



NA and DNA strand

Molecular Characterization of *Hepatozoon* (Apicomplexa: Adeleorina) Blood Parasite Within the Threatened *Gopherus polyphemus* species



Johanna Eusse¹, Brian Cooney¹ and Evelyn Frazier¹

¹Department of Biological Sciences, Charles E. Schmidt College of Science Florida Atlantic University



Abstract

Gopherus polyphemus is a burrowing chelonian endemic to the Southeastern United States. Gopher tortoises are considered a keystone species due to their construction of extensive burrows which provide shelter to approximately 362 commensal species. Haemogregarines are the most common species rich with hemoparasites of reptiles. A previous study has shown that the genus Hepatozoon (Apicomplexa: Adeleorina) has been discovered within Jonathan Dickinson State Park gopher tortoise population. We will isolate hemoparasite genomic DNA from blood samples from the gopher tortoises and compare 18s rRNA gene fragment to other Hepatozoon species with similar evolution. This project will allow us to better understand the phylogenetic diversity of this parasite in comparison to other Hepatozoon species previously discovered.

Objective

Conduct molecular analysis on the blood of infected gopher tortoise to isolate and identify the 18S rRNA gene & compare to other *Hepatozoon* species with similar phylogeny

Introduction

Gopher Tortoise: Gopherus polyphemus

- · Endemic to SE United States
- · Digs extensive burrows
- · Keystone species
- · Threatened status



Step 5:

3)

Study Site (2014):

Jonathan Dickinson State Park (JDSP)

• 11,500 acres





Haemogregarines

- · Most common reptilian hemoparasite
- Three Families
 - Hepatozoida
 - Karvolvsidae
 - Haemogregarinidae

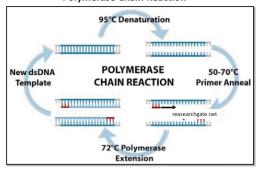
Genus Hepatozoon (1932)

- · Potential health effects:
 - Nondetrimental in natural hosts
 - · Life threatening in unnatural host

Method

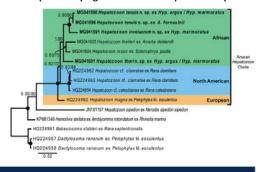
1) In 2014, blood was collected from three adult gopher tortoises at Jonathan Dickinson State Park.

Polymerase Chain Reaction



Anticipated Results

Example of Phylogenetic Tree of Hepatozoon sp.



Future work

- 1) Order 2 sets of forward and reverse primers to conduct PCR reaction
- Collaborate with Bracken-Grissom laboratory at Florida International University for sequencing and phylogenetic analysis.

References

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