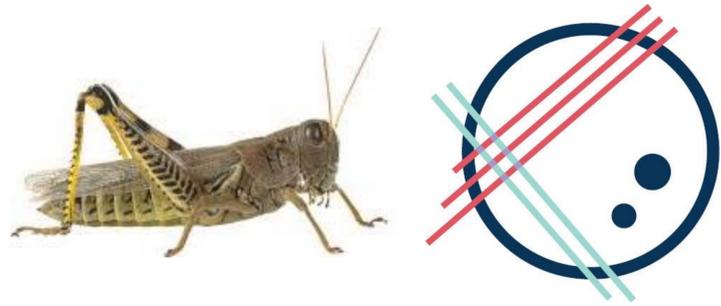


Patience, Young Grasshopper: Analyzing the fungal composition of the grasshopper gut microbiome



Staci Seitz and Melani Muratore
Advisors: Drs. Yvonne Sun, Ying-Ju Chen,
and Chelse Prather



Introduction

- Microbes are found all around us.
- Intestinal microbes, or gut microbiome, play an important role in overall animal health.
- The fungal communities within the gut microbiome of insects are important in insect physiology and thus, insect control.

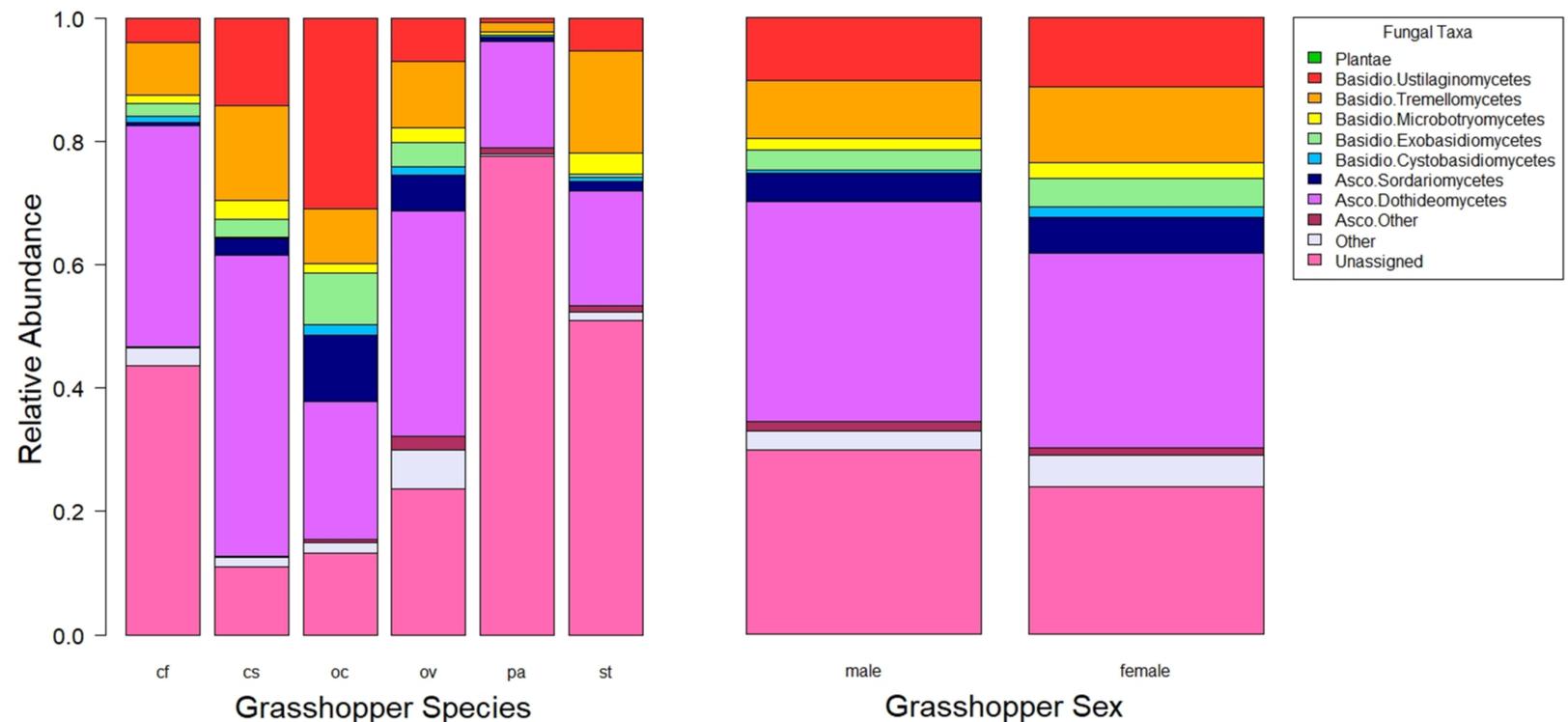
Main Questions

- What is the composition of the fungal communities in the microbiome of grasshoppers?
- What drives the composition of fungal communities within the microbiome of grasshoppers?
 - Species? Sex?
- Is there a “core” fungal microbiome in grasshoppers?

Methods

- The grasshoppers were collected in the summer of 2017 from a Texas prairie (multifactorial micronutrient experiment in which plots of the prairie were treated with various micronutrients)
- After collection and dissection, DNA was extracted from the grasshopper gut and submitted for sequencing by Zymo Research.
- Subsequently, the fungal and bacterial groups present in the sample were identified and analyzed using R programming.

Results



Summary

- No significant difference between males and females; The grasshopper species, *P. atlantica*, varied from the other grasshopper species
- The fungal phyla Ascomycota and Basidiomycota were most prevalent, while other phyla were scarcely found within the samples.
- Within Ascomycota → class Dothideomycetes most prevalent
- Within Basidiomycota → classes Tremellomycetes and Ustilaginomycetes most prevalent
 - Classes are directly associated with plant matter → food sources, presence in soil

Conclusions

- There exists a “core” fungal microbiome in grasshoppers.
- Diet and phylogeny are potential drivers of the fungal composition within the grasshopper gut microbiome.

Acknowledgements

I would like to thank Melani Muratore for allowing me to take part in the research project she started and always being willing to answer questions, Dr. Prather for the opportunity to perform research, and Drs. Sun and Chen for their mentorship on my Honors Thesis and senior Capstone.

