

#### Context

<u>Fusarium head blight:</u> Fungal disease → lot of damages (yield / food safety / added value)



Caused by different species of Fusarium  $\rightarrow$  different ecological niche with climate change?



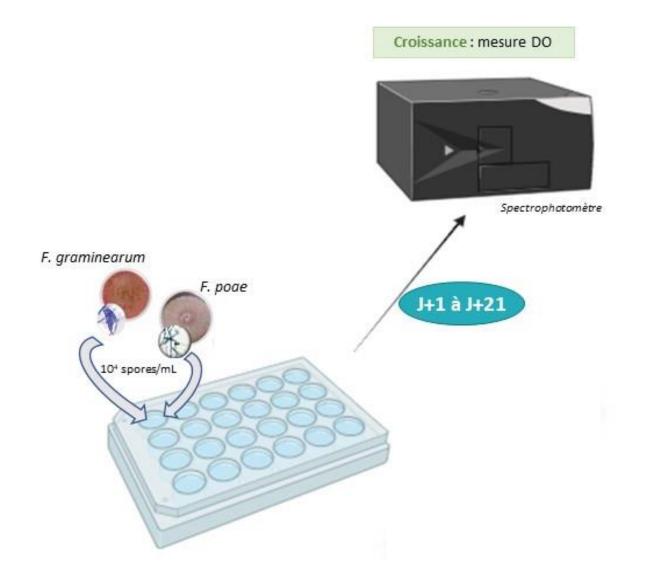


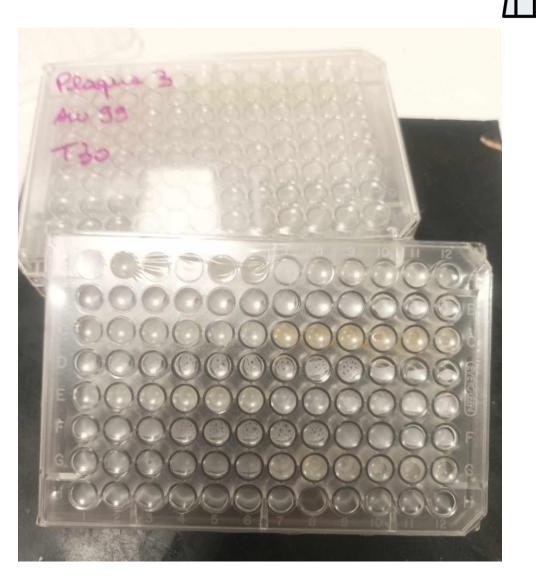
### How will the repartition of the different species of Fusarium evolve with climate change?

- 1. Does it grow ? Characterize the probability of growth of 25 strains of Fusarium under ≠ T° x aw (Water activity)
- 2. How does it grow? Characterize the kinetics of growth of 25 strains of Fusarium under ≠ T° x aw
- 3. What happens when it grows? Study the mycotoxins emissions (\$\$\$) 10/06/2025

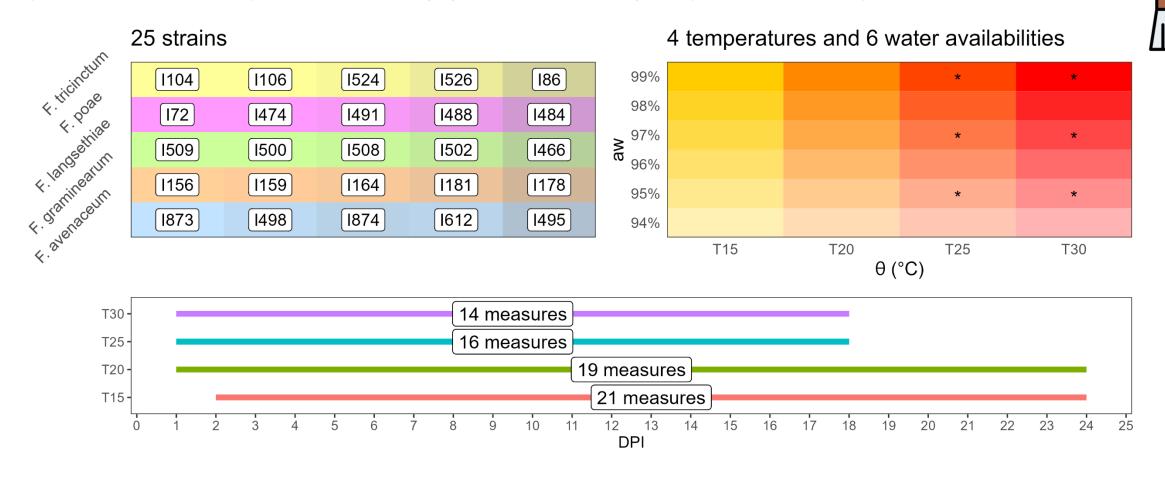


Experimental setup: monitoring growth through optical density measurements (OD)





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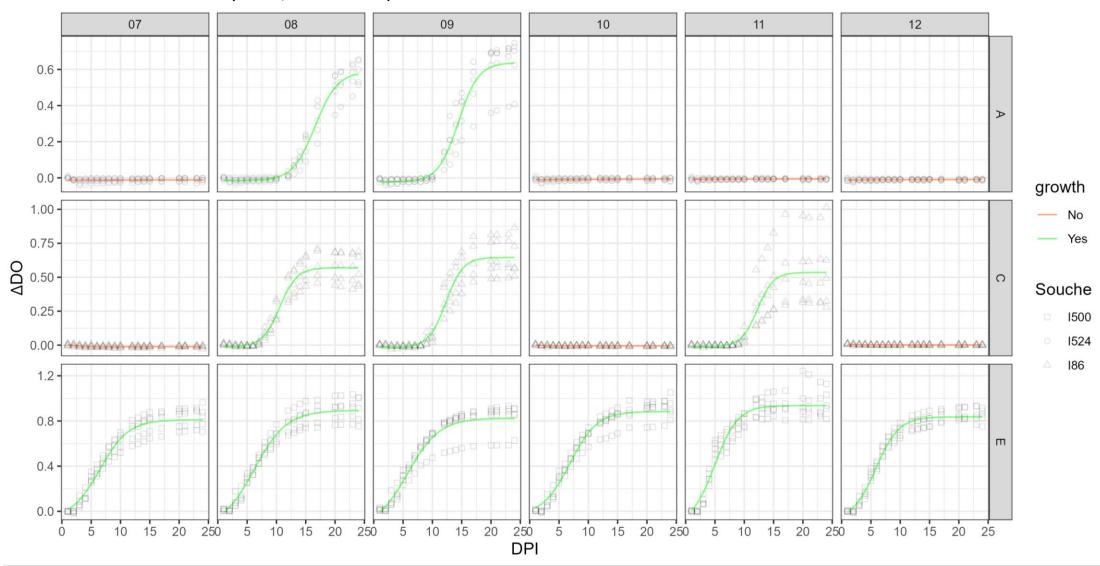


600 treatments, 6 replicates/treatments = 3600 samples (statistical individuals)
120 000 measures of DO



# All the samples did not grow

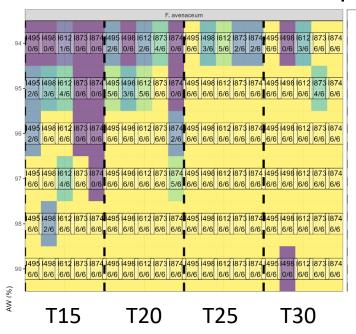
Evolution of OD (20 $^{\circ}$ C, aw = 98%)

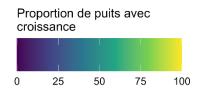


## Proportion of wells with growth, given temperature and aw



## Different behavior depending on the species?

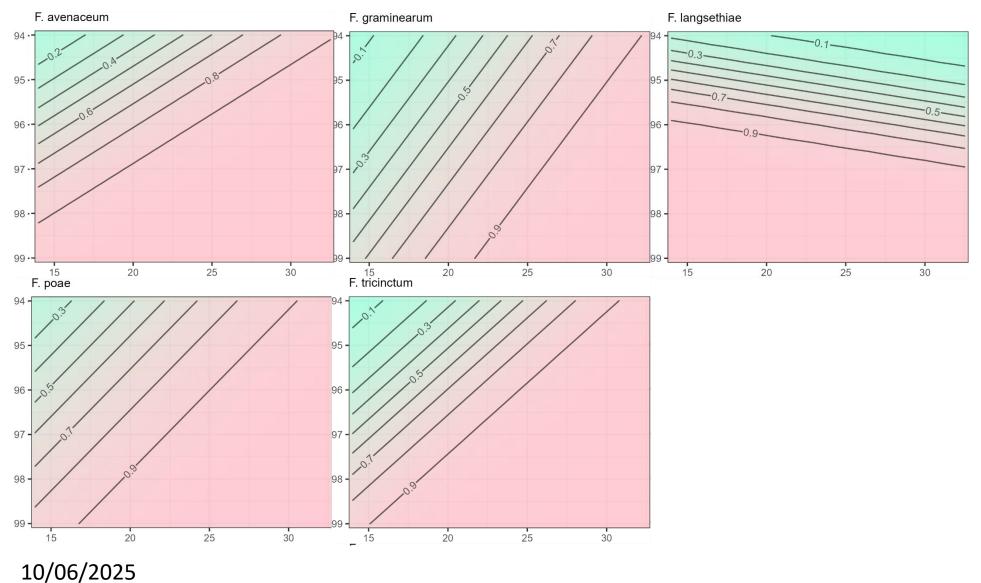




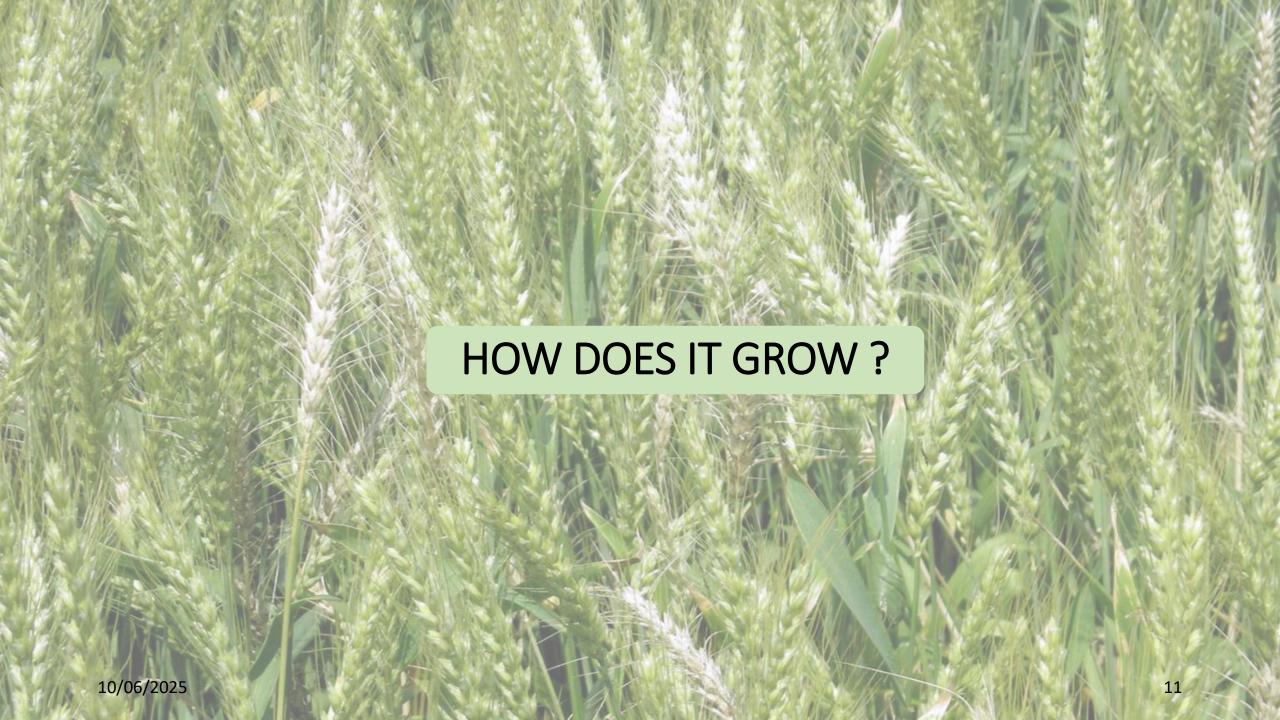


# Logistic regression to model the probability of growth

$$logit(P(growth)_i) = \beta_0 + \beta_{1i}Species_i + (\beta_2 + \beta_{2i})Temp + (\beta_3 + \beta_{3i})aw, i = 1, \dots, N$$

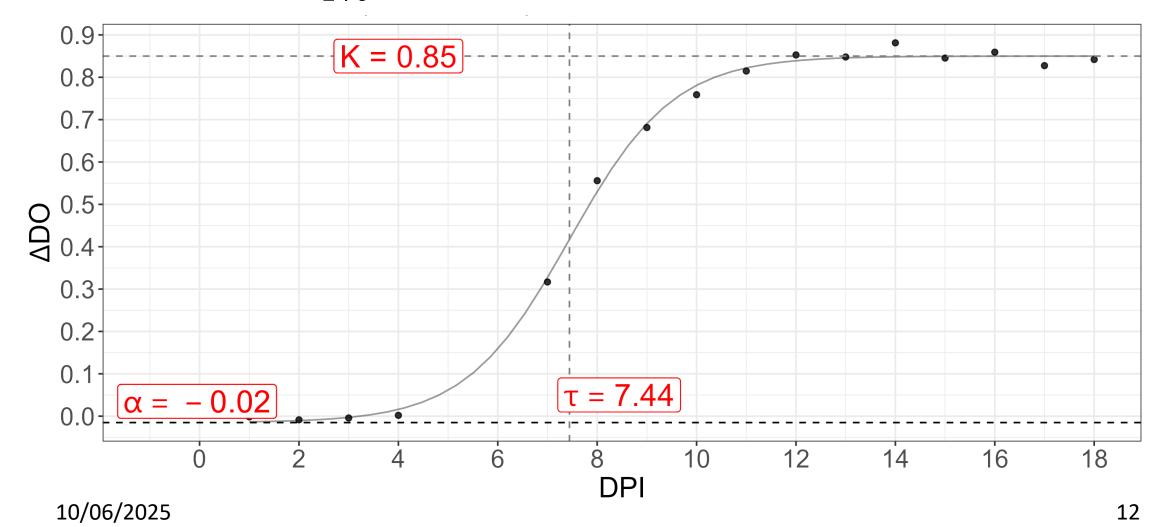


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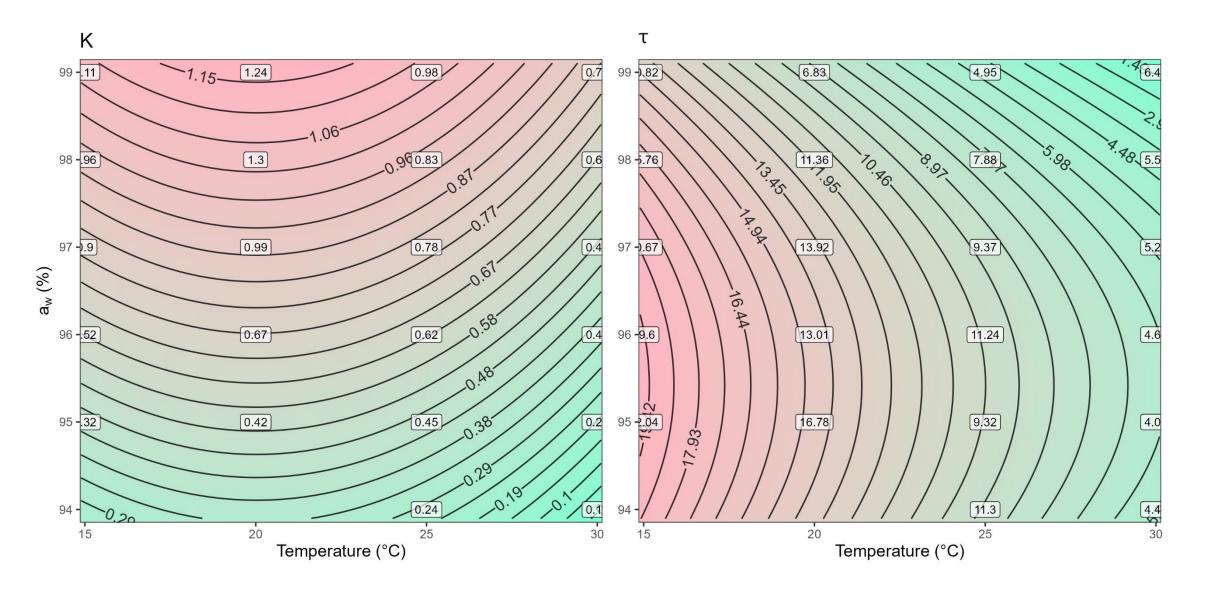


## Model the kinetics of OD

OD(t) = 
$$f_{\theta}(t) = \alpha + \frac{K}{1 + e^{r(\tau - t)}}$$
, where  $\theta = (\alpha, r, \tau, K)$ 

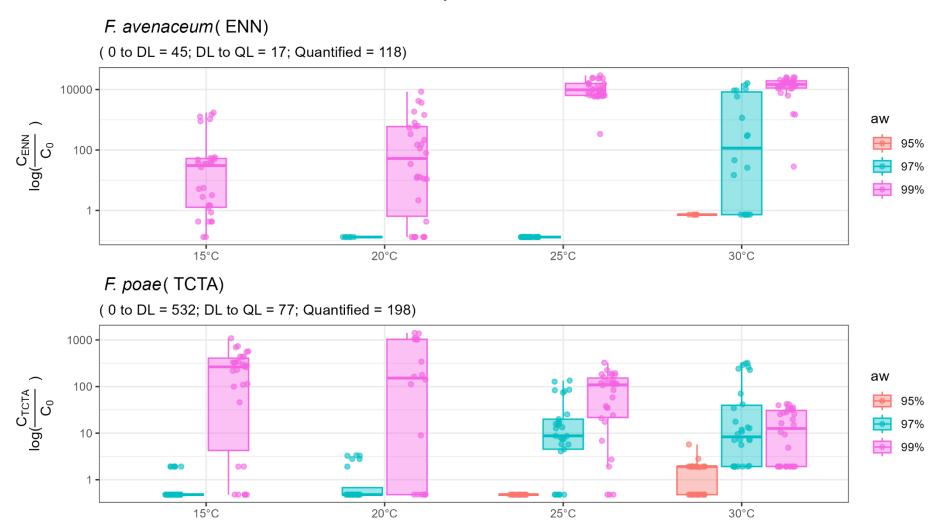


## Parameters as a function of T° and aw





# Toxins, different behaviour vs Temp x aw



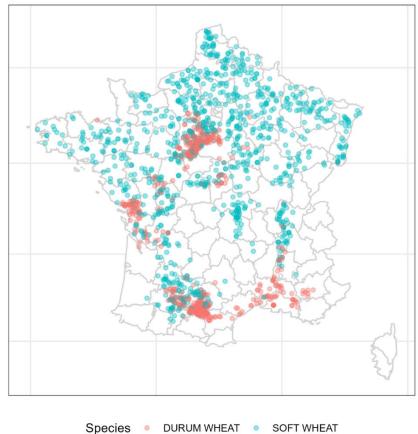
#### → What happens in vivo?

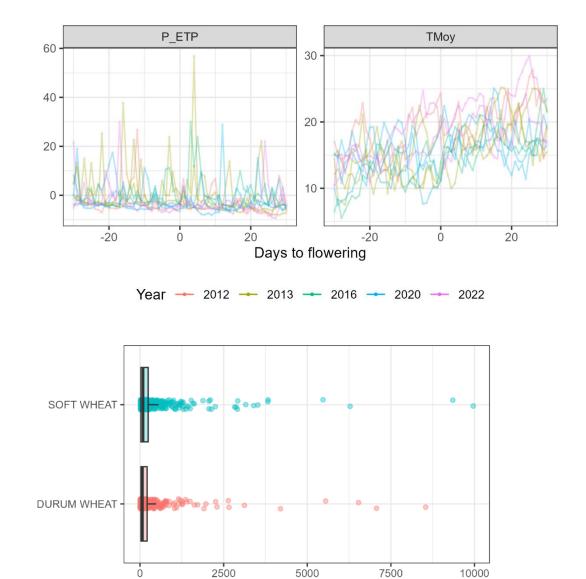


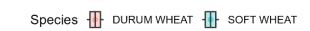
## Toxins = f(climate)?

#### Repartition of the samples

(575 durum wheat,1006 soft wheat)





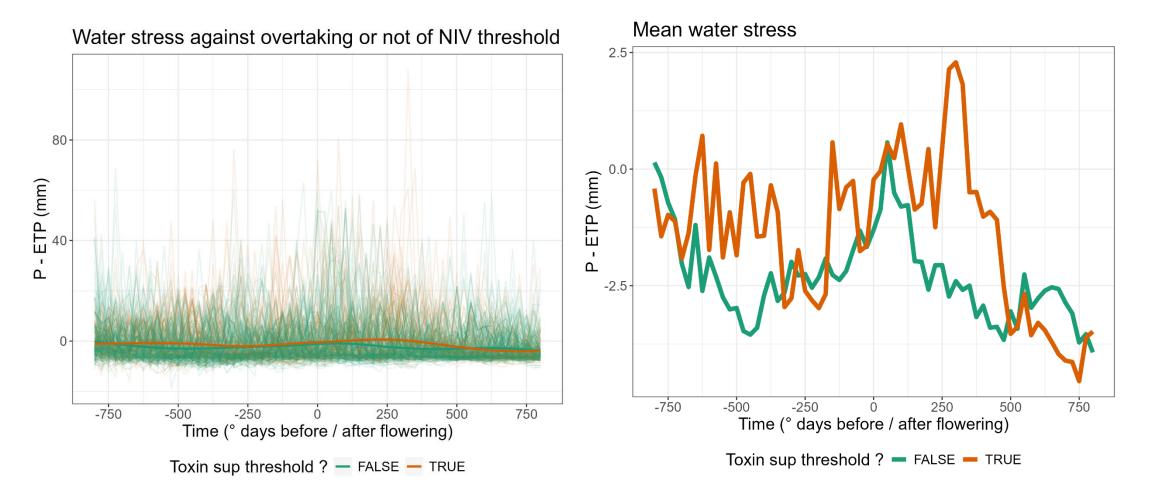


Concentration DON

## A running example

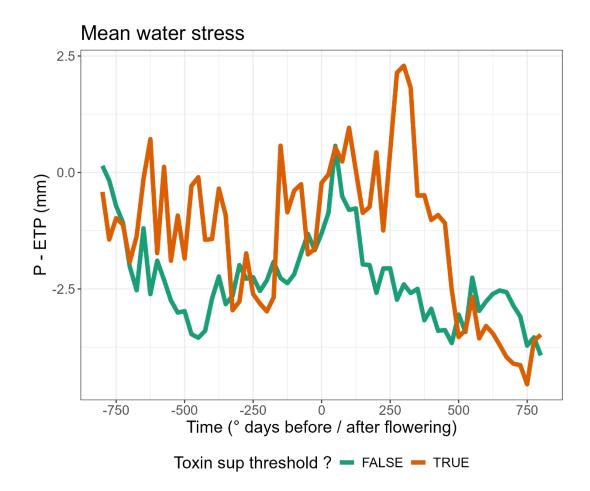
# WORK IN PROGRESS

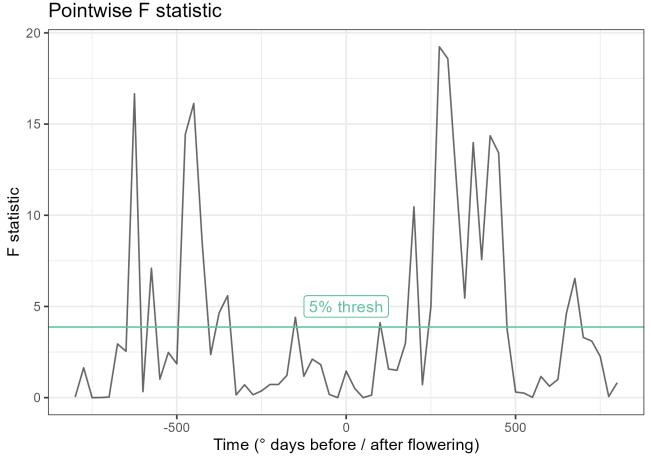
#### How to find climatic zones related to the overtaking of a given mycotoxin



## Pointwise F tests





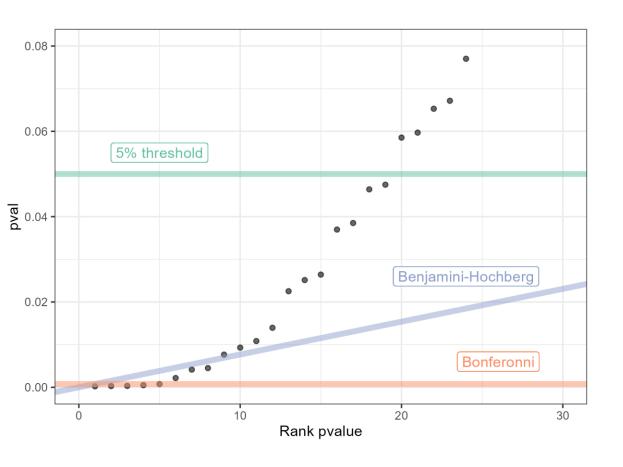


→ Just a multiple comparison problem?

## Two possibilities to consider



1) Perform pointwise Ftests and correct the pvalues with a Benjamini-Hochberg procedure



- 2) Perform a functional ANOVA (Fanova)
- a) Compute pointwise F statistics
- ) Aggregate the F statistics (sum for ex.)
- c) Distribution of the aggregated F statistics under H0 computed by random permutation technics

**Problems**: Fanova results in a **yes/no** answer, no local information and BH procedure leads to no / noisy results → Need for a compromise

#### Local Fanova: a compromise between the two approaches (on going work!)



**Idea**: find the largest intervals at which no significant effect can be detected with the functional ANOVA test

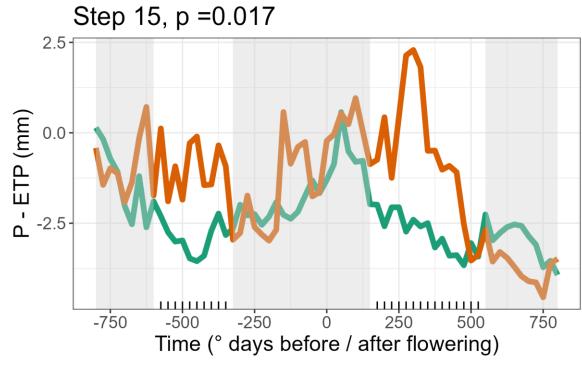
#### Inputs:

k = window size

n = number of intervals to consider

#### Algorithm:

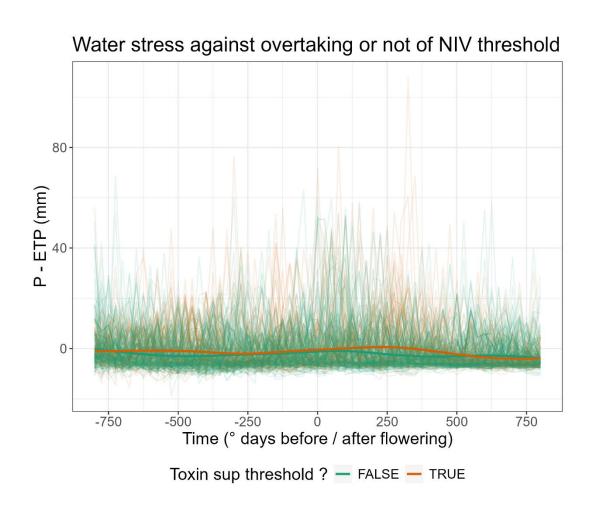
- 1. For each interval i in 1 to n:
  - a. Define a time frame centered on interval i with a window size k
  - b. Perform a Fanova on these time frames
- 2. Remove all intervals from the time frame that are significant.
- 3. While the Fanova on the remaining whole time frame is not significant:
  - a. Add more intervals to the time frame
  - b. Perform a Fanova
- c. Stop when the largest possible interval is found for which the test is not significant

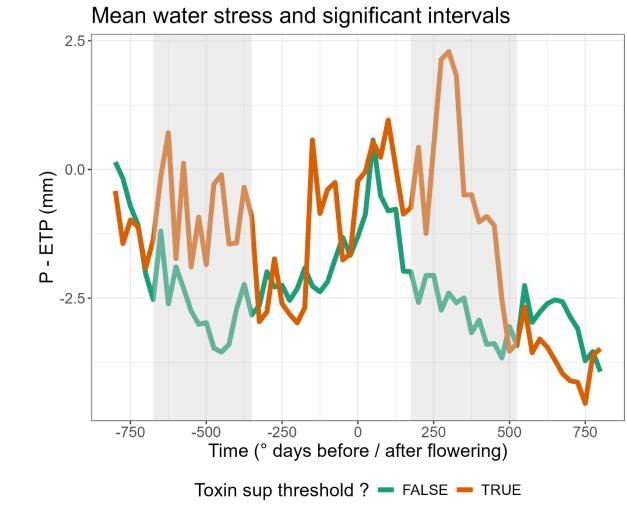


Toxin sup threshold? — FALSE — TRUE

## Results







## Perspectives





- How to set hyperparameters (window size, number of intervals)?
- Comparison with other methods (Servien et al. 2019 (SISIR/SFCB), Grollemund et al. 2018 (BLISS): function-on-scalar regression)
- Simulation study
- Link with in vitro results?

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