

UNIVERSITY **OF SOUTHERN** QUEENSLAND

Pathogenicity and aggressiveness of Macrophomina phaseolina isolates to sorghum in Australia's northern grains region

<u>DL Adorada</u>, EE Adorada, PV Gonzales and AH Sparks University of Southern Queensland, Centre for Crop Health, Toowoomba QLD 4350, Australia E: <u>dante.adorada@usq.edu.au</u>

What's the issue?

- Causes major sorghum stalk rotting, which can lead to plant lodging
- *M. phaseolina*, a soilborne pathogen, causing charcoal rot in more than 500 crop species
- Splitting sorghum stalks will show ash grey tissue or microsclerotia, the survival structure of the fungus, giving the internal stalk tissue a peppered look



Withhold

irrigation

stress

Internal tissue rot

Peppered look

Lodging

- Common during seasons with prolonged hot, dry weather or when other unfavourable environmental conditions stress the plant.
- Despite the lack of formal quantification in Australia, significant yield losses have been associated to prevailing hot dry conditions resulting to widespread high incidences of charcoal rot and subsequent lodging
- The present work aims to compare pathogenicity and aggressiveness of isolates, from sorghum and other hosts from the northern region, to sorghum

Methodology

Ash grey tissue







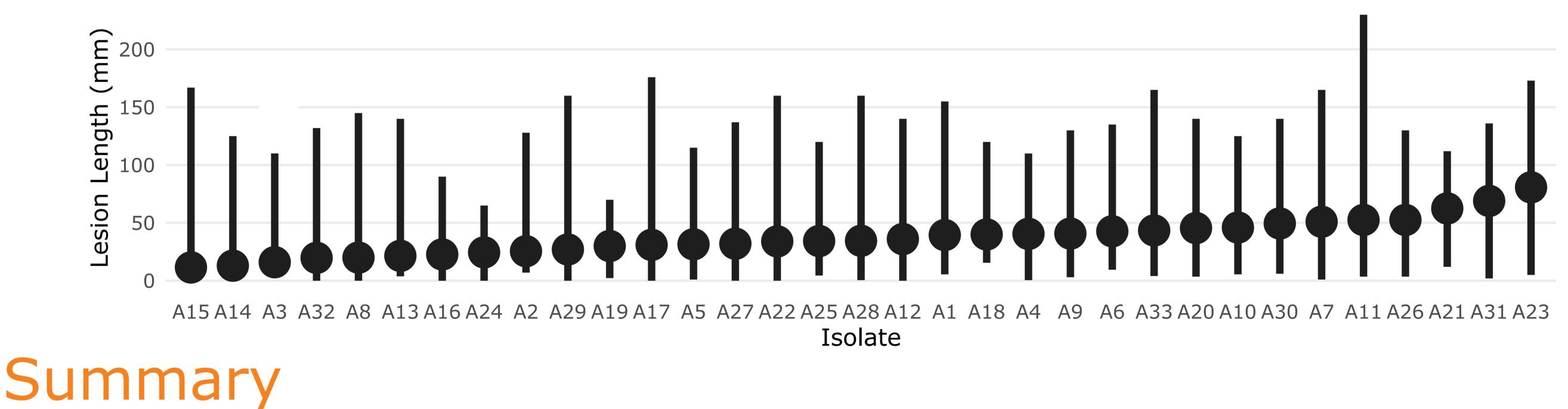


Grow sorghum cv. MR-Bazley in sterilised soil

Results

M. phaseolina-infested toothpick inoculation (33 isolates from sorghum and other hosts from CQ, SQ & NNSW) in 4 replications, repeated trial

Split open each stalk and measure charcoal rot lesion length



• In both trials, all isolates were pathogenic and capable of causing disease.



- In both trials, there are no statistically detectable differences
 - in lesion length due to the **effects of**:
 - Isolate, ullet
 - **Region** that the isolate originates from, or
 - **Host** that the isolate originates from.

This result has implications in the identification of sources of resistance to the charcoal rot disease, as well as in crop rotation decision-making in an integrated disease management programme.

Acknowledgements

We thank Sriram Padmanaban, Peter Buyoyu, Laura Baartz, and the agronomists from CQ, SQ and NNSW, for their contributions in this investigation, and GRDC Project DAQ00186 for the funding support.

