### Color pattern polymorphism in Eleutherodactylus coqui: selection in Puerto Rico and founder effects in Hawai'i



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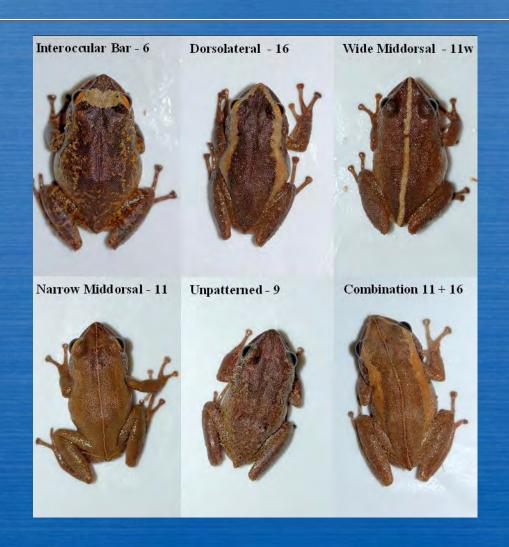
## Background



- Color patterns polymorphisms have historically been instrumental in studies of evolution
- E. coqui exhibits distinct color patterns (Woolbright, 2005)
- E. coqui was introduced to Hawai'i in the late 1980s and appears to have undergone a loss of genetic diversity in mtDNA (Velo-Antón et al., 2007)



# Color patterns in E. coqui





### Goals



- Determine the mode of inheritance
- Characterize color patterns in native and introduced ranges
- Test for founder effects in Hawai'i
- Test for selection in Puerto Rico



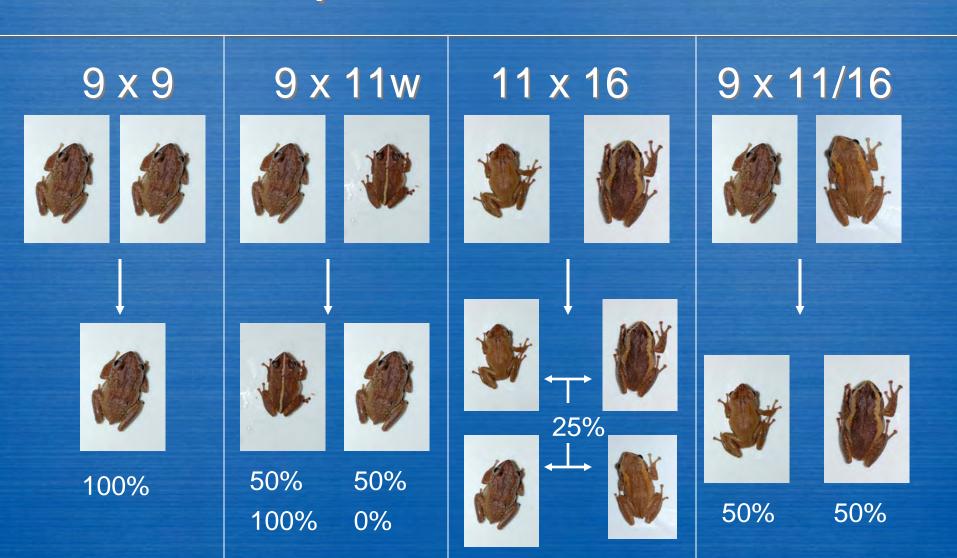
## Mode of Inheritance



- In nature
  - Do frogs have multiple color patterns?
  - Yes, but only 2
  - Are color patterns sex limited?
  - No, all patterns seen in males and females
- In lab
  - Mendelian crosses
  - Number of loci and alleles
  - Dominance / recessive relationships



## Crosses performed in the Lab





#### Results



- Lab data
  - all stripe patterns dominant to unpatterned
  - all stripe patterns are codominant with each other
  - $\chi^2$  tests indicate single autosomal locus, with 5 alleles

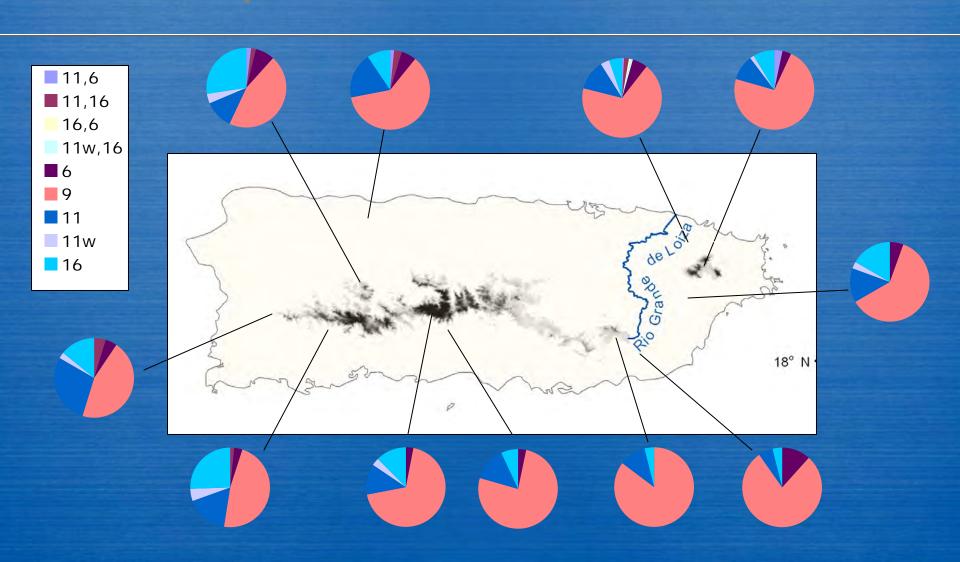


# Color patterns in native and introduced ranges

- Sampled multiple populations from across both ranges
  - PR: n=11
  - HI: n=14
- Counted number of frogs with each color pattern
- Used results of breeding experiment to determine allelic richness
- Compared allelic richness among PR and HI



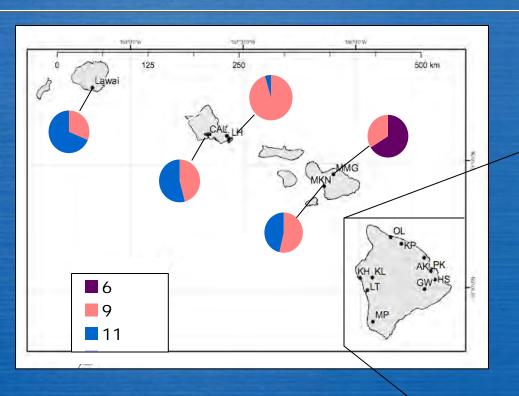
## Color patterns in Puerto Rico

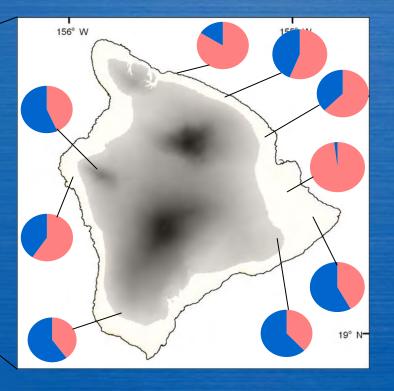




# Color patterns in Hawai'i









## Results



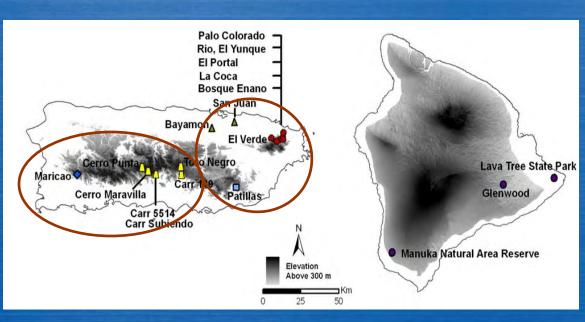
- Puerto Rico
  - $\bar{x} = 4.3 \pm 0.2 \text{ SE}$
- Hawai'i
  - $\bar{x} = 2.0 \pm 0.0 SE$
- Loss of color patterns may be result of founder effects

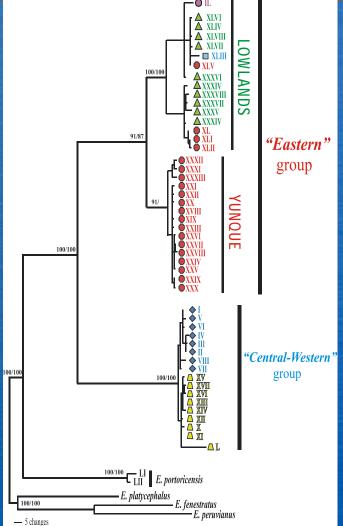


# Test for selection on color patterns in Puerto Rico

- Locus comparison approach
  - Compare population differentiation (φ<sub>ST</sub>) among neutral and test loci
  - differentiate between drift/migration and selection
- Neutral: mtDNA from Velo-Anton et al. (2007)
- Test: Color pattern allele frequencies estimated from field data, single locus, 5 allele
- Compare estimates of φ<sub>ST</sub> using Arlequin 3.1

# mtDNA phylogeography of *E. coqui*



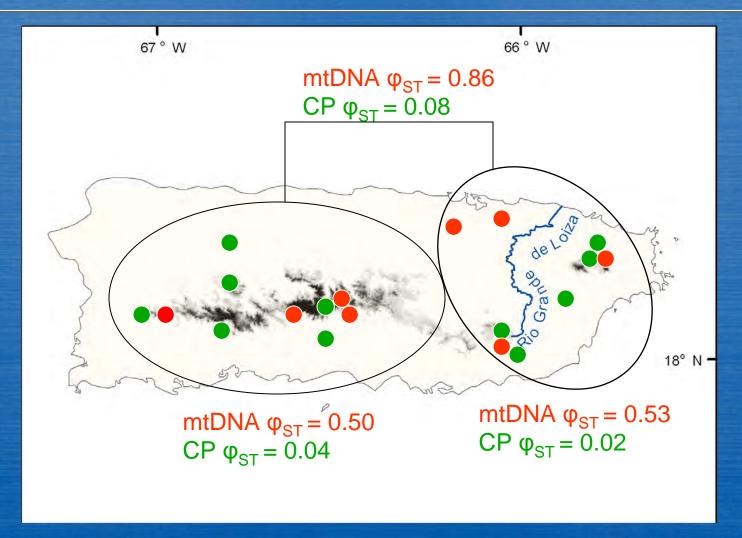


G. Velo-Antón et al. (2007) Molecular Phylogenetics and Evolution



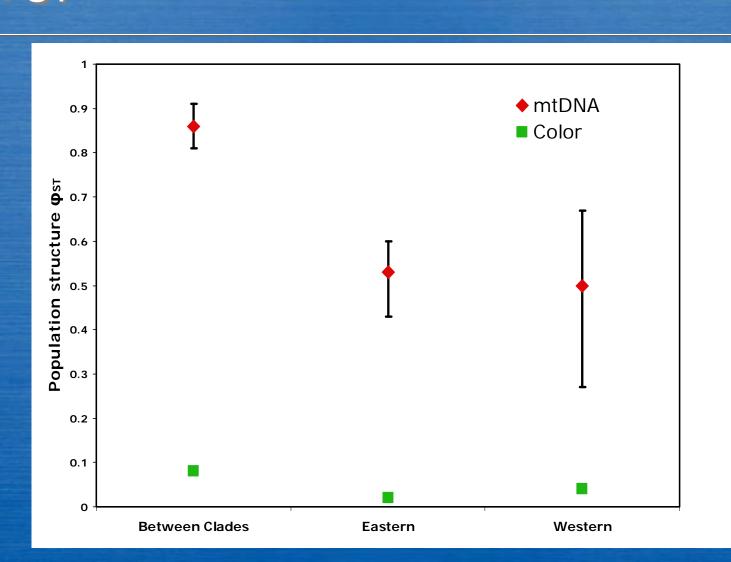
# φ<sub>ST</sub> across Puerto Rico







# φ<sub>ST</sub> Estimates for Puerto Rico





### Conclusions



- Color patterns result from a single autosomal locus with 5 alleles
  - Stripes are codominant, unpatterned is recessive
- Fewer alleles in Hawai'i may be result of founder effects
- Color patterns are under balancing selection in Puerto Rico



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