

# Sticking together: Inter-species aggregation of bacteria isolated from iron snow is controlled by chemical signaling

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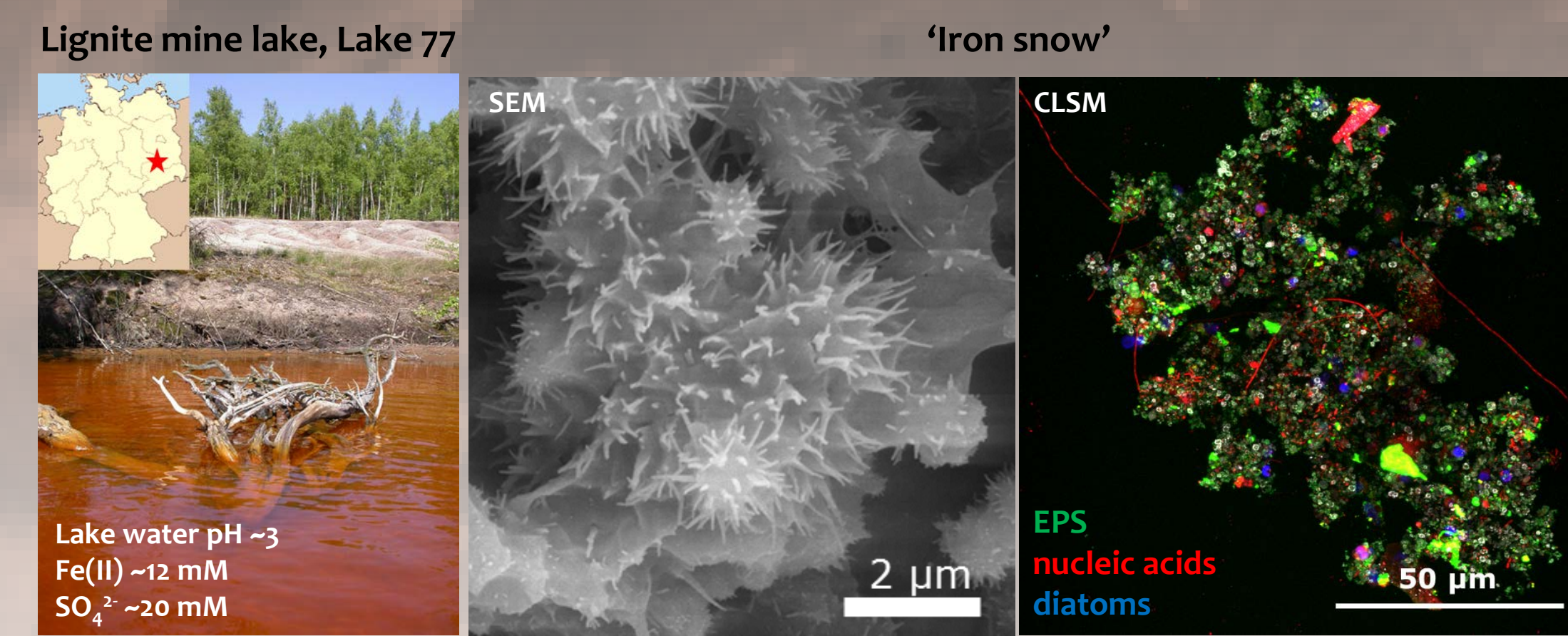
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## Background

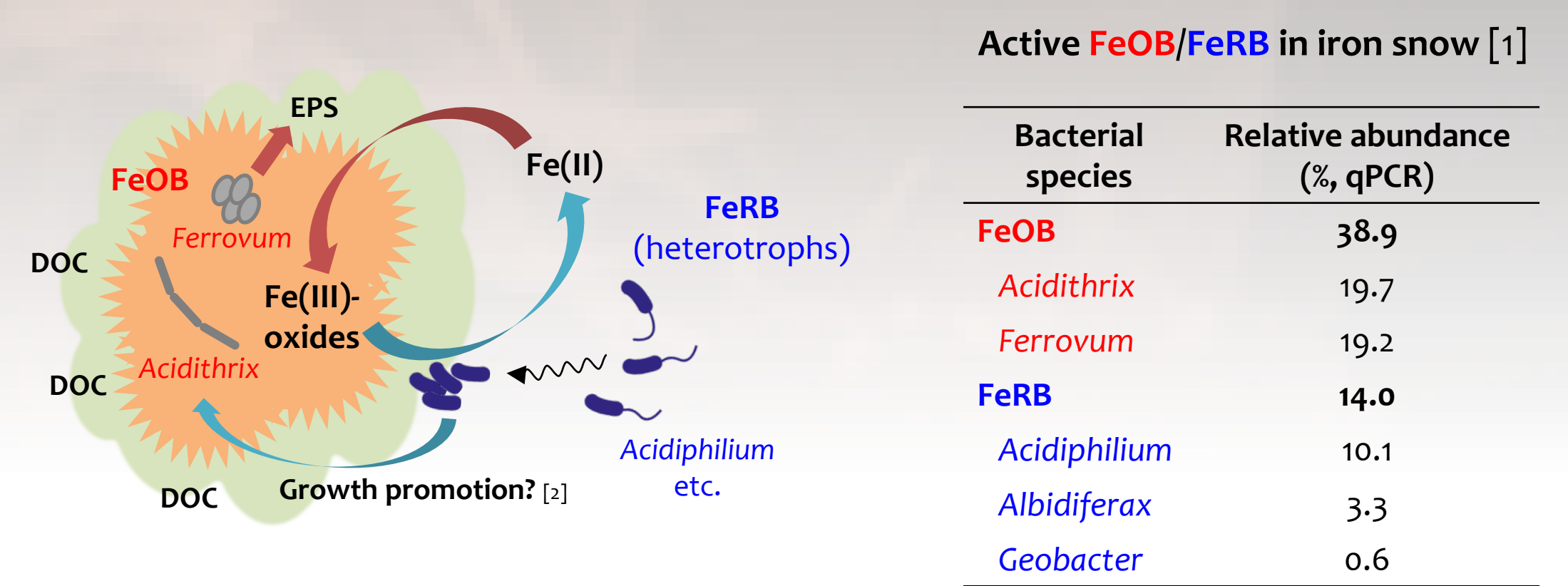
Mixed organic and inorganic pelagic aggregates (marine/lake snow) which form in the water column are considered local hot spots for microbial interactions. However, the exact mechanisms for their interactions is not well characterized due to the high microbial and chemical complexity.

Interestingly, Iron-rich pelagic aggregates (**'iron snow'**) found in acidic mine lake are formed and colonized by small dominant groups of Fe(II)-oxidizing bacteria (**FeOB**) and Fe(III)-reducing bacteria (**FeRB**), thus providing a simplified model to study mechanisms of interaction and communication among key bacterial players within pelagic aggregates.

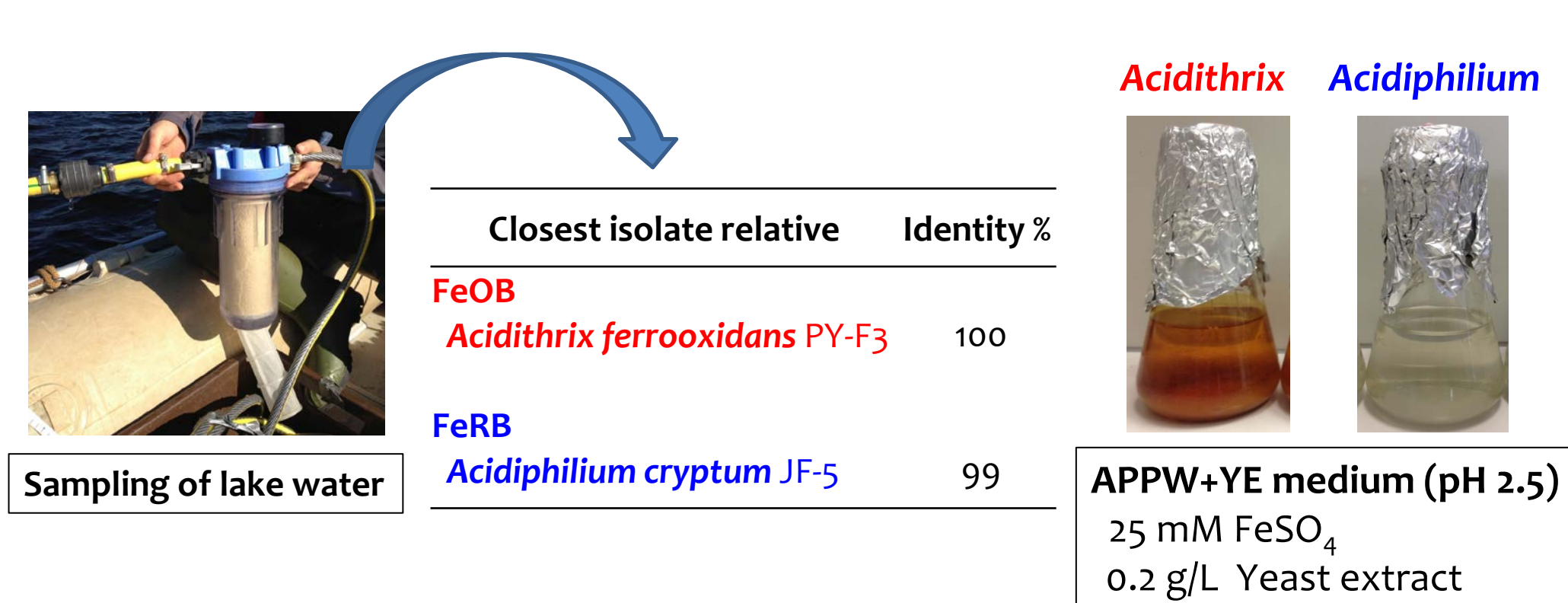


## Research objective

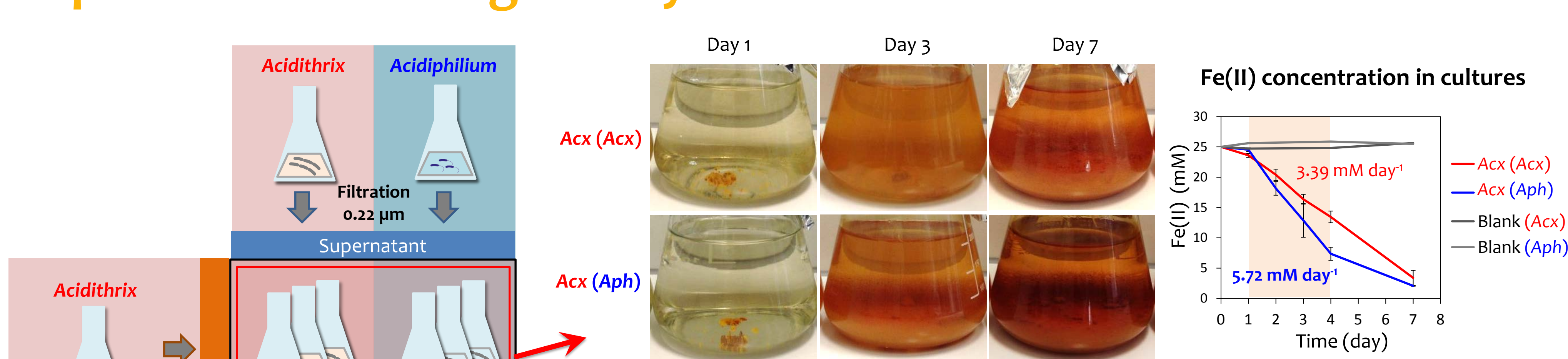
Detection and identification of **infochemicals** that signify potential chemical communication between **FeOB** & **FeRB** within the iron snow.



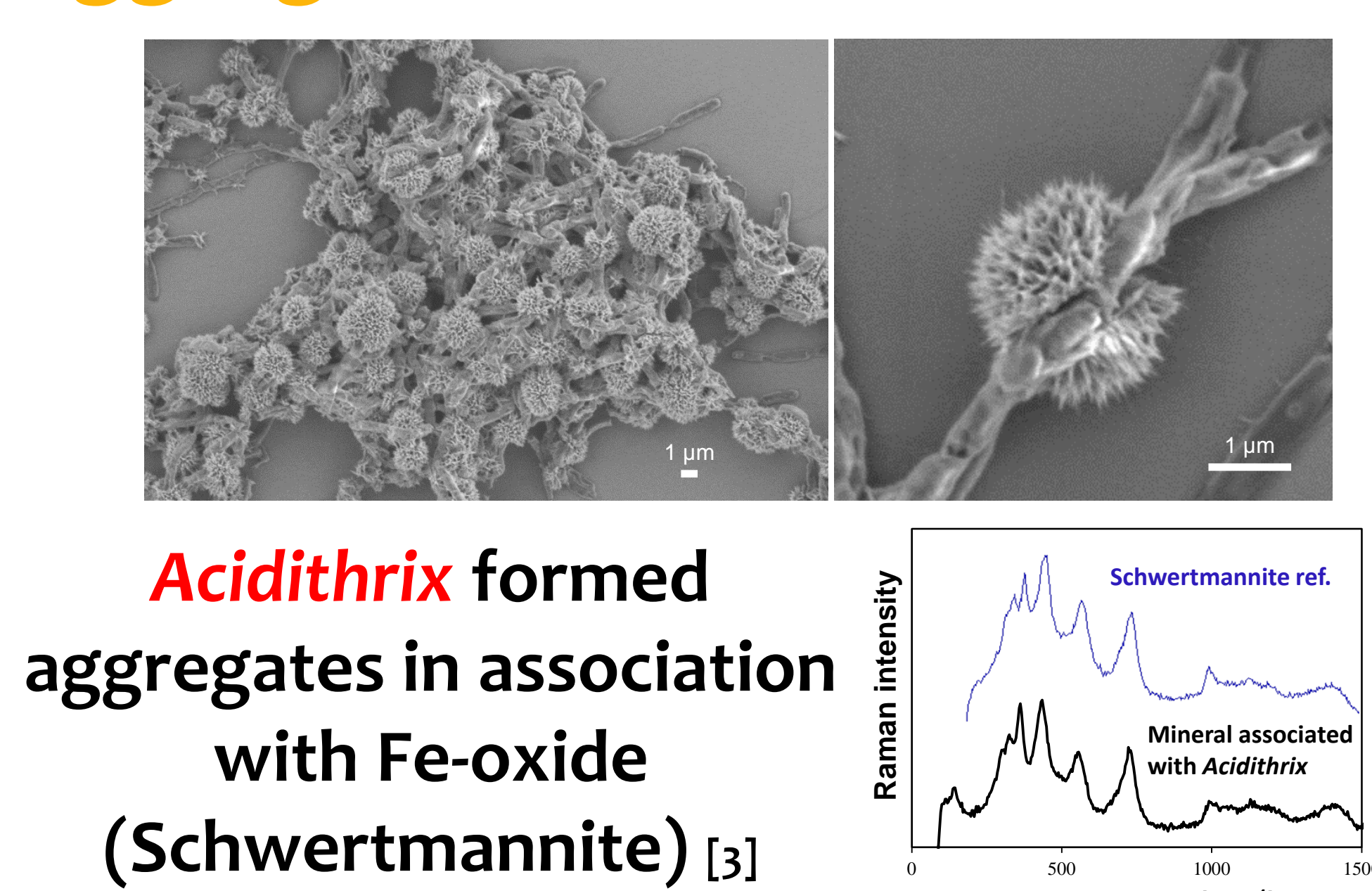
## Isolation of Bacteria



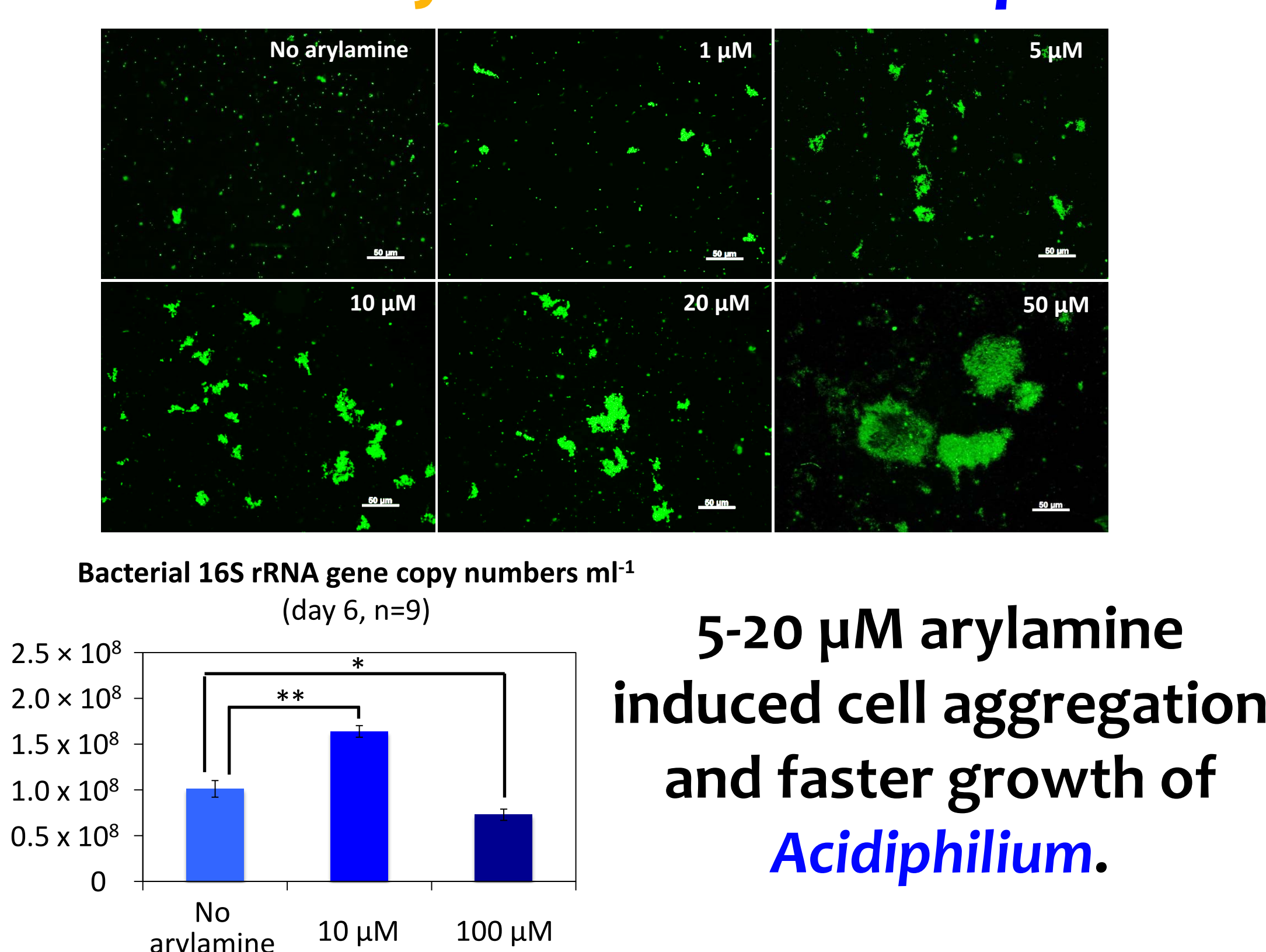
## Supernatant exchange assay



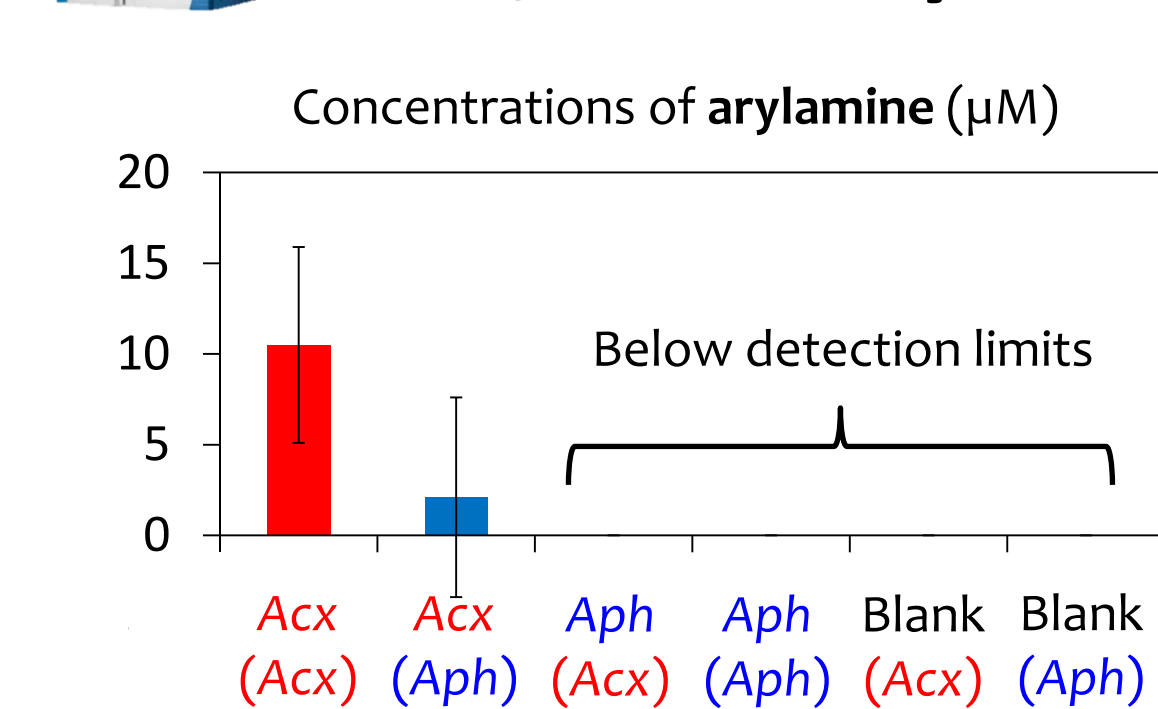
## Aggregation of *Acidithrix*



## Effect of arylamine on *Acidiphilium*

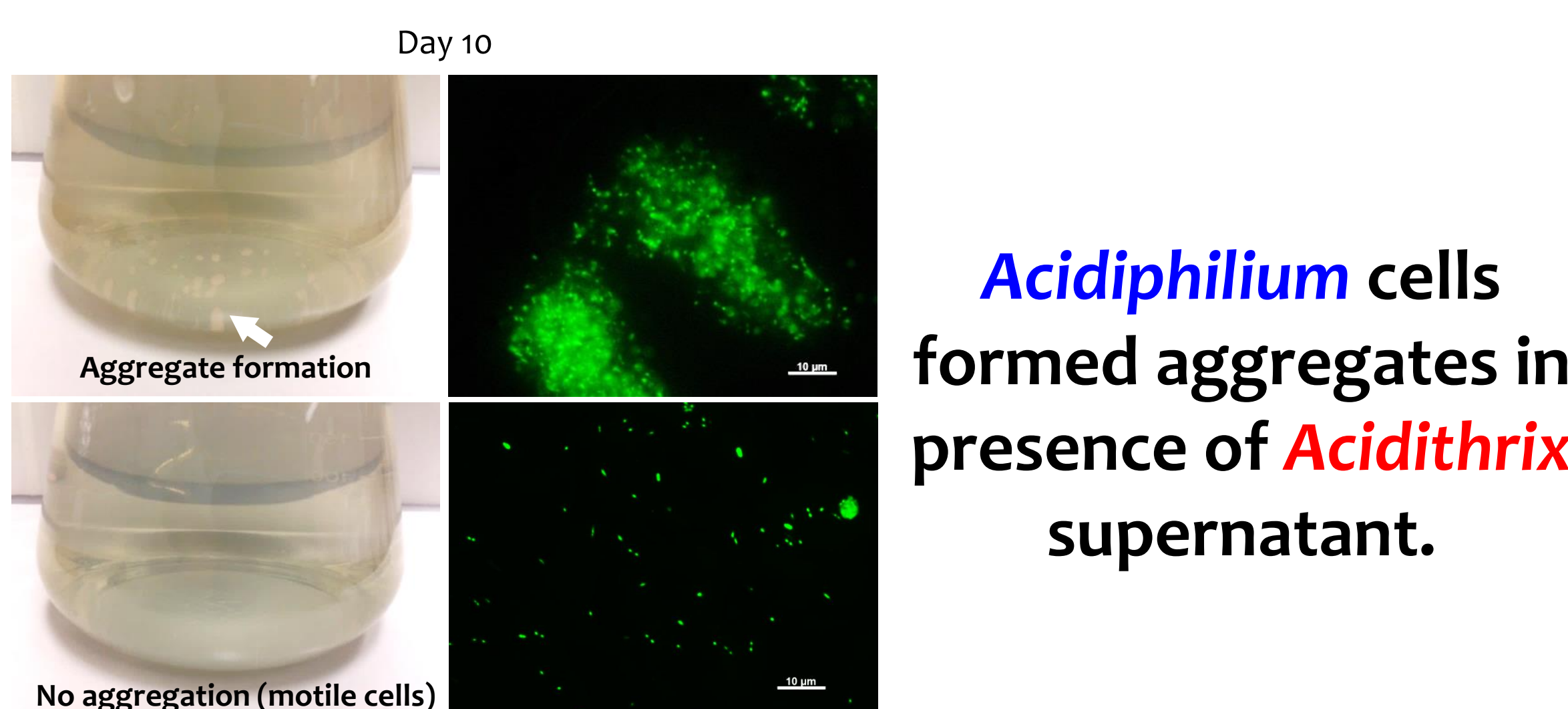


## GC/MS analysis

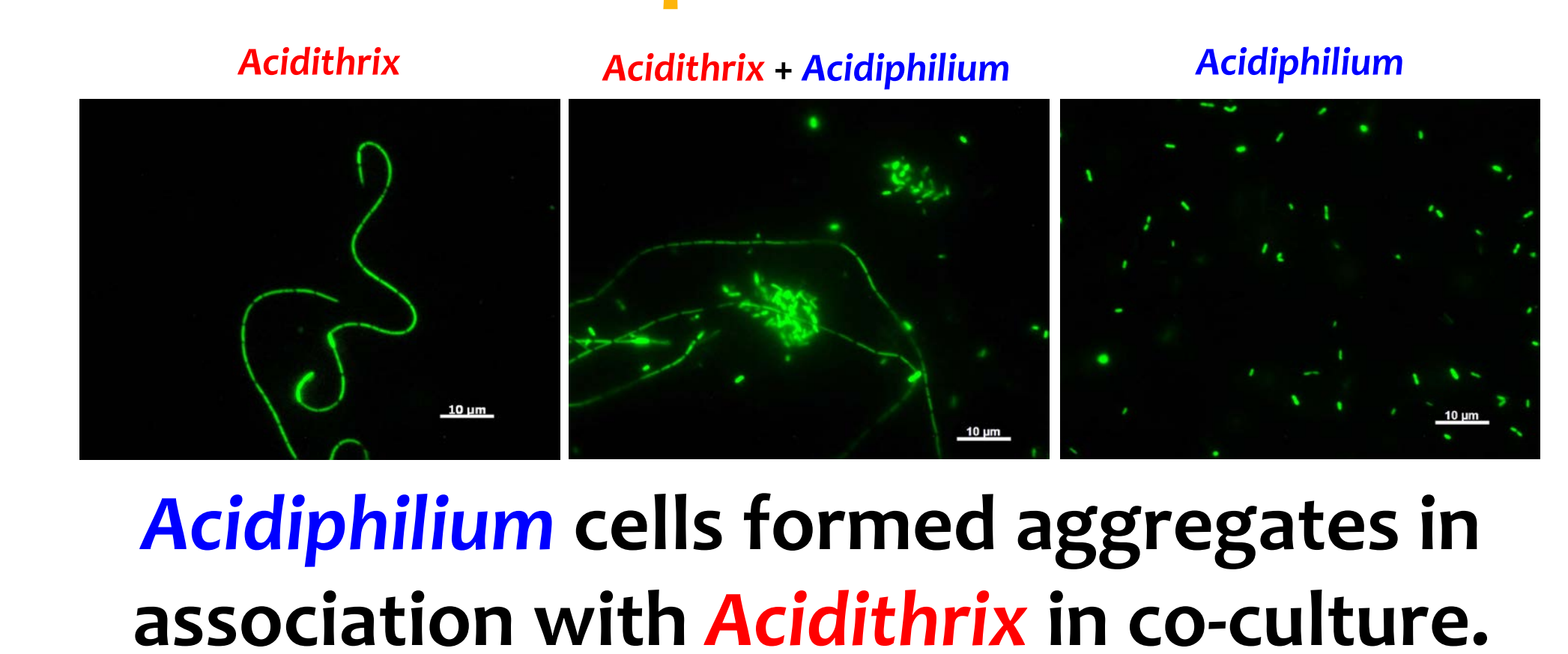


arylamine : a possible cell consolidation signal molecule for gram negative bacterium which suppress cell motility

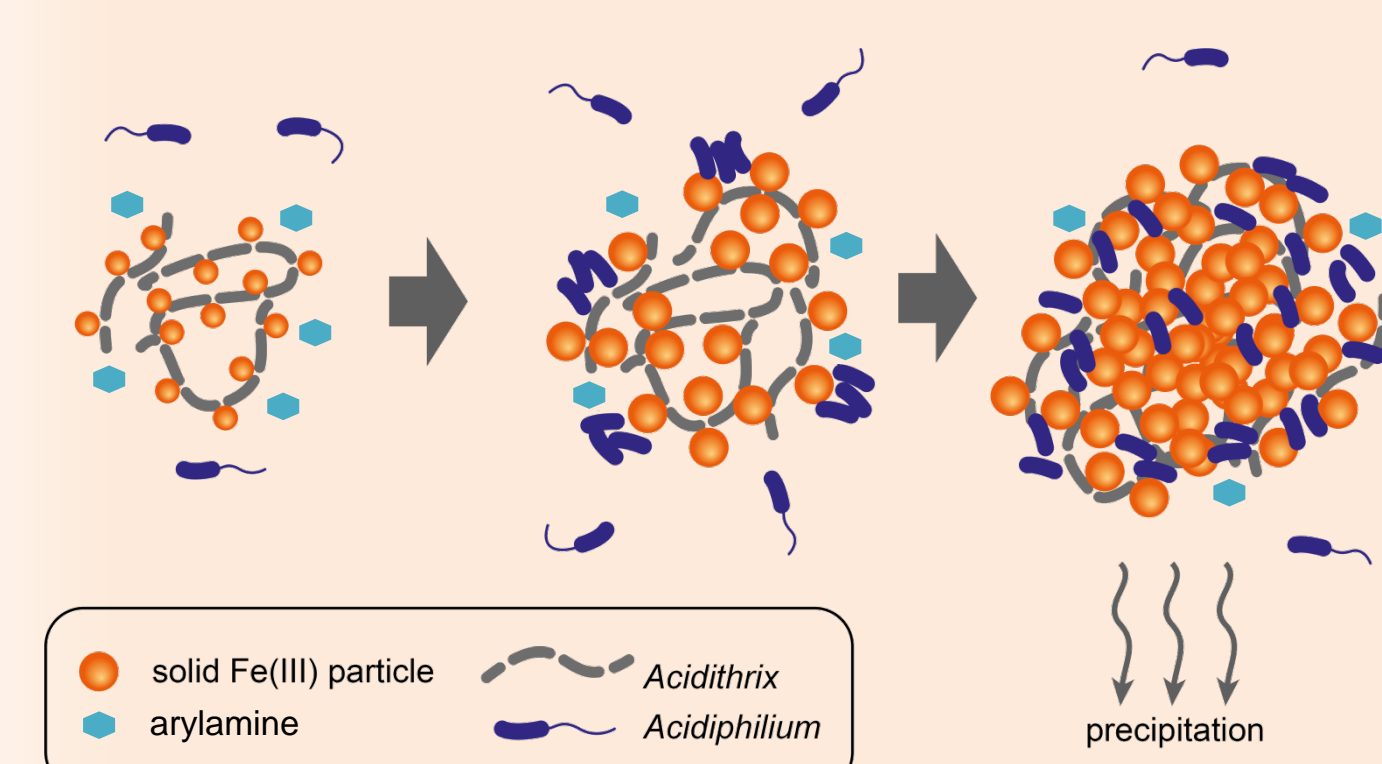
Rates of Fe-oxidation by **Acidithrix** were accelerated in presence of **Acidiphilium** supernatant.



## Co-culture experiment



## Summary



- **Acidithrix** (**FeOB**) rapidly forms iron-rich aggregates and plays a central role in iron snow formation.
- Metabolites of **Acidiphilium** (**FeRB**) accelerates growth and Fe(II) oxidation of **Acidithrix**.
- Not only the shuttling of resources, but an active control by chemical signals shapes the association of the iron snow consortia and their behavior.

## References

- [1] Lu, S. et al., *Applied and Environmental Microbiology* (2013)  
 [2] Liu, H. et al., *Archives of Microbiology* (2011)  
 [3] Mori, J. F. et al., *Microbiology* (2016)

## Acknowledgements

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