

The Gulf Coast's Local Star: Exploring the differences between Gulf, Atlantic, and Pacific Coast Starlet Sea Anemones

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Background

- *Nematostella vectensis* is a mud-dwelling, **model cnidarian organism** found living in salt marshes and frequently studied due to its high tolerance of temperature, salinity, and starvation.
- Our lab **rediscovered** the presence of *Nematostella vectensis* in the northern **Gulf of Mexico** in a salt marsh on Dauphin Island, AL.
- Before its closure, Florida's prehistoric **Suwannee Canal** connected the Gulf of Mexico to the Atlantic Coast.
- The Suwannee Canal **closed 1.8 mya** disconnecting Gulf Coast salt marsh habitats from those on the Atlantic Coast, disrupting a possible avenue for gene flow.

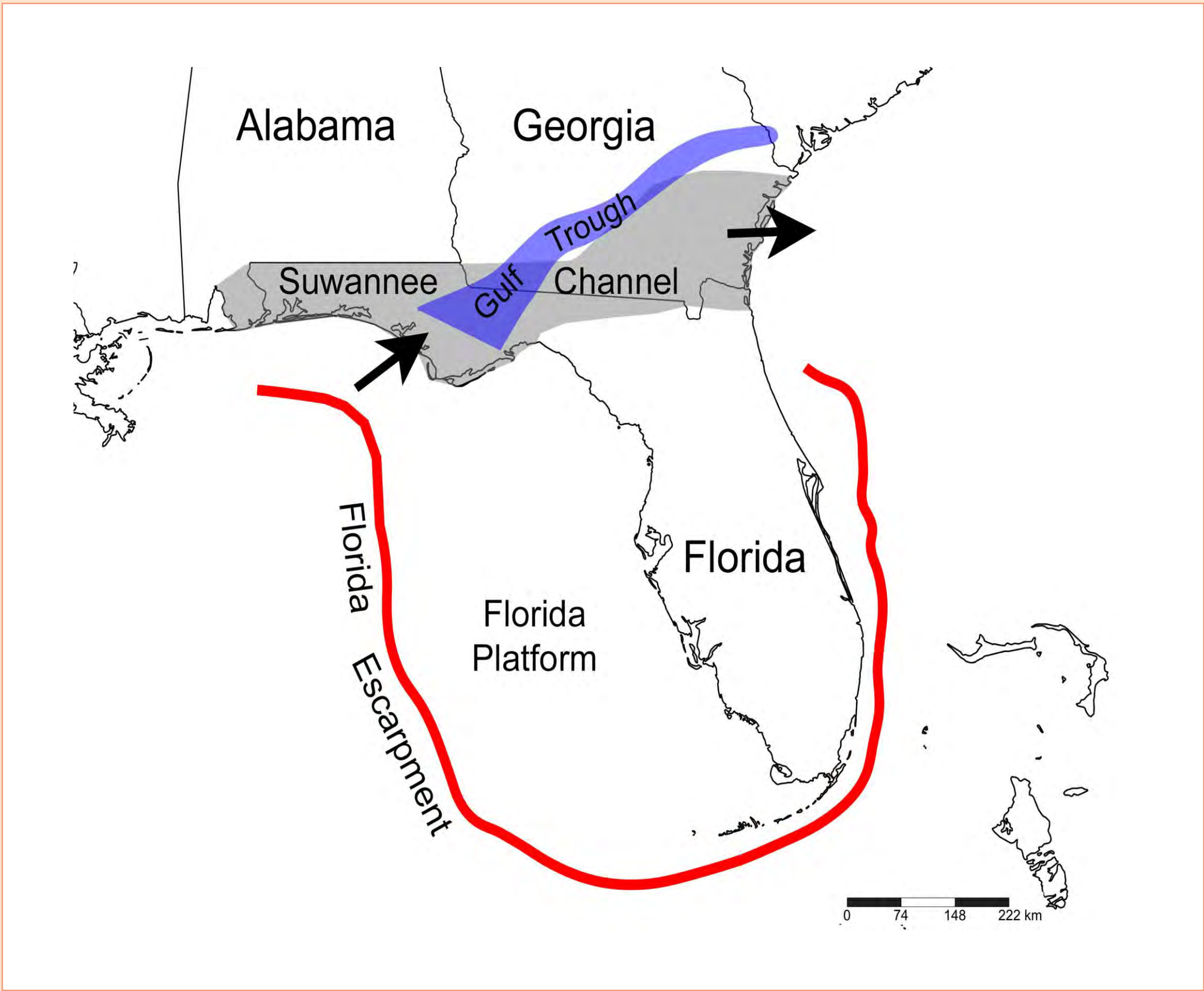


Fig. 1: Map of the prehistoric Suwannee Channel.

Questions

- Are Gulf Coast *Nematostella vectensis* **significantly different** than those along the Pacific and Atlantic coasts?
- How might the **Suwannee Canal** and its closure have impacted **gene flow** between Gulf Coast and Lower Atlantic individuals?

Results

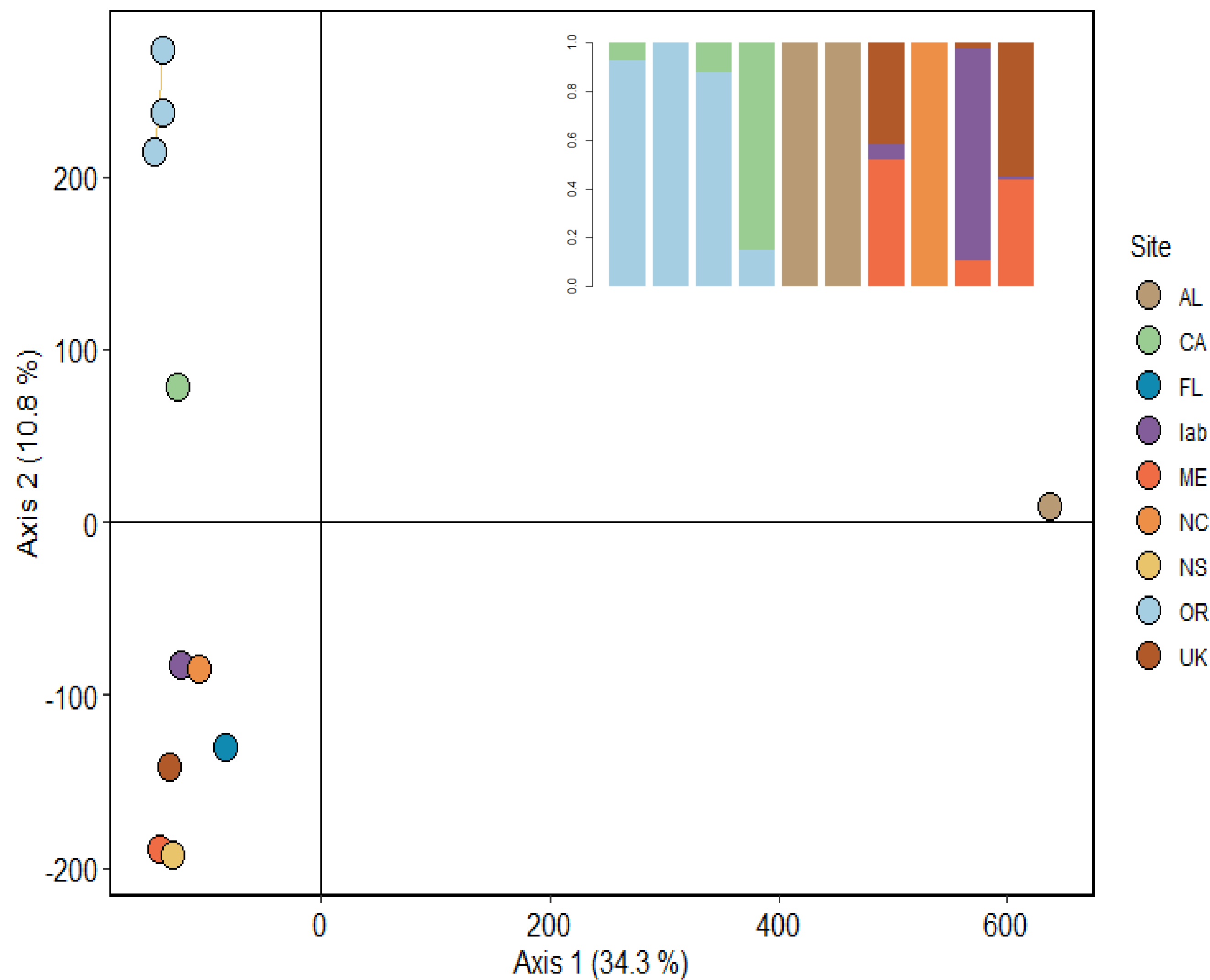


Fig. 2 PCA plot displaying overall pattern of Single Nucleotide Polymorphism (SNP) differences between *Nematostella vectensis* individuals and an integrated membership probability plot.

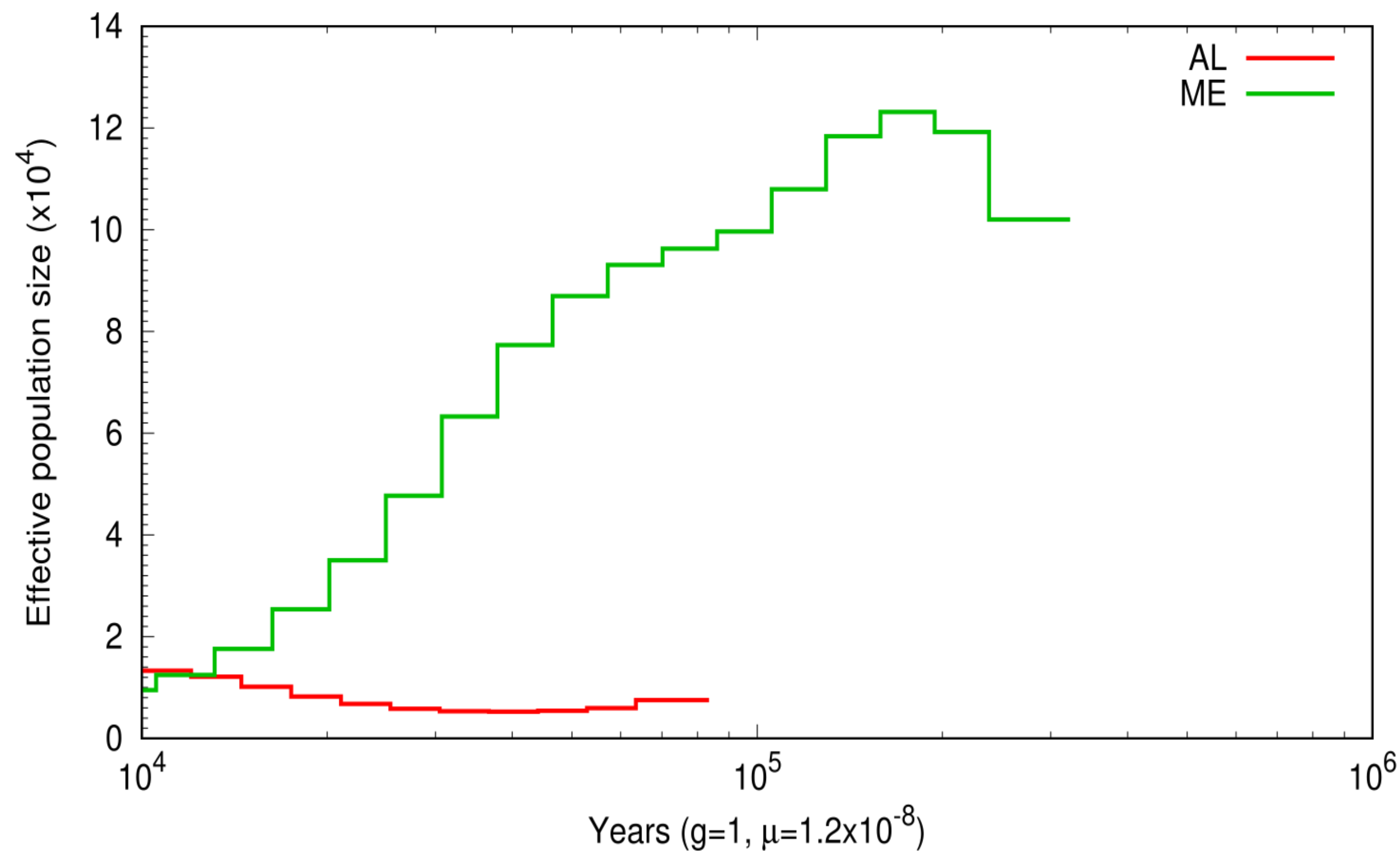
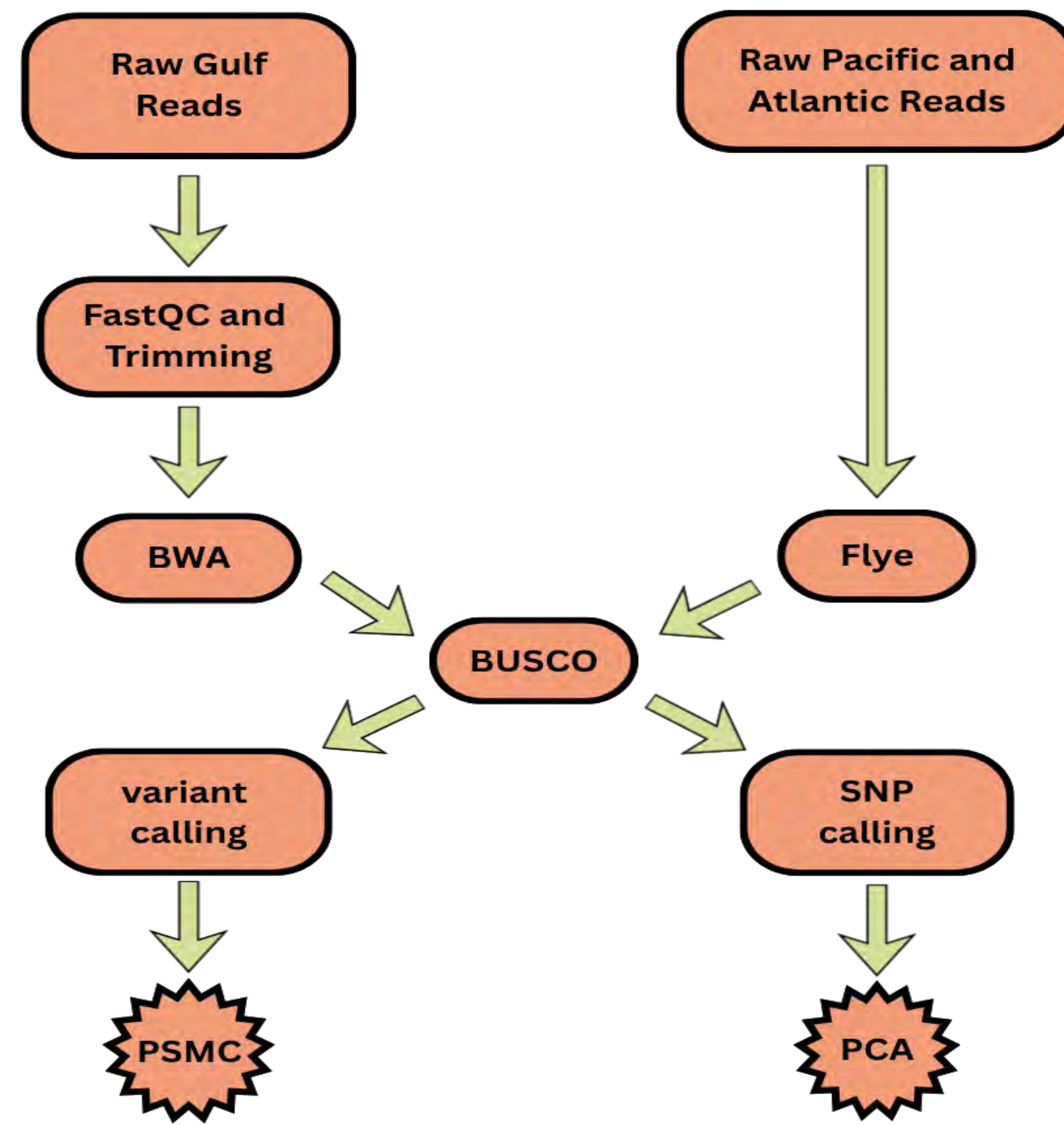


Fig. 3 Pairwise Sequentially Markovian Coalescent (PSMC) plot displaying reconstructed effective population size values of *Nematostella vectensis* individuals.

Methods



Conclusions

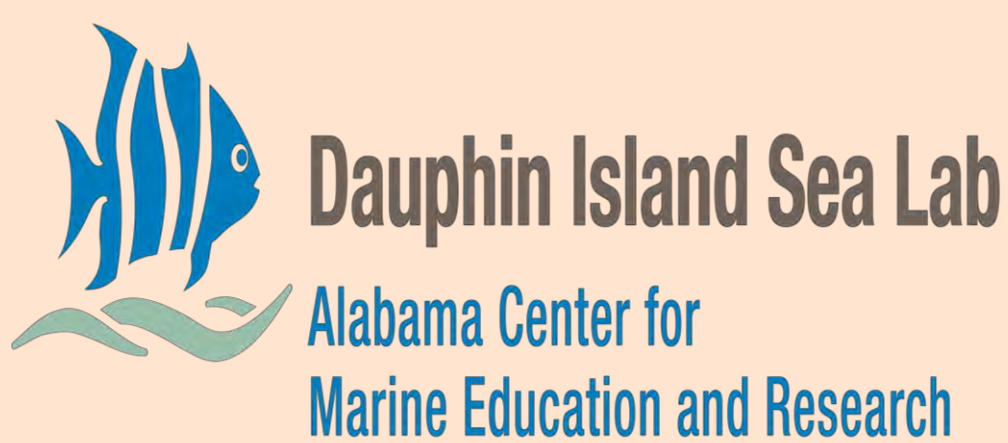
- The two Gulf Coast *Nematostella vectensis* are **not** a part of **the same population** as the Pacific or Atlantic individuals and may represent a **native Gulf population**.
- Alabama and Florida individuals are not closely related which suggests the **Suwannee Seaway did not play a role in facilitating gene flow**.
- Results from **PSMC** are **preliminary** but support the conclusion that individuals from the Gulf are from a separate native population disjunct from the Atlantic.
- PSMC will continue to be used to compare Atlantic effective populations and **reconstruct** Atlantic and Gulf **phylogenies**.

Acknowledgements

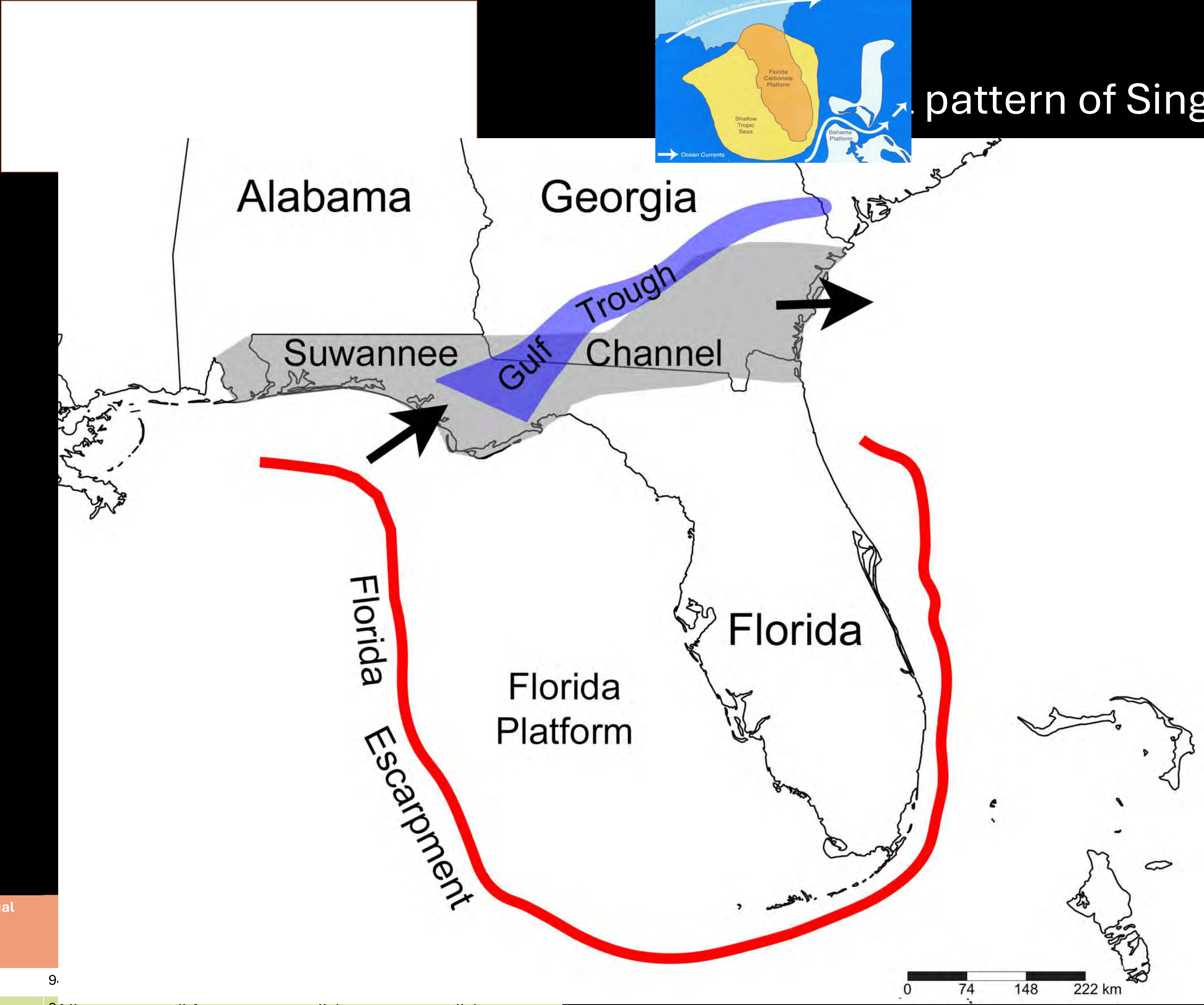
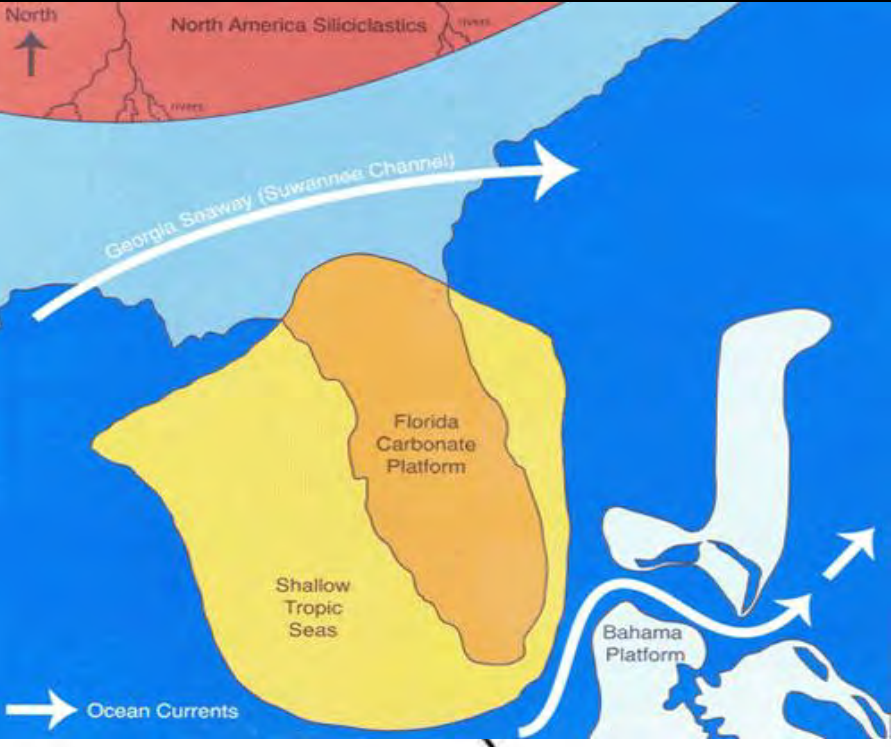
This work was funded by the National Science Foundation award number 2318922. Photo Credit to Benjamin Titus. Atlantic and West Coast DNA sequences provided by Auston Rutledge in the UNC Charlotte Reitzel Lab



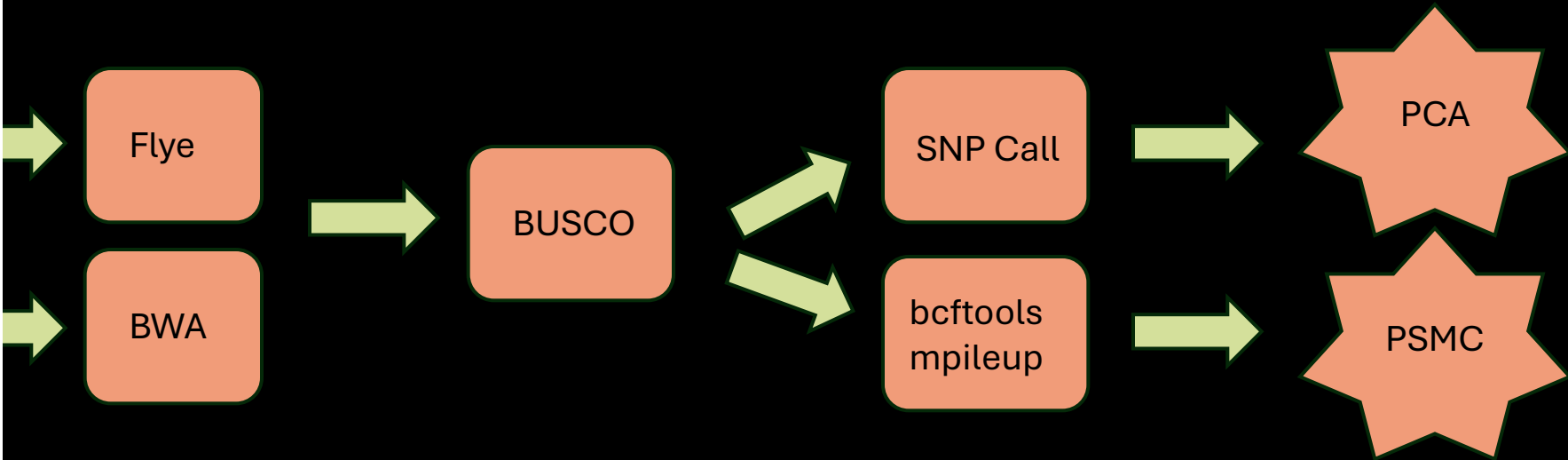
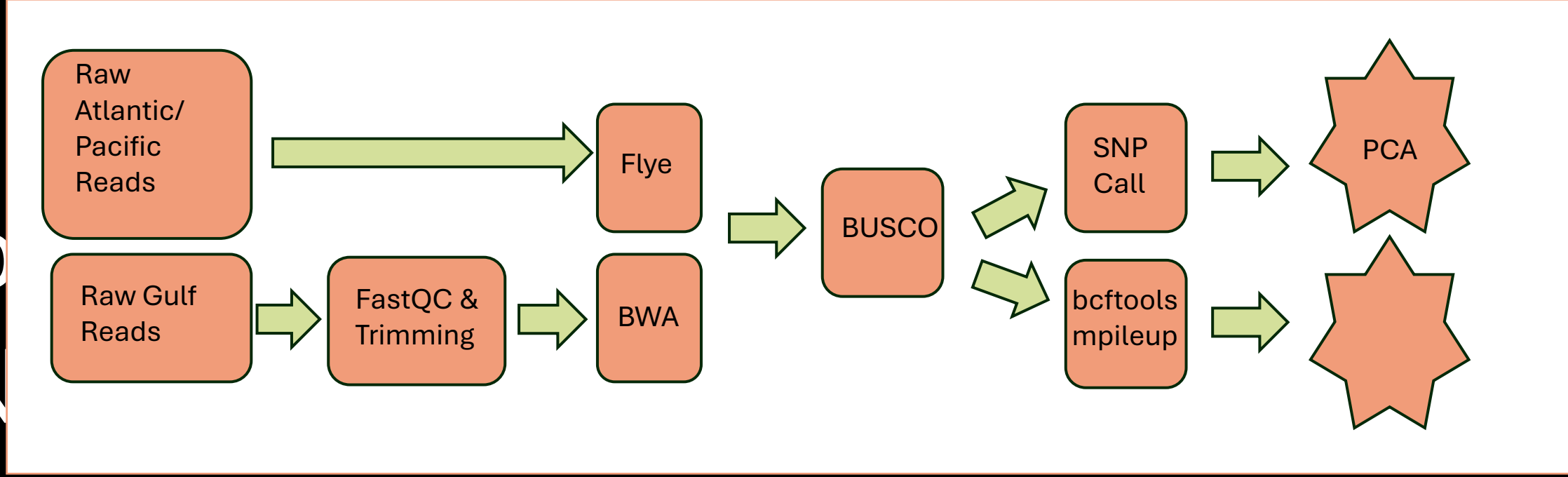
References



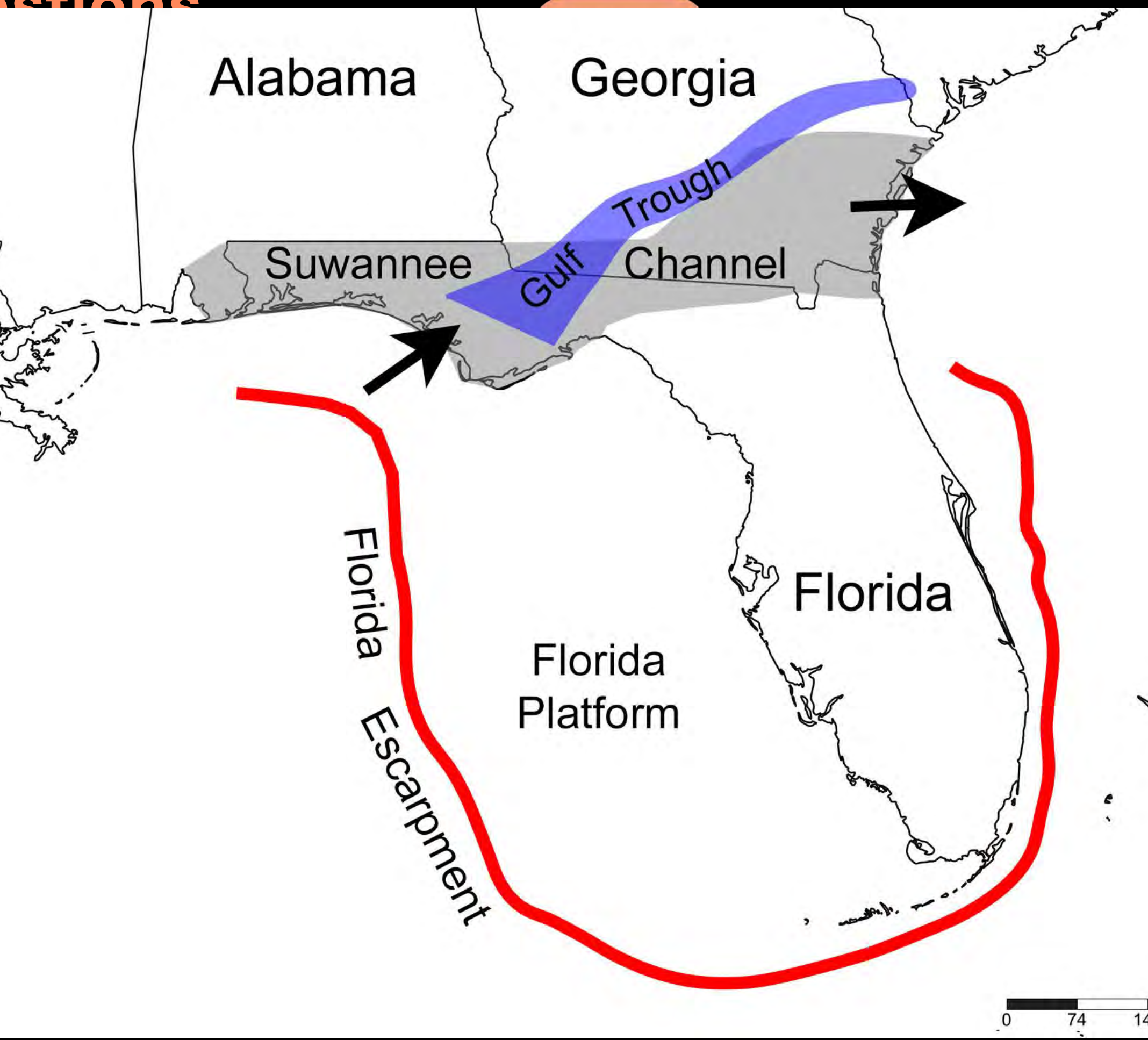
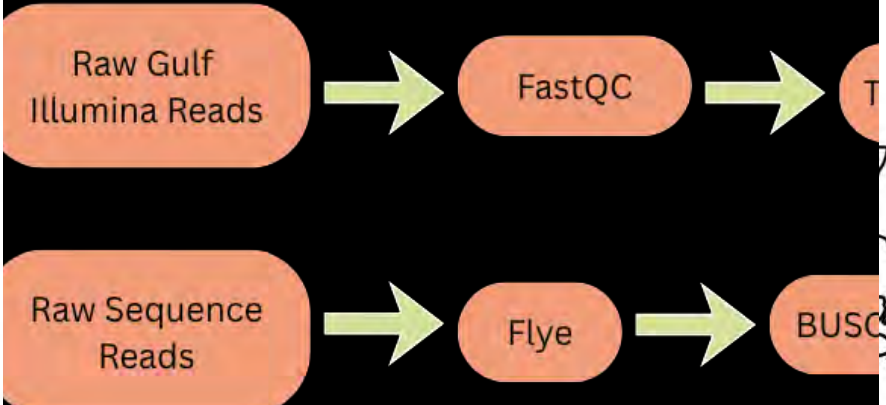
Individual				
AL1	9.			
AL2	94.2	0.4	2.7	2.7
ME	92.9	0.9	4.4	2.5
NC	89.5	3.9	4	2.6
FL	91	3.7	3	2.3
NS	93.5	1.5	2.8	2.2



pattern of Single Nucleotide Polymorphisms between N



Questions



Markovian Co-constructed eff

population size values