

# Role of Seed-Borne Infection in Rhynchosporium Epidemics in Barley

Amarnath Thirugnanasambandam<sup>1</sup>, Adrian Newton<sup>1</sup>, Steve Whisson<sup>1</sup>, Kath Wright<sup>1</sup>, Neil Havis<sup>2</sup> and Simon Atkins<sup>3</sup>.

<sup>1</sup>Scottish Crop Research Institute, Invergowrie, Dundee, DD2 5DA.  
<sup>2</sup>Scottish Agricultural College, West Mains Road, Edinburgh EH9 3JG.  
<sup>3</sup>Rothamsted Research, Harpenden, Herts, AL5 2JQ.

Email: athiru@scri.ac.uk



Figure 1 Typical Rhynchosporium lesions.

## Background

Rhynchosporium is the most damaging disease of barley. *Rhynchosporium secalis* can complete its life cycle within a host without producing any visible symptoms. A greater understanding of the epidemiology of this pathogen, how it spreads during the symptomless phases of its life cycle, the role of seed-borne infection and subsequent proliferation and spread, will assist resistance breeding, improve guidance to farmers for control measures and help optimize fungicide application.

## Transformation

*Agrobacterium tumefaciens*-mediated transformation of *R. secalis* using two pathogenicity-characterised isolates was done by co-cultivating high concentration of *R. secalis* spores with *A. tumefaciens* in the presence of acetosyringone. The transformed colonies were selected on minimal media containing hygromycin.

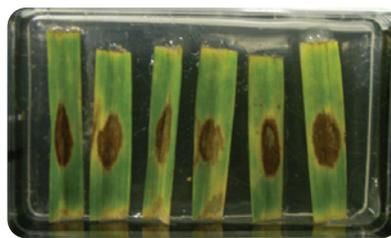


Figure 2a Symptoms produced by transformed *R. secalis* on a susceptible cultivar.

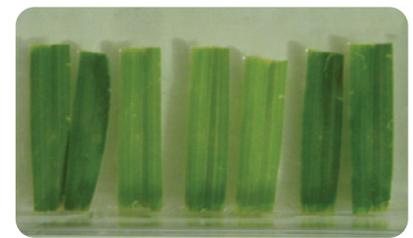


Figure 2b Symptomless infection on a resistant cultivar.

## Time Series Experiment

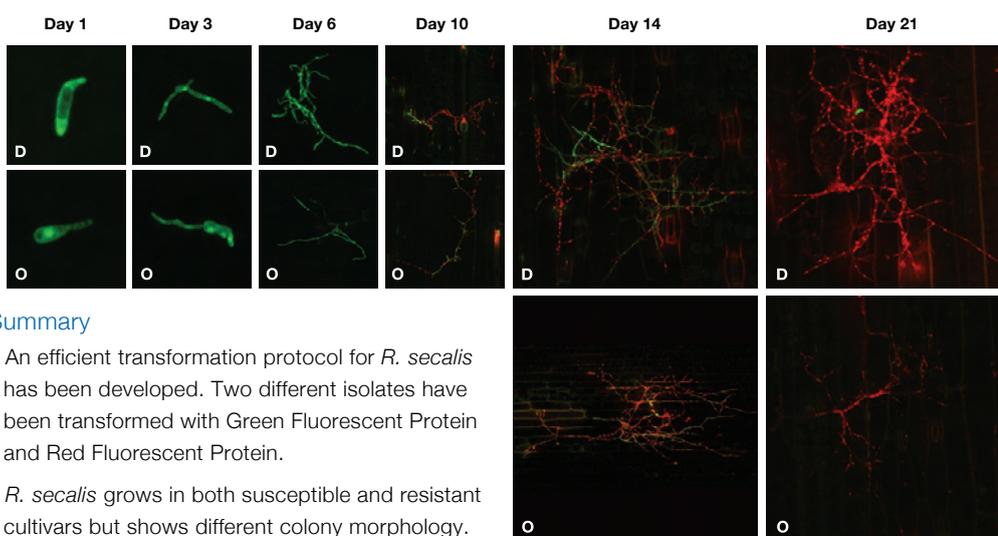


Figure 3  
 D= Digger (Susceptible)  
 O= Osiris (Resistant)

### Summary

- An efficient transformation protocol for *R. secalis* has been developed. Two different isolates have been transformed with Green Fluorescent Protein and Red Fluorescent Protein.
- *R. secalis* grows in both susceptible and resistant cultivars but shows different colony morphology.
- Use of propidium iodide during imaging helps distinguish living and senescent mycelia in infections.

## Future Prospects

- Observations made during the asymptomatic phase will help to identify and understand different Rhynchosporium resistance genes in barley and their potential use in breeding.
- Tools developed in this project will aid in identifying the mechanisms of seed-plant-seed infection.
- This study will be helpful for devising variety-specific crop protection strategies (fungicide application and timing).