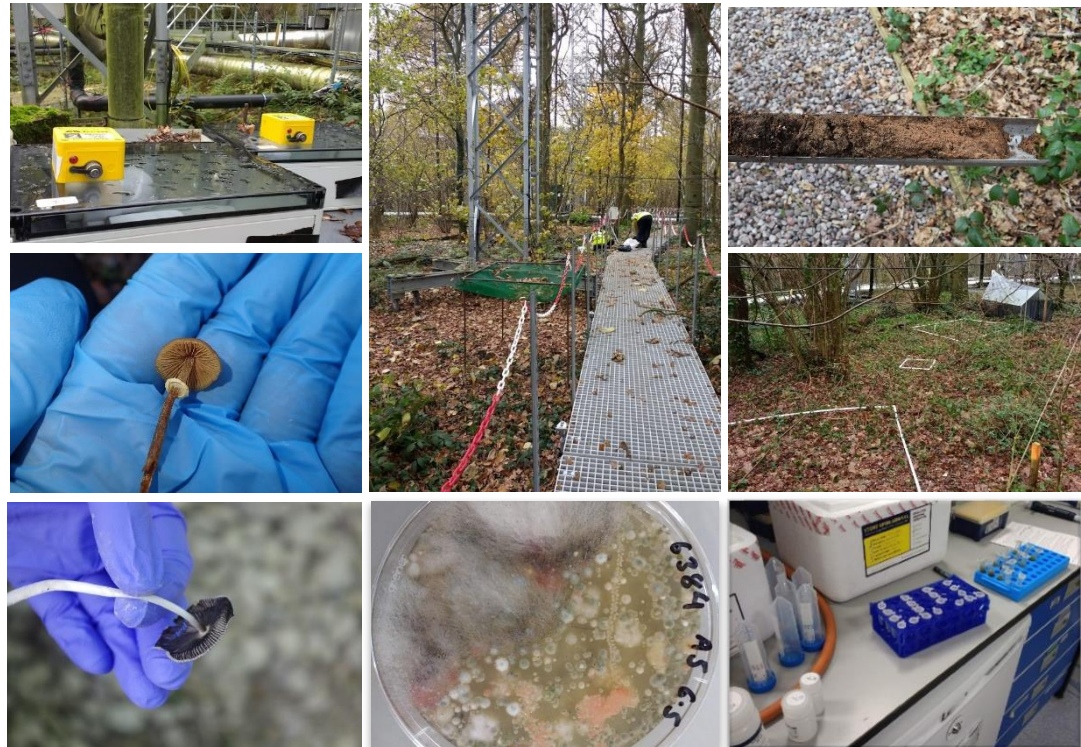


Fungi of the future:

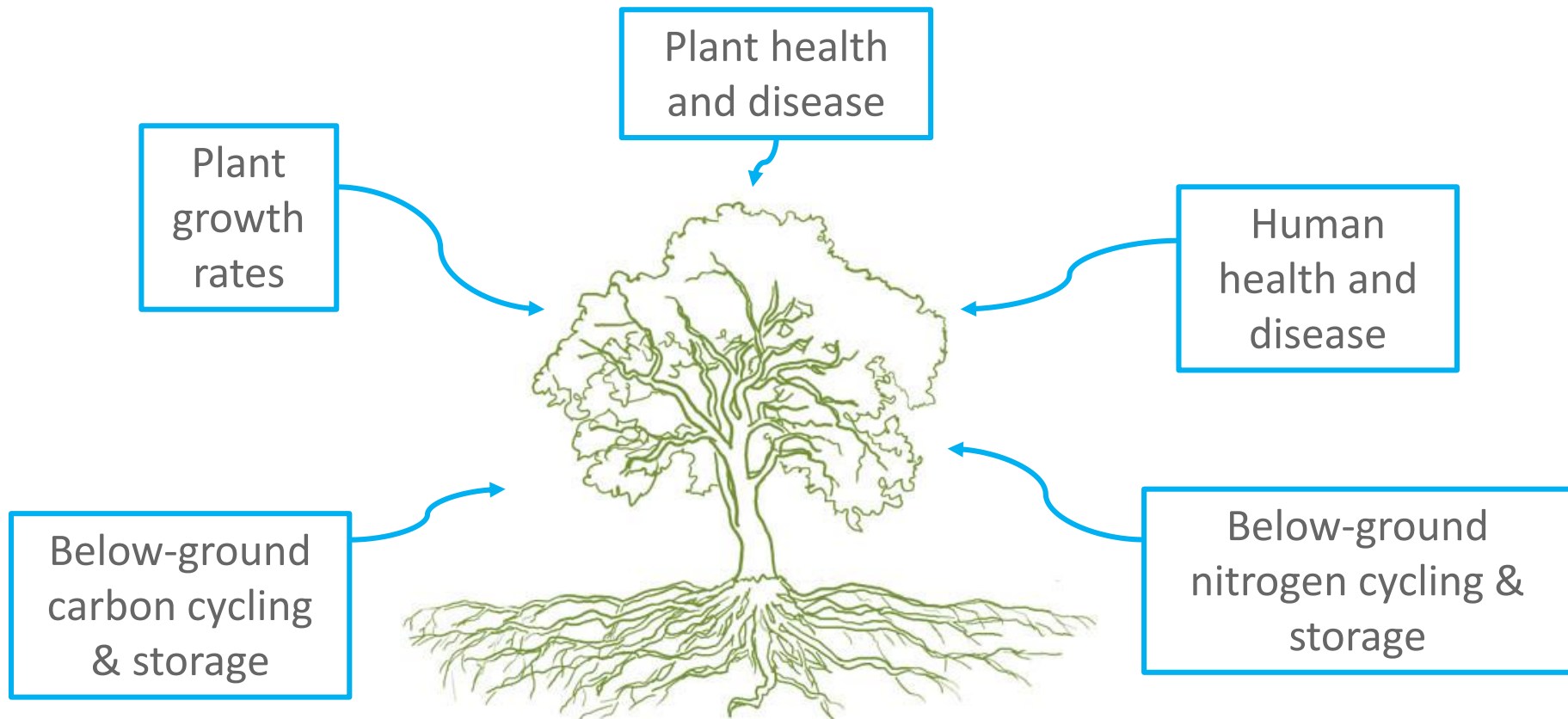
Assessing the effects of elevated CO₂ on forest fungal communities



BIFoR FACE

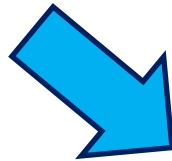


Why are fungi important in forest systems?



Rationale

Fungi could have a significant impact in how a forest responds to eCO_2 .



Environmental fungal populations are largely unstudied.



Measuring the fungal populations will significantly contribute to our understanding of forest responses to eCO_2 .

Aims



1. Characterise the fungal populations (and their spatial and temporal variation) in a temperate forest (BIFoR Mill Haft).
2. Investigate whether fungal community composition at BIFoR FACE is affected by $e\text{CO}_2$.

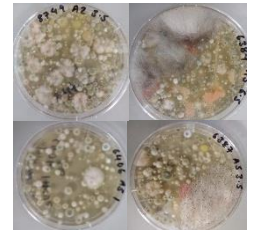


Experimental strategies



Counting & measuring bioaerosols

Sampling & culturing bioaerosols



Surveying fungal fruiting bodies

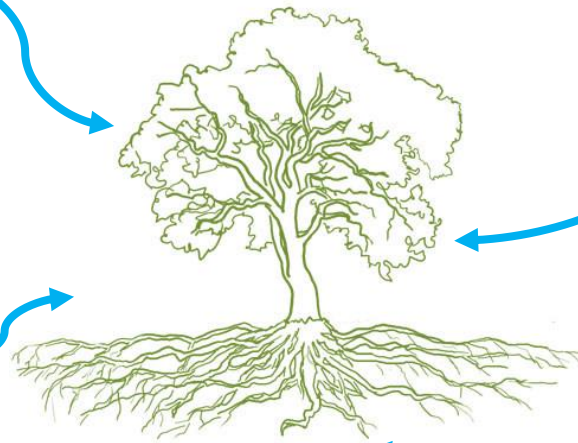


Extracting and analysing fungal DNA from soil



All work is completed in 6 arrays at BIFoR:

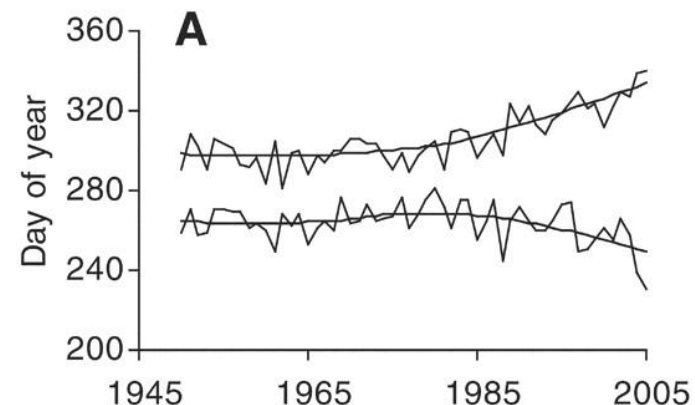
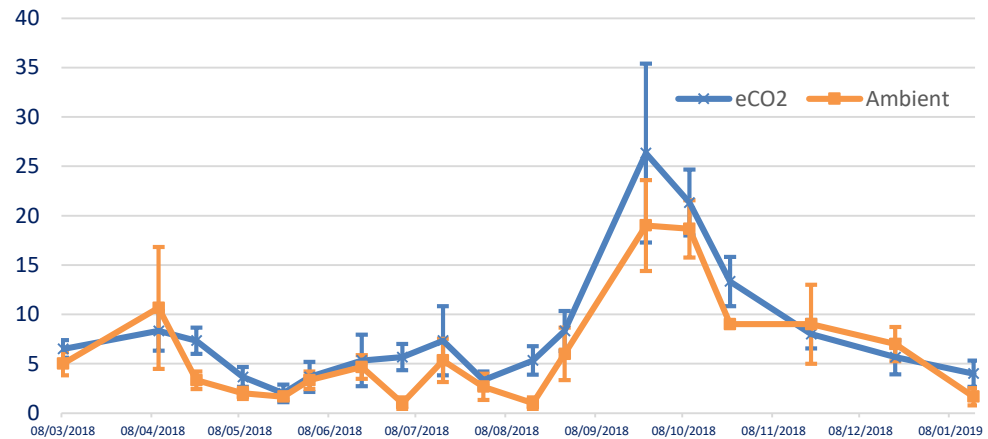
- 3 eCO₂ arrays
- 3 ambient control arrays



Strategy number 1: Traditional fungal survey techniques

Methods

- Manual survey of fruiting bodies of defined areas in 6 arrays
- Photographs and samples of all species found.

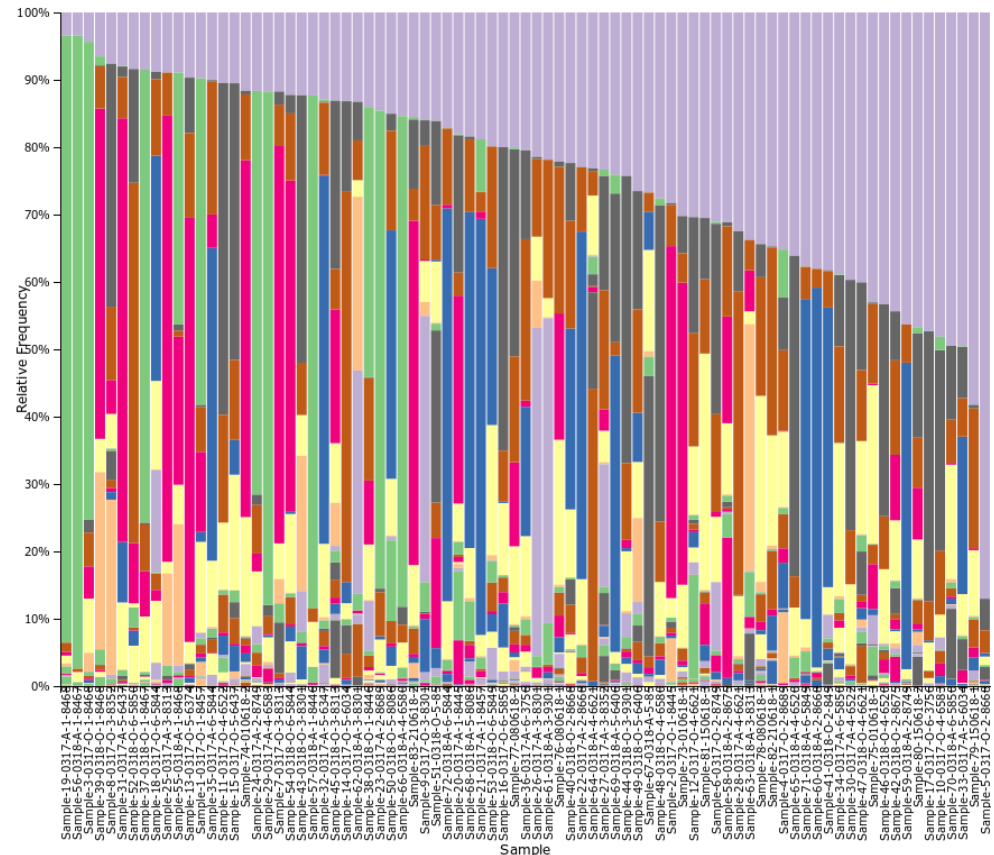


Gange et al., 2007

Strategy number 2: High-throughput DNA sequencing

Methods

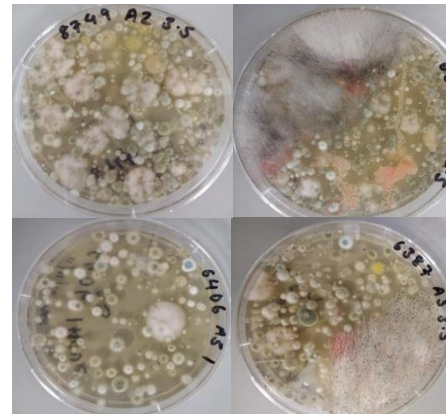
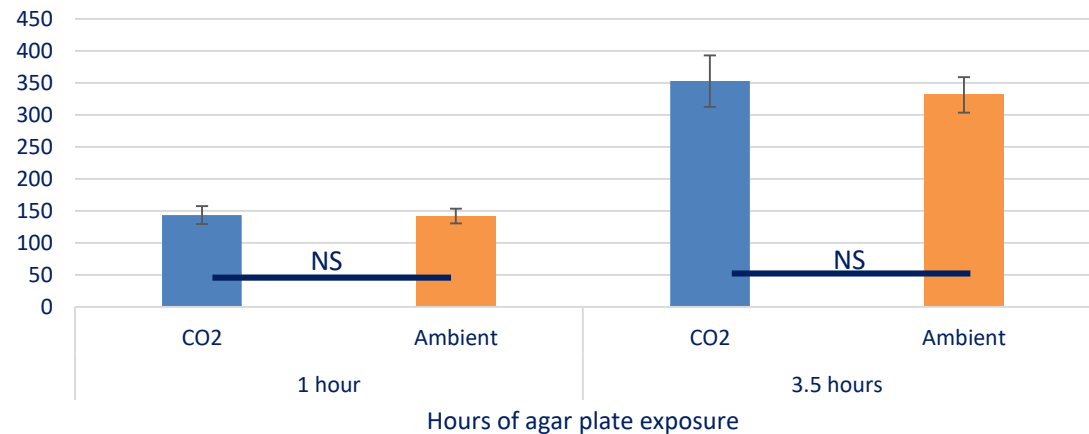
- Soil samples from March 2017 & March 2018
- Extract total DNA
- Isolate and sequence a fungi-specific section of the genome : ITS1
- Compare ITS1 sequences with ITS1 sequences of known fungi to collate a list of fungal species in each soil sample



Strategy number 3: Bioaerosol sampling & measurements

Methods

- Sampled & cultured fungal aerosols over a 3 week period in Autumn 2018.
- We have also collected 2 additional datasets not presented here which use Optical Particle Counters (OPCs)
 - Throughout 2016 (pre-switch on)
 - Autumn 2018



Plans for 2019



- A. In-depth analysis of soil metagenomics data set
- B. Integration of current data with BIFoR environmental data
- C. 2019 sampling season



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- Members of the HAPI lab
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- David Tubbs
- Bioinformatics team @ University of Liverpool

