



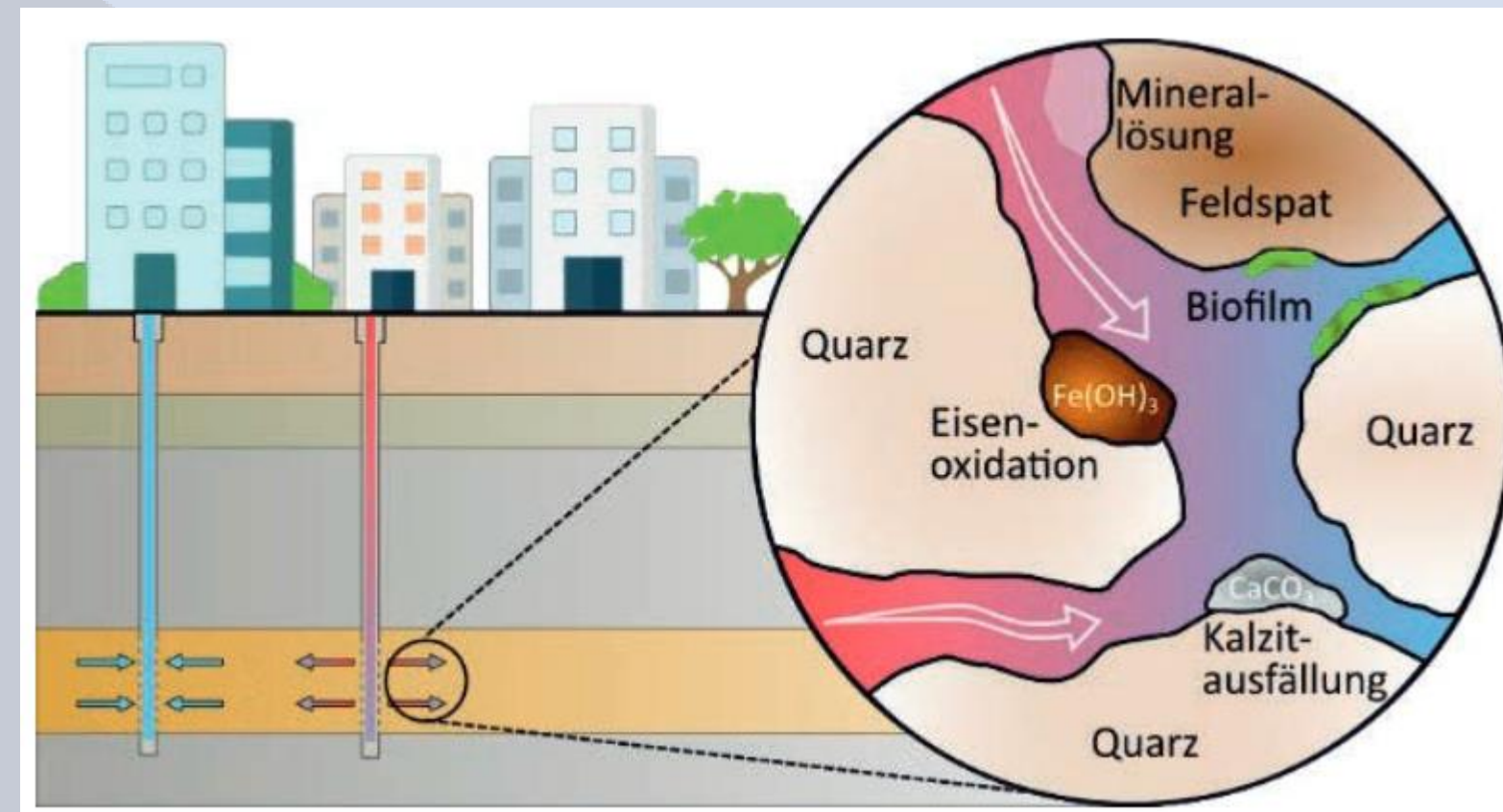
# Microbial diversity in a saline siliciclastic aquifer at the ATEs exploration site Berlin-Adlershof

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## Motivation

- **Aquifer Thermal Energy Storages (ATES)** can store excess energy (heat) and provide it when required
- **microbial activity** can impair the efficiency and integrity of ATEs facilities in different ways
- **crucial to identify microbial key players and monitor their activity**



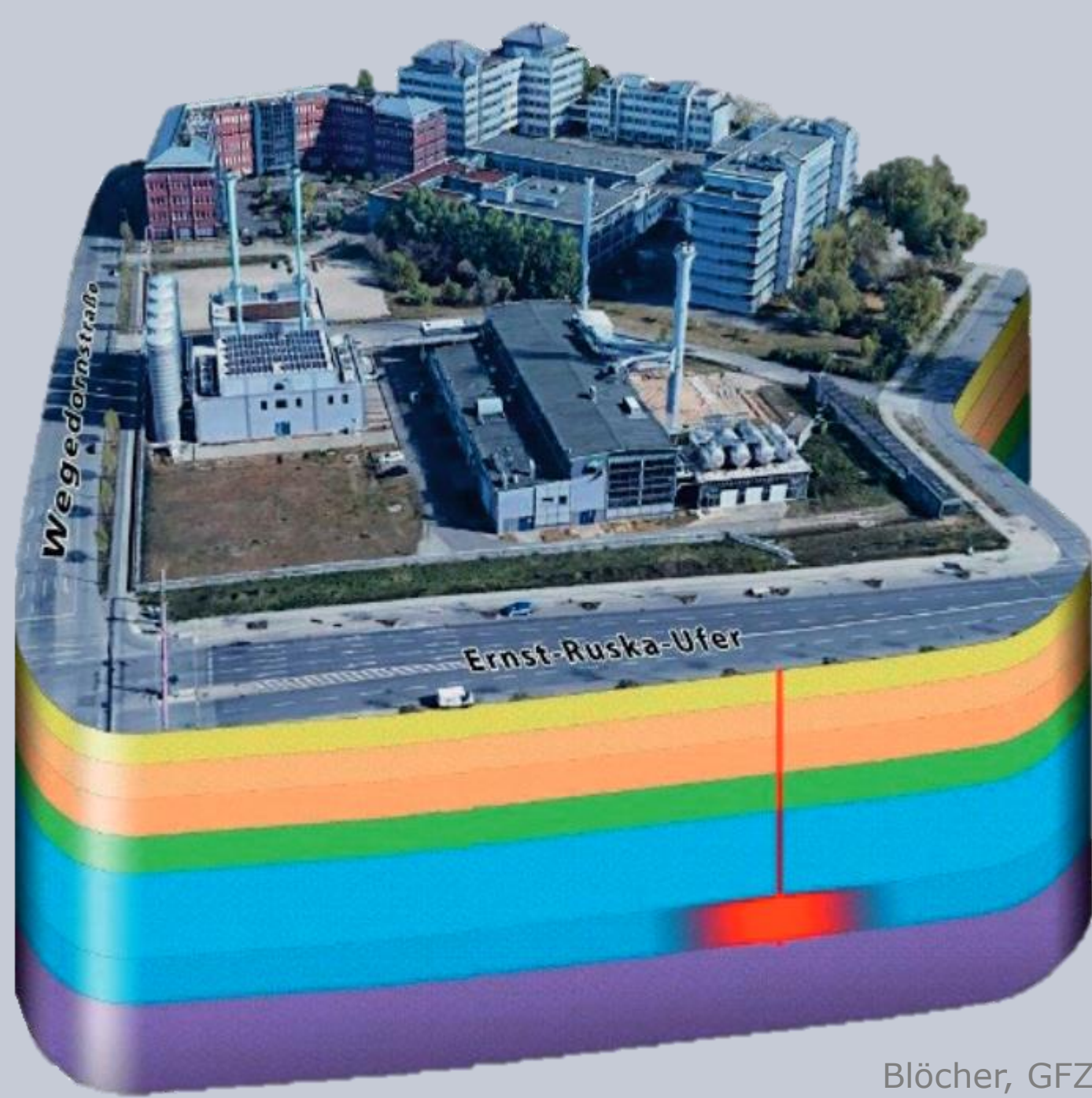
## Objectives

- **explore microbial community** composition, behaviour and potential functions in a siliciclastic aquifer at the **ATES exploration site** over 2 years after drilling

## Take home

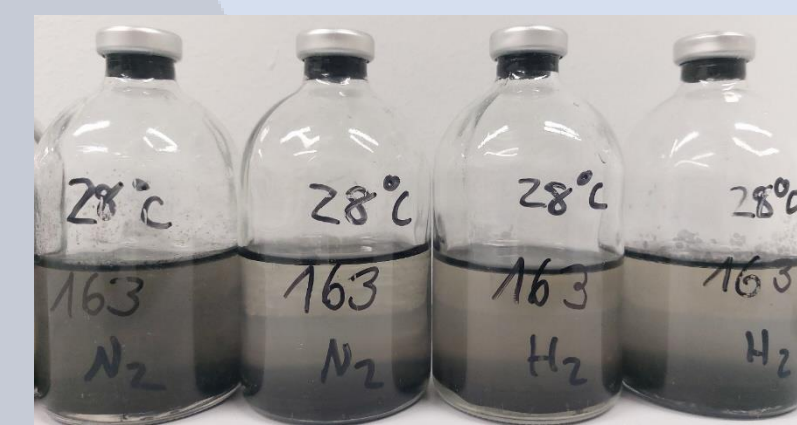
- **Aquifer microbial community** is characterized by **syntrophic** relationships between **fermentative** and acetogenic bacteria with **sulfate reducing bacteria (SRB)** and **methanogens**
- Community undergoes **succession over time** – fermentation → sulfate reduction → methanogenesis
- **Higher temperature** selects for **SRB, in situ temp.** for **acetogens**
- **Potential key consequences on ATEs:**
  - H<sub>2</sub>S formation → **corrosion** of metal components
  - Iron sulfide scales → pipe **clogging**
  - CH<sub>4</sub> formation → **gas buildup**/pressure changes/safety hazard
  - Biofilm formation → permeability/**efficiency reduction**

## Study Site & Methodology

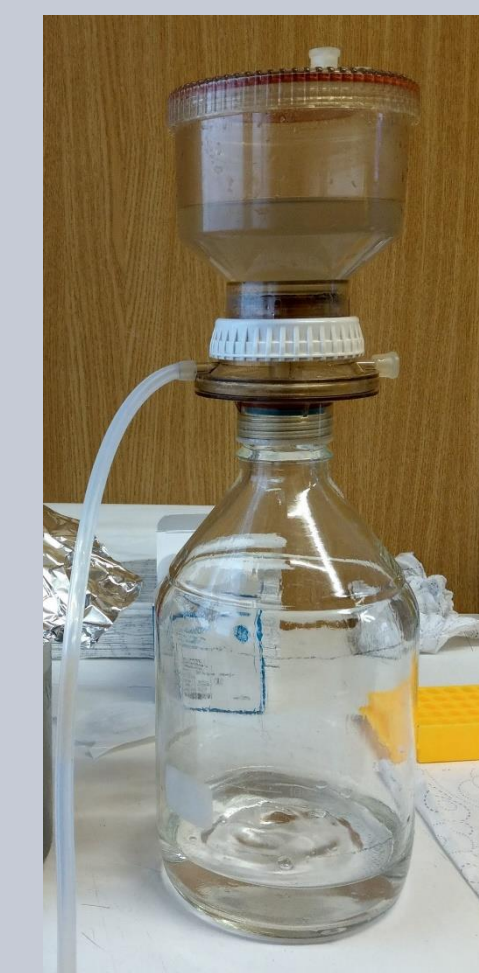


- Water sampling**
- 2 months
  - 14 months
  - 24 months
  - 28 months after drilling

Origin of groundwater at ~225 mbs



**Enrichment for sulfate reducers (SRB):** +/- NaCl, N<sub>2</sub> or H<sub>2</sub> at 16/28/55 °C



**Microbial community analysis based on amplicon sequencing of 16S rRNA gene**

## Hydrochemistry

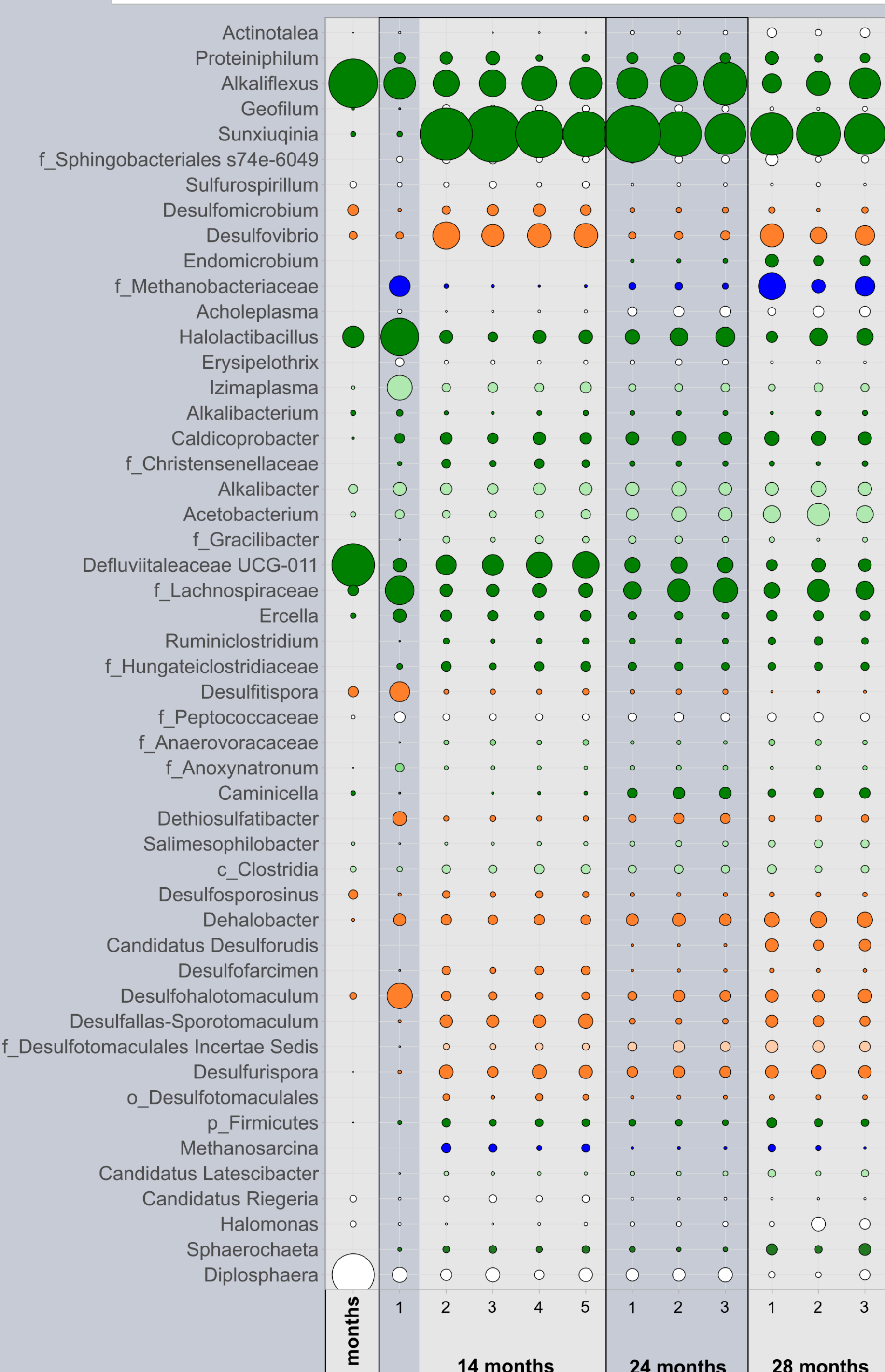
**Table 1: Groundwater characteristics.** Saline water (~2%) is dominated by Na and Cl. Sulfate can act as electron acceptor for anaerobic respiration by SRB. The presence of organic acids indicates fermentative microbial metabolism. \*Standing water in the annulus of the production well as indicated by higher pH, lower temperature and different microbial community (see Fig. 1 & 2A).

sample no.	pH	EC [mS cm <sup>-1</sup> ]	Temp [°C]	[mg L <sup>-1</sup> ]							Lactic acid			
				Na	K	Mg	Ca	F	Cl	SO <sub>4</sub>	acid	Acetate	Propionate	Valerate
2 months 1	n.m.	31.9	15.9	6970	220	146	166	1	11688	270	n.a.	0.4	n.a.	n.a.
14 months 1*	9.5	26.4	11.8	6615	259	109	71	1	11163	270	<1	1.8	n.a.	<1
2-5	7.7	31.9	13.6	6656	283	130	139	1	11060	250	3.6	4.0	0.6	0.9
24 months 1-3	7.6	31.4	14.1	5401	110	60	54	1	10457	235	0.3	<0.1	<0.1	n.a.
28 months 1-3	7.7	31.7	15.3	6843	152	151	190	n.a.	10770	225	<0.1	0.8	0.1	n.a.

**Table 2: Dissolved gas composition** – dominated by H<sub>2</sub> 2 months after drilling, but N<sub>2</sub> 14 months after drilling.

	[%] H <sub>2</sub>	N <sub>2</sub>	CO <sub>2</sub>	Ar	O <sub>2</sub>	He	CH <sub>4</sub>
2 months	91.6	7.0	1.3	n.a.	n.a.	n.a.	n.a.
14 months	0.1	95.8	0.5	2.6	0.7	0.0	0.2

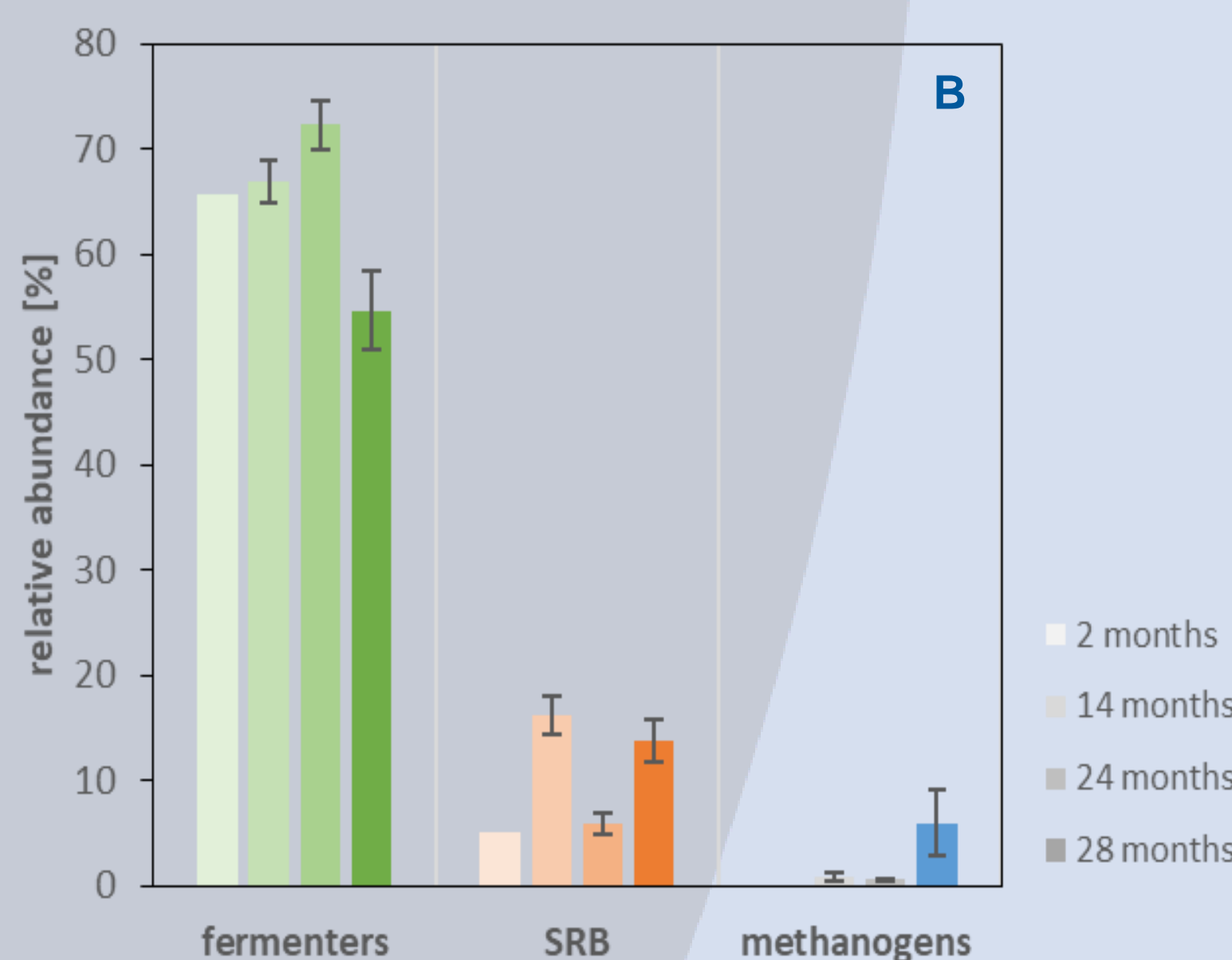
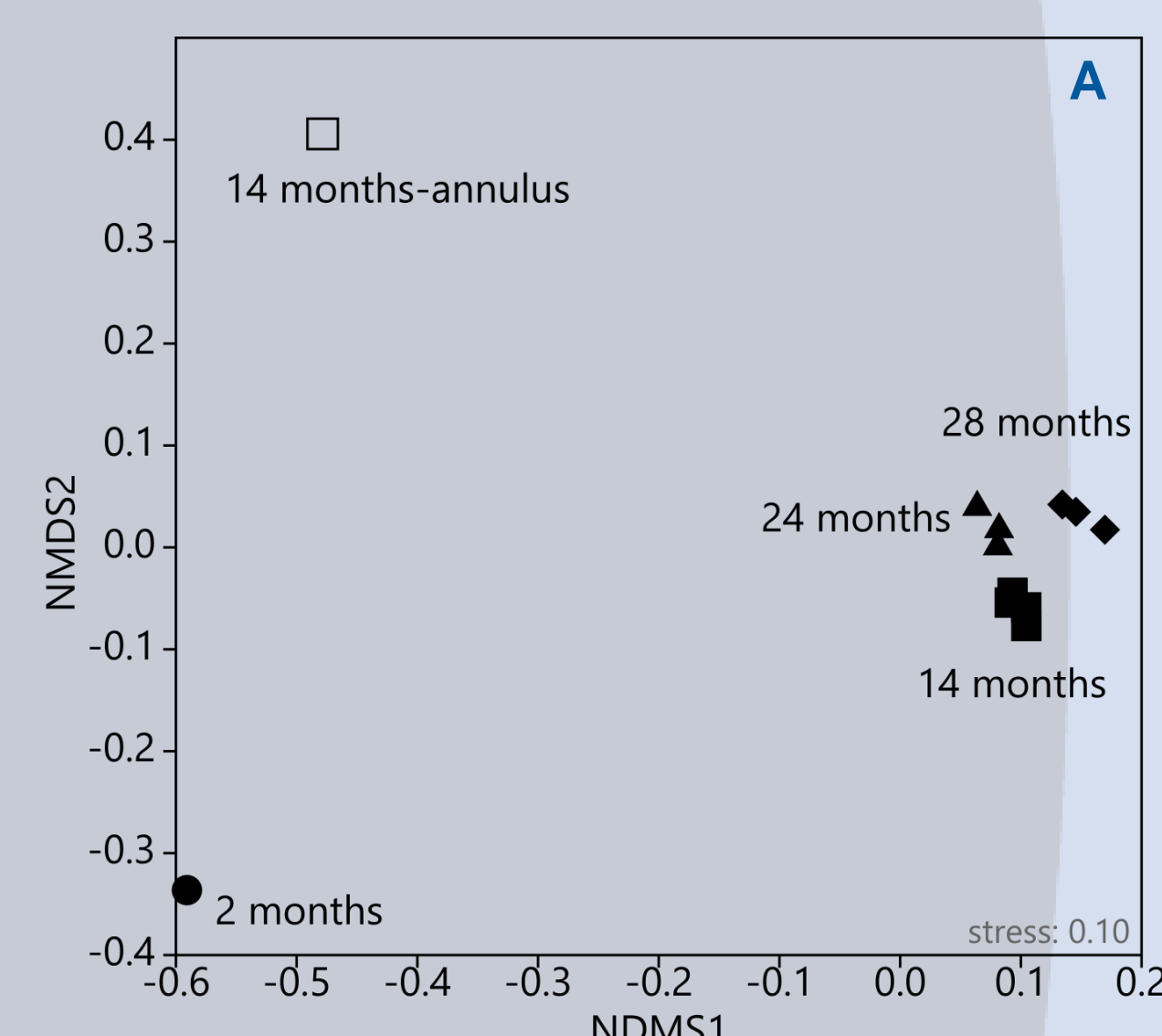
## Aquifer Microbial Community Composition and Dynamics



**Fig. 1: Top 50 most abundant genera and their classification** into fermentative, sulfate reducing and methane producing microorganisms.

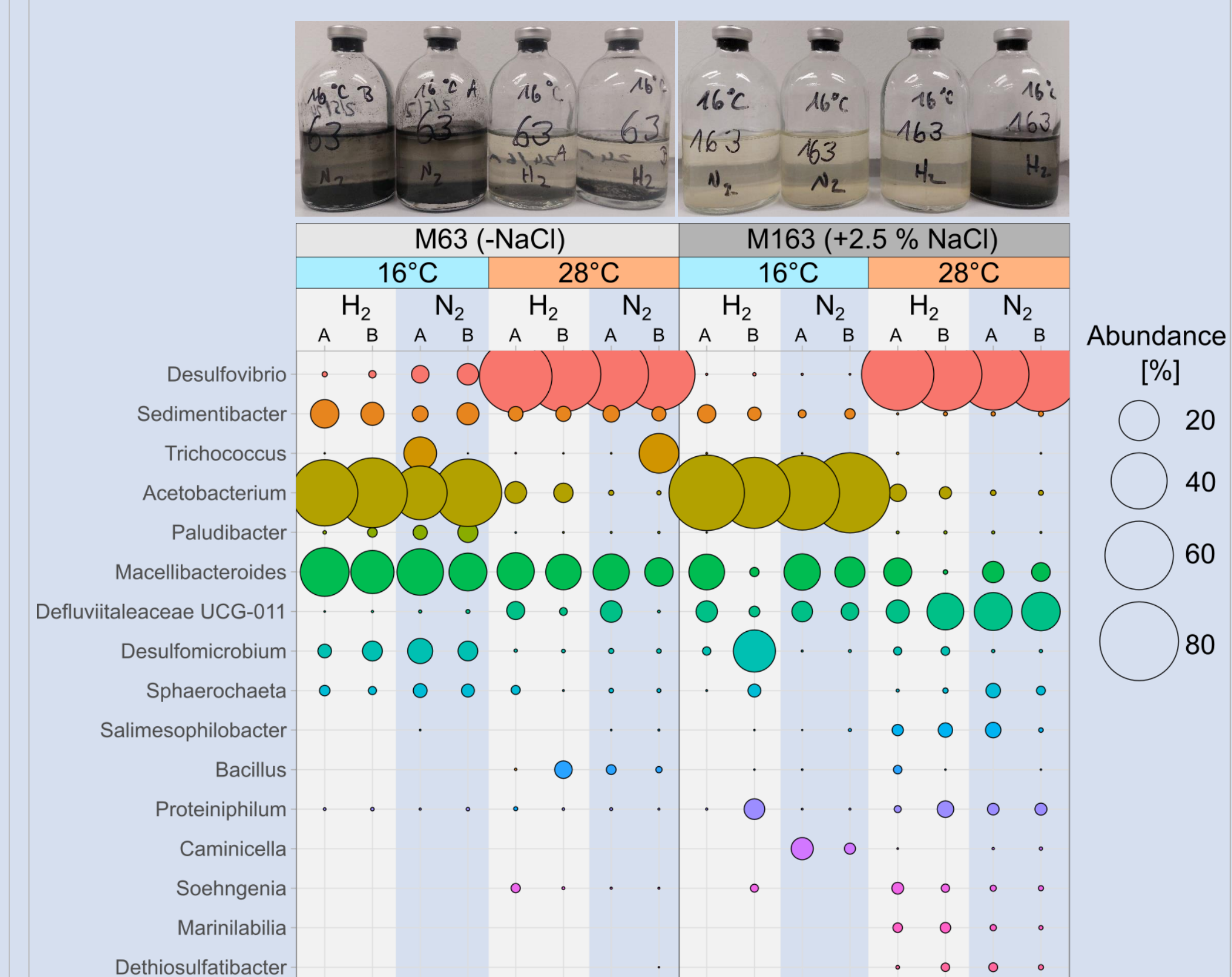
Abundance  
● 10  
● 20  
● 30  
● 40

● fermenters  
● sulfate reducers (SRB)  
● methanogens



**Fig. 2A: NMDS analysis** – microbial community shift after > 2 months and stabilization after > 14 months. **2B: Relative abundance** changes of fermenters, SRB and methanogens over time.

## Enrichments



**Fig. 3: Composition and relative abundance of taxa after enrichment** for sulfate reducers at different temperatures and head spaces with and without NaCl.

- Abundance of SRB correlates with formation of black precipitates (most likely Fe-sulfide)
- Temperature controls the type of organisms being enriched
- No enrichment at 55 °C
- Salinity and head space composition have no significant impact