Exploring combined stress incited disease dynamics of chickpea x dry root rot interaction

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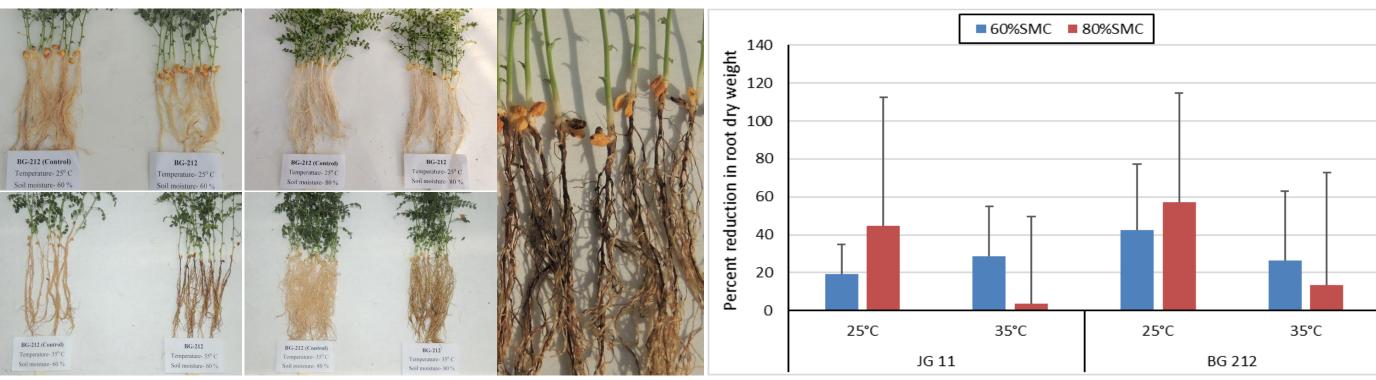
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Background

The incidence of dry root rot (DRR) caused by *Rhizoctonia* bataticola (Taub.) Butler (Rb) [Syn: Macrophomina phaseolina (Tassi.) Goid] is very high towards later stages of the crop, which incidentally in the semitropics, coincides with arid higher atmospheric and soil



temperatures and a decrease in **A DRR infested chickpea field**



Root symptoms of DRR at 28 DAS

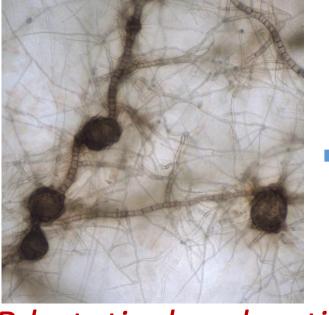
Reduction in root dry weight

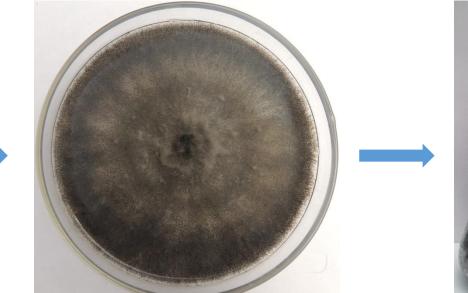
- Rb colonization was observed to increase with increasing time points in both cultivars
- Colonization of Rb was realized in both cultivars irrespective of temperature and SMC

soil moisture content. In the near future, increase in average temperature and inconsistent rainfall patterns resultant of changing climatic scenario is strongly believed to exacerbate the disease to epidemic proportions. The present study aims to quantify the collective role of temperature and soil moisture on disease progression in chickpea under simulated environmental conditions.

Controlled environmental setup

1. Development of sick soil





R.bataticola sclerotia



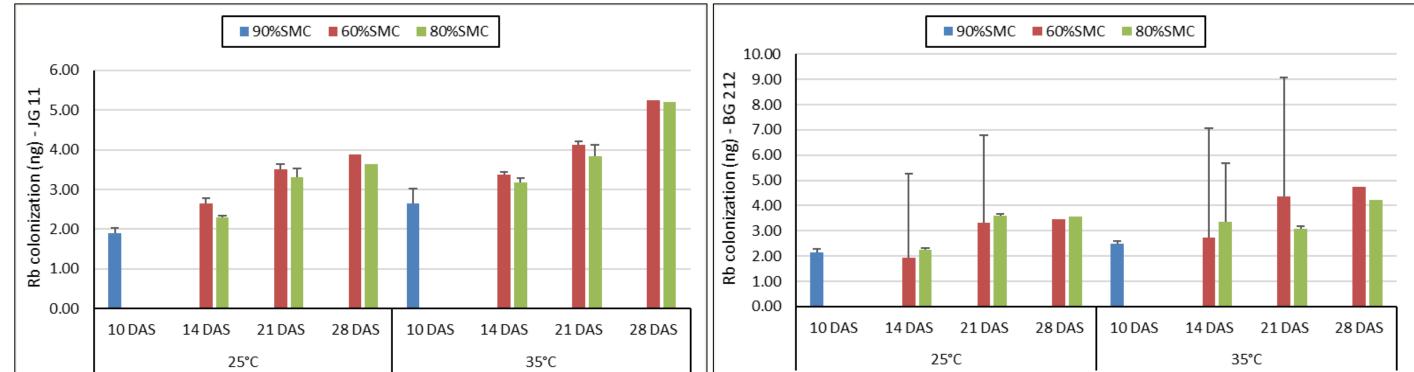
Mass multiplication Inoculum mixing

PGC 2

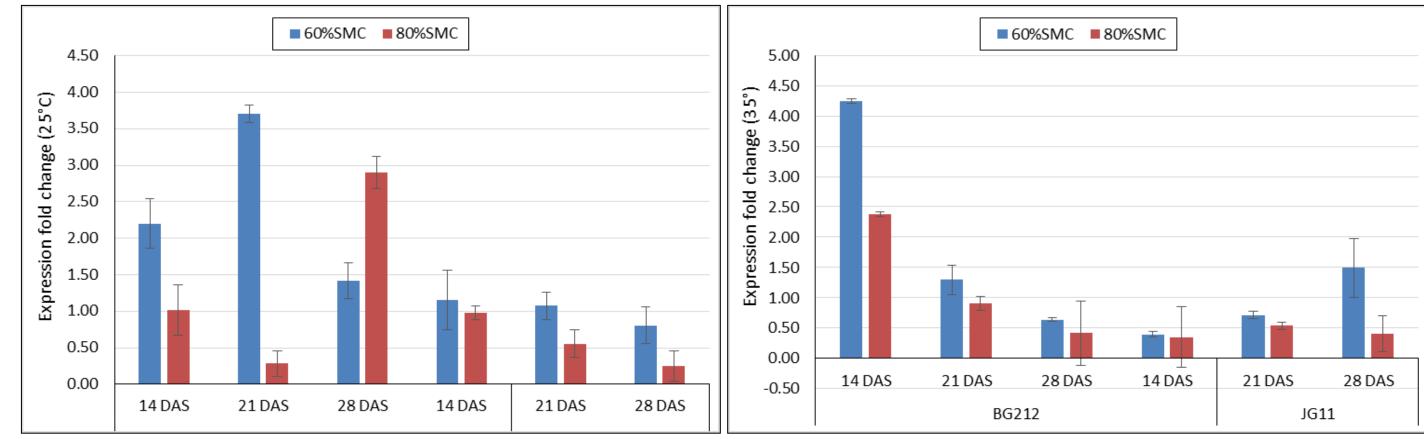
- Soil moisture condition (SMC) 60% and 80%
- Plant growth chambers (PGC) of Conviron having gradients for 3. temperature and relative humidity (RH) - Controlled environment research facility (CERF), ICRISAT.

Time

Pathogenesis-related (PR) genes responded with significant difference under different temperatures and soil moisture



R.bataticola colonization in host root tissue (JG 11 and BG 212)



Differential gene expression of β -1,3 endoglucanse (PR-2 gene) at 25°C and 35°C





					2 5
00.00	22	60	12	60	4 change
06.00	25	60	18	60	
10.00	30	50	22	60	on fold
12.00	35	50	25	60	essi
15.00	30	50	22	60	EX DI
18.00	26	60	18	60	0
23.59	22	60	12	60	
radiants sat in DCC					

RH (%)

Conviron Plant Growth Chamber

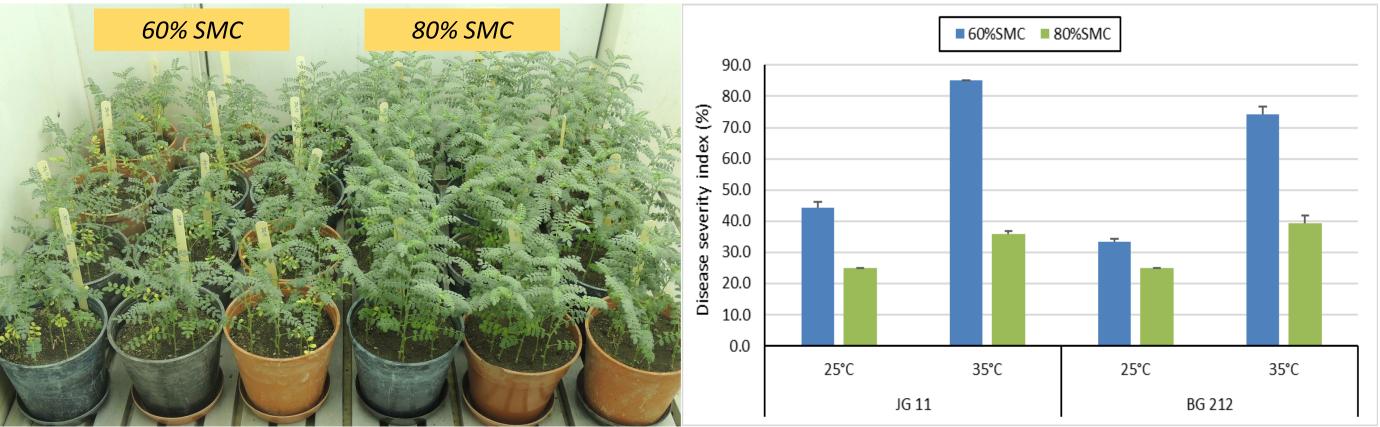
Gradients set in PGC

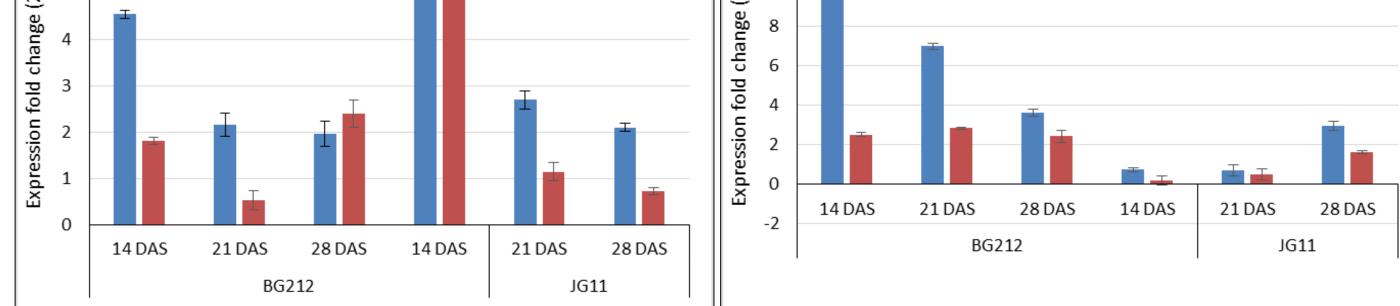
PGC 1

Temp (°C)

Results

- A high positive correlation (R²= 0.68) was found between disease severity and temperature
- * The disease severity showed high negative correlation (R^2 = -0.71) with SMC
- A significant reduction in root dry weight was also observed in infected plants





Differential gene expression of chitinase, CH III (PR-3 type) at 25°C and 35°C

Conclusion

- As evident from the disease severity index, a high temperature and low soil moisture plays a crucial role in pre-disposing chickpea to DRR
- Since Rb colonization occurred at all the temperatures and soil moisture range tested, it is to be assumed that regulation of different PR genes is responsible for the differential expression of disease symptoms
- Climate change incited disease epidemic scenario is possible in near future due to the nature of disease dynamics. Hence, further studies in patho-genomics and transcriptomics is necessary

Acknowledgment

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Foliar symptoms of DRR at 28 DAS Disease severity index

