INVESTIGATING THE INTERACTION BETWEEN CROP AGE AND TIMING OF COTTON LEAFROLL DWARF VIRUS INOCULATION ON DISEASE SEVERITY AND YIELD LOSS

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Introduction

Cotton leafroll dwarf virus (CLRDV) (genus Polerovirus, family Luteoviridae) is an emerging plant virus in the U.S. cotton belt that was first identified in samples collected from Alabama in 2017 (Alevar et al., 2018). CLRDV was first identified in Africa in 1949 (Cauquil, 1977.) and has since been found in Argentina, Brazil, India, Thailand, and Timor-Leste (Corrêa et al. 2005, Mukherjee et al. 2012., Sharman et al. 2015., Ray et. Al 2016). In the U.S., CLRDV has been reported in NC, SC, GA, AL, MS, and TX (Aboughanem-Sabanadzovic et al. 2019, Alabi et al. 2020, Ayelar et al. 2018, Huseth et al. 2019, Tabassum et al. 2019, Wang et al. 2020). Further testing has confirmed this to be genetically distinct from CLRDV isolates in South America (Avelar et al. 2020). CLRDV is a positive single stranded RNA virus. The known vector of this virus is the cotton aphid (Aphis gossypii). The cotton aphid transmits the virus in a persistent, circulative manner. Alate aphids are reported to transmit in 40 seconds and can retain the virus for up to 12 days (Michelotto and Busoli, 2003, 2009). The geographic range of the cotton aphid in the U.S. extends across the cotton belt, and infestations occur annually (Abney et al. 2008, Gore et al. 2013, Kerns et al. 2015). The symptoms attributed to the disease caused by CLRDV in the U.S. are highly variable and include: leaf distortion, red tinting of the leaves, greenish-blue leaf color, red veins, red petioles, red stem, upward cupping, downward cupping, drooping, vein clearing, shortening of internodes, abnormal top growth, square abortion, and reduced size of bolls (Brown et al. 2020, Hagan et al. 2019). There is some data showing variation of CLRDV symptoms among cultivars (Brown et al. 2019), but based on our understanding of plant viruses we know that there are a variety of factors that can influence symptom development including plant age at the time of infection, interactions with other abiotic and biotic plant stress factors, and environmental conditions. The objective of this study was to compare disease severity and yield loss among cotton plots that were infected with CLRDV at three different growth stages.

Methods

In 2019, a field study was conducted at the E.V. Smith Research Center in Shorter, Alabama. Cages made of thrips-proof netting were used to cover cotton plots to confine insects released into the plots, and exclude natural infestations of early-season insects. Two-row plots that were 20' long were planted with DP1646 on May 30. The day after plant, plots were covered with insect cages, and all plots were caged for the same amount of time to reduce cage-related effects on symptom expression and yield across treatments. A randomized complete block design with 4 replicates was used to investigate the following treatments: 1) Control plots that were not infested with aphids; 2) Control plots in which non-viruliferous aphids were released (aphids only, no CLRDV); 3) Plots infected with CLRDV by releasing viruliferous aphids 1 week after emergence at the 3-4 true-leaf stage; 4) Plots infected with CLRDV by releasing viruliferous aphids 2 weeks after emergence at the 5-7 true-leaf stage; 5) Plots infected with CLRDV by releasing viruliferous aphids 3 weeks after emergence at the first-pinhead growth stage. Viruliferous aphids that were released into plots were generated by infesting CLRDV-infected cotton plants with *Aphis gossypii* 2-3 weeks prior to release. Plants infested with aphids were grown in a greenhouse and in cages made from thrips-proof screen until they were

placed in field cages. In field cages alates were allowed to disperse for one week, and infested leaves were distributed across the plot to promote rapid colonization and virus transmission. Non-viruliferous aphids reared on healthy cotton in a separate greenhouse were reared and released the same. Four border rows that were not covered in cages were planted to monitor natural infection of CLRDV at this location. Aphids were eliminated from infested plots by spraying insecticide 2 weeks after infestation, and once the week before cage removal to prevent unintended virus spread among plots. Cages were then removed 8 weeks after plant. Ten plants were marked in each plot and monitored weekly for aphid numbers, plant mapping, and symptom presence. Plant mapping consisted of plant heights, presence/absence of 1st and 2nd position bolls, first fruiting node, and number of total nodes. Weekly spectral imaging was performed to determine early detection of disease. Virus was confirmed by Auburn's plant disease diagnostic lab in August, and yield was evaluated for each plot.

Results

PCR testing confirmed virus transmission in these plots, but not all plants tested positive for the virus. Analyses are underway to examine multispectral imaging data, plant mapping data, CLRDV symptoms, and aphid numbers. Preliminary analyses conducted on plot yield showed a trend in which the earlier a plot was infested with viruliferous aphids, the lower the yield. Non-infested plots and plots infested with non-viruliferous aphids (control plots) were numerically higher than plots infested with viruliferous aphids. Yield was significantly lower in plots infested with viruliferous aphids on week one, than in control plots (P=0.05) (approx. 400 lb. difference). Preliminary analyses of plant mapping data show that there are no significant differences in the number of first and second position bolls between plants testing positive for CLRDV and those that tested negative for CLRDV (P=0.05). The preliminary data from this one-year study show that plant age at time of infection impacts yield, and suggests that early CLRDV infection impacts lint production. This study will be repeated in 2020.

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