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# Integrated response of plant, microbial and N-cycling interactions to precipitation patterns

**Ilonka Engelhardt, Laurent Philippot, Romain Barnard** 

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# INTRODUCTION

Climate is changing towards more extreme precipitation patterns

	Dry period	Rewetting after dry period
SOIL	Decreased water saturation of pore spaces	<ul> <li>Increased saturation of pore spaces</li> <li>Breaking up of soil aggregates</li> <li>Release of trapped nutrients</li> <li>Increased respiration and release of greenhouse gases</li> </ul>
MICROBES	<ul> <li>Decreased mobility of microbes and nutrients</li> <li>Accumulation of intracellular osmolytes to prevent dehydration</li> <li>Increased number of dormant and resistant forms</li> <li>Increased fungal:bacterial ratio</li> <li>Decreased N and C cycling rates</li> </ul>	<ul> <li>Increase in competitive and fast growing species</li> <li>Release of accumulated osmolytes to prevent cell lysis</li> <li>Increase in microbial activity including N and C cycling</li> </ul>
PLANT	<ul> <li>Reduced stomatal conductance to avoid water loss</li> <li>Decreased photosynthesis rate</li> <li>Access deeper water sources if possible</li> </ul>	<ul> <li>Increased photosynthesis rate</li> <li>Resumed transfer of rhizodeposits to the soil</li> </ul>

# **OBJECTIVES**

Gain insight into the dynamics of the interactions involved in the coupling between water and N cycles, as affected by changes in precipitations patterns. A multidisciplinary approach is used to evaluate the response of soil microbial community, plant, and soil N cycle to repeated cycles of drying-rewetting patterns of contrasted frequency and magnitude.



# <sup>18</sup>O Stable Isotope Probing (<sup>18</sup>O-SIP)

Capture the temporal dynamics of metabolically active soil microorganisms upon rewetting after contrasted water availability.

## Aim

Identify bacterial strategies

Relate to soil N cycling and N<sub>2</sub>O emissions

## Principle

<sup>18</sup>O-labelled water allows identification of metabolically active communities in soil without the addition of external nutrient sources (Blazewicz et al. 2014).



#### Approach

- Isolate metabolically active community – SIP
- Identify community  $\bullet$ composition – sequencing
- Evaluate abundance of N cycling genes - qPCR

Fig. 2. Dynamics of soil water content in the precipitation pattern treatments and controlled wet-up (adapted from Barnard et al. 2015).

## **GREENHOUSE EXPERIMENT**

Evaluate integrated response of system as a whole (plant, microbiota, soil) to rewetting after different precipitation regimes



Fig. 1. Conceptual diagram of plant-soil interactions during and after a dry period. Arrow widths are proportional to flux rate.

# REFERENCES

Barnard RL, Osborne CA, Firestone MK. 2015. Changing precipitation pattern alters soil microbial community response to wet-up under a Mediterranean-type climate. ISME J. in press.

Blazewicz SJ, Schwartz E. 2014. Growth and death of bacteria and fungi underlie rainfallinduced carbon dioxide pulses from seasonally dried soil. Ecology **95**:1162–1172.

- Aim
- Establish fate of N by tracking <sup>15</sup>N-NH<sub>4</sub> tracer as plants and microbes compete for resources
- Link to interaction between plant performance and soil microbial activity

### Approach



- Soil microbial community: metagenomics, taxonomic diversity lacksquare
- Plants: high thoughput phenotyping platform
- Soil N cycling: nitrification, denitrification, N<sub>2</sub>O efflux rate