







Journée Scientifique BASC 28 janvier 2021

Axe "comprendre et gérer la dynamique de la diversité biologique"

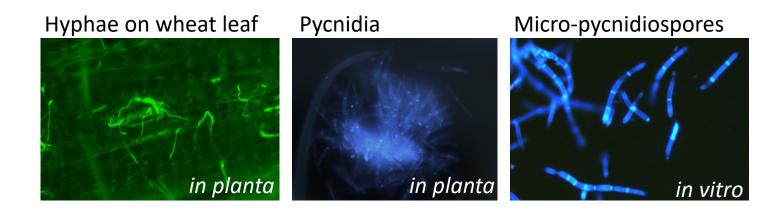
Evofungi (2013-2016)

Partner 1: Anne GENISSEL, UMR BIOGER, INRAE

Partner 2: Arnaud LE ROUZIC, UMR EGCE, CNRS

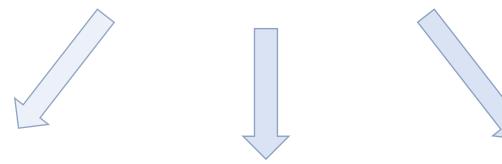
Assessing the evolvability of a pathogenic fungus in the context of global change

- Context: Important challenge is to understand the effects of environmental variation on pathogen evolution
- Questions: What is the role of phenotypic plasticity in adaptation? Is fluctuating environment a driver of plasticity?
- Model: Zymoseptoria tritici





Evofungi (2013-2016)





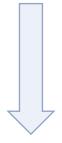
Evofungi (2013-2016)



 First experimental evolution in fungal pathogen

PROJECT

Tackle fundamental questions

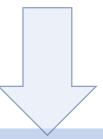






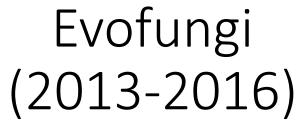
Evofungi (2013-2016)

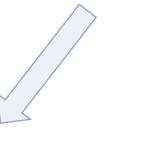


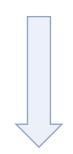










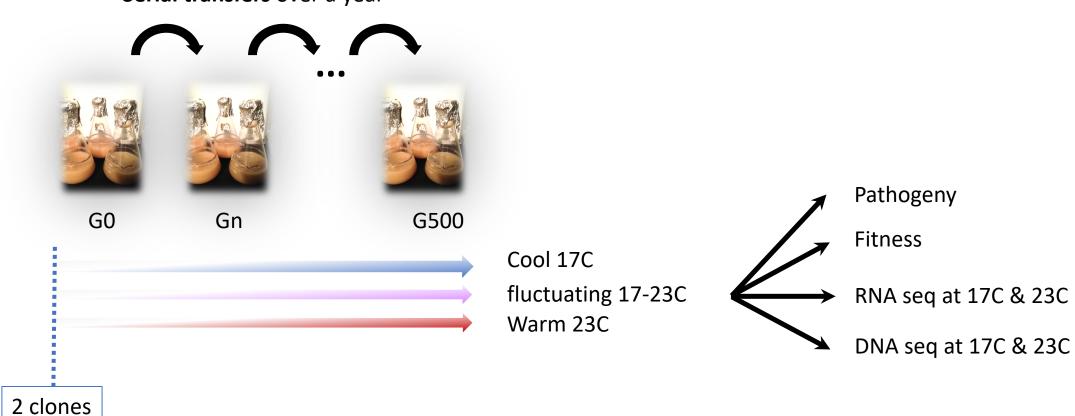


3 replicates



Experimental design for Evofungi

Serial transfers over a year



Results and interpretations: A. Jallet's PhD

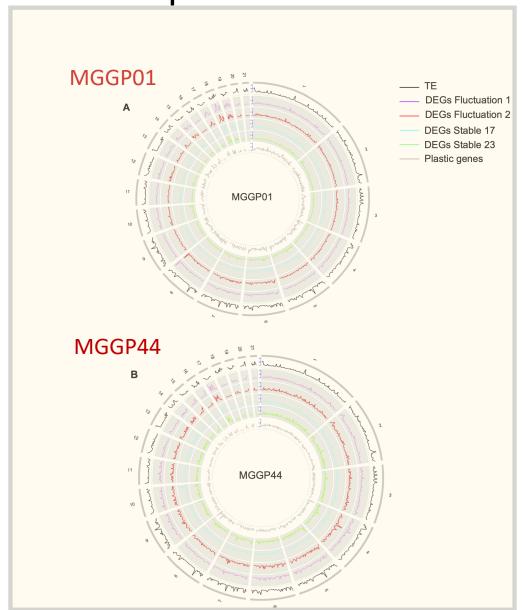
 Thorough description of gene expression variation due to temperature using Experimental Evolution

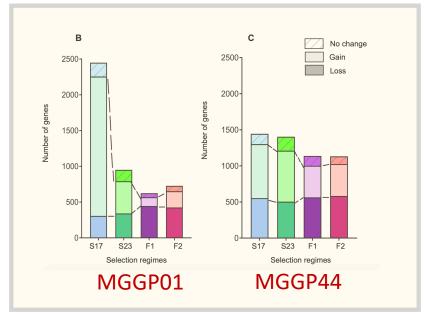


Arthur Jallet, now Postdoc at I. Bravo's lab, Montpellier

AgroParisTech

Transcriptome evolution under fluctuations

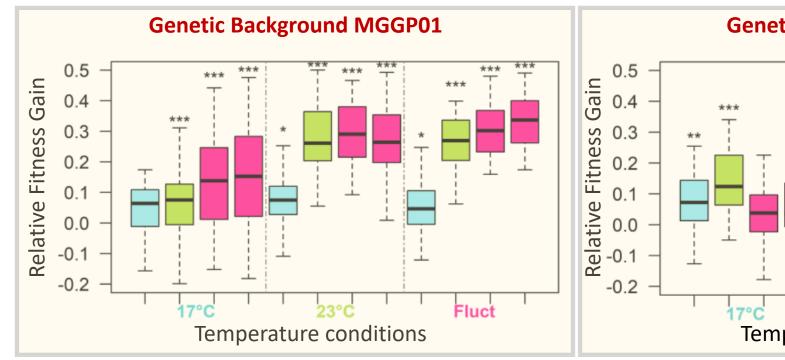


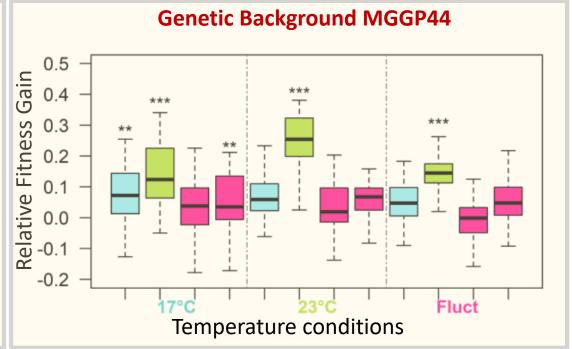


- > ~ 10% of transcriptome responded to the fluctuating regime
- > often localized in genomic regions enriched in TE
- > Evolution towards a loss of plasticity under fluctuations



Fitness evolution

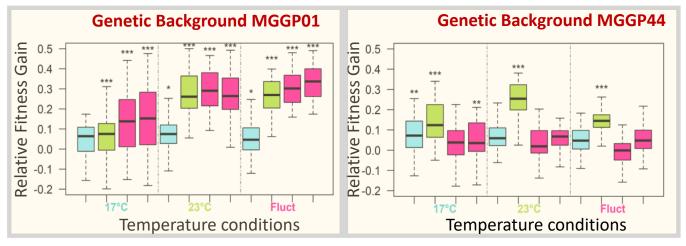


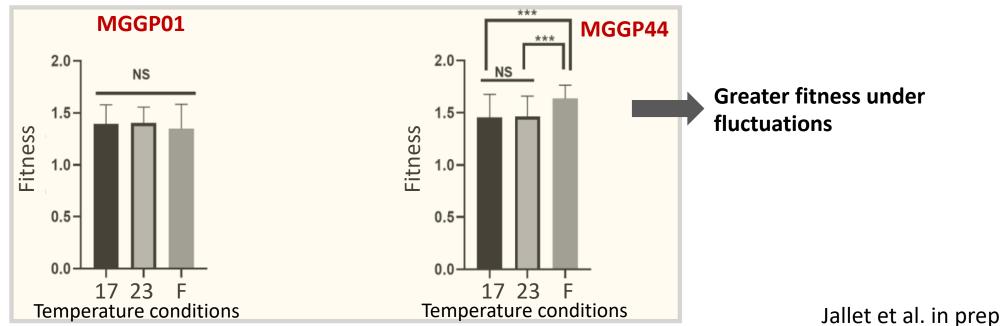


> Evolved generalists: gain of fitness regardless of the condition tested



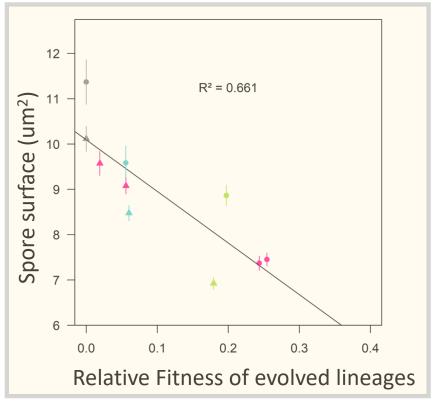
Fitness evolution







Body size matters

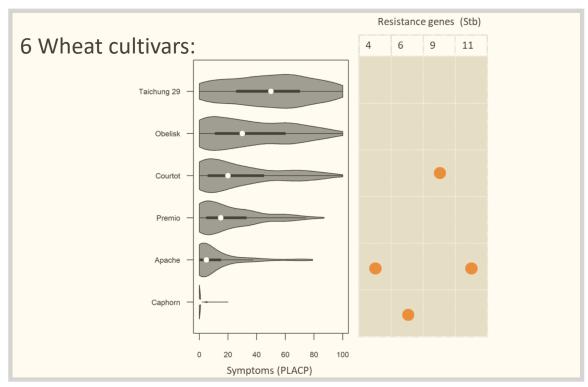


Adaptive evolution with a reduction of spore size

- Is there a common genetic basis or different evolutionary changes led to the same phenotypic evolution? ...Some candidates are currently tested
- Pleiotropic effects or direct selection on the trait?

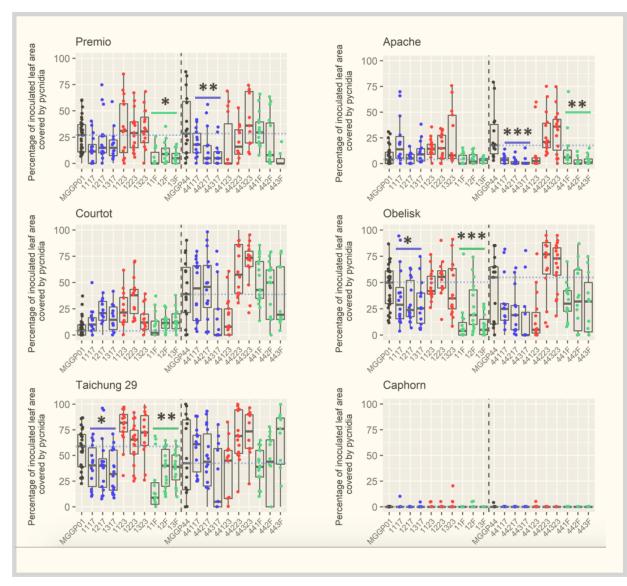
Consequences on disease symptoms

- After a year of evolution with no host we expect that pathogeny may decrease due to relaxed selection pressure
- Are different regimes influencing mutation accumulation and what is their effect on pathogeny?



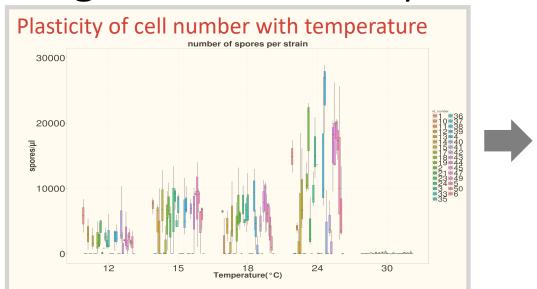
> Differential of disease symptoms among wheat genotypes

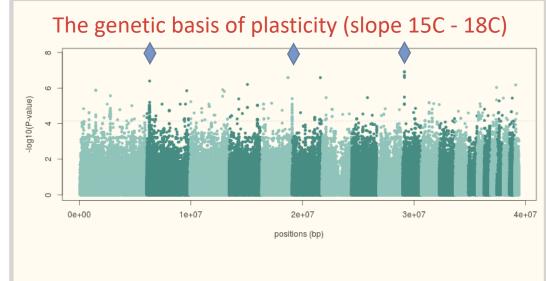
Decrease of disease symptoms in evolved lineages



- > Some significant decrease of symptoms, no evidence for a link with the selection regime
- ➤ No correlation with fitness gain measured in vitro
- > Trans-lineage segregating mutations in the genome, notably mutations in effector proteins.
- No loss of dispensable chromosomes occurred during the experimental evolution after 500 generations

More projects linked to Evofungi: Finding the heritability for the phenotypic plasticity

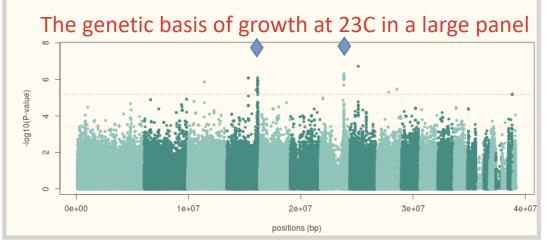






Maroua Bouzid (2013)

- Genetic basis of plasticity is hard to detect
- Need to develop high-throughput and precise phenotyping

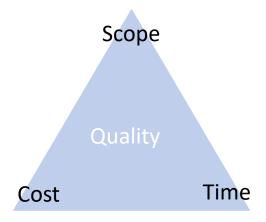




Amandine Bonnet (2015)

Labex Grant: unprecedented stimulus

- Excellent for the establishment of new research
- Positive long term effects



Thank you for attention