Understanding selection of fungicide resistance using blackleg of canola as a model

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What we do and don't know about fungicide resistance

What we do know:

- Wide-spread DMI resistance across Australia
- Frequency within populations ranges from <0.05-32%

What we don't know:

- What frequency leads to field failure?
- What practices lead to fungicide resistance?

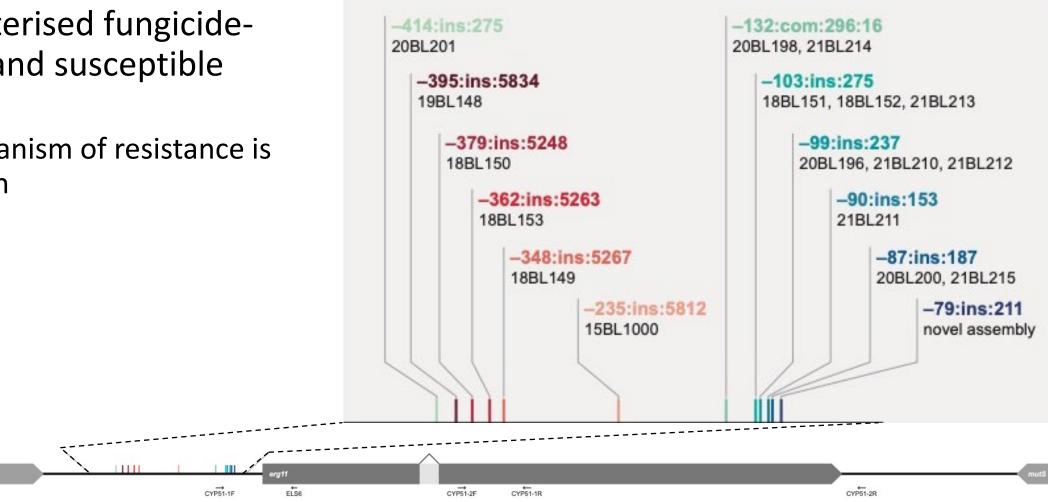
Flutriafol 44 80 All regions 188 NSW Central 2 **NSW Vic Slopes** 31 24 Vic High Rainfall 9 26 7 SA Vic Mallee 3 SA Vic Bordertown-Wimmera 10 20 43 SA Midnorth-Lower Yorke Eyre 10 21 48 WA Sandplain 3 4 20 WA Central 16 4 2 25 50 75 100 0

Blackleg as a model for understanding fungicide resistance

1. Characterised fungicideresistant and susceptible isolates

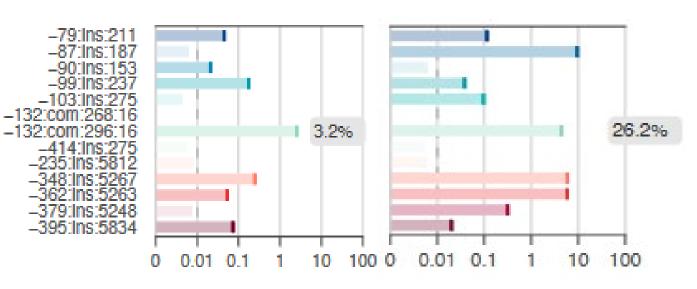
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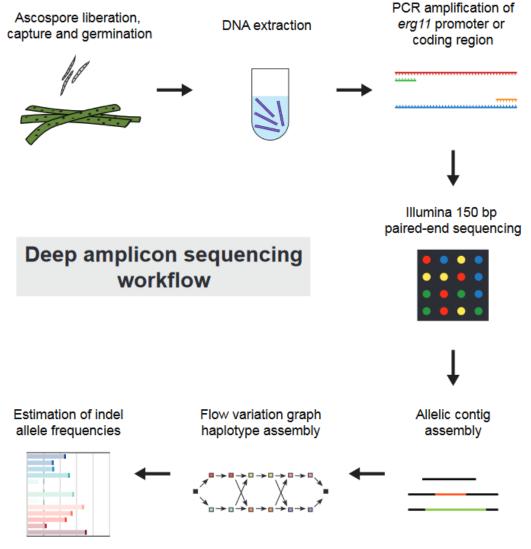
 Mechanism of resistance is known



Blackleg as a model for understanding fungicide resistance

- 2. Molecular markers for tracking changes in populations
 - Markers are applied to whole populations, not individual isolates
 - Captures all mutations





Blackleg as a model for understanding fungicide resistance

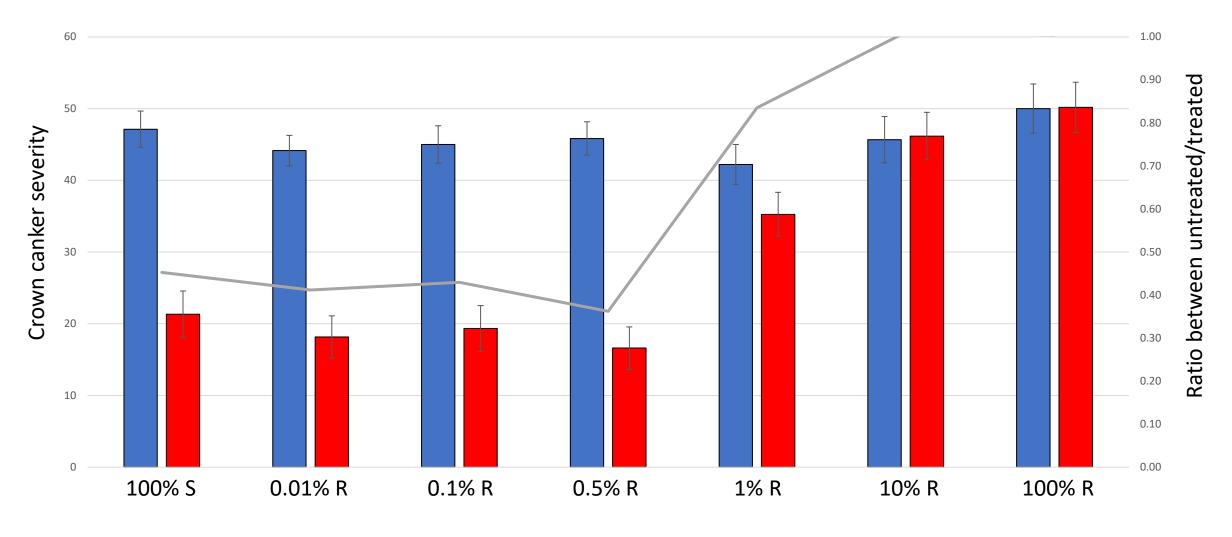
- 3. *in planta* assays to simulate different selection regimes
 - Use ratios of different isolates
 - Inoculations at different growth stages
 - Grow through to maturity and allow sexual reproduction to occur on stubble



What frequency of resistance is needed to render the fungicide ineffective?

- Untreated and Jockey-treated plants
- Inoculated with populations of isolates with different ratios of fungicide resistance
- Inoculated at multiple growth stages to simulate field conditions
- Assessed disease severity at the end of the year
- Looked at changes in allele frequency following sexual reproduction

Only 1% of the population needs to be resistant for loss of fungicide efficacy



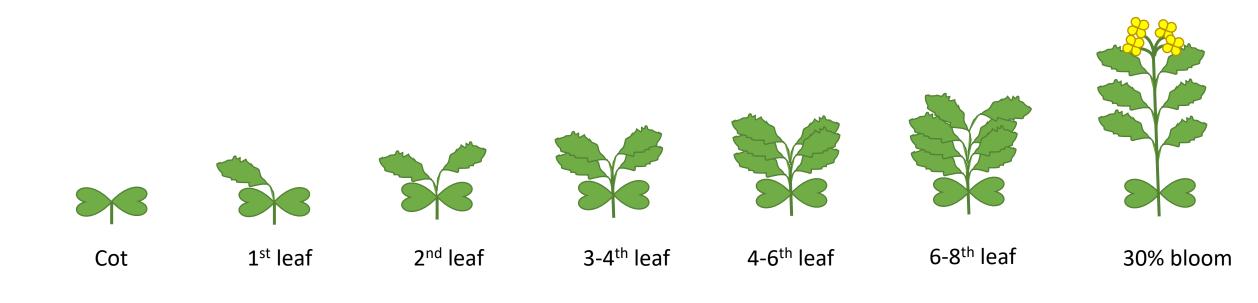
🔳 Untreated 📕 Jockey — Ratio

Populations change dramatically after selection



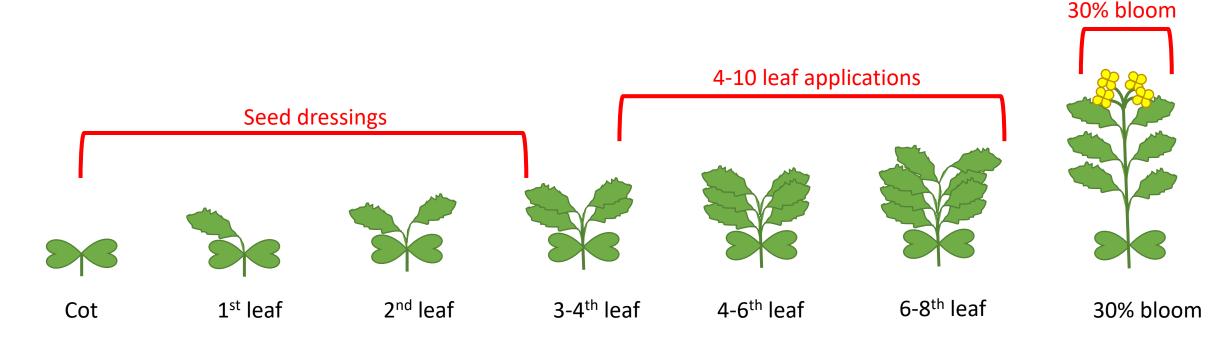
How does timing of infection contribute to fungicide evolution?

• Do later infections have time to grow in the plant and contribute to the next generation?



How does timing of infection contribute to fungicide evolution?

- Do later infections have time to grow in the plant and contribute to the next generation?
 - If not, then do later fungicide applications matter for fungicide resistance management?



How does timing of infection contribute to fungicide evolution?

 Plants inoculated with two different populations at different growth stages

3-4th leaf

4-6th leaf

6-8th leaf

30% bloom

- Fungicide susceptible •
- Fungicide resistant

1st leaf

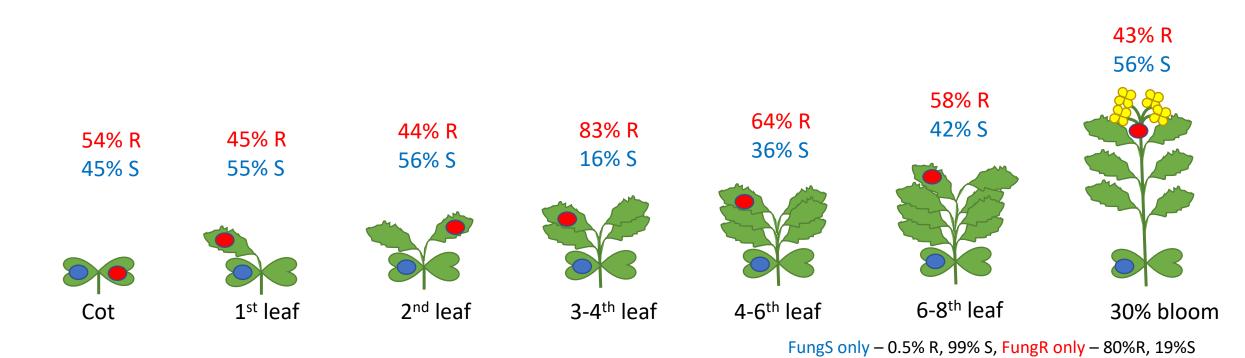
Cot

2nd leaf

• Plants grown to maturity, assessed for disease and stubble kept

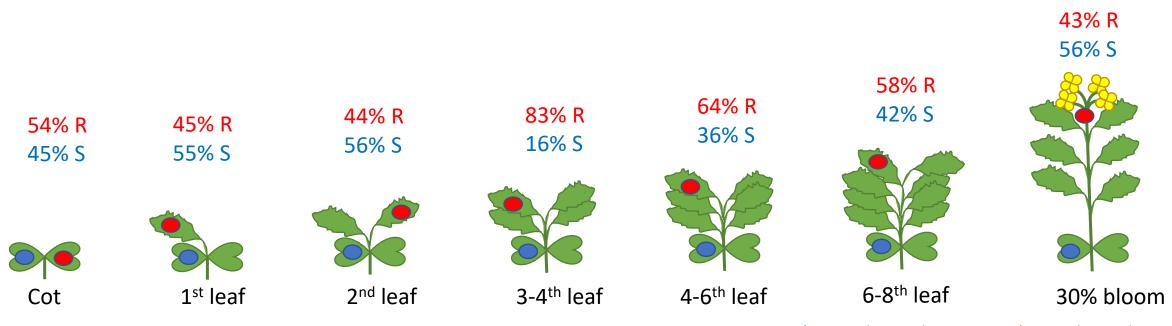
Sexual reproduction detected from all inoculation timings

- No fungicides applied to this experiment
 - Fungicide resistance used as a marker for tracking populations



Sexual reproduction detected from all inoculation timings

- Populations were designed to be able to have sex within itself
 - Positive control for experimental design



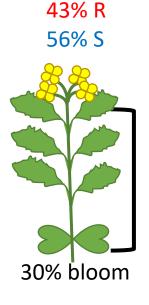
FungS only – 0.5% R, 99% S, FungR only – 80%R, 19%S

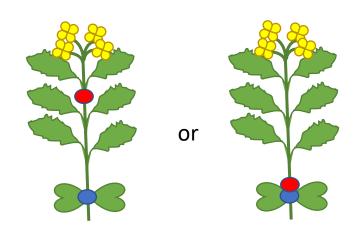
Have the populations mated together or individually?

• Capturing of spores was done with entire stem

Stubble kept

 Has the 30% bloom FungR population had sex with itself or with the FungS population?





Findings

- Only 1% of the population needs to be resistant for field failure to occur
 - Monitoring strategies need to be sensitive
- One year of selection increases the frequency of resistance dramatically
- More work around timing of infection is required to determine where the sexual reproduction is occurring







