

College of Veterinary Medicine UNIVERSITY OF GEORGIA

Associations between biotic and abiotic factors and Chagas disease vector abundance in palm trees across different habitat types

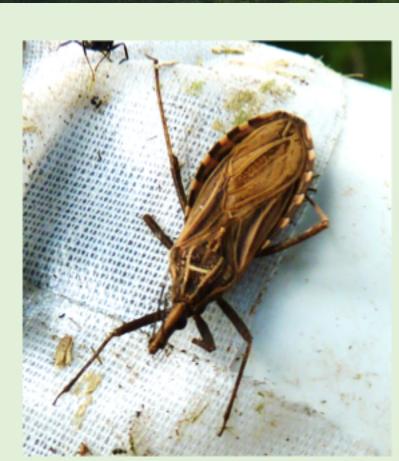
INTRODUCTION

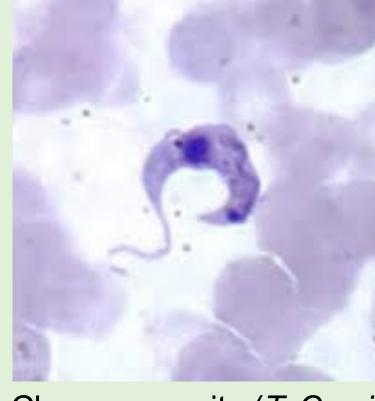
vector-borne disease is Chagas а infectious disease the caused by protozoan Trypanosoma cruzi, transmitted by triatomine insect vectors known as "kissing bugs" (*Rhodnius pallescens*).

Around 8 million people are infected with Chagas disease in the Americas, which can be life threatening if not treated.

In Panama, sylvatic transmission of T. cruzi commonly occurs in the crown of the Attalea butyracea palm, where kissing bugs live, but can often spillover into human populations when infected vectors come into contact with humans.

Previous studies^[1,3] demonstrate that land use change (e.g. deforestation) increase *pallescens* abundance, but the R. underlying mechanisms driving vector abundance are unknown.





Chagas parasite (*T. Cruzi*)



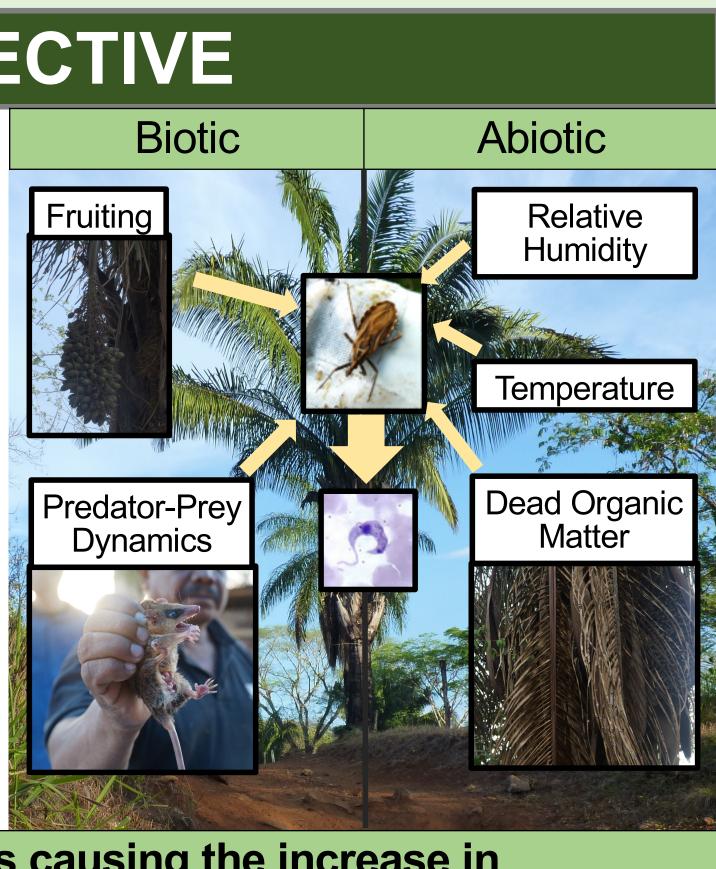
OBJECTIVE

To evaluate how biotic and abiotic factors influence R. pallescens abundance in palm trees across different types land cover and locations geographic in Panama.

(Right) Interaction diagram of mechanisms that play a role in determining kissing bug abundance and ultimately Chagas disease transmission.

Potential mechanisms causing the increase in *R. pallescens* abundance across habitat types

Mechanism	Scale		Expected effect on abundance
1. Dead Organic Matter (DOM)	Microenvironment		More space
2. Relative Humidity (%)	Microenvironment	~	Maximizes
3. Temperature (C°)	Microenvironment	~	Maximizes at
4. No. of Connected Trees	Microenvironment		More space, attrac
5. Fruiting	Microenvironment		Attrac
6. Predator biomass (g)	Community	₩	Increa
7. Prey-insect biomass (g)	Community		Increases food supply
8. Species Richness	Community	₩	Biodiversity regula
9. Habitat Type	Ecosystem		Increases anthropogen
10. Location	Ecosystem	2	Varie



Jason Soriano^{1,2}, Christina Pilar Varian¹, Nicole Gottdenker¹ ¹Department of Pathology, College of Veterinary Medicine, University of Georgia ²University of California, Berkeley

Kissing Bug (*R. pallescens*)

characteristic marker of acute Chagas disease

n vector

ce, attracts prey s at optimal RH t optimal Temp acts vertebrates acts vertebrates eases predation y for predators ates population nic disturbance es with location



- Collected dry weight biomass measurements of hundreds of individual invertebrate specimens caught directly from palm tree crowns.
- Assessed species diversity of 50 palms.
- the species relationships (predator/prey) within the palm community.

RESULTS



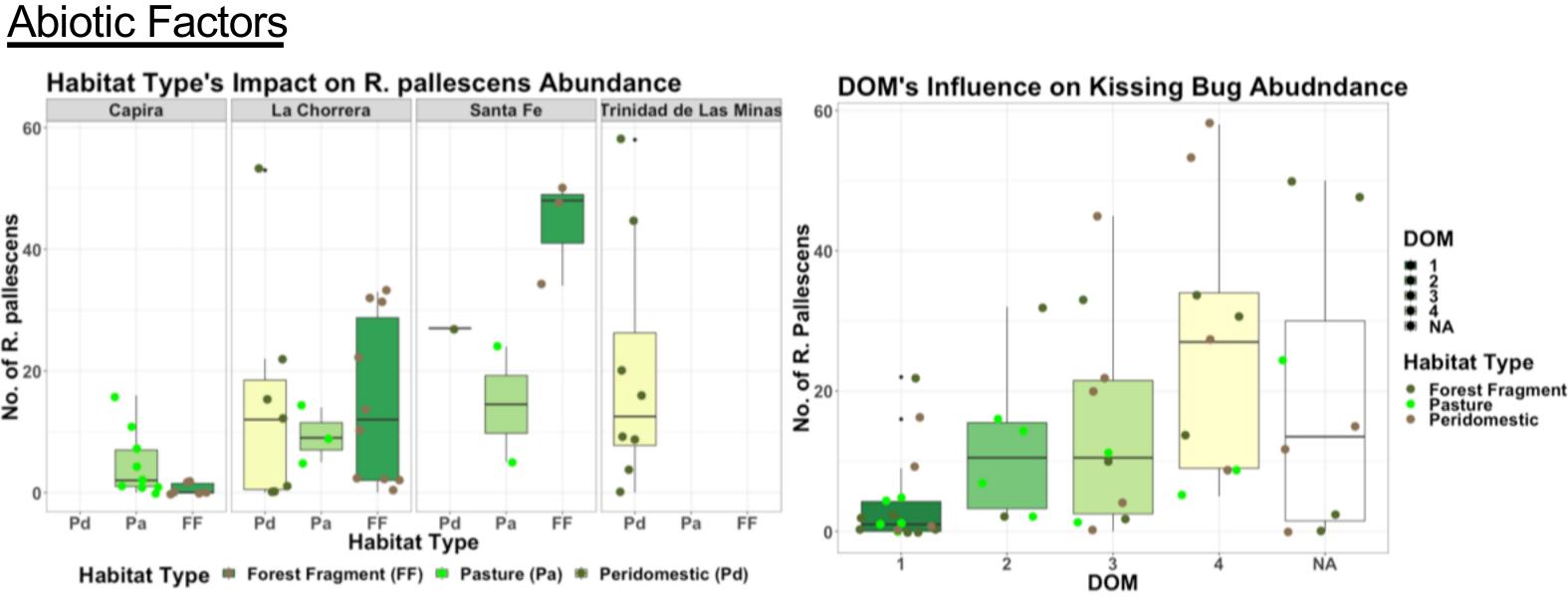
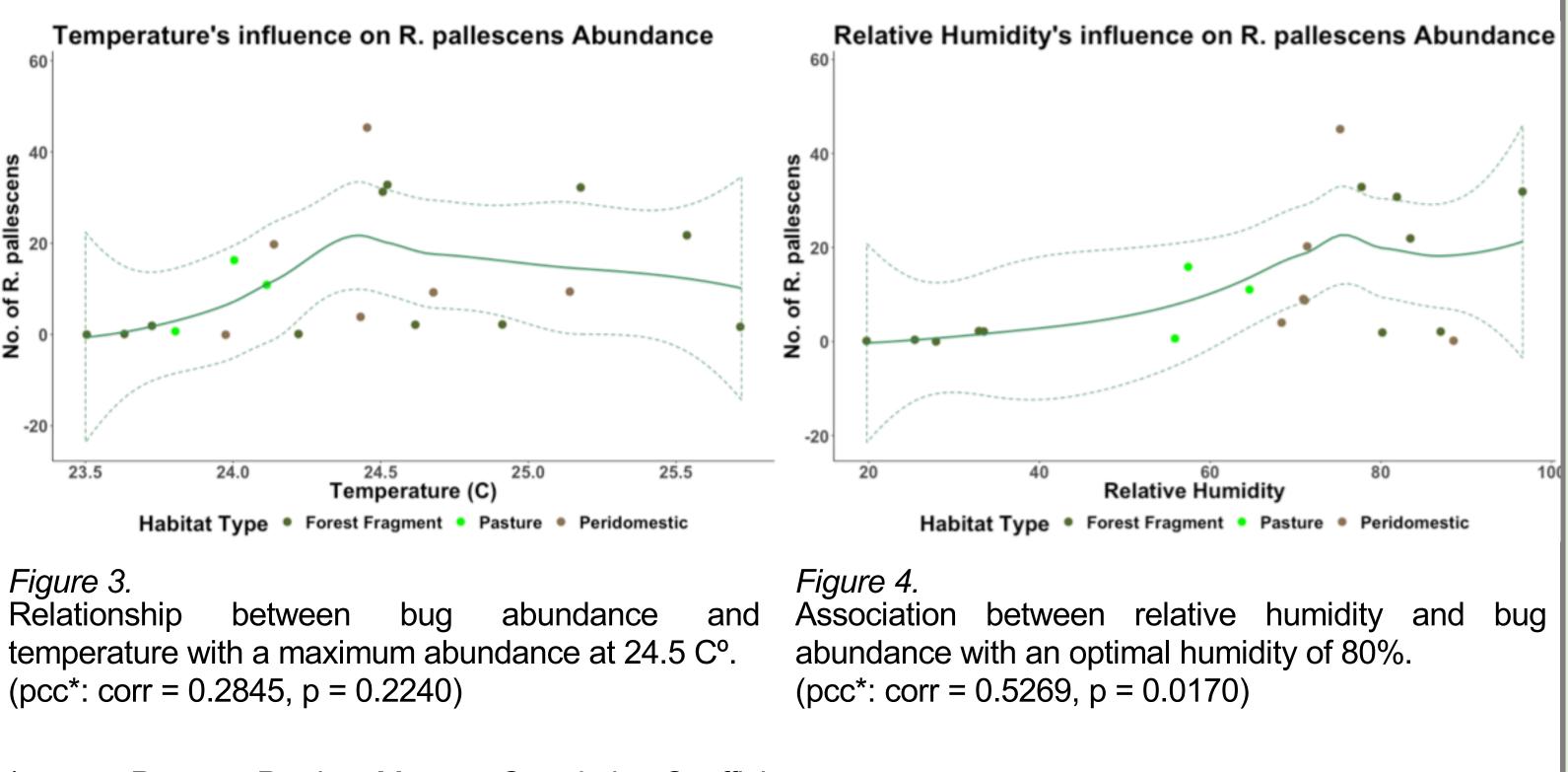


Figure 1.

Ŕ

Relationship between habitat type, location, and number of *R. pallescens* found per palm tree. Conditions in peridomestic habitat are positively vector abundance with (glmm' associated p<0.005) while pasture habitats are negatively vector associated abundance with (glmm* p<0.00001).



pcc – Pearson Product-Moment Correlation Coefficient ** glmm – Generalized Linear Mixed Model Fit by Maximum Likelihood (Laplace Approximation)



(Left) Map of Panama zoomed in where field data was collected.

(Right) Collection of a few specimen brought back to the lab.

Explored and modeled field data regarding microhabitat/climate and

Figure 2

Kissing bug abundance is positively associated with the dead organic matter score in the palm crown (1=low, 4=high). P-value below suggests that the R. pallescens abundance and DOM have a significant relationship.

(Kruskal Wallace test, p = 0.00166)

Biotic Factors Figure 5. Kissing bug abundance and invertebrate species richness within the palm crown. $(pcc^*: corr = 0.1725, p = 0.2359)$

- from removal

(1) Gottdenker, Nicole L., et al., *Am J Trop Med Hyg* 84.1 (2011): 70; (2) Gottdenker, Nicole L., et al., *EcoHealth* 11, 619–632 (2014); (3) Abad-Franch, et al., *Acta Tropica* (2015)

ACKNOWLEDGEMENTS

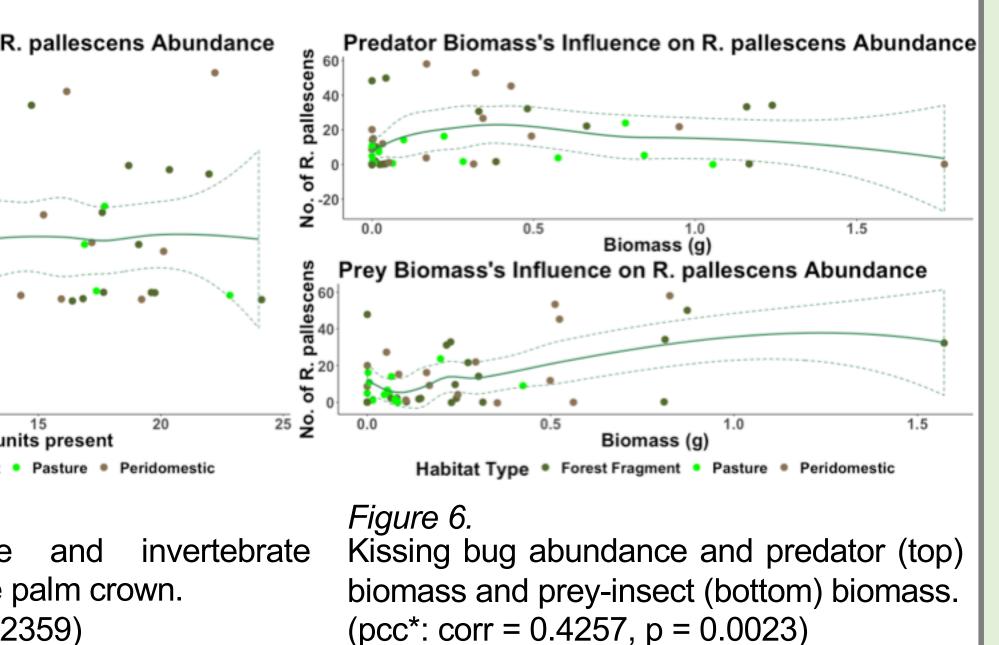
Thank you to the Population Biology of Infectious Diseases REU program at the University of Georgia for their support, and to the members of the Gottdenker Lab for their guidance and mentorship. Funding by the National Institutes of Health (NIH 1U54GM111274-01). Population Biology



Contact Information:



RESULTS CONT'D



CONCLUSIONS

Significant factors associated with vector abundance were microenvironment-related: Dead Organic Matter, No. of Connected Trees, and Relative Humidity. Community mechanisms, such as species diversity, was not significantly associated with vector abundance.

Habitat type also has a significant effect on R. pallescens abundance, but different degrees of disturbance have varying levels of effect, suggesting that the microclimate/habitat drive abundance over location.

This knowledge will influence the design and testing of novel pest control strategies to reduce Chagas disease infection rates, incorporating selective palm management.

FUTURE DIRECTIONS

Ecological experiments^[2] evaluating effectiveness of DOM palm crowns and monitoring crown temperature and relative humidity on vector abundance.

Evaluate relationships between the presence of specific predator taxa and traits that might be additional mechanisms influencing kissing bug abundance.

REFERENCES

National Institutes of Health

of Infectious Diseases REU site @ UGA

Jason Soriano