

## Introduction

- Flax (*Linum usitatissimum*) cultivation in Europe, counts for **80% of worldwide flax production**. This place is known for **appropriate climate conditions** to grow flax properly and **ancestral skills** in terms of cultural practices. Moreover, France and Belgium produce the **best quality fibers**.
- For the last 15 years, the observation of *Verticillium* wilt has increased in French and Belgian fields, which leads to significant **economic losses** because of the **alteration of bast fibers** by the pathogenic fungus.
- Verticillium* wilt of flax is caused by a **vascular fungal pathogen**: *Verticillium dahliae*. Long-lasting surviving structures of this fungus (**microsclerotia**) are present in natural soils. **Root exudates** of potential host plants stimulate the **germination** of microsclerotia. Resulting **hyphae** can enter the **root tissue** in the **intercellular space** (Fig. 1). Then, *V. dahliae* reaches the **xylem vessels** and **spreads systemically**.

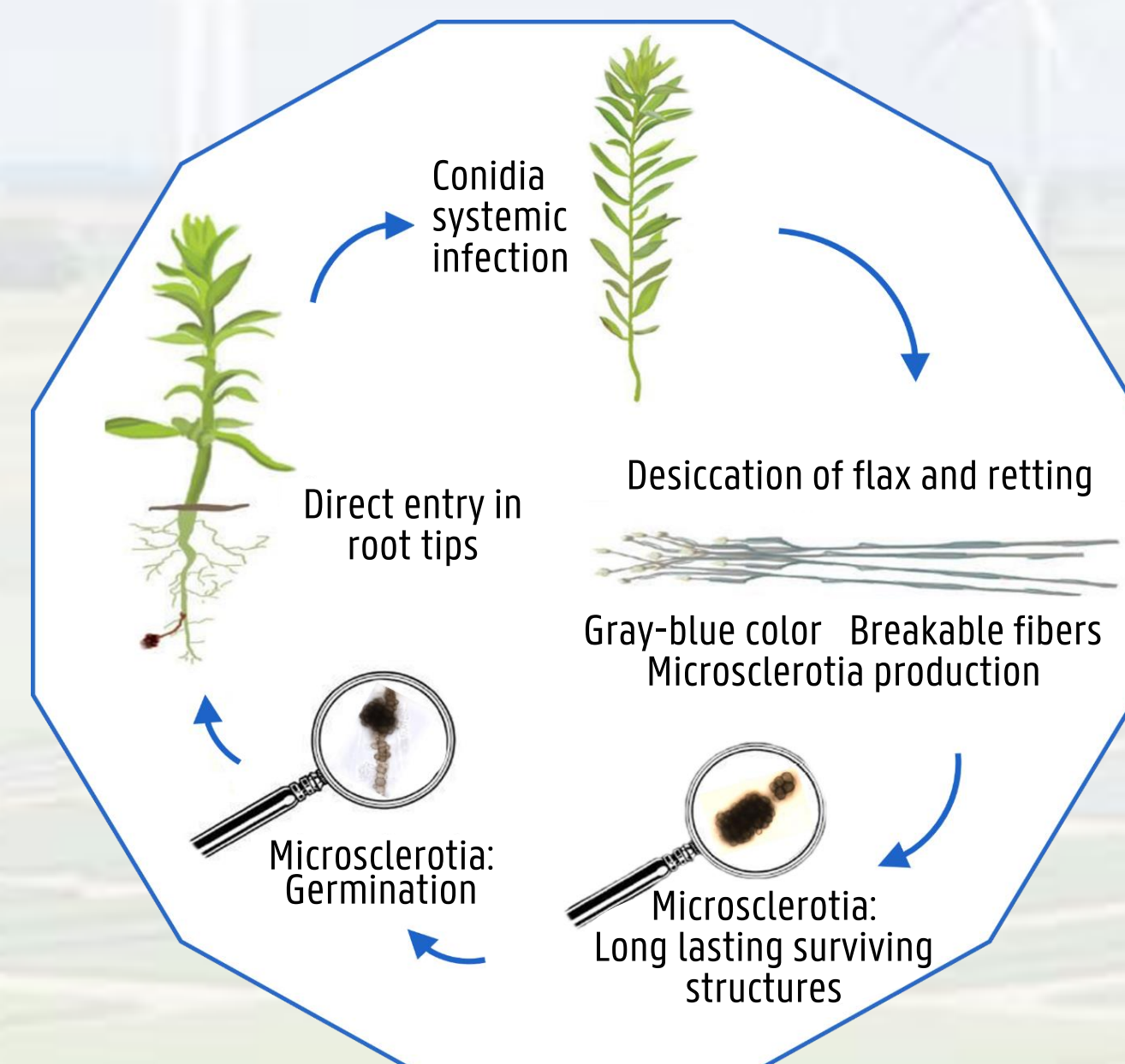


Figure 1: Infection cycle of *Verticillium dahliae* on flax (Modified from ARVALIS-Institut du végétal)

## PATHOFLAX project

This **INTERREG** project brings together **11 partners** from Hauts-de-France, Wallonia and Flanders regions, involved either in **study** or **control** of **fungal plant diseases** or **study of flax**. Launched in **January 2019** for **4 years**, its aims are to:

- Realize a **state of progress** of the disease on flax in the studied area
- Test **biological control strategies** to control the disease
- Map** *Verticillium* **occurrence** in fields soils
- Assess the **efficiency** of **elicitors** and **antagonistic strains**
- Select suitable **cultivars** and crop **management techniques**
- Organize a **laboratories network** to help farmers facing *Verticillium* wilt



## PhD thesis work: Studying the pathosystem and finding a potential biocontrol product

**INFECTION:** Setting up a **protocol of infection** of flax (cultivar Lisette, Wiersum Plantbreeding BV) with *Verticillium dahliae* microsclerotia to be **as close as possible to what happens in soil in the field**:

- Produce
- Harvest
- Test the viability of
- Mix with soil

microsclerotia (MS)

Flax seeds are sown in a mix of soil + MS powder.

### *V. dahliae*: ID card

*Verticillium dahliae* isolated in Amiens (France) (unpublished)  
**Pathogenic strain** from **symptomatic flax plants** from the field (Oise, France)  
Causal agent of *Verticillium* wilt of flax

**BIOCONTROL:** Testing a potential biocontrol strain: *V. isaacii* (Vt305) to decrease the impact of *Verticillium* wilt on flax

- Protocol:** *V. isaacii* isolate Vt305 MS (powder) + clean flax seeds (250 MS / seed)  
With *V. dahliae* microsclerotia in the soil or not

- Measuring **heights** (30 plants / condition) : (Fig. 4) : the significant **reduction in plant height** due to *V. dahliae* infection is counteracted in the presence of *V. isaacii*.
- Scoring** plants for the intensity of their symptoms : (Fig. 5)  
The addition of *V. isaacii* seems to **decrease** the **symptoms** of infected flax plants.

### *V. isaacii* / Vt305: ID card

*Verticillium isaacii* isolated in Ugent (Tyvaert *et al.*, 2014)  
Endophyte from **suppressive soil** in **cauliflower fields**  
Shows abilities to **control** *Verticillium* wilt on **cauliflower** (Deketelaere *et al.*, 2020)

**MOLECULAR QUANTIFICATION:** qPCR assays have been done on **infected** and **inoculated** or not with *V. isaacii* plants to follow the **amount** of *V. dahliae* DNA *in planta*

Early results showed a **decrease** of *V. dahliae* DNA amount *in planta* when *V. isaacii* is present (data not shown)

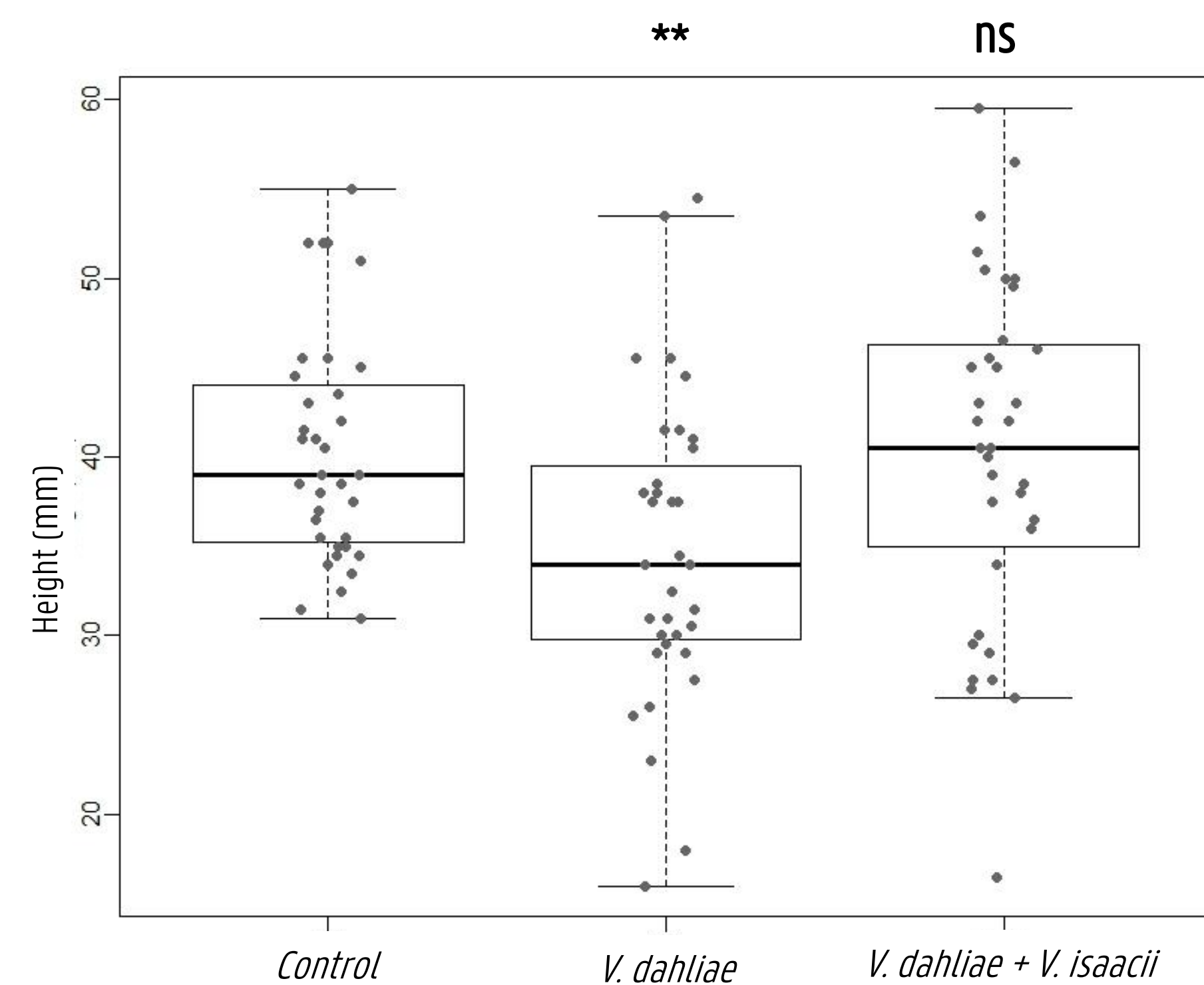


Figure 4: Heights of flax plants per condition  
ns: no significant difference  
Every dot is a flax plant

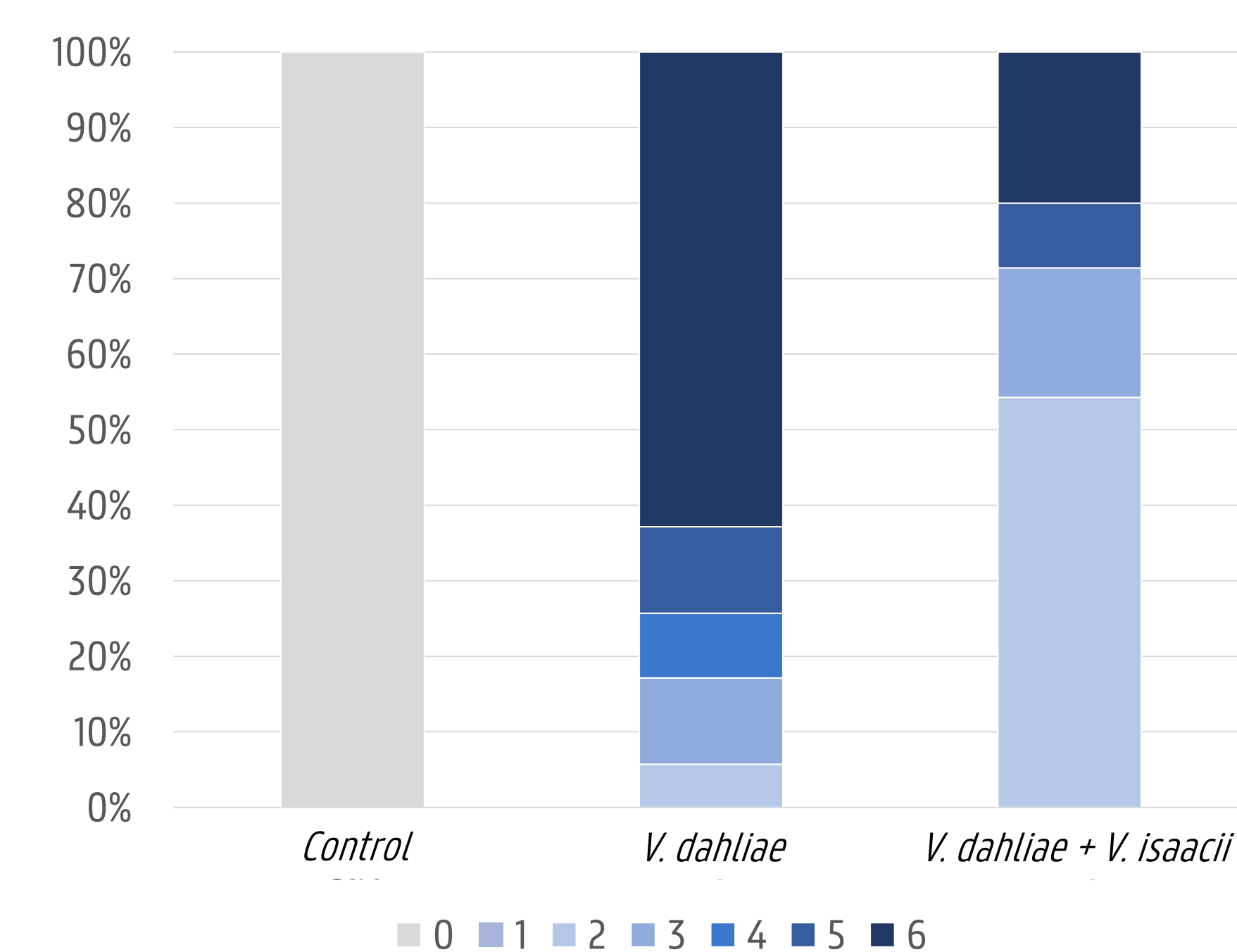


Figure 5: Disease scores of plant per condition  
*V. isaacii* alone on flax plants did not show any symptoms  
0 = no symptoms  
3 = a third of the plant presents chlorosis/necrosis  
6 = the entire plant shows chlorosis/necrosis

**SCREENING:** Searching for **new markers** of the **disease** in a **phenotyping robot** (Fig. 2):

- Automatization of watering and measures
- Fluorescence camera
- Visible top and side cameras (Fig. 3)
- Hyperspectral visible and near infrared (350 to 900 nm) and short-wave infrared (900 to 1700 nm) top cameras



Figure 2: Insight of the phenotyping robot in CRRBM of Amiens



Figure 3: Side visible camera pictures  
Left : Control flax plants  
Right : *V. dahliae* Infected flax plants

**METABOLOMICS:** Xylem sap of **infected** and **control** plants have been harvested. *Verticillium* wilt seems to have an impact on xylem sap flow and composition (unpublished)

- GC-MS** analysis has been performed, looking for **polar metabolites**
- Searching** for potential **differences** in the amounts of those metabolites between **infected** and **control** plants (Fig. 6): **white bars** mean that metabolites are **less present** in infected plants and **blue bars** mean that metabolites are **more present** in infected plants

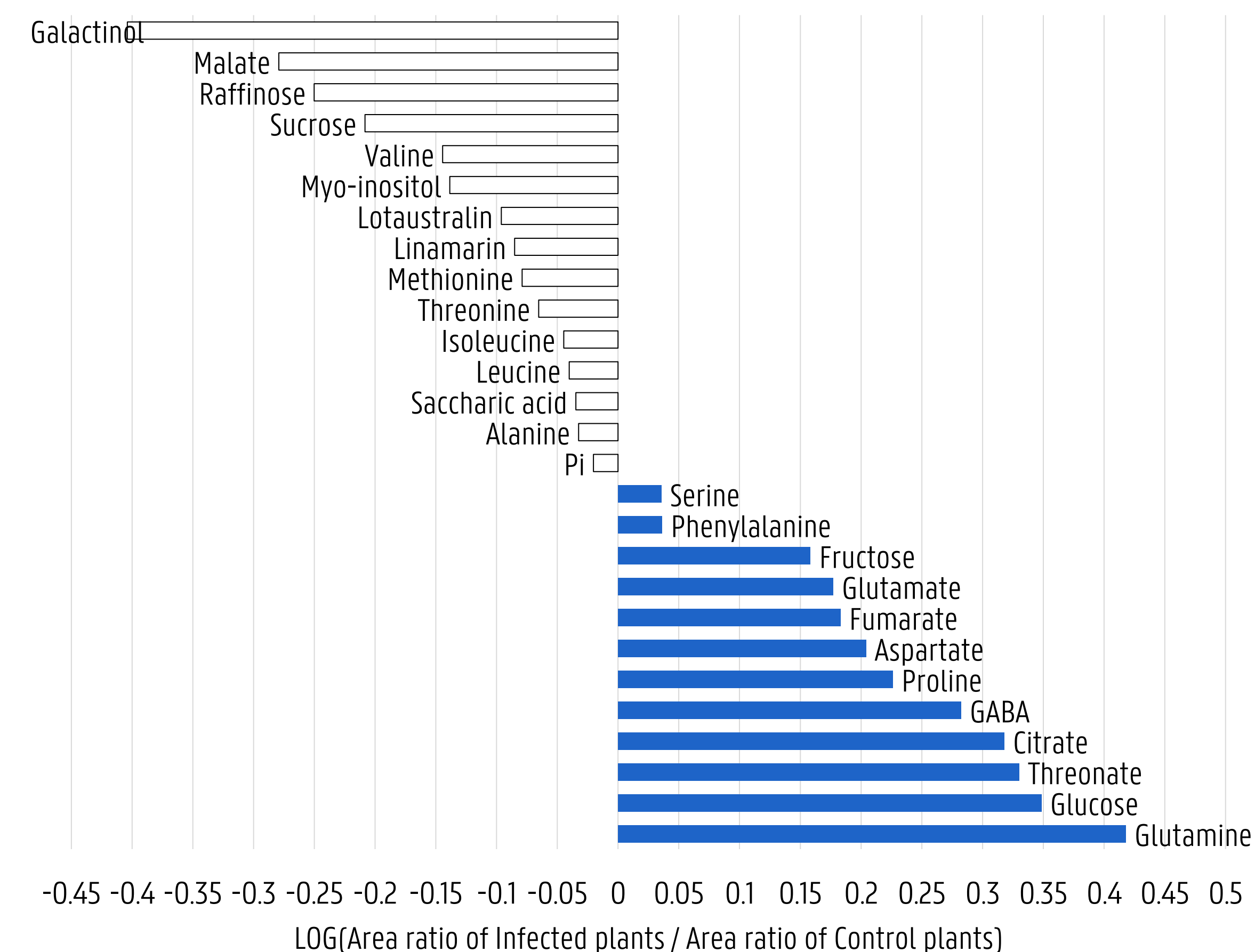


Figure 6: Metabolite accumulation in flax xylem sap after infection with *V. dahliae*  
Differences between **area ratios** of **control** and **infected** plants for each analyzed metabolite (6 control samples and 8 infected samples)

## Perspectives

- Testing *V. isaacii* efficiency in the field: → **Plants** are currently growing
- Testing *V. isaacii* efficiency on other cultivars : → **Plants** are growing in controlled conditions

- Carry on metabolomics: → **Confirm results**  
→ **Other cultivars, biocontrol conditions**