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INVESTIGATING THE DROUGHT-PRONE BIOLOGICAL INTERPLAY OF SOIL MICROBIAL COMMUNITIES AND SCOTS PINE TREES

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Website of the project





Acknowledgements: We thank the Swiss National Science Foundation for funding the project PZOOP2_180030. We are thankful for the help and advice of Matti Barthel, Britta Jahn-Humphrey, Brigitta Herzog, Manon Longepierre, Roman Hüppi, Rafaela Feola Conz.



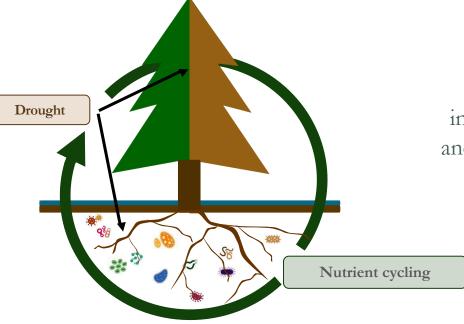




Background

Investigating the effects of drought on the biological interplay between Scots pine trees and soil microbial communities.

For Scots pine (*Pinus sylvestris* L.) high mortality rates have been observed in Switzerland^[1]. This dieback appears to be primarily caused by effects of drought^[1].

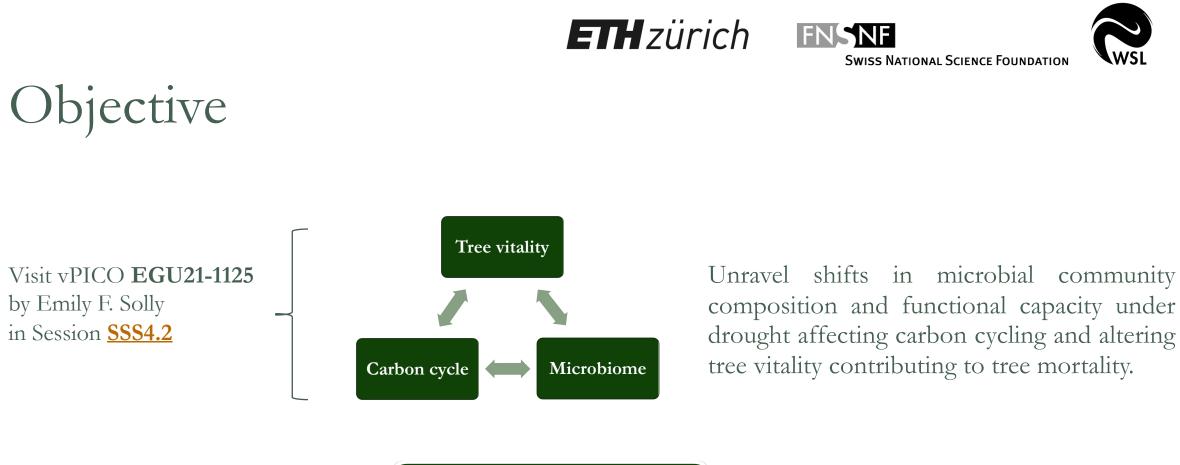


Drought induces a series of interconnected effects between trees and the associated soil microbiome^[2]. → This could strongly alter carbon cycling in forests

Bigler, Bräker, Bugmann, Dobbertin, Rigling. Ecosystems. 2006;9(3):330–43.
Classen AT, Sundqvist MK, Henning JA, Newman GS, Moore JAM, Cregger MA, et al. Ecosphere. 2015;6(8).









Activity	Basal Respiration	Abundance	qPCR of taxonomic markers (16S, 18S)
Biomass	CFE \rightarrow C _{mic} , N _{mic} , δ^{13} C enrichment	Diversity	Metabarcoding
Structure	Extractable PLFAs $\rightarrow \delta^{13}$ C enrichment	Function	Metagenomics









Experimental Set-up

18 Scots pine- soil mesocosms Origin of trees & soil: Pfyn forest, CH Controlled seasonal temperatures

> 1st year 3 drought treatments

2nd year + 4 °C in growing season → drought + heat stress



Control field capacity



Dry 40 % reduction



Drought 75 % reduction







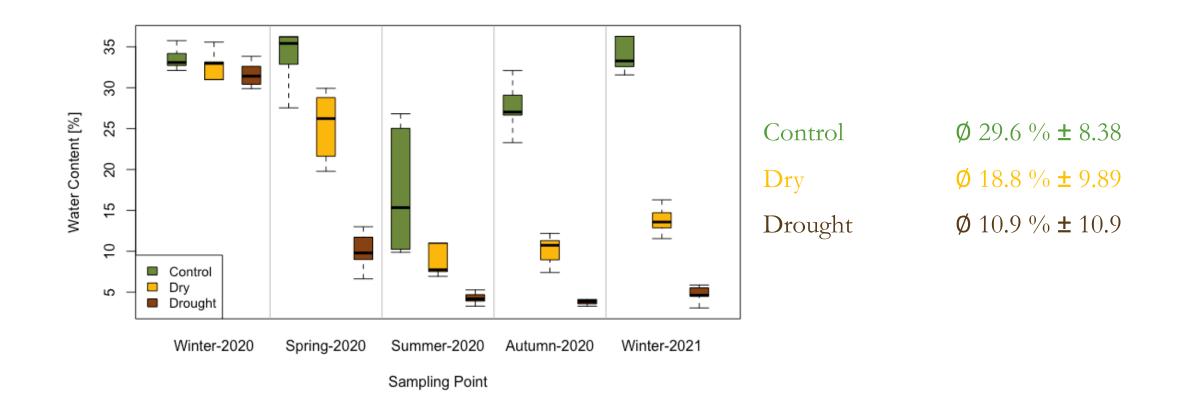






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One year of experimental drought



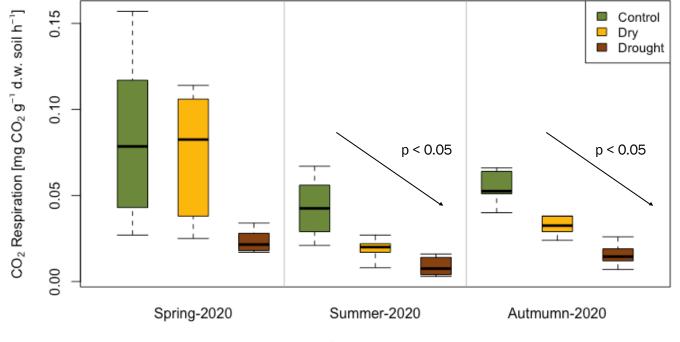








Basal Respiration during growing season



Sampling Point

 \rightarrow Basal respiration used as indicator for microbial activity

→ Experimental drought influenced basal respiration in soils

→ Drought condition showed significantly lower basal respiration during growing season







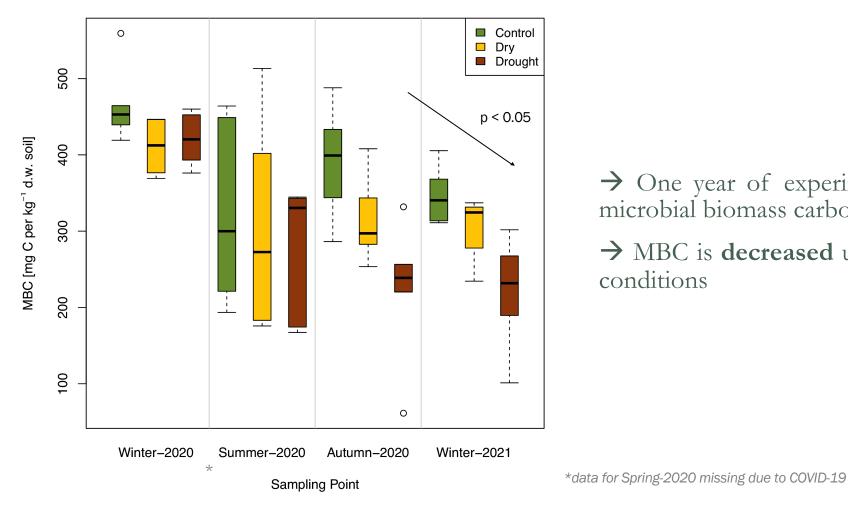






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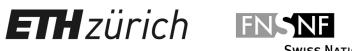
Microbial Biomass Carbon (MBC)



 \rightarrow One year of experimental drought altered microbial biomass carbon (MBC) in soils

→ MBC is **decreased** under **dry** and **drought** conditions









Outlook



Tracing the natural pathway of photosynthetic CO₂ assimilates into microbial groups under drought

¹³C-CO₂ pulse-labelling of9 mesocosms

1000 ppm of 99.9 % $^{13}\text{C-CO}_2$

45 min per Tree

Sampling: 0h, 24h, 48h, 72h, 7d

Structure	Extractable PLFAs $\rightarrow \delta^{13}$ C enrichment
Biomass	CFE \rightarrow C _{mic} , N _{mic} , δ^{13} C enrichment



