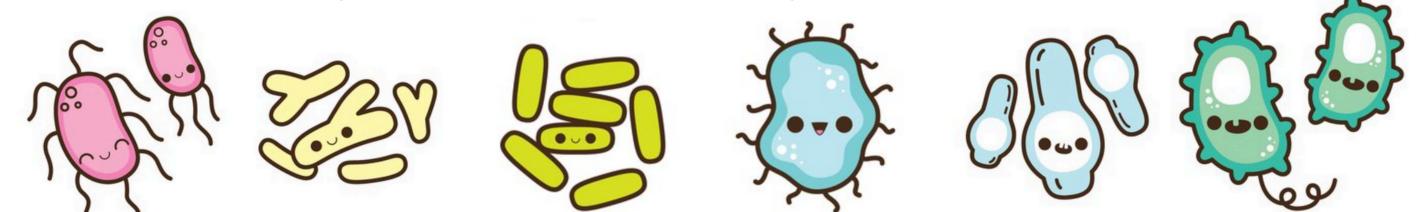
In literature:

- Community level
- Absorbance directly measured over time → quantitative assessment

Community-Level Physiological Profiling (CLPP):

Characterization and classification of heterotrophic microbial communities based on **sole carbon source utilization patterns**

→ Functional diversity relates to the activity of the microflora



→ Quickly evaluate impact of practices (pollutant, fertilizer etc) on soil microflora



Community-Level Physiological Profiling (CLPP):

Metrics derived:

• **AWCD** (Average Well Color Development): High AWCD: High **microbial activity + diversity**

$$AWCD = \frac{\sum_{i=1}^{N} OD_{i}}{N}$$

• Shannon-Weaver Index (H):

Considers both Richness and Evenness

High Shannon–Weaver Index = the community uses many carbon sources

equally well -> greater diversity

$$H = -\sum_{i=1}^{OD_i} P_i \cdot \ln P_i \text{ , } P_i = \frac{OD_i}{\sum_{i=1}^{N} OD_i}$$

Limitations of Ecoplates

Microbial • Only cultivable bacteria • Bias toward fast growing bacteria

- Incubation

 High substrate concentrations
 No combination of substrate

- Measurements should be done as fast as possible after sampling
 Choice of the part of the soil to use

The Biolog EcoPlate™ Technique for Assessing the Effect of Metal Oxide Nanoparticles on Freshwater Microbial Communities

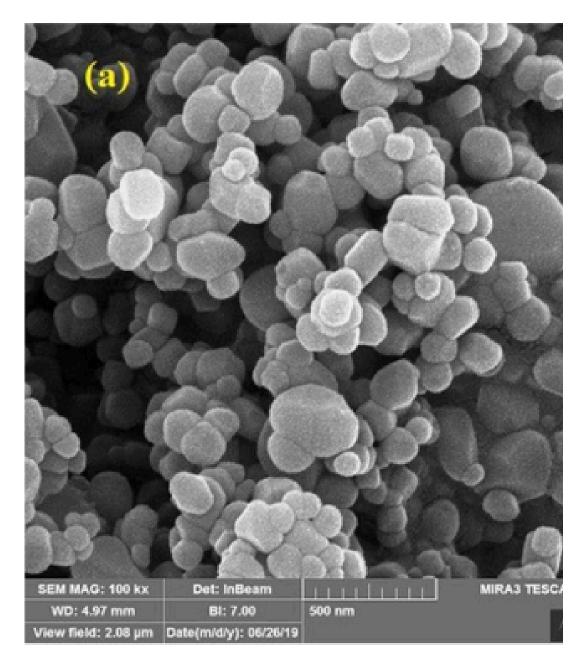
<u>Imre Németh</u> ¹, <u>Szabina Molnár</u> ¹, <u>Emese Vaszita</u> ¹, <u>Mónika Molnár</u> ^{1,*}



Research Question:

Assessing the impact of nTiO2 and nZnO released in freshwater ecosystems on local microbial communities

What are nTiO2 and nZnO?



https://www.nanoshel.com/product/tio2-nanoparticles

Nano titanium dioxide (nTiO2) and nano zinc oxide (nZnO):

- Anthropogenic engineered metal oxide nanoparticles
- Used in medicine, electronics, cosmetic and textile industry
- Widely produced and transported
 - → risk of release in the environment



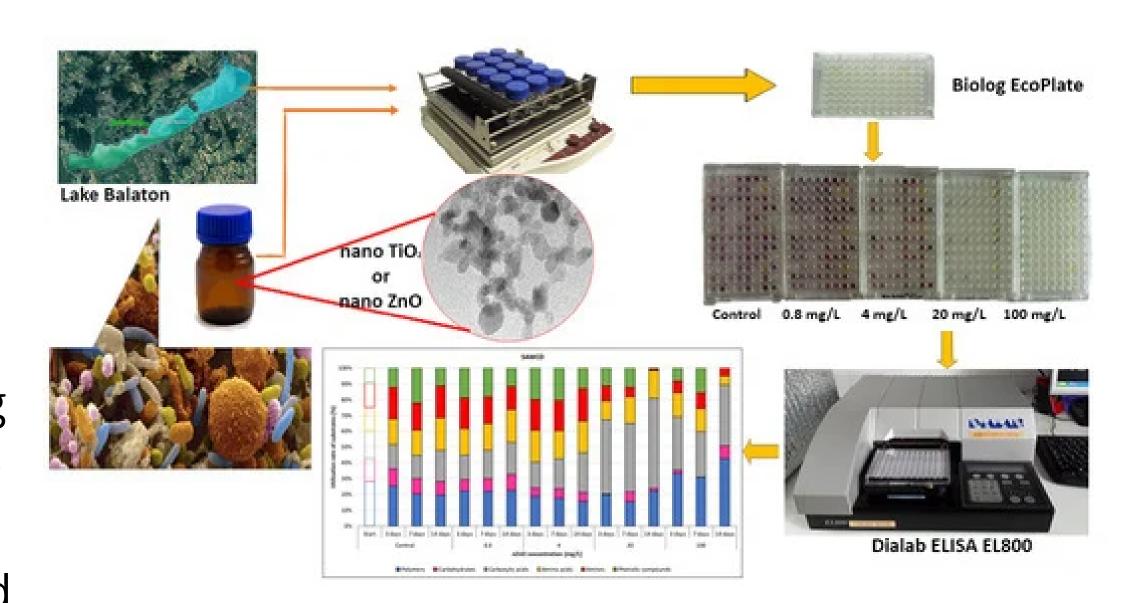
Impact of nanoparticles on bacteria:

Can interact with various component in the cells (membrane, proteins, genes expression etc)

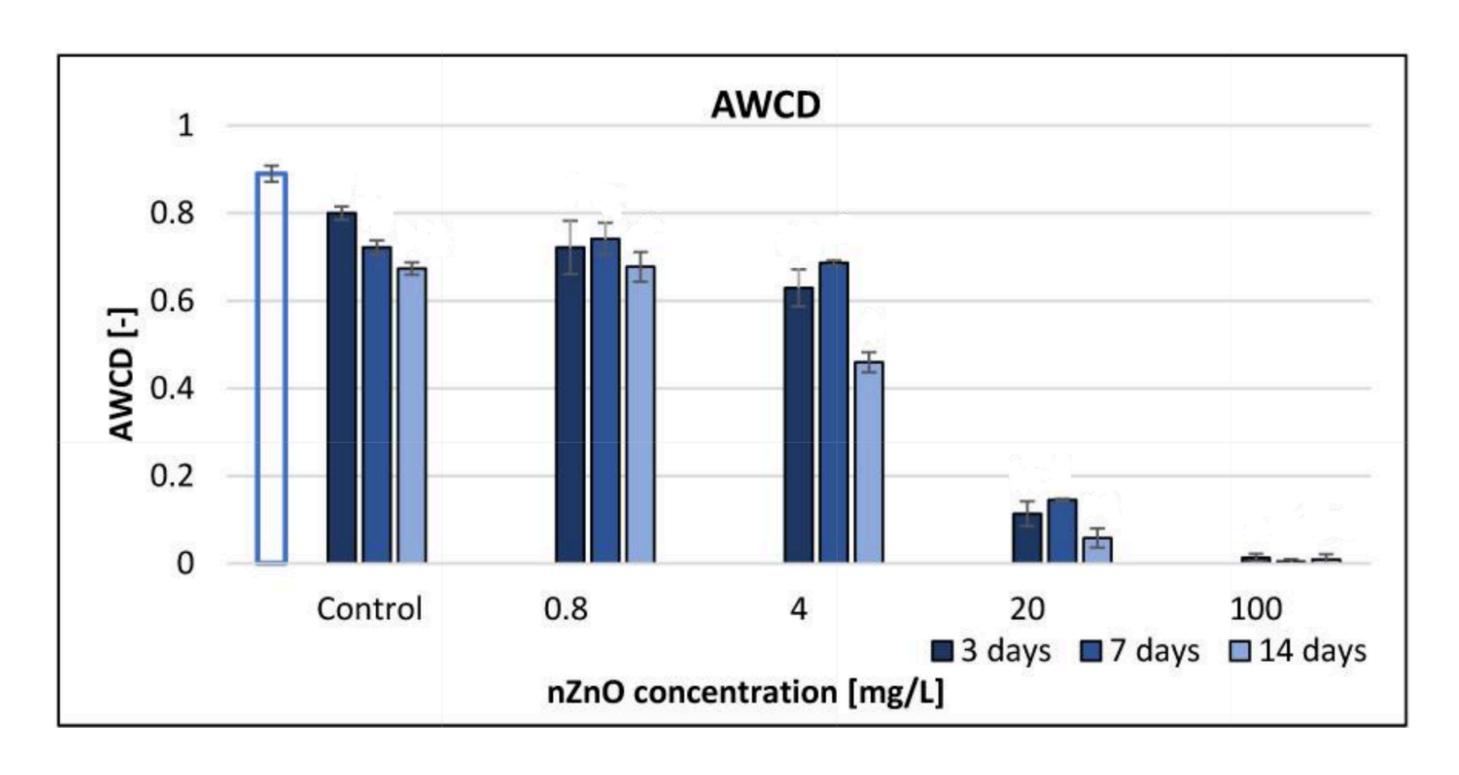
→ reduction of enzymatic activity, reproduction, or increase morbidity

Methods

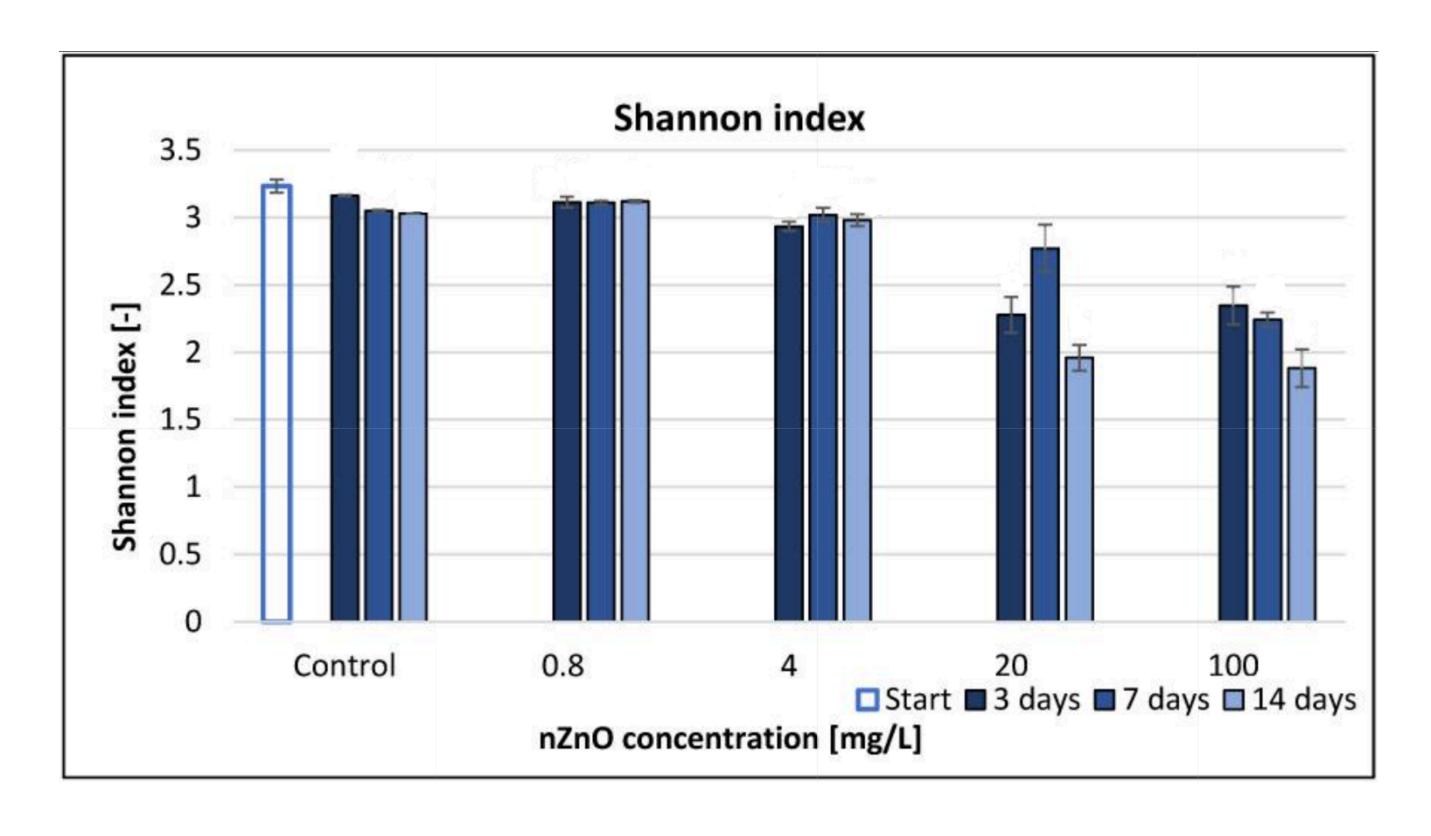
- Water samples collected in Lake Balaton, Hungary in June and October 2020
- **Inoculation** with the two nanoparticles suspension
- Community-level
 physiological profiling using
 Biolog Ecoplates → Substrate
 utilization over time
- Metrics analyzed : AWCD and Shannon diversity index



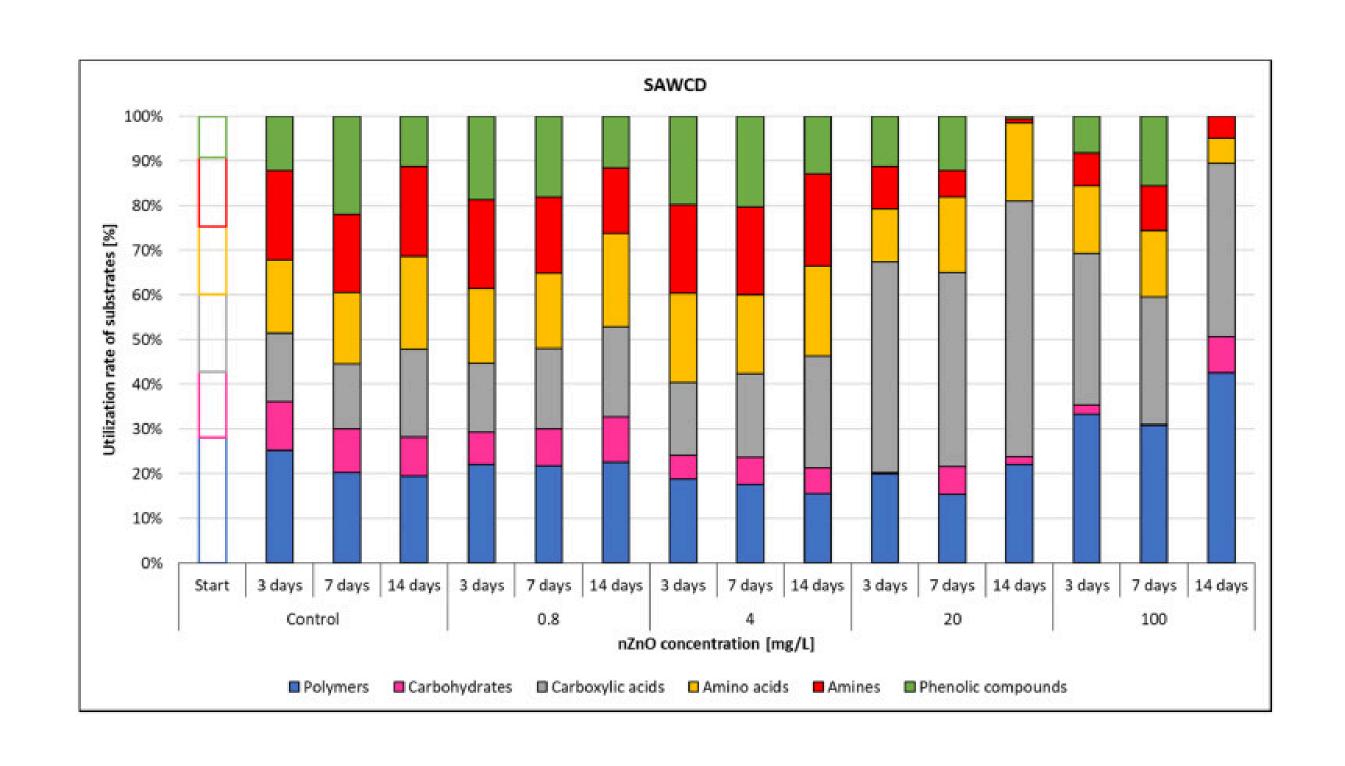
Results: Average Well Color Development



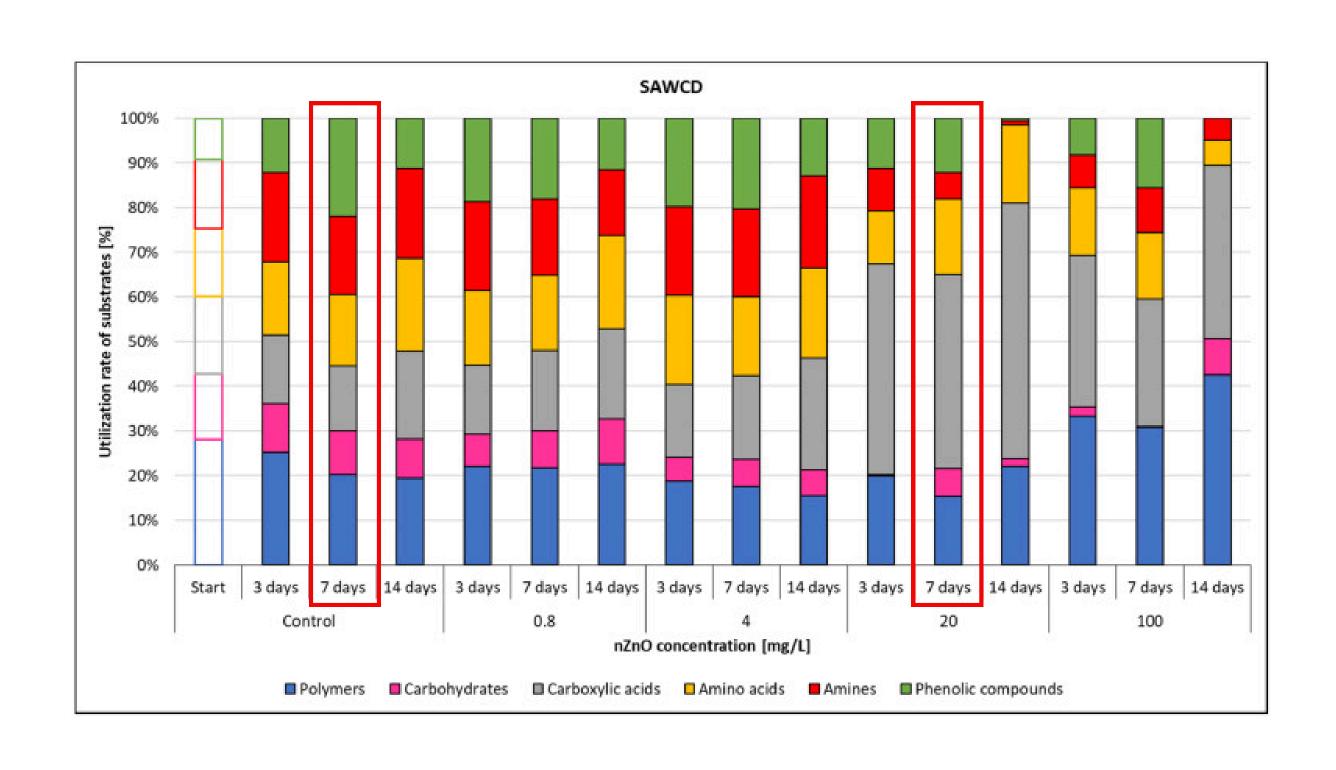
Results: Shannon diversity index



Results: AWCD by substrate category



Results: AWCD by substrate category



Conclusion

Main advantages:

- Simple, rapid and cost-effective method
- Allows for the comparison of community level response against changing/disturbed environments

One step further:

- Ecoplates give only a first insight on the community profile
- Pair its utilization with enzymatic activity measurement or genomics methods