



## Optimisation of deployment strategies of plant resistance to pathogens

Loup Rimbaud, Julien Papaix, Luke G. Barrett, J.J. Burdon, Peter H. Thrall

### ► To cite this version:

Loup Rimbaud, Julien Papaix, Luke G. Barrett, J.J. Burdon, Peter H. Thrall. Optimisation of deployment strategies of plant resistance to pathogens. Convergence 2017 - CSIRO Postdoctoral Fellow Conference, Mar 2017, Melbourne, Australia. , 2017. hal-02787790

HAL Id: hal-02787790

<https://hal.inrae.fr/hal-02787790>

Submitted on 15 Feb 2022

**HAL** is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

# Optimisation of deployment strategies of plant resistance to pathogens

Loup Rimbaud<sup>1</sup>, Julien Papaïx<sup>2</sup>, Luke G. Barrett<sup>1</sup>, Jeremy J. Burdon<sup>1</sup>, Peter H. Thrall<sup>1</sup>

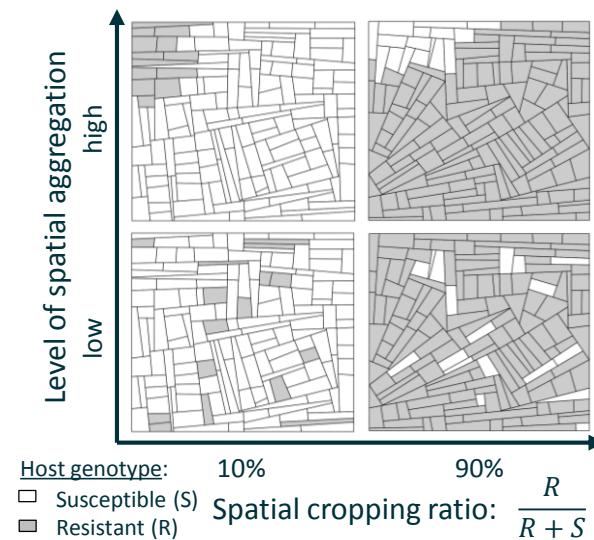
CSIRO AGRICULTURE AND FOOD  
www.csiro.au



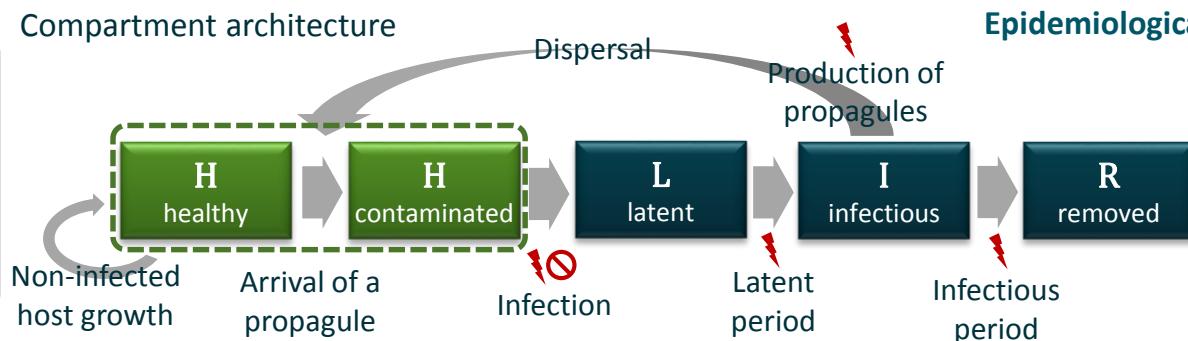
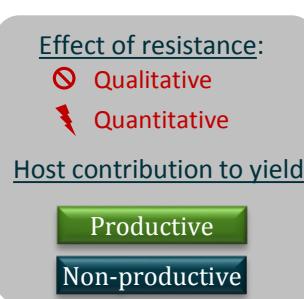
Genetically controlled plant resistance allows reduction of damage caused by pathogens. However, pathogens have the ability to evolve and overcome such resistance, resulting in significant crop losses. We developed a spatially explicit model to investigate the potential for different resistance gene deployment strategies to provide both efficient (ability to reduce disease impact) and durable (ability to limit pathogen adaptation) disease control<sup>a</sup>.

## Explicit landscape

1. Generation of a landscape of 155 paddocks using a T-tessellation algorithm
2. Allocation of different cultivars in controlled proportions and spatial aggregation

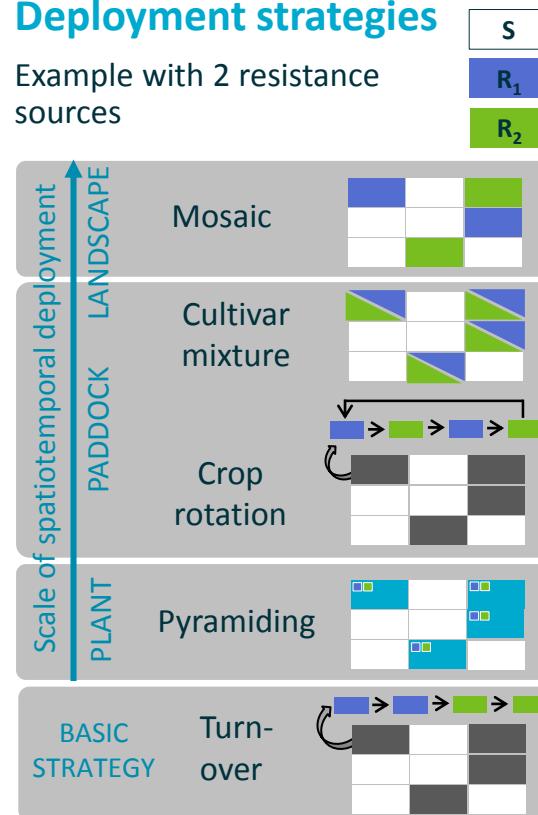


## Host response to disease



## Deployment strategies

Example with 2 resistance sources



## Pathogen adaptation

Breakdown of qualitative resistance (🚫)

| Infectivity matrix | Host genotype |            |
|--------------------|---------------|------------|
|                    | S             | R          |
| non-infective      | 1             | $1 - \rho$ |
| Infective          | $1 - \theta$  | 1          |

Cost of infectivity  
Major gene efficiency

Erosion of quantitative resistance (⚡)

| Aggressiveness matrix | Host genotype |            |
|-----------------------|---------------|------------|
|                       | S             | R          |
| not aggressive        | 1             | $1 - \eta$ |
| ...                   | ...           | ...        |
| ...                   | ...           | ...        |
| Fully aggressive      | $1 - \eta$    | 1          |

Trade-off strength:  $\beta$   
QR efficiency

Key parameters estimated through literature or laboratory experiments

## Evaluation criteria

Example of a mosaic with 1 resistant cultivar<sup>b</sup>

- Evolutionary**  
**Epidemiological**
- Durability (D)
  - Short-term control (STC)
  - Long-term control (LTC)

