

# DECRyPT Workshop **Ecological Microbiome Studies**5-7 October, Tübingen

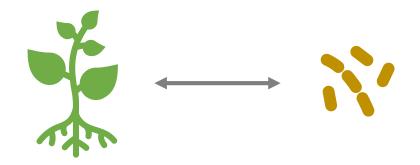
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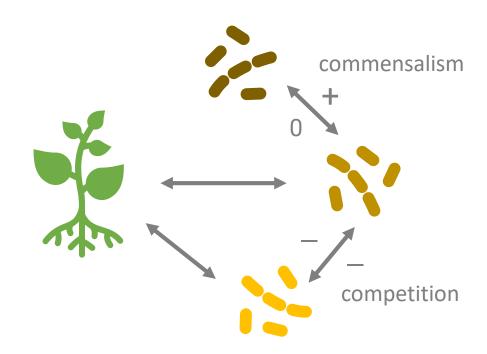
### Ecological microbiome studies?

**Ecology** = Study of relationships between organisms and their environments

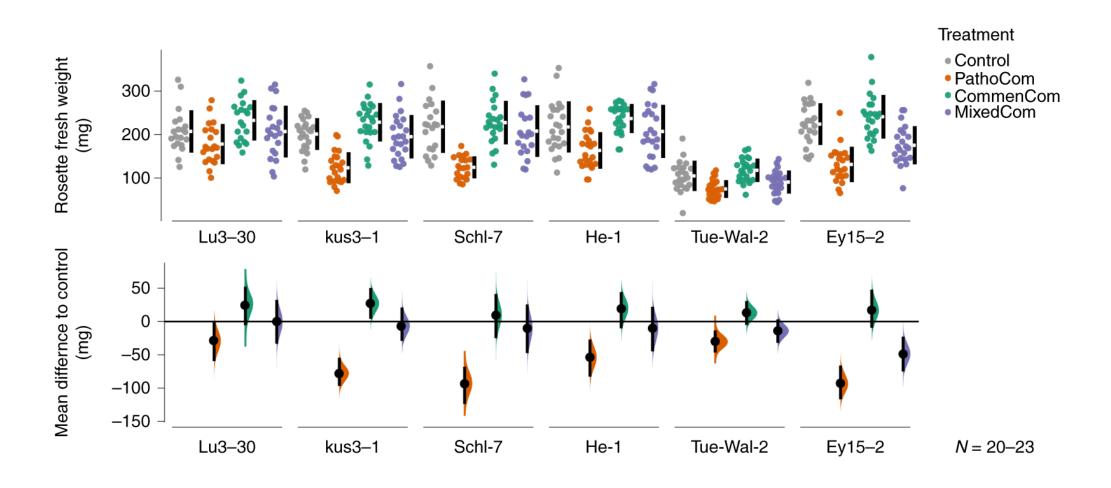


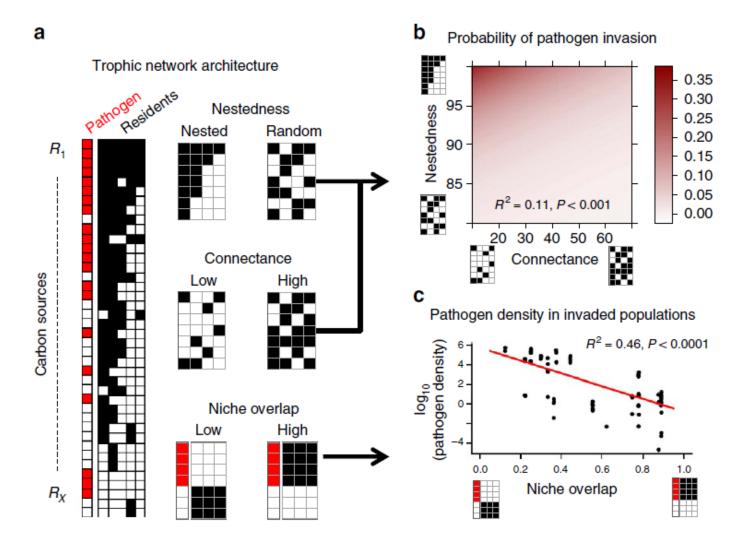
#### Microbe-microbe interactions

→ Direct and indirect effects of microbes on plants!

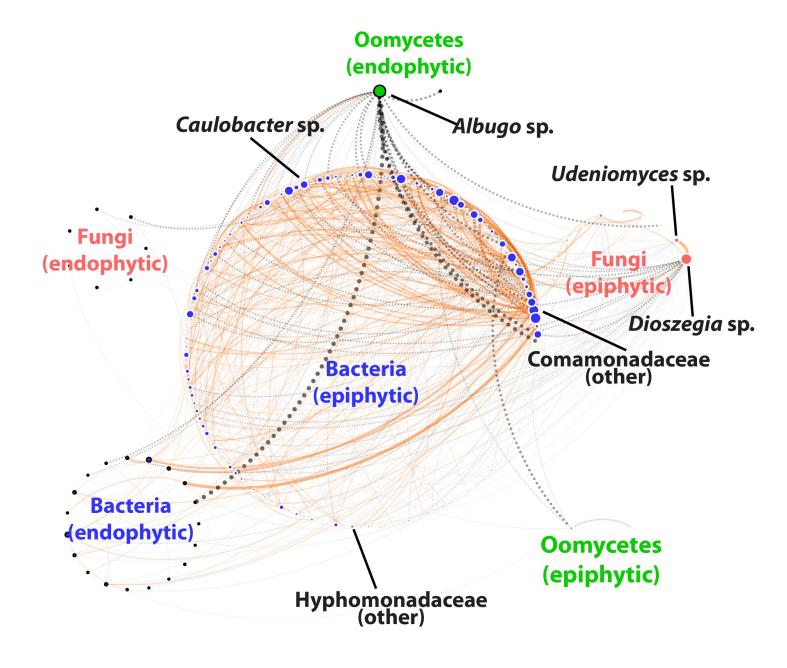


#### Commensal *Pseudomonas* strains protect *Arabidopsis* from pathogens

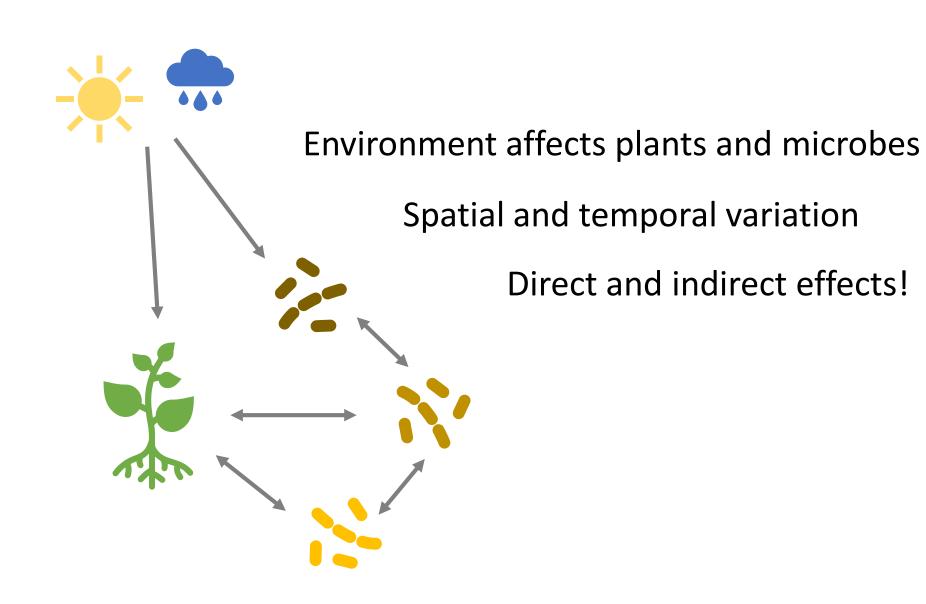




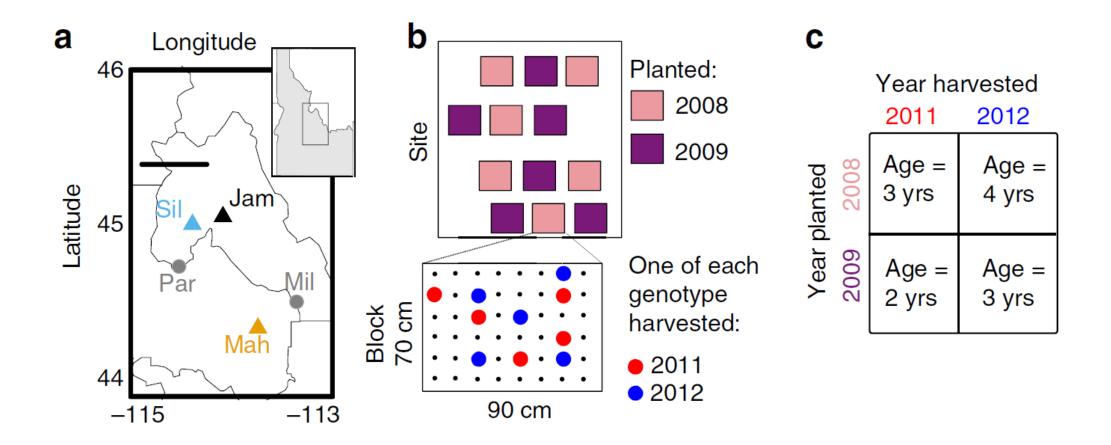
Bacterial community structure determines tomato pathogen resistance

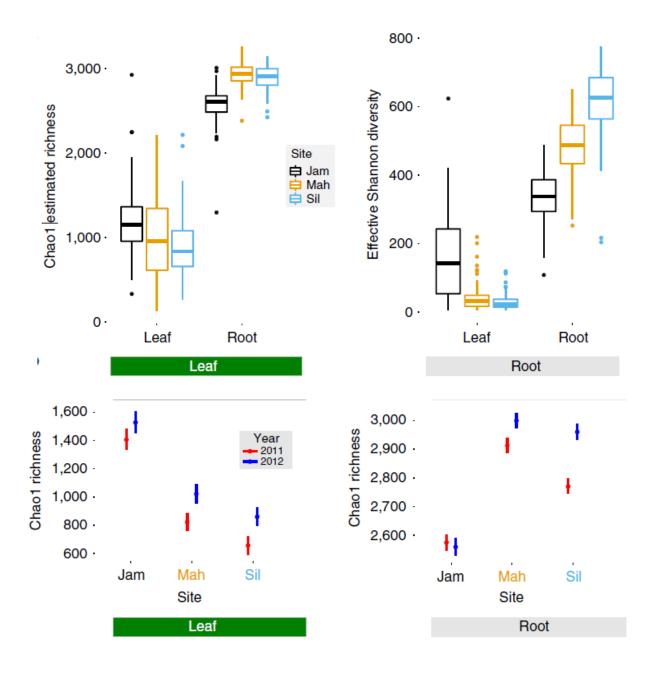


Network analyses try to understand microbe-microbe interactions from patterns of co-occurrence

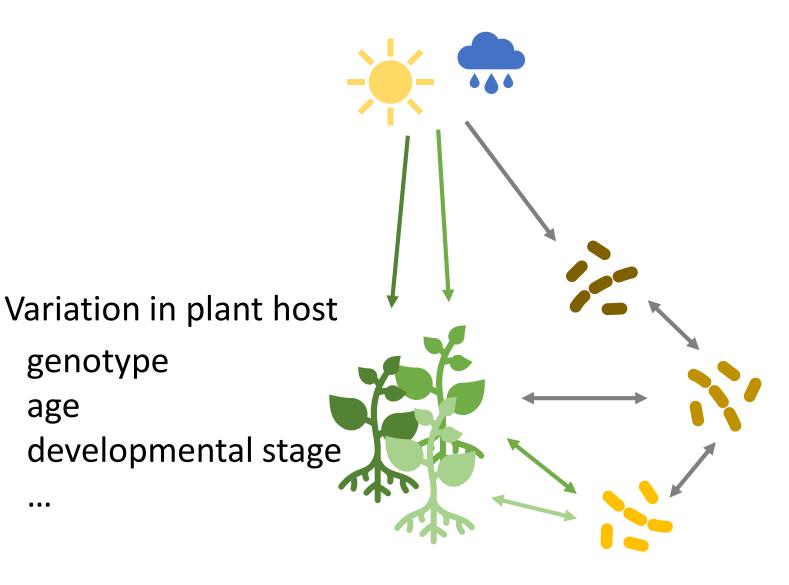


A multi-site field experiment to test environmental and host genotype effects on *Arabidopsis* microbiome



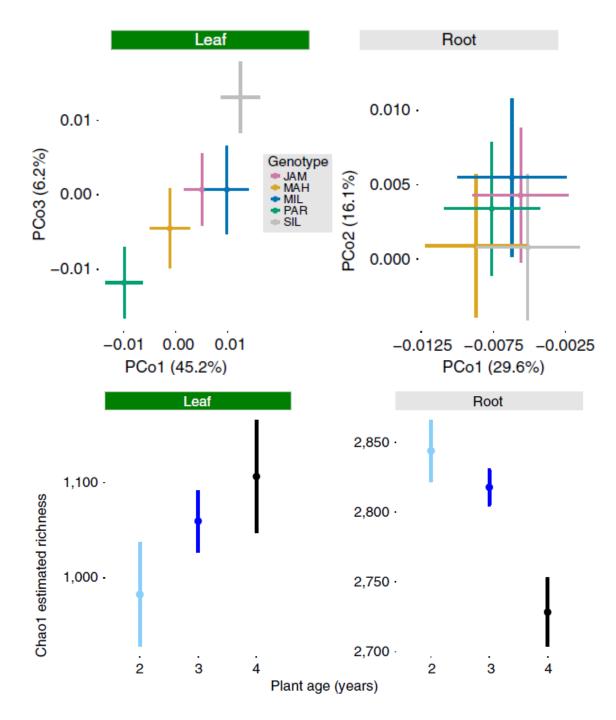


Differences in microbiome diversity and composition between sites and years!

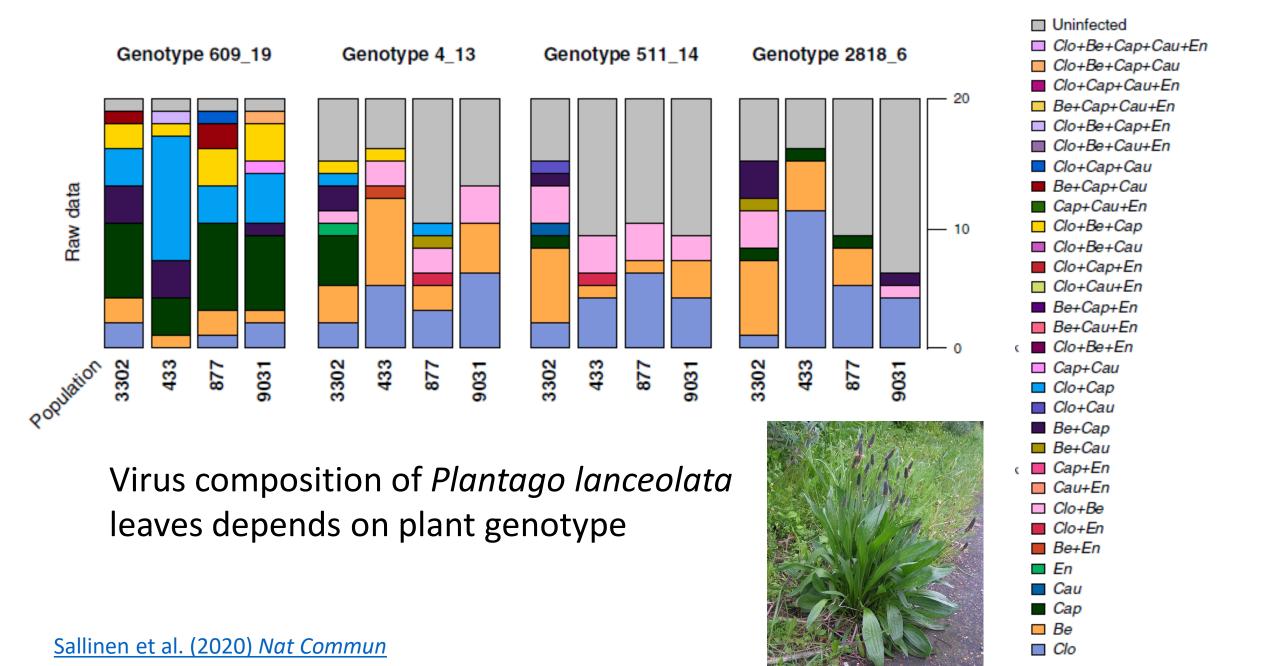


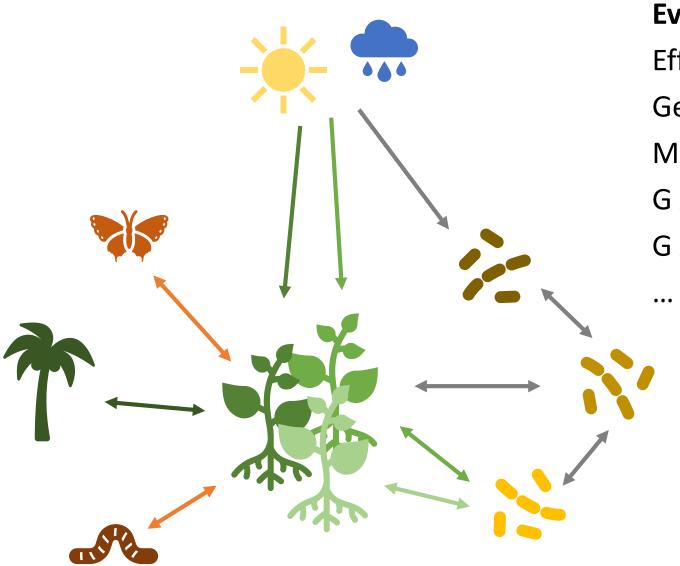
genotype

age



Arabidopsis microbiome differs between plant genotypes and plant age





#### **Even more complexity:**

Effects of other organisms

Genotype x environment (G x E)

Microbe x environment (M x E)

GxMxE

GxMxMxE

## Realism

Complex communities
Natural environments

Controlled environments
Sophisticated methods

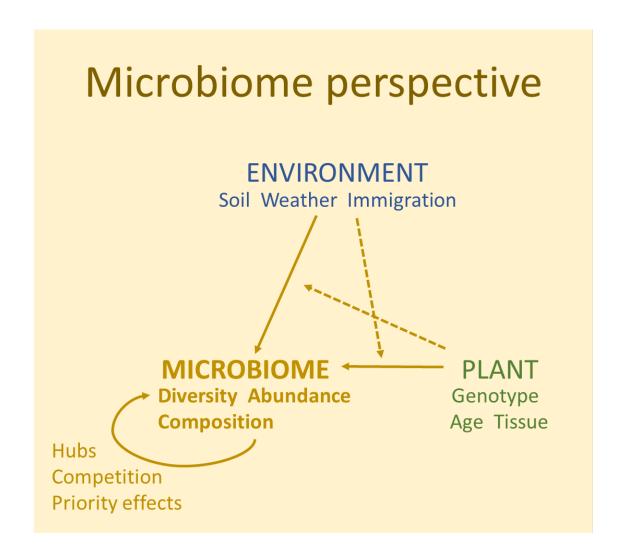
Many genotypes

Many environments

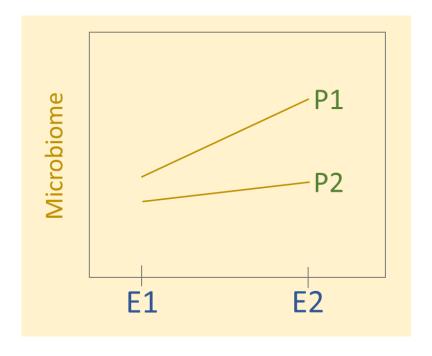
Precision

Generality

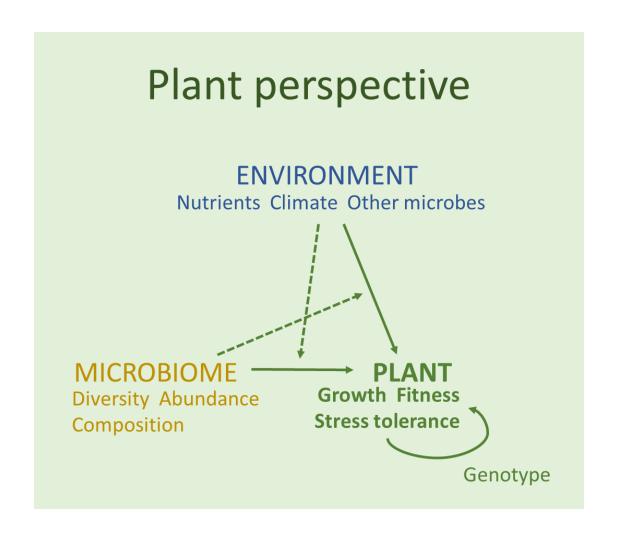
# Two types of ecological plant-microbiome studies



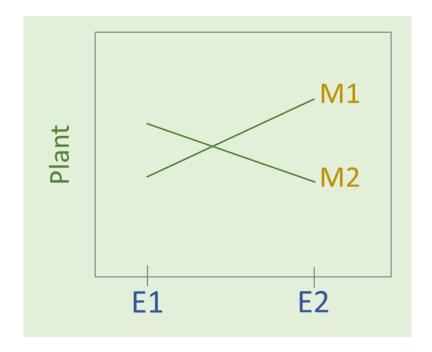
Microbiome is dependant variable
Often analysis of sequencing data

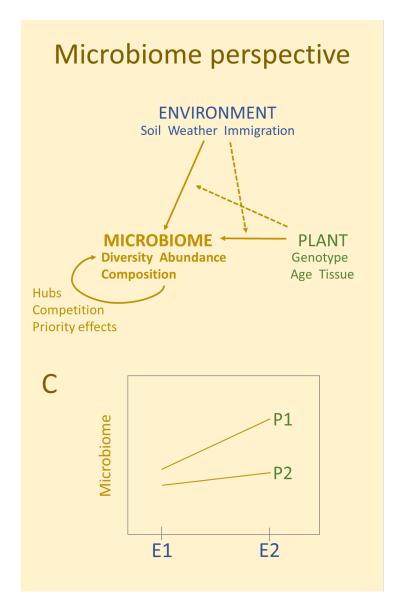


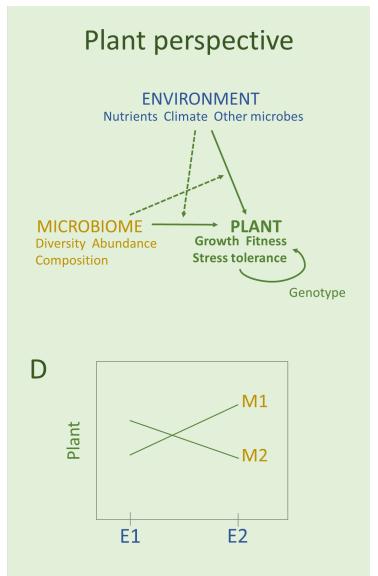
# Two types of ecological plant-microbiome studies



**Plant** is dependant variable
Analysis of plant fitness/physiology







Jung et al. (2021) Understanding plant microbiomes requires a genotype x environment framework. *Am J Bot* 

# Goals of this workshop

- Think about ecological perspectives
- Learn some computational & statistical methods
- Get to know each other and have fun

## Who are we?



And who are you?

5 October		6 October	
08:30	Room open, with coffee & croissants	08:30	Room open, with coffee & croissants
09:00	Welcome & introduction What are ecological plant-microbiome studies? Getting to know each other	09:00	Analyzing microbial community data III  Network statistics, hubs, etc. continued  Using machine learning to understand microbial communities
10:00	Amplicon sequencing data Setting up OTU/ASV tables/metadata. Evaluating data quality. Jupyter/Google colabs notebooks Short intro to R	12:00	Lunch
		13:00	Analyzing inoculation studies I Basic logic of GLM Environmental influences Plant species/genotype effects
12:30	Lunch	15:00	Coffee & cake
13:30	Analyzing microbial community data I R packages for microbiome analyses; normalizing data Simple stats: comparing samples, diversity metrics, etc.	15:30	Analyzing inoculation studies II G x E interactions Visualizing multifactorial results
15:30	Coffee & cake	17:00	Group work – connect with own project(s)
16:00	Analyzing microbial community data II Analysing & visualising networks Network statistics, hubs, etc.	18:00	
		19:30	Dinner at <u>Die Wurstküche</u> , Am Lustnauer Tor 8 (Bus stop "Nonne
18:00	End	7 October	
19:30	Dinner at <u>An An</u> , Aixer Strasse 46 (Bus 13 to "Aixer Strasse)	08:30	Room open, with coffee & croissants
		09:00	Group work – connect with own project(s)
		10:00	Analyzing inoculation studies III Functional biodiversity Analysing microbial diversity experiments
		12:00	Farewell