University of Hertfordshire

Life and Medical Science

Genetic basis of partial resistance against Pyrenopeziza brassicae in oilseed rape



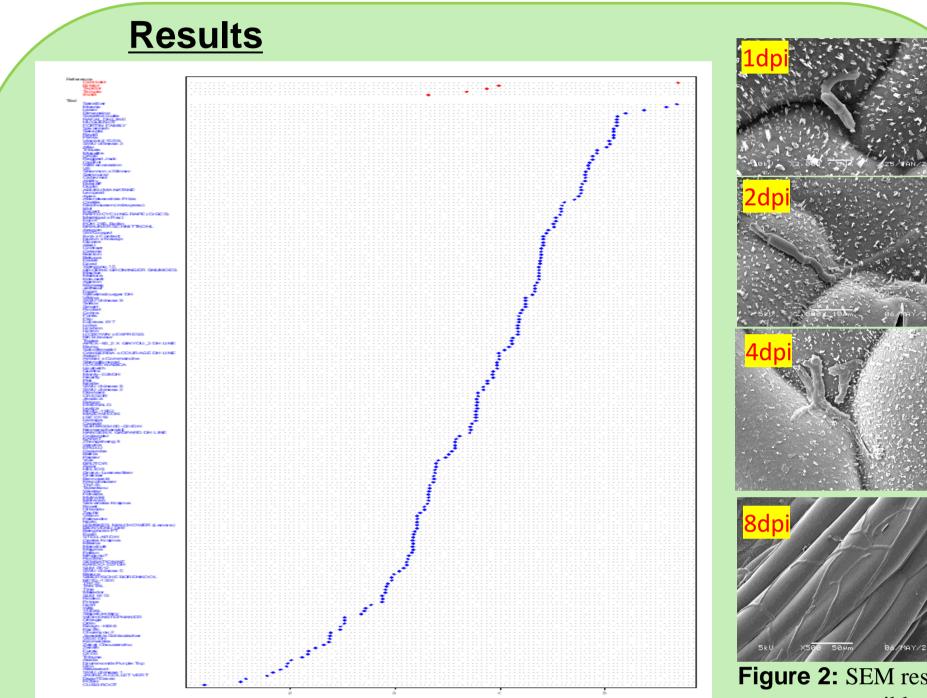
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Introduction

Light leaf spot (LLS), caused by the pathogen Pyrenopeziza brassicae, is the most damaging disease of oilseed rape in the UK.



- LLS accounts for up to £160M annual yield loss despite expenditure of £20M on fungicides.
- This disease is controlled by fungicide applications and partially resistant cultivars (AHDB recommendation lists).
- ✤ P. brassicae has developed fungicide insensitivity.
- Better understanding of quantitative resistance (QR) against *P. brassicae* in oilseed rape is important in designing improved and durable control strategy against LLS.

Aim

- To study genes involved in QR in oilseed rape against P. brassicae.
- To understand the contribution of host wax/cuticle to this pathosystem.

Objectives

- To screen *B. napus* accessions and *B. rapa* wax mutants under glasshouse conditions.
- To assess cuticle permeability of *B. rapa* wax mutants using toluidine blue staining.

Figure 2: SEM results on susceptible

Figure 1: LLS disease scoring of 195 *B.napus* accessions

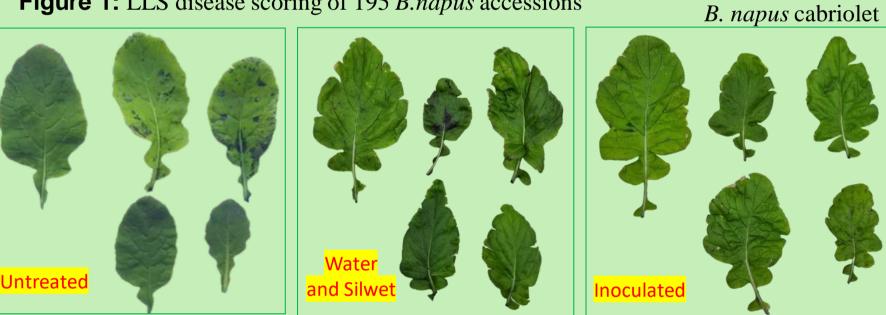


Figure 3: Toluidine blue staining with wild type R-O-18, two glaucous (GL) and non-glaucous (NG)mutants of *B. rapa* from top left to bottom

Discussion

- The four most susceptible (Dwarf Essex, Laser, Sansibar and Cabriolet) and resistant (POSH, SWU Chinese1, Cubs Root and Moana) cultivars were chosen from glasshouse results for further experiments.
- ✤ R-O-18 and NG (more wax) showed more permeability and susceptibility.
- GL (less wax) was less permeable and more resistant.

Materials and Methods

- ✤ 195 B. napus cultivars were spray inoculated with a *P. brassicae* spore suspension. Disease were scored on a scale of 1-6 (Fig.1).
- Associative transcriptomics was used to identify 8 genes that were differently expressed in partially resistant and susceptible accessions.
- Scanning electron microscopy was done to track pathogen and decide future gene expression works.
- Toluidine blue staining were done before and after inoculation with pathogen on *B. rapa* wax mutants.

Works in progress

- Gene expression using multiplex Taqman qPCR to study the expression of GEMs during the interaction with P. brassicae.
- ✤ KASP primer analysis of *B. rapa* TILLING mutants to confirm the involvement of GEMs in partial resistance against P. brassicae.
- Quantification of pathogen DNA.
- Analysis of wax and cutin quantities and compositions in pathogen challenged *B. napus* leaves with differences in susceptibility to P. brassicae and in and B. rapa wax mutants.





