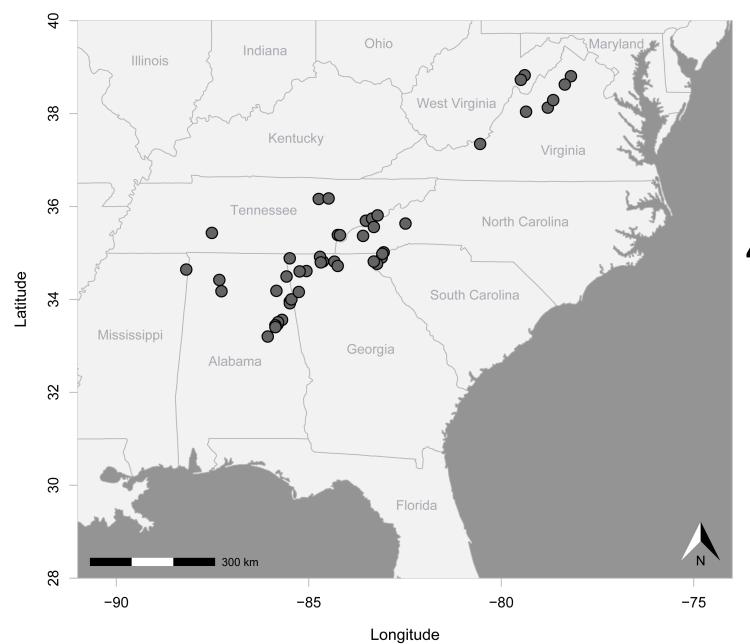




Sampling



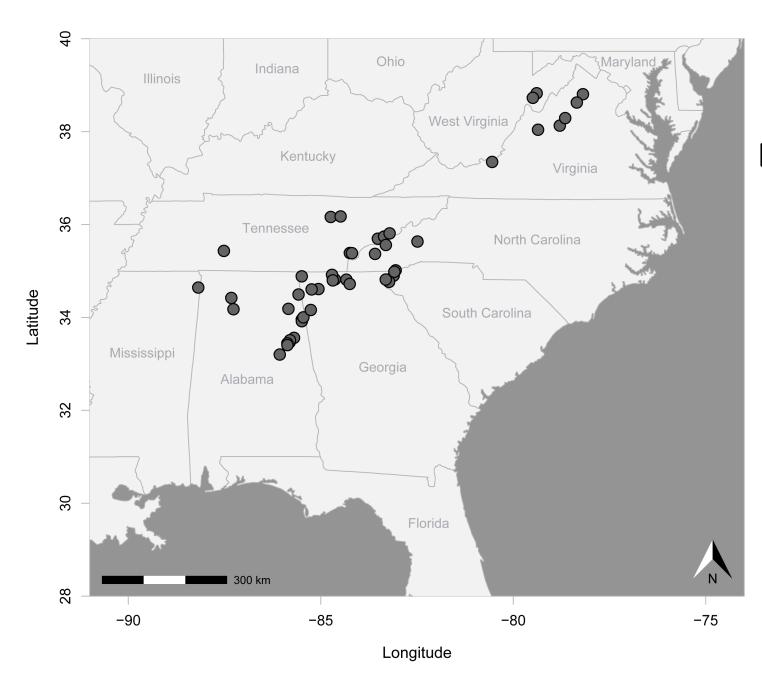


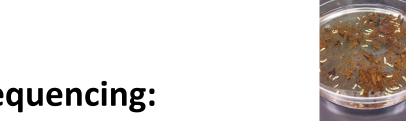
46 sites:

50 logs:

128 individuals

Genetic Data Collection





DNA sequencing:

2 markers:

1 nuclear: endo-beta-1,4-glucanase (seq. length: 251 bp)

1 mitochondrial: cytochrome c oxidase I and II (seq. length: 1,117 bp)

Evolutionary History of *Reticulitermes flavipes*

Divergence of genetic lineages \rightarrow dealing with environmental change

Population size dynamics \rightarrow is population size increasing?

Geographic distribution \rightarrow is geographic distribution shifting/expanding?

Evolutionary History of *flavipes*: Questions

How many lineages of *flavipes*?

When did these lineages arise and how did they change through time?

Evolutionary History of *flavipes*: Questions

How many lineages of *flavipes*?

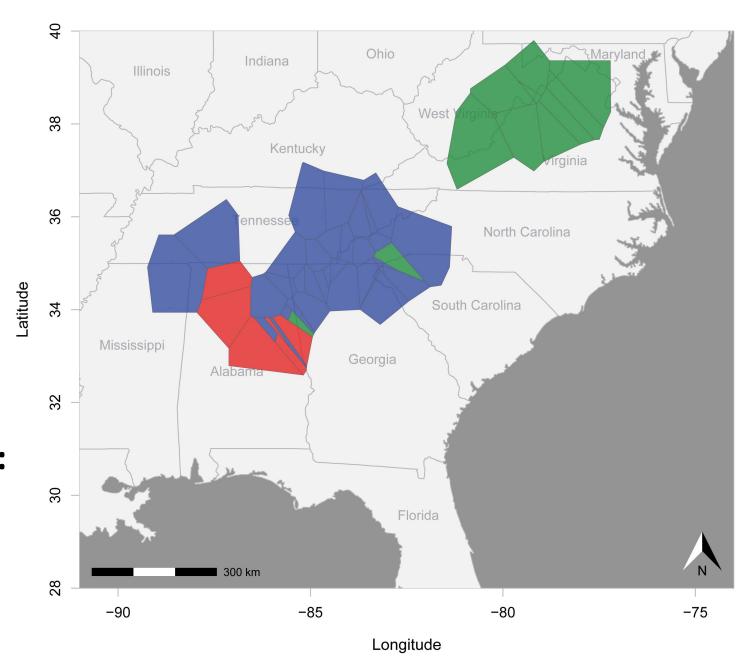
When did these lineages arise and how did they change through time?

Evolutionary History of *flavipes*: Questions

How many lineages of *flavipes*?

When did these lineages arise and how did they change through time?

How many lineages?



Three Lineages:

N = Northern

C = Central

S = Southern

When did *flavipes* lineages arise and how did they change through time?

Approximate Bayesian Computation

Competing evolutionary history hypotheses characterized using:

Divergence time

Population

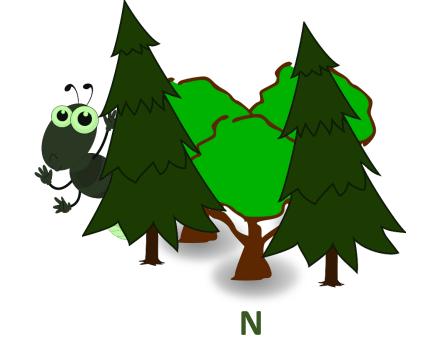
Best-fit hypothesis: closest match between simulated and empirical data:

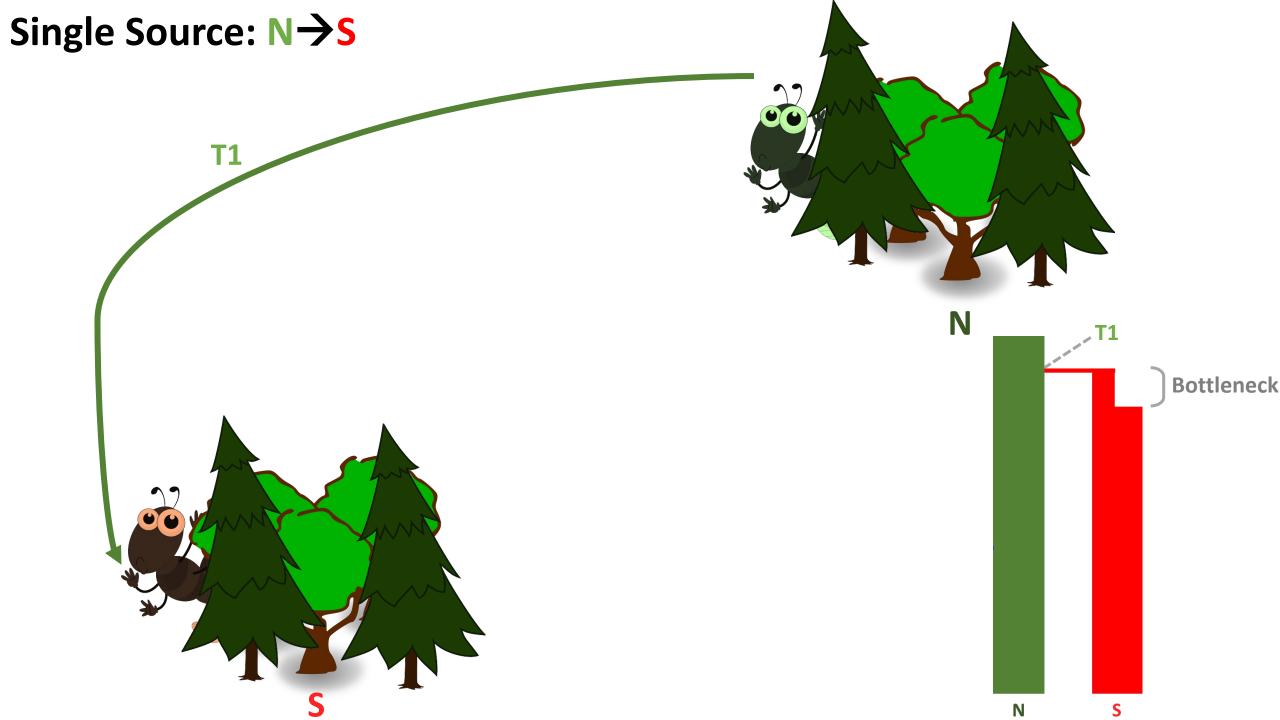
1 million simulated datasets for each hypothesis

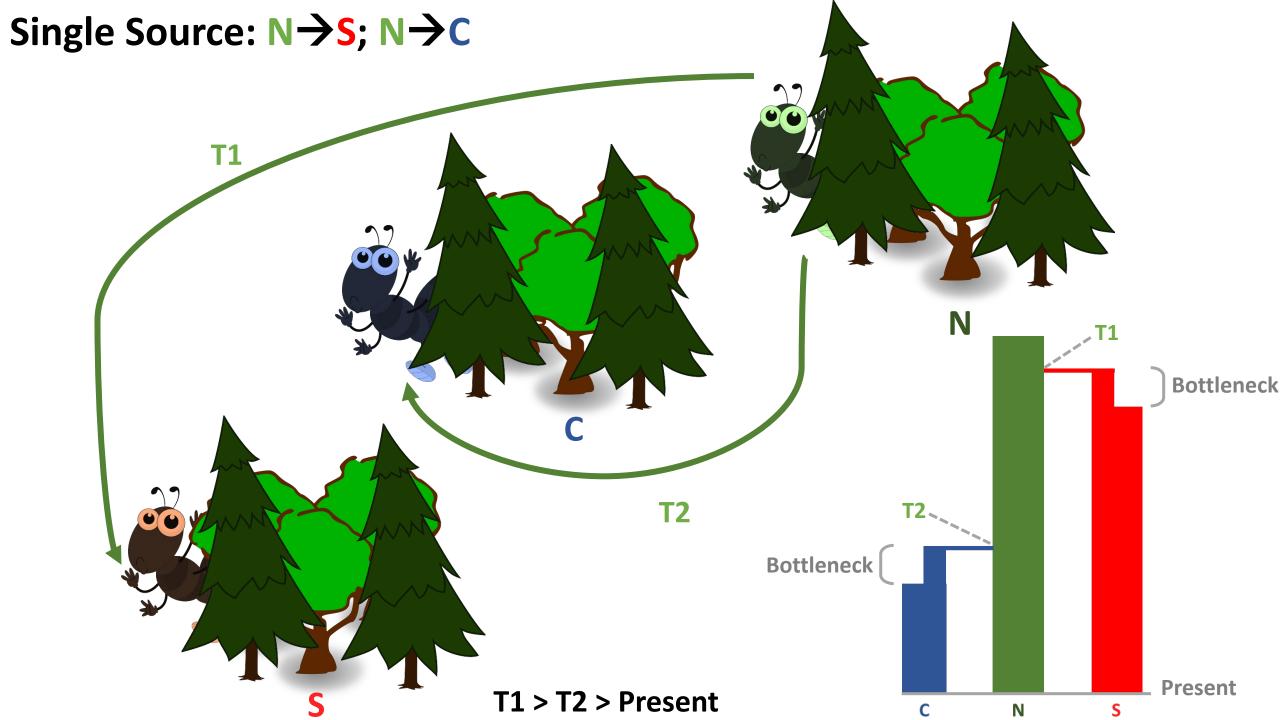
Compare simulated data with empirical data

Evolutionary History Hypotheses: Single Source

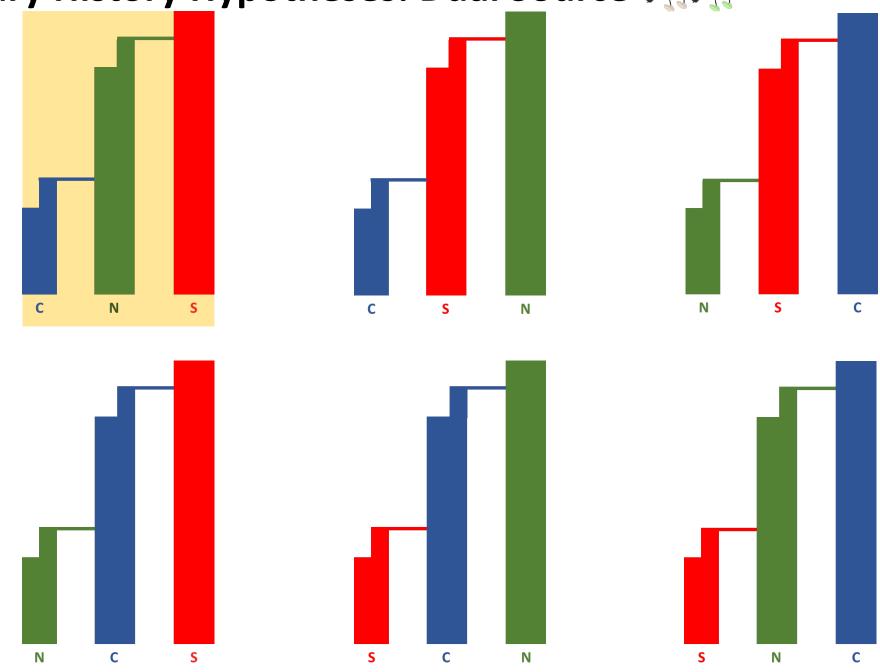
Single Source: N

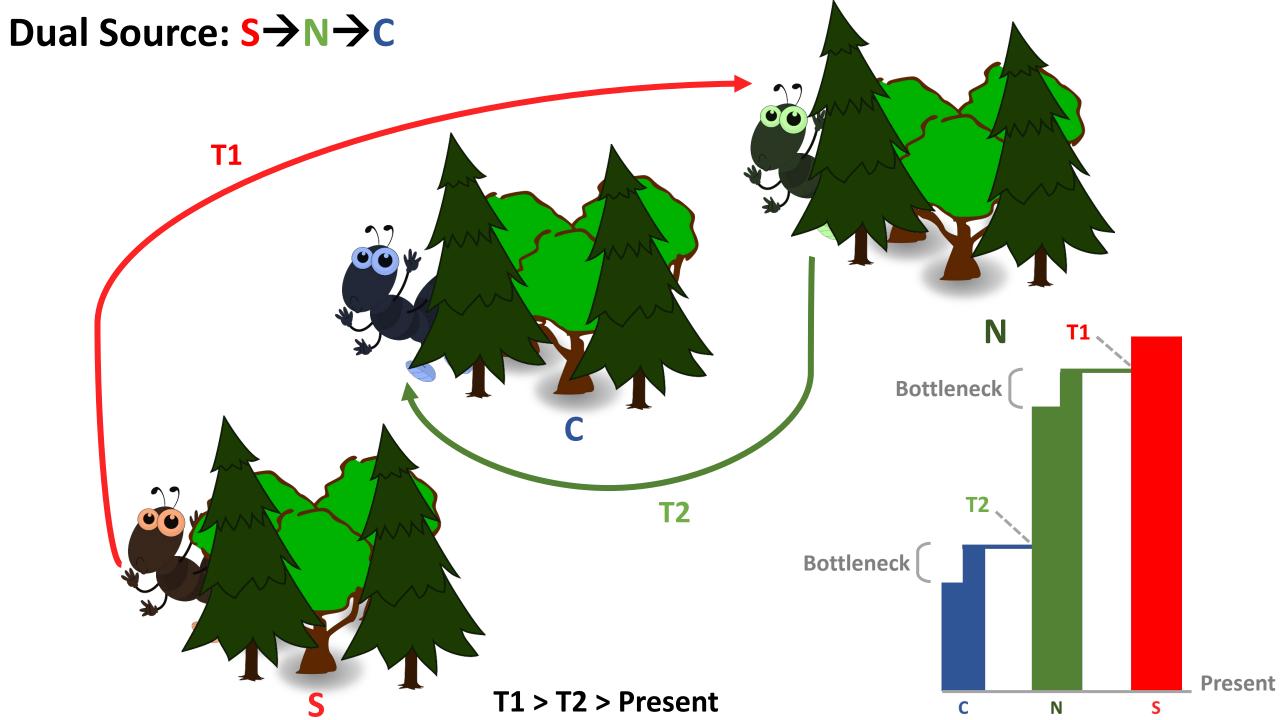






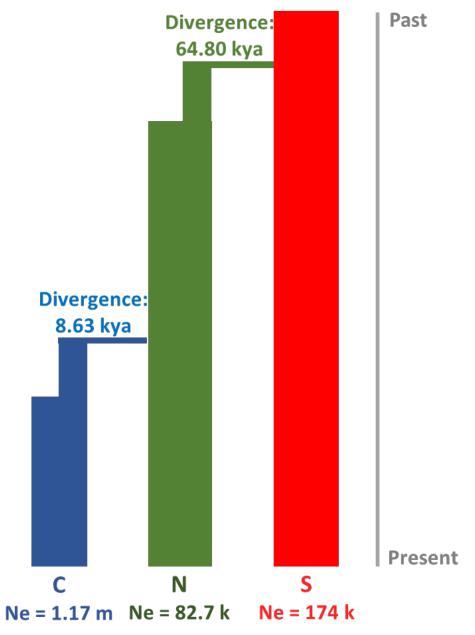
Evolutionary History Hypotheses: Dual Source

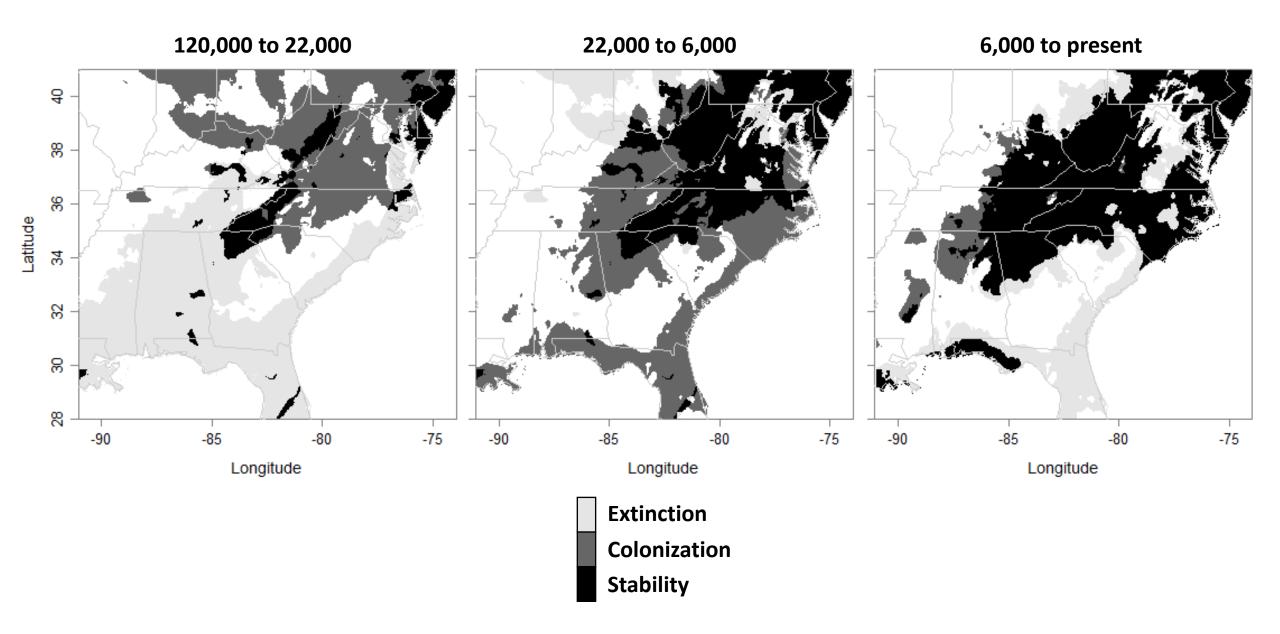




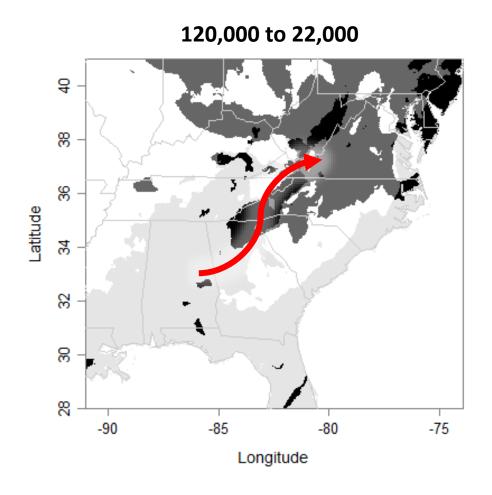
Best-Fit Hypothesis

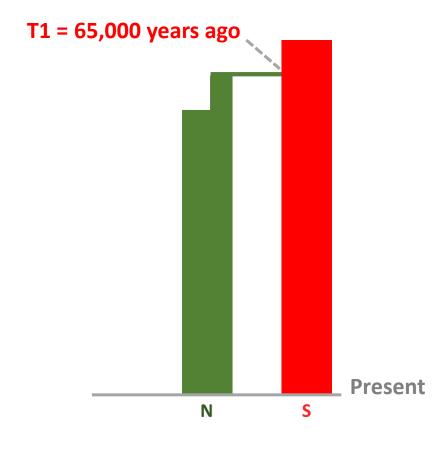




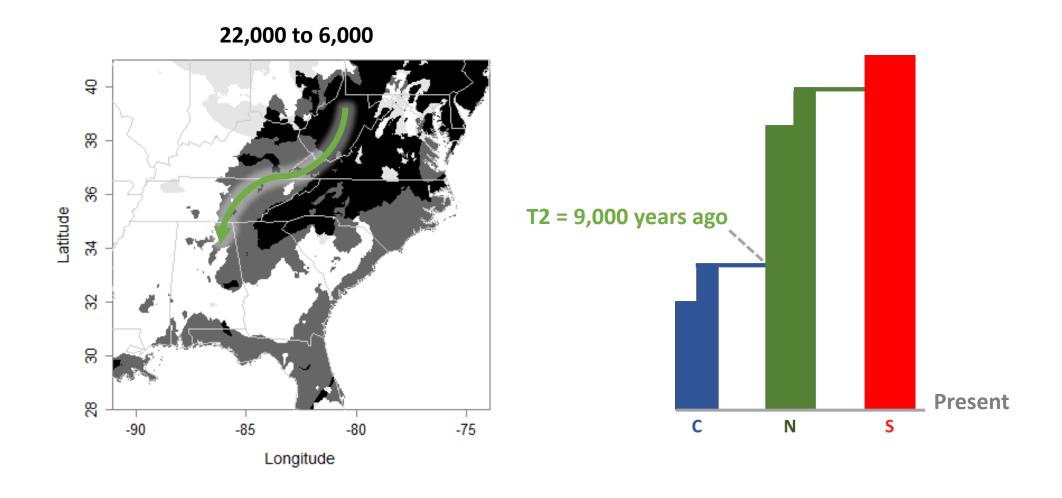


Distributional Shift: South-to-North

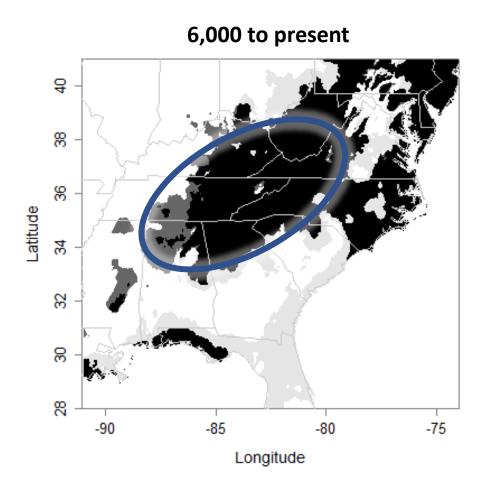


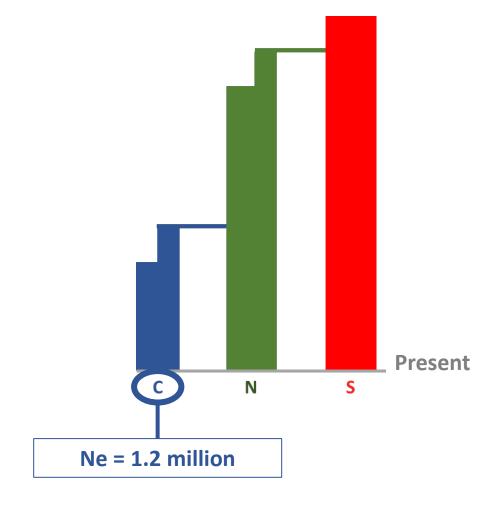


Distributional Shift: North-to-Center



Central Expansion





Correspondence between distributional change and genetic divergence

