

EVOLUTION OF MICROBIAL COMMUNITIES IN AEROBIC GRANULAR SLUDGE DURING CHANGES OF THE WASTEWATER COMPOSITION

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SWISS NATIONAL SCIENCE FOUNDATION



ÉCOLE POLYTECHNIQUE
FÉDÉRALE DE LAUSANNE

Microbial processes for wastewater treatment

Activated sludge

- Conventional microbial process in wastewater treatment plants
- Flocular structures, settle slowly



Aerobic granular sludge

- Granular structures, settle fast
 - Allows high sludge concentration
 - Suited for biological phosphorus removal
 - > chemicals saving
- Space and time saving

After 1
minute
settling

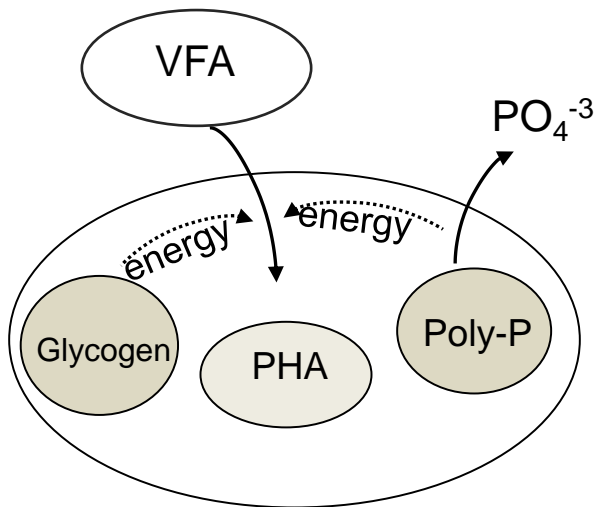


Phosphate accumulating organisms (PAO)

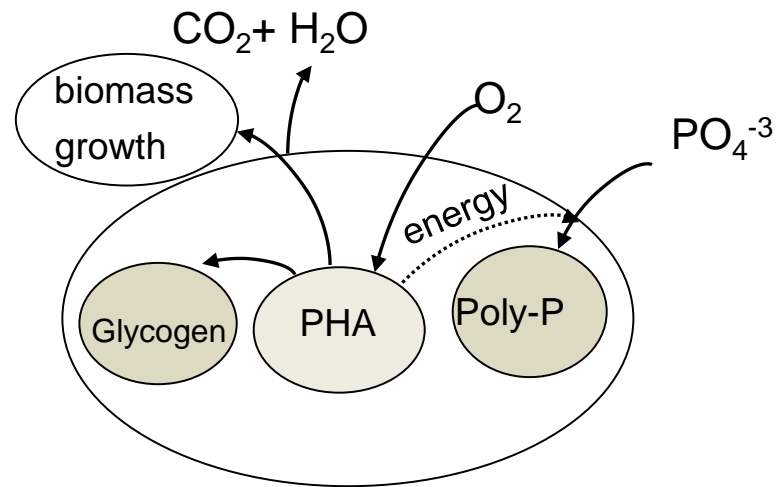
PAO model organism

Accumulibacter

Anaerobic



Aerobic

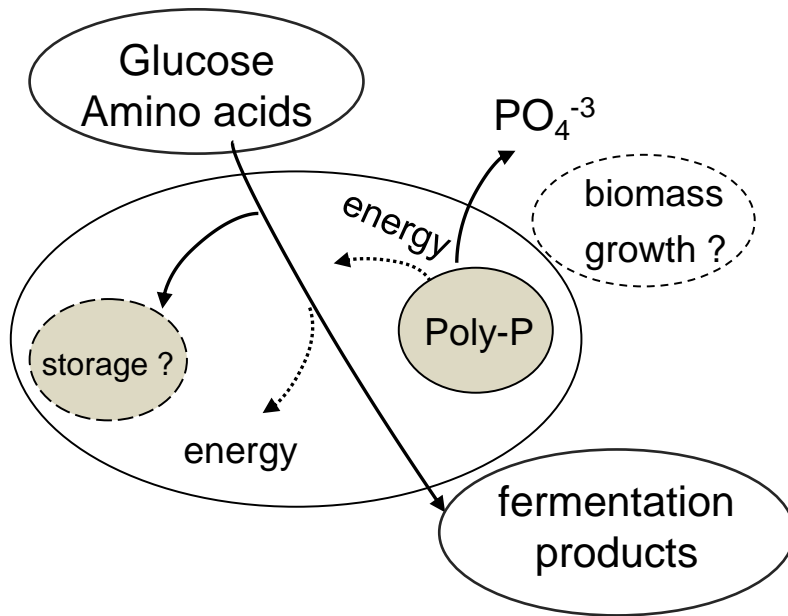


Phosphate accumulating organisms (PAO)

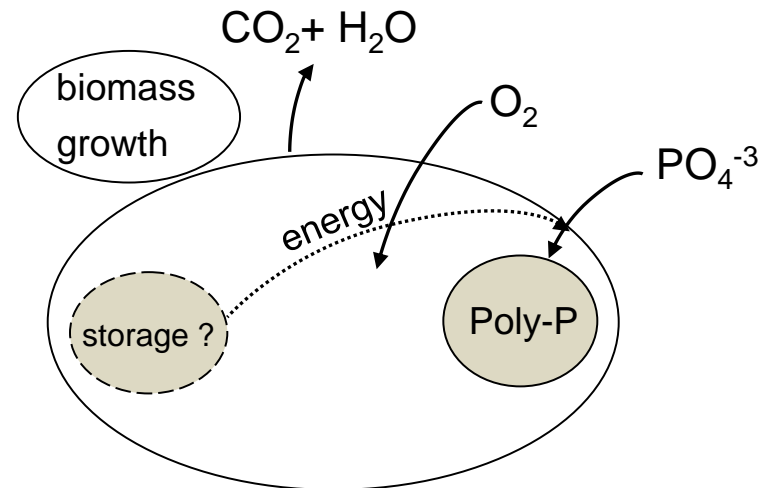
Fermentative PAO

Tetrasphaera

Anaerobic

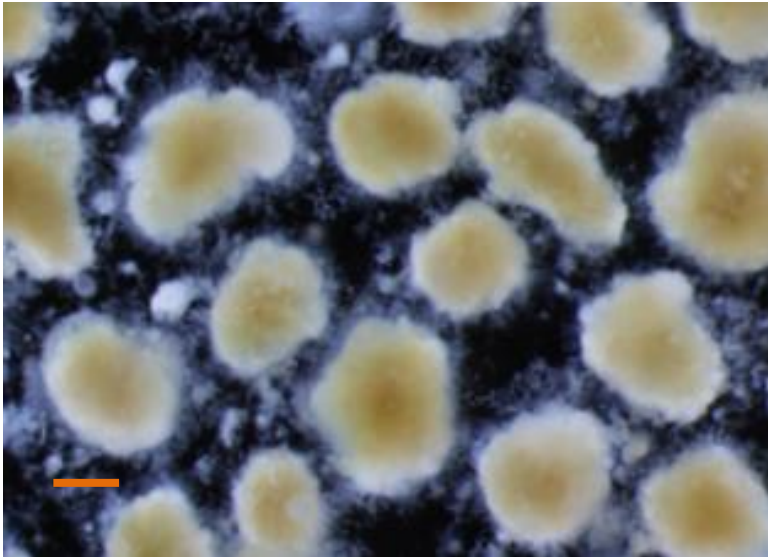


Aerobic

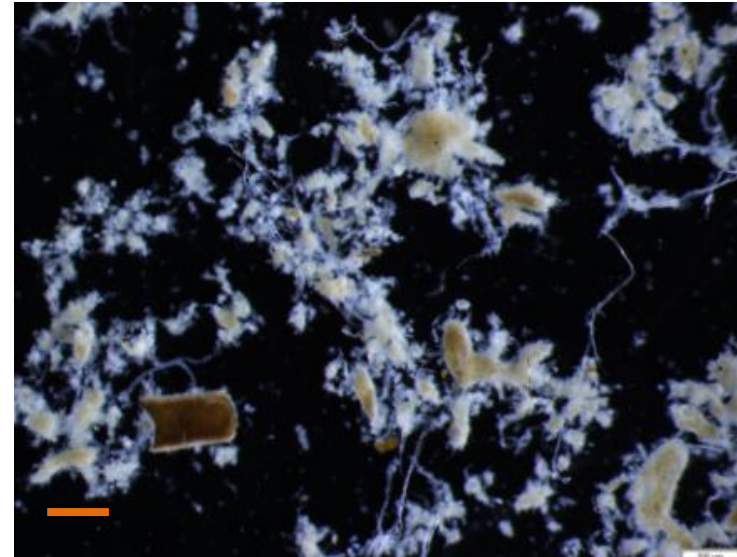


Aerobic granular sludge (AGS) for wastewater treatment

AGS fed with simple synthetic wastewater ...



... and with raw municipal wastewater

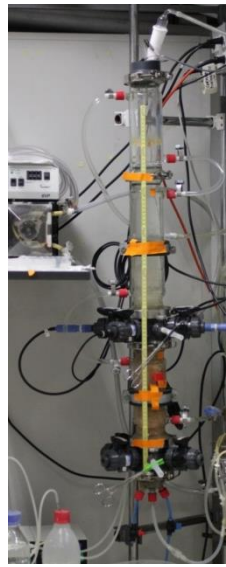


— 500 μ m

Eawag- LTE

- Are **fermentable** or **polymeric** compounds responsible for this difference ?
- How do they impact - the settling characteristics,
 - the nutrient removal,
 - the microbial communities of the AGS?

Progressive changes of the wastewater composition in two lab-scale reactors



lab-scale reactor

1 reactor ← 2 reactors → 1 reactor

Transition to
**simple
monomeric
wastewater**

AGS fed with
**complex
monomeric
wastewater**

Transition to
**complex
polymeric
wastewater**

VFA

VFA, glucose,
amino acids

VFA, glucose,
amino acids,
starch, peptone

Introduction

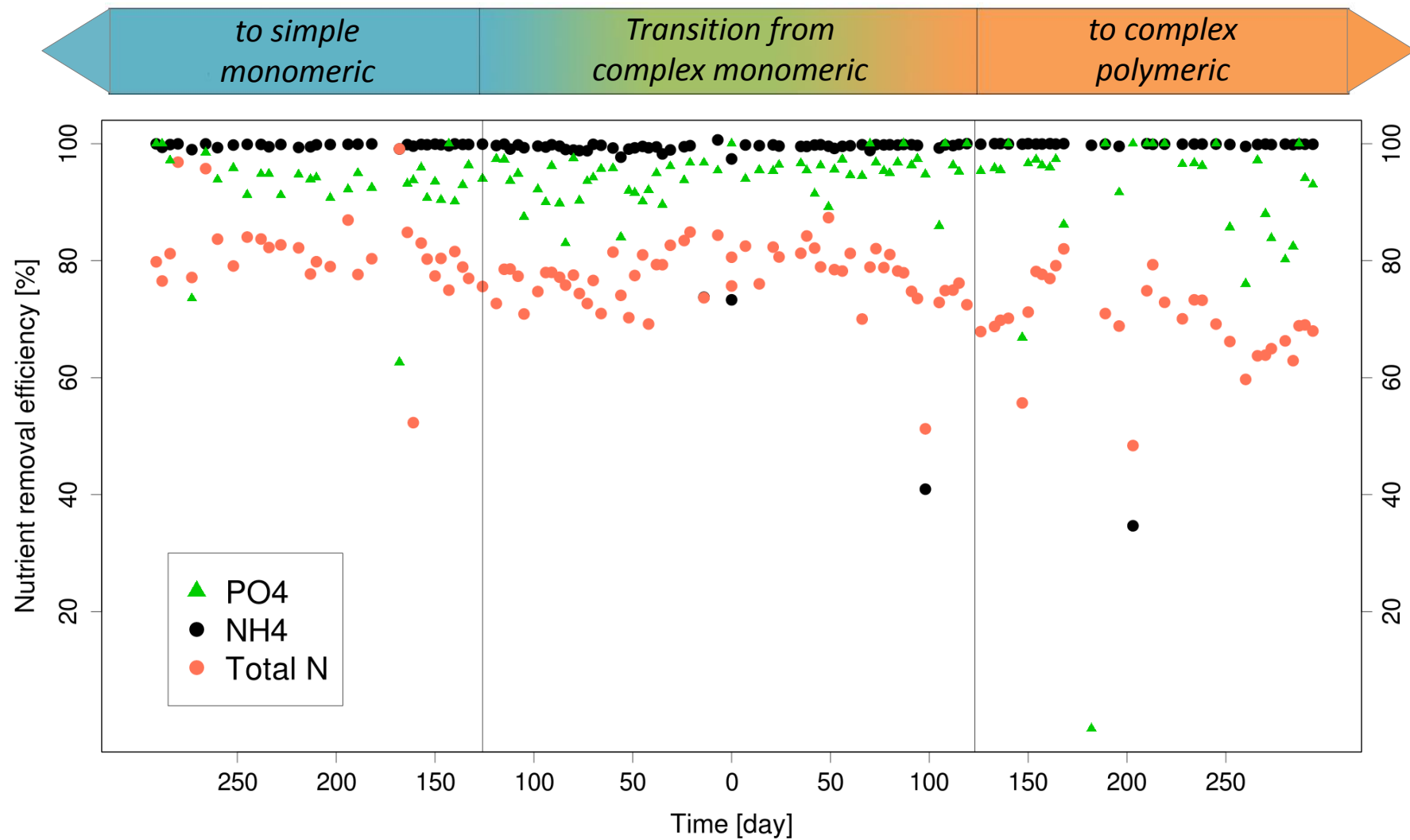
Objectives

Method

Results

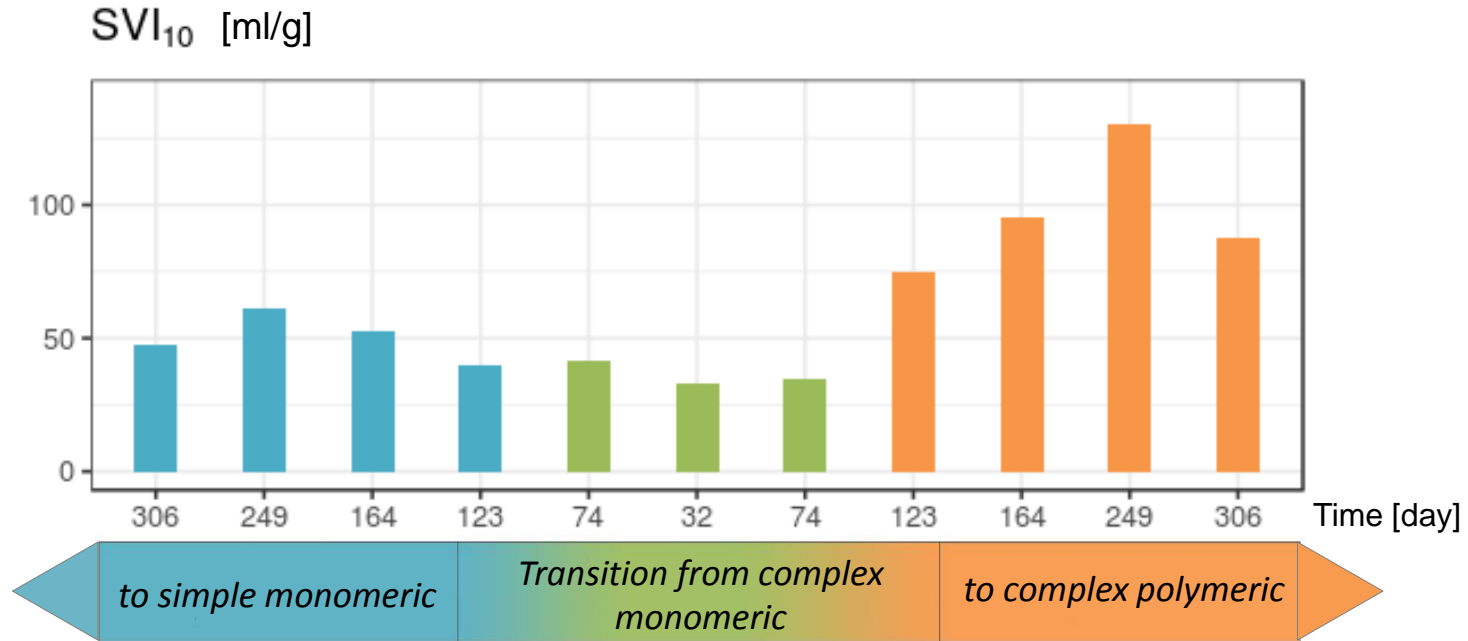
Conclusion

Evolution of nutrient removal efficiency



Introduction
Objectives
Method
Results
Conclusion

Evolution of the settling properties



Introduction

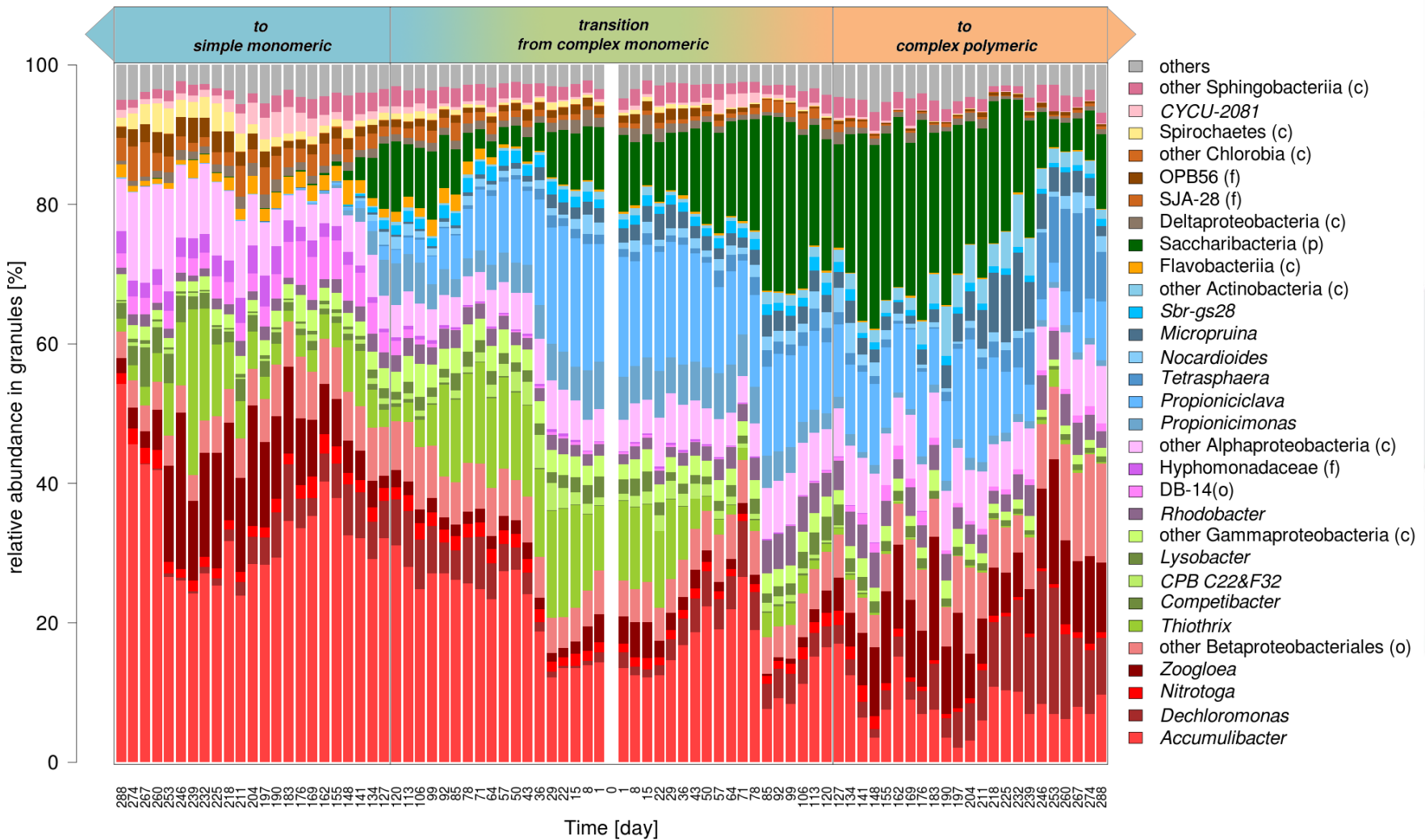
Objectives

Method

Results

Conclusion

Evolution of the structure of the bacterial communities



Introduction
Objectives
Method
Results
Conclusion

phosphate accumulating organisms (PAO)

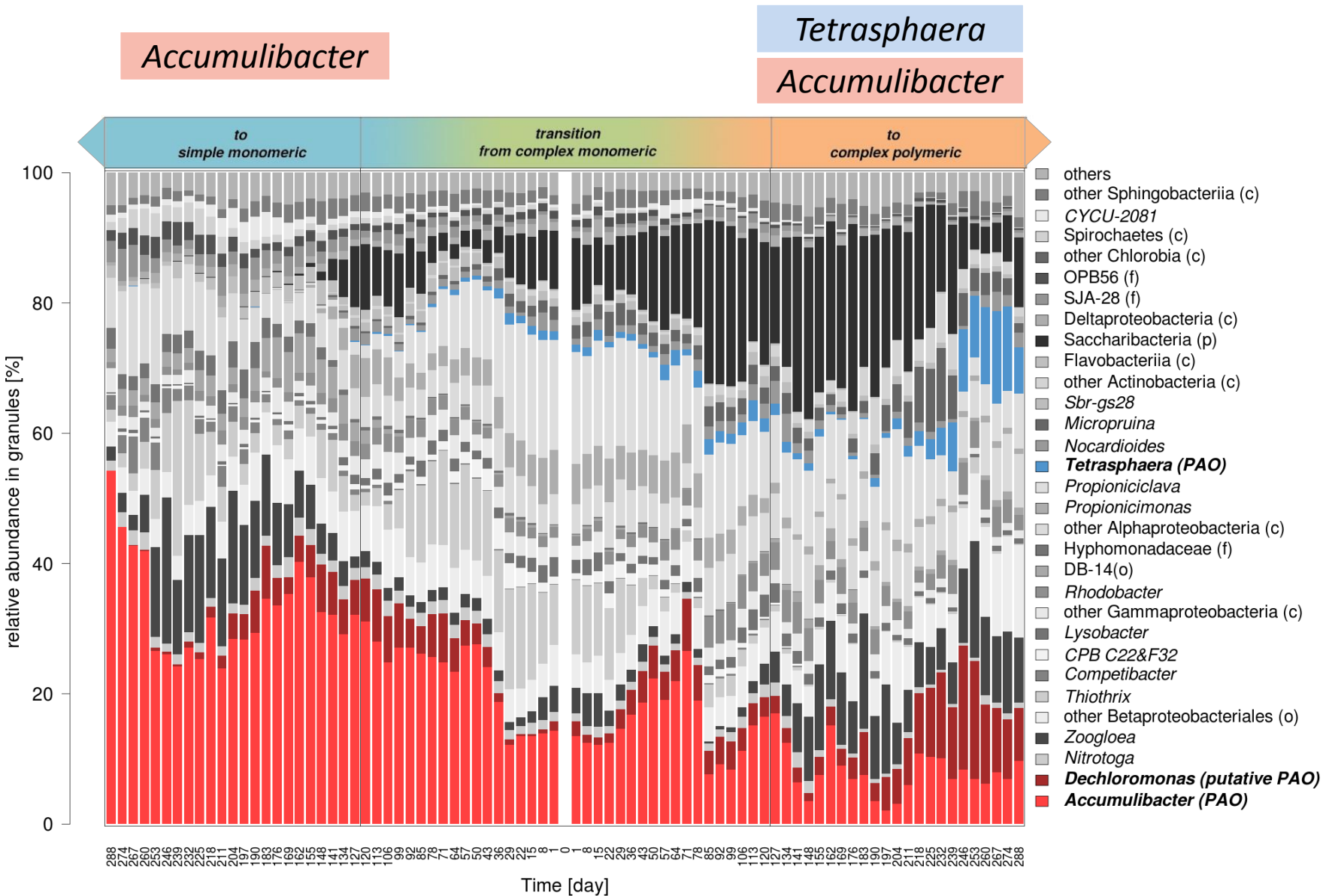
Introduction

Objectives

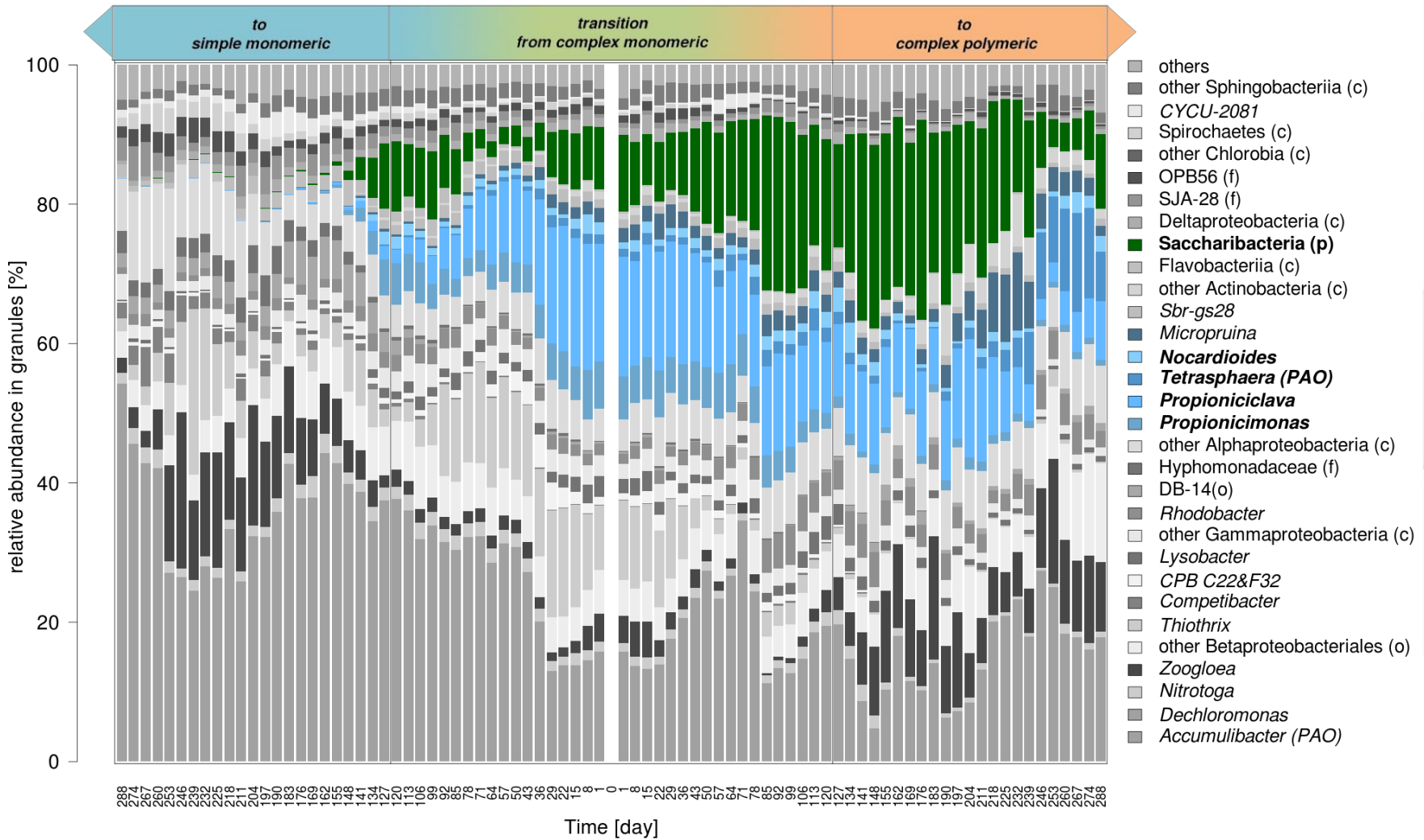
Method

Results

Conclusion

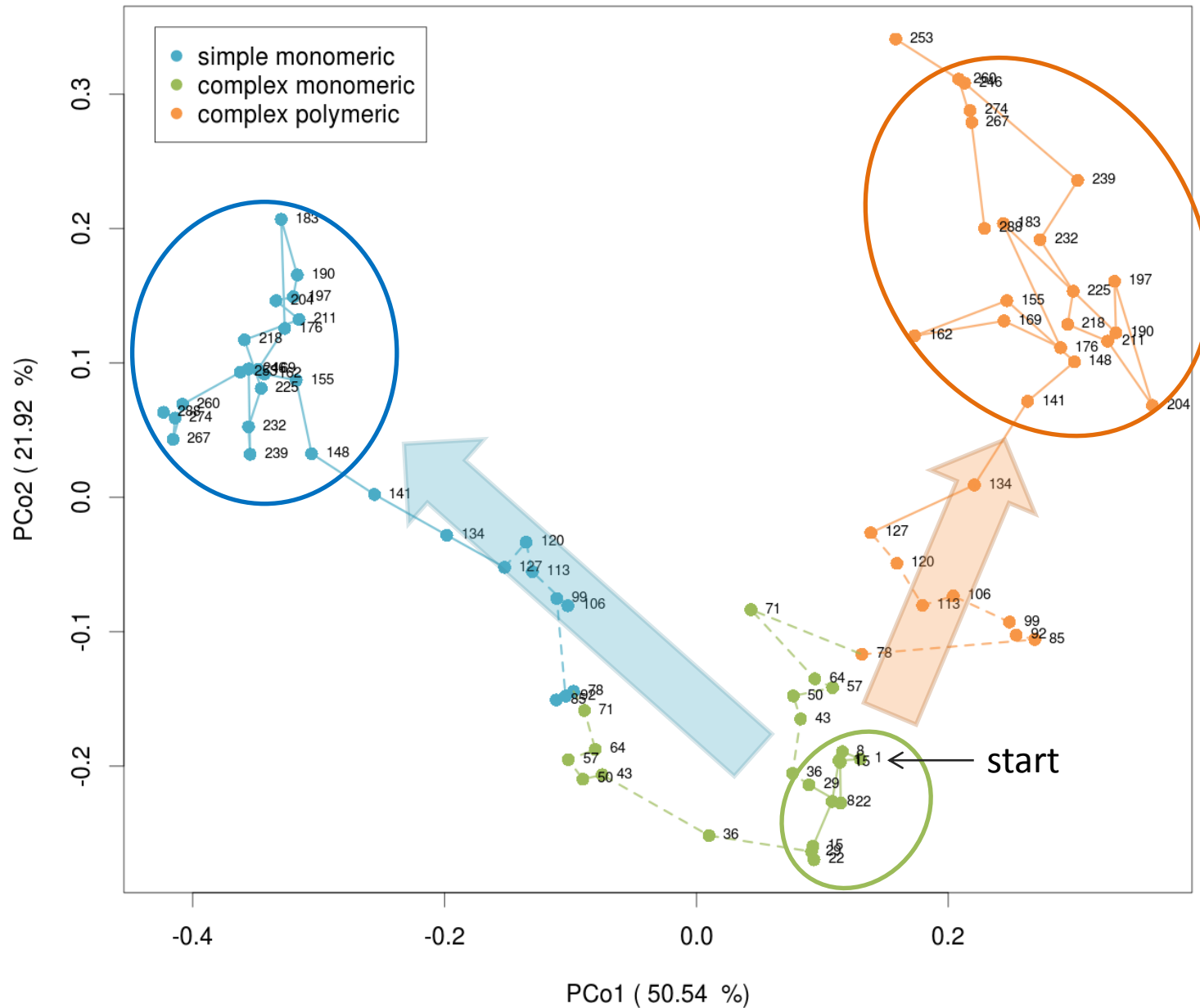


fermenting bacteria



- Introduction
- Objectives
- Method
- Results
- Conclusion

Overall evolution of the microbial communities



Conclusions

- The AGS settleability and the nutrient removal performances were good with the simple and the complex monomeric wastewaters.
- With the introduction of polymeric compounds, a significant proportion of flocs appeared and the N-removal decreased.
- Accumulibacter was the predominant PAO with the simple wastewater. With the fermentable and polymeric compounds, the guild of PAO was more diverse.

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Christof Holliger

DPE - Eawag

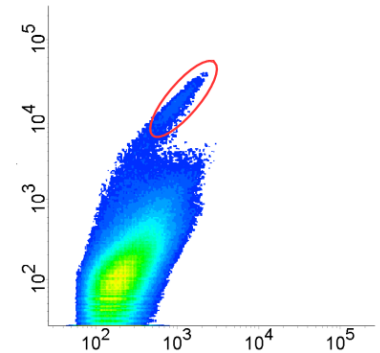
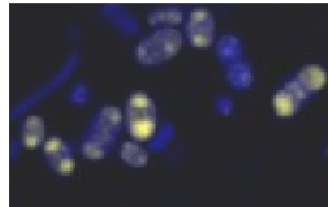
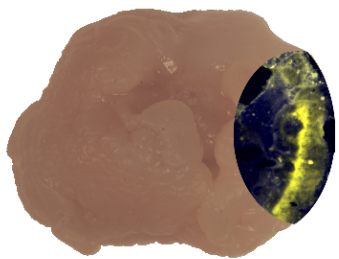
Manuel Layer

Nicolas Derlon

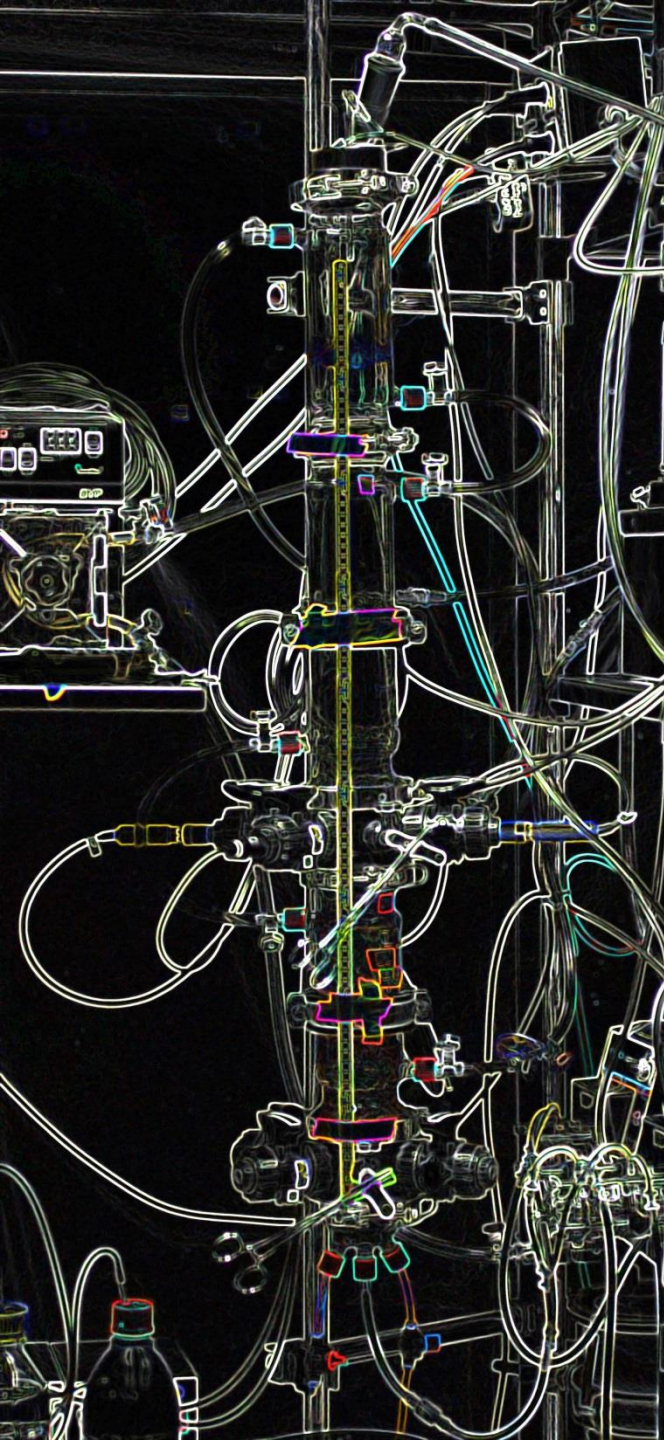
Eberhard Morgenroth

Arnaud Gelb

Identification of microorganisms in aerobic granular sludge actively involved in biological phosphorus removal



Poster n°51



Thank you for your
attention !

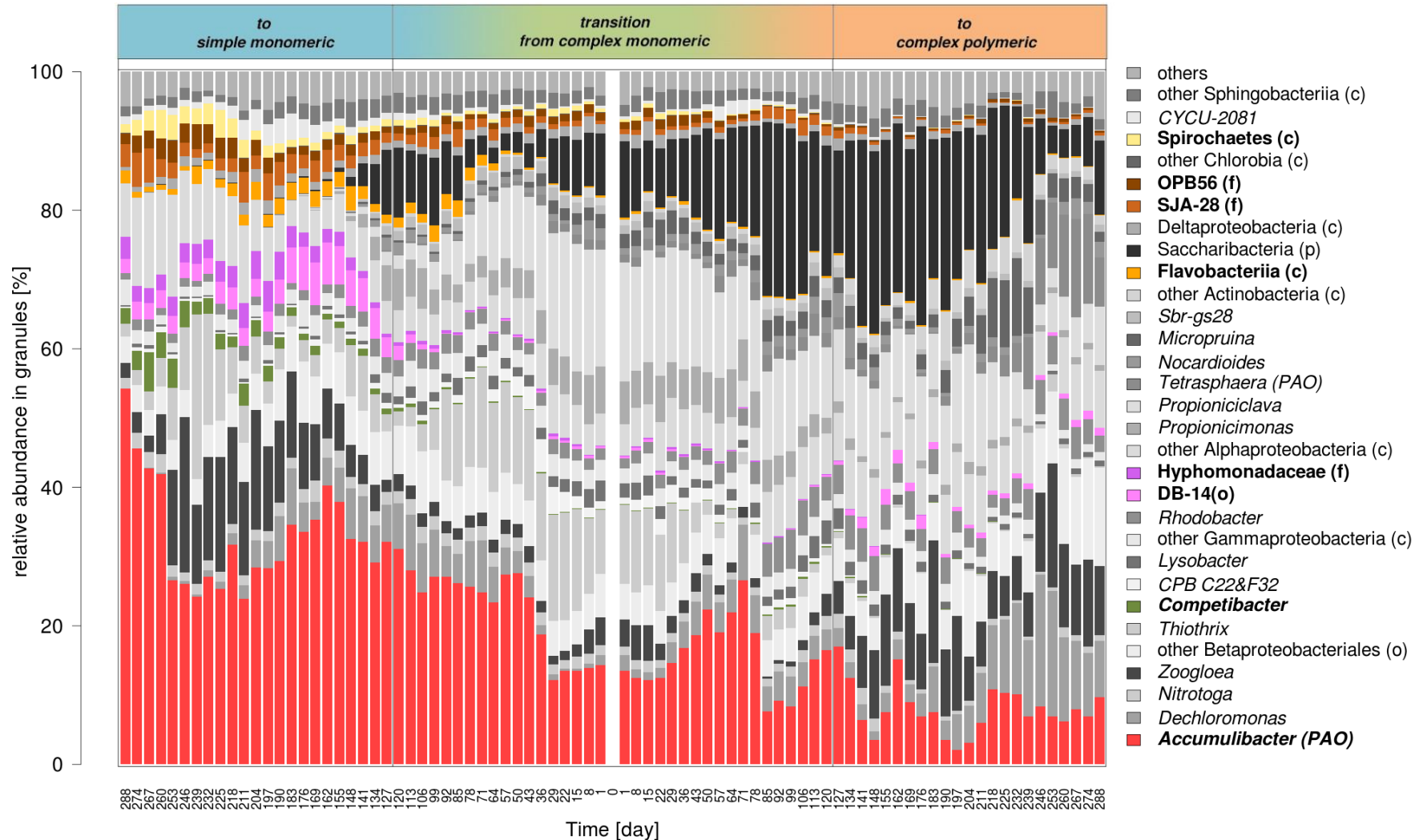
Synthetic wastewater compositions

<i>Medium</i>	<i>COD [mgO₂/L]</i>	<i>VFA</i>	<i>Glucose and amino acids</i>	<i>Starch and peptones</i>
Simple monomeric	450	100 %	-	-
Complex monomeric	600	33 %	66 %	-
Complex polymeric	600	33 %	33 %	33 %

Phosphorus : 22 [mg/L]

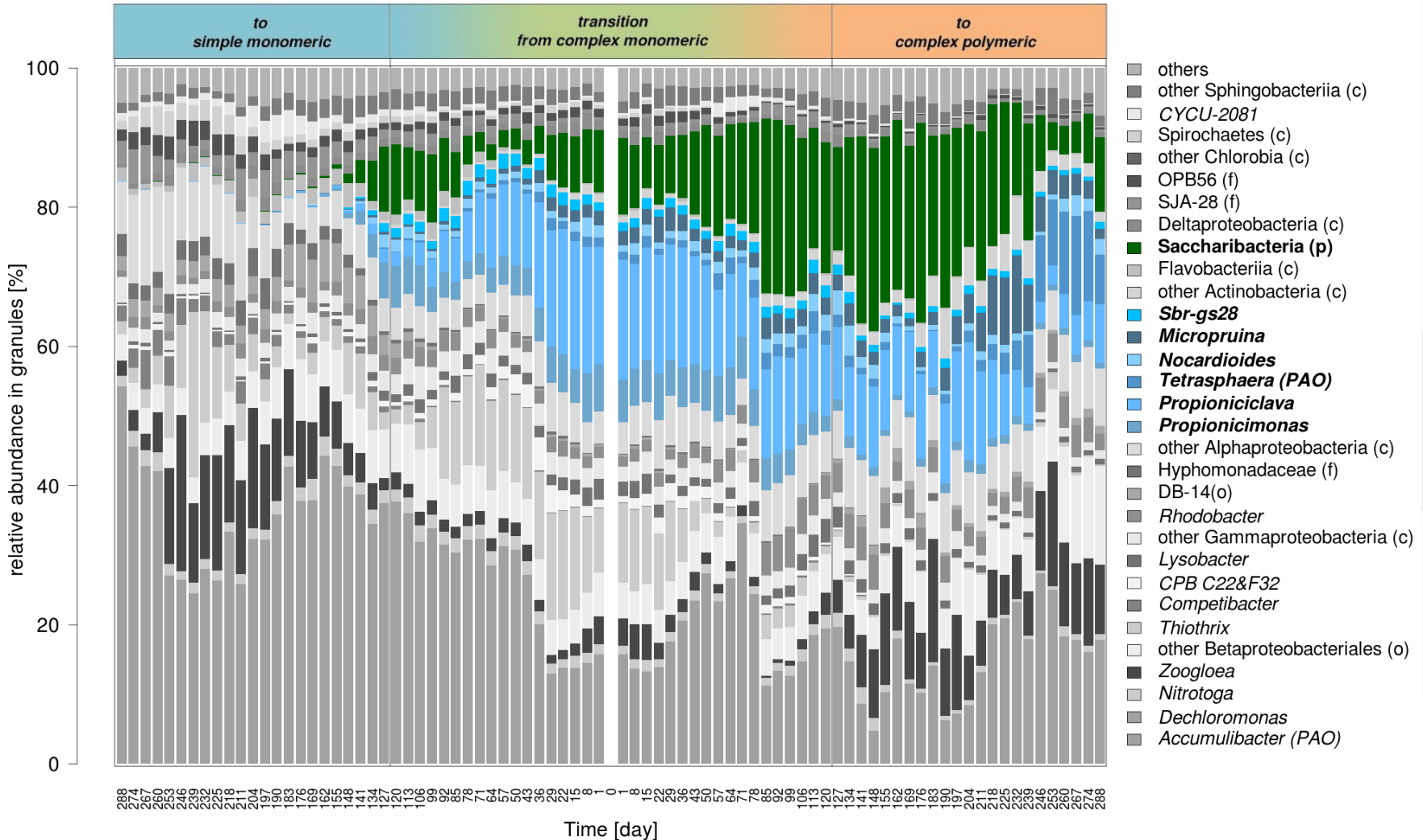
Nitrogen : 56 [mg/L]

Some taxa are found in higher abundance with the simple wastewater



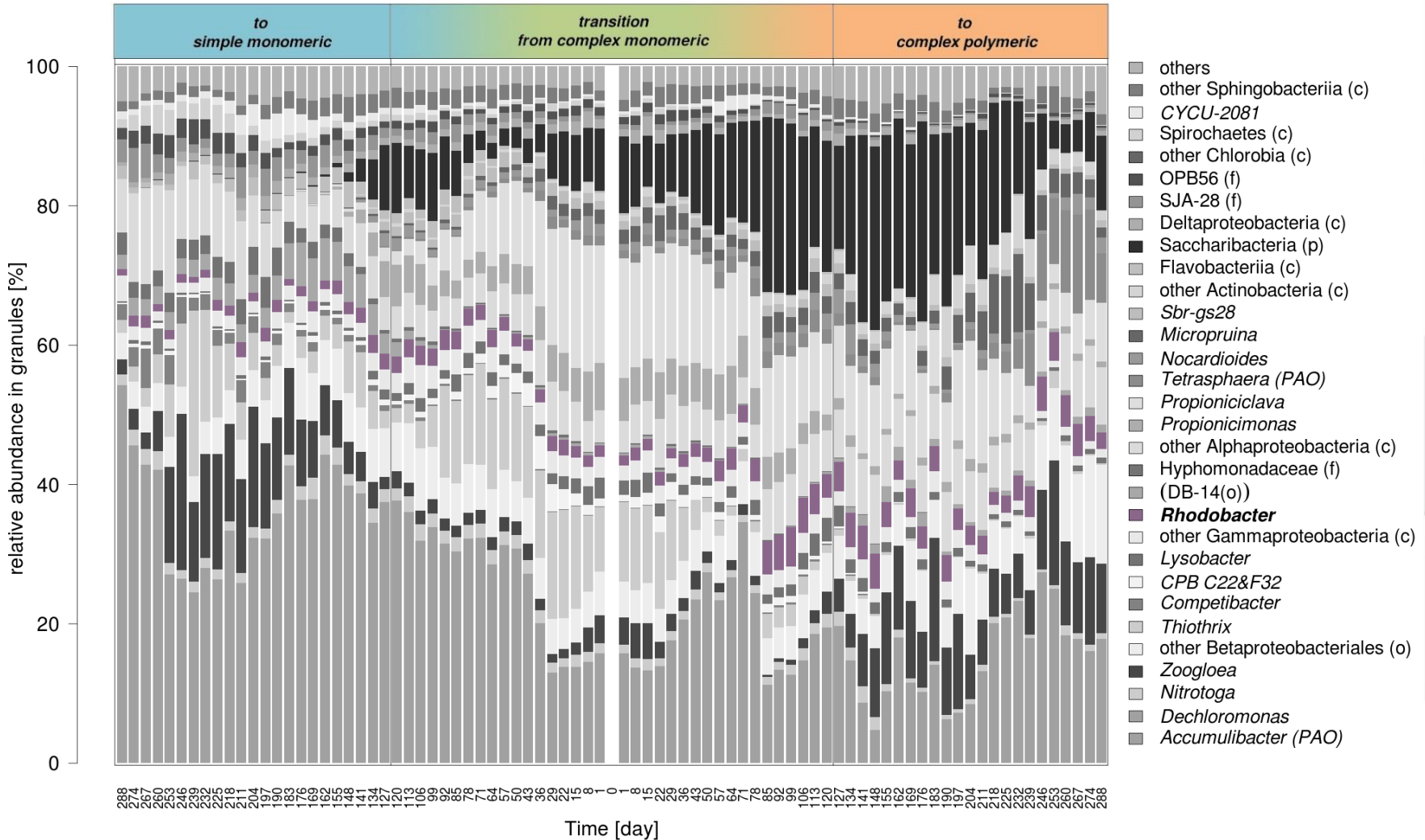
- Introduction
- Objectives
- Method
- Results
- Conclusion

Some taxa are found in higher abundance with the complex wastewater



Introduction
Objectives
Method
Results
Conclusion

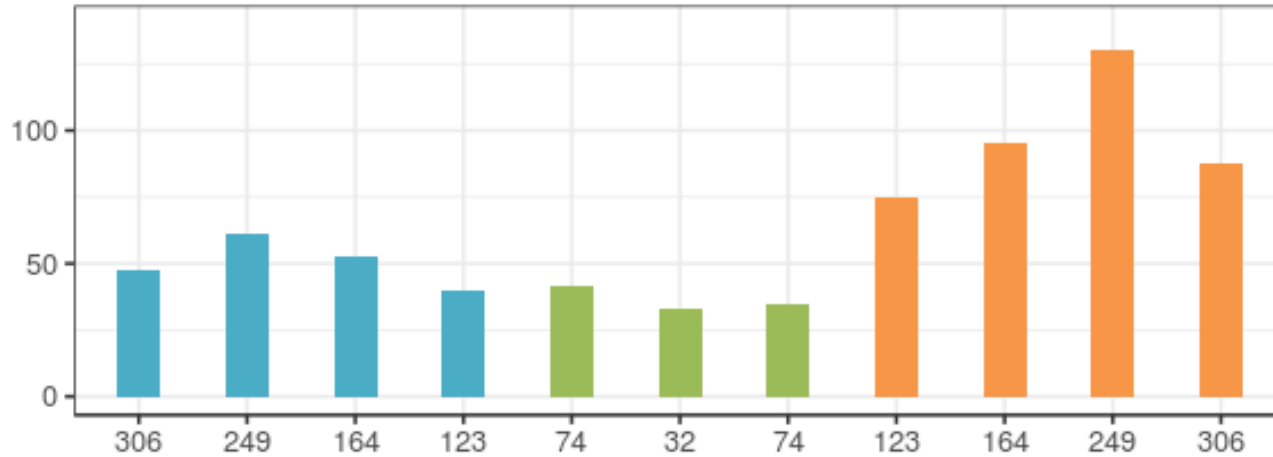
Rhodobacter was found in higher abundance with the polymeric wastewater



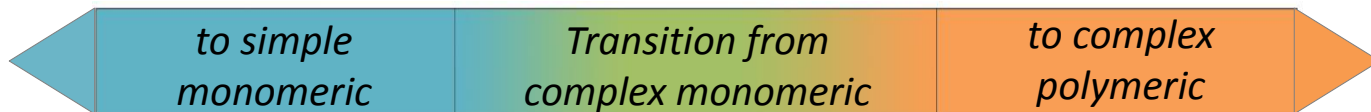
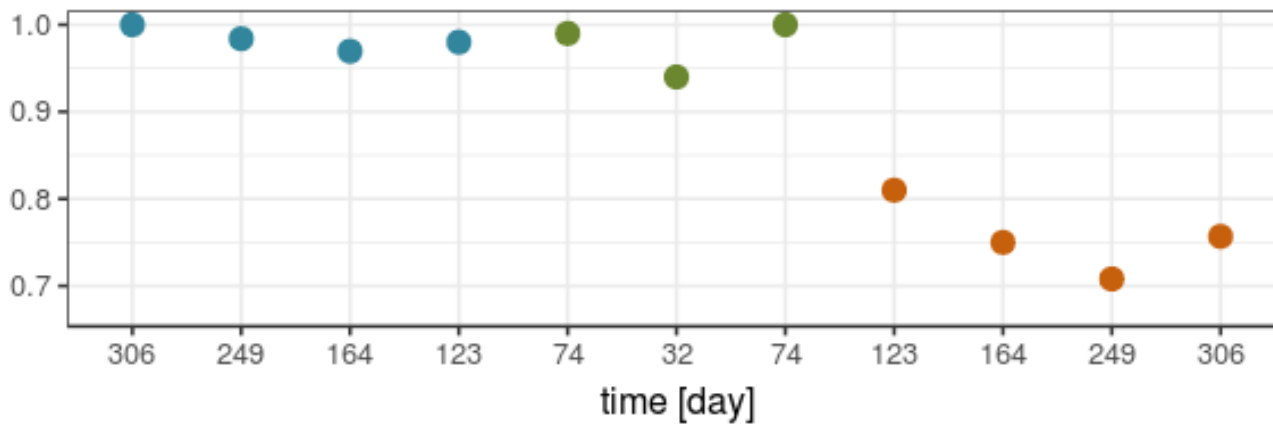
Introduction
Objectives
Method
Results
Conclusion

Sludge volume index (SVI)

SVI₁₀ [ml/g]



ratio SVI₁₀/SVI₃₀



Introduction

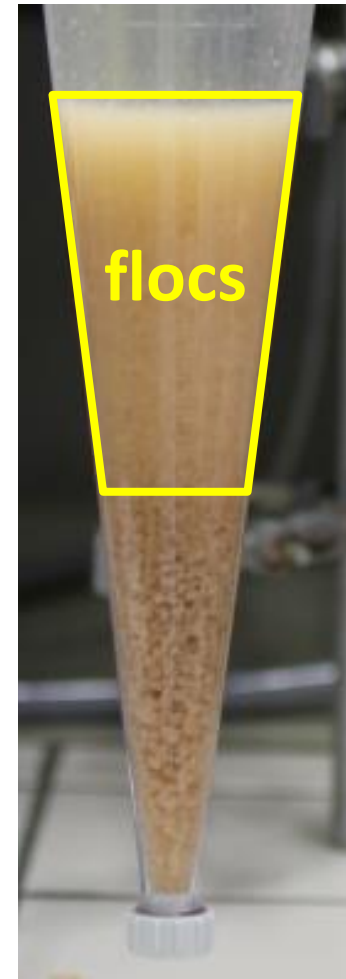
Objectives

Method

Results

Conclusion

Evolution of the settling capability of the AGS



Transition to **simple monomeric**

Complex monomeric

Transition to **complex polymeric**

Introduction

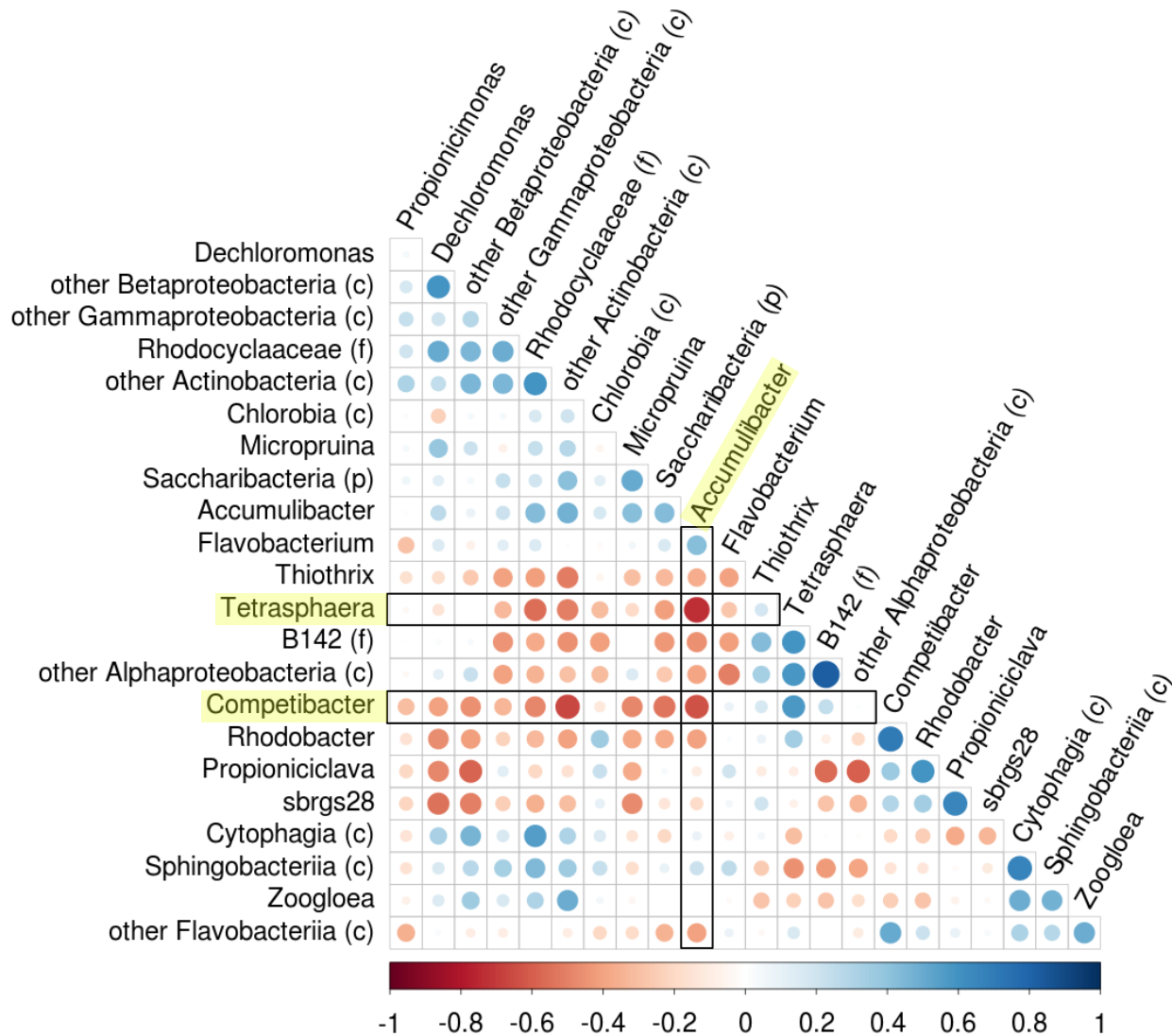
Objectives

Method

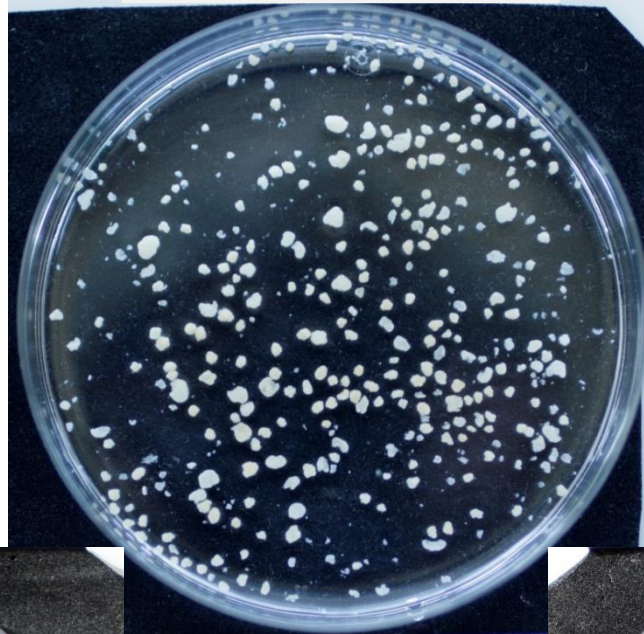
Results

Conclusion

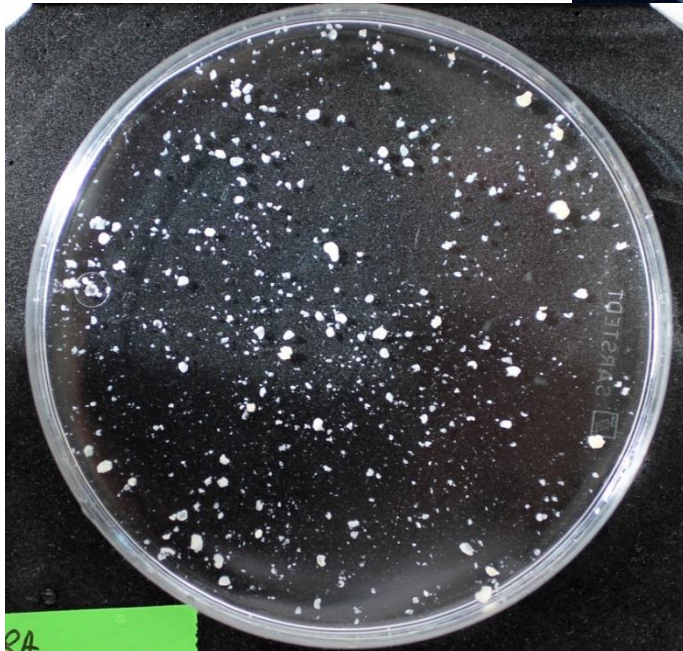
Correlation between the relative abundance of main taxa with complex monomeric wastewater



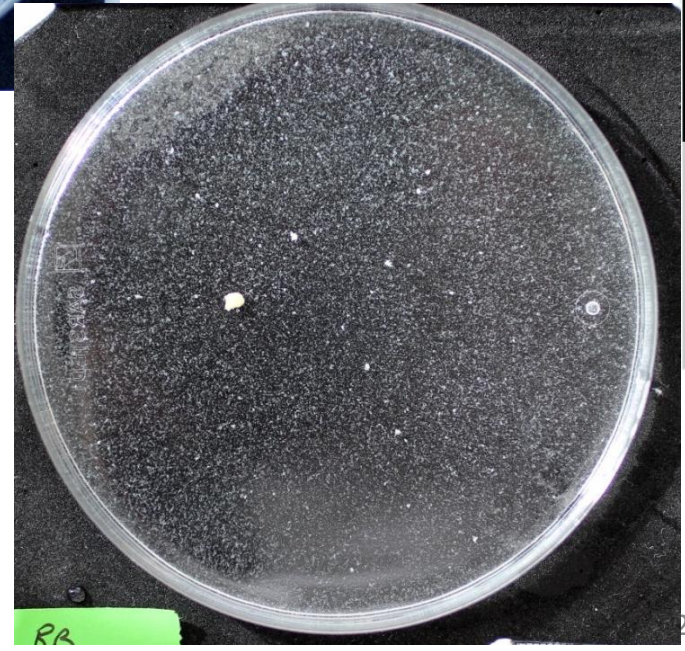
Complex
monomeric



Simple
monomeric



Complex
polymeric



Introduction

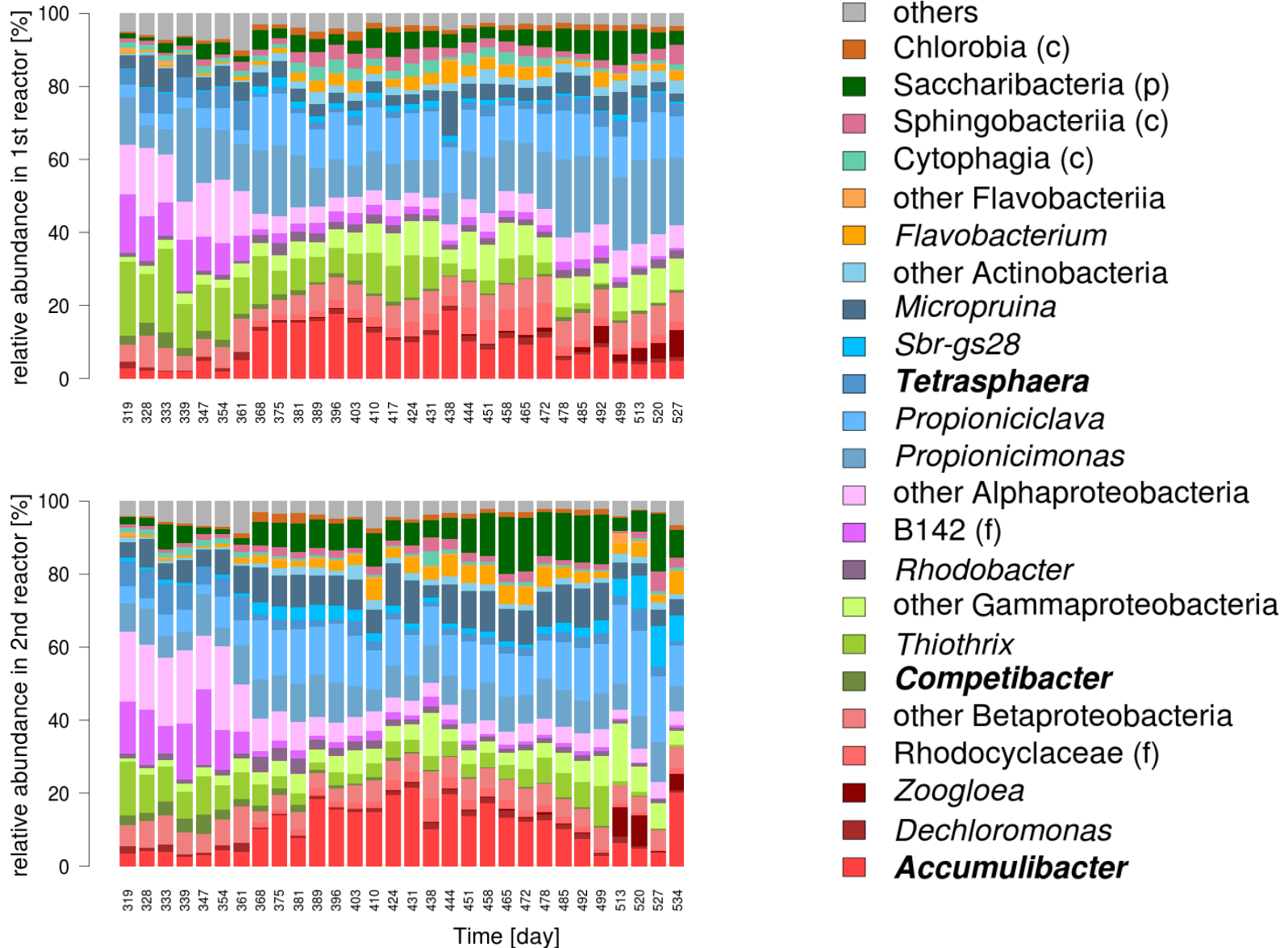
Objectives

Method

Results

Conclusion

Microbial communities evolution in two reactor fed with complex monomeric wastewater



- Introduction
- Objectives
- Method
- Results**
- Conclusion