

Evaluating threats and providing effective tools to conserve Coquí Guajón (Eleutherodactylus cooki)

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#### Cooperative Agreement - Proyecto Coquí & USFWS

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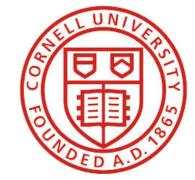
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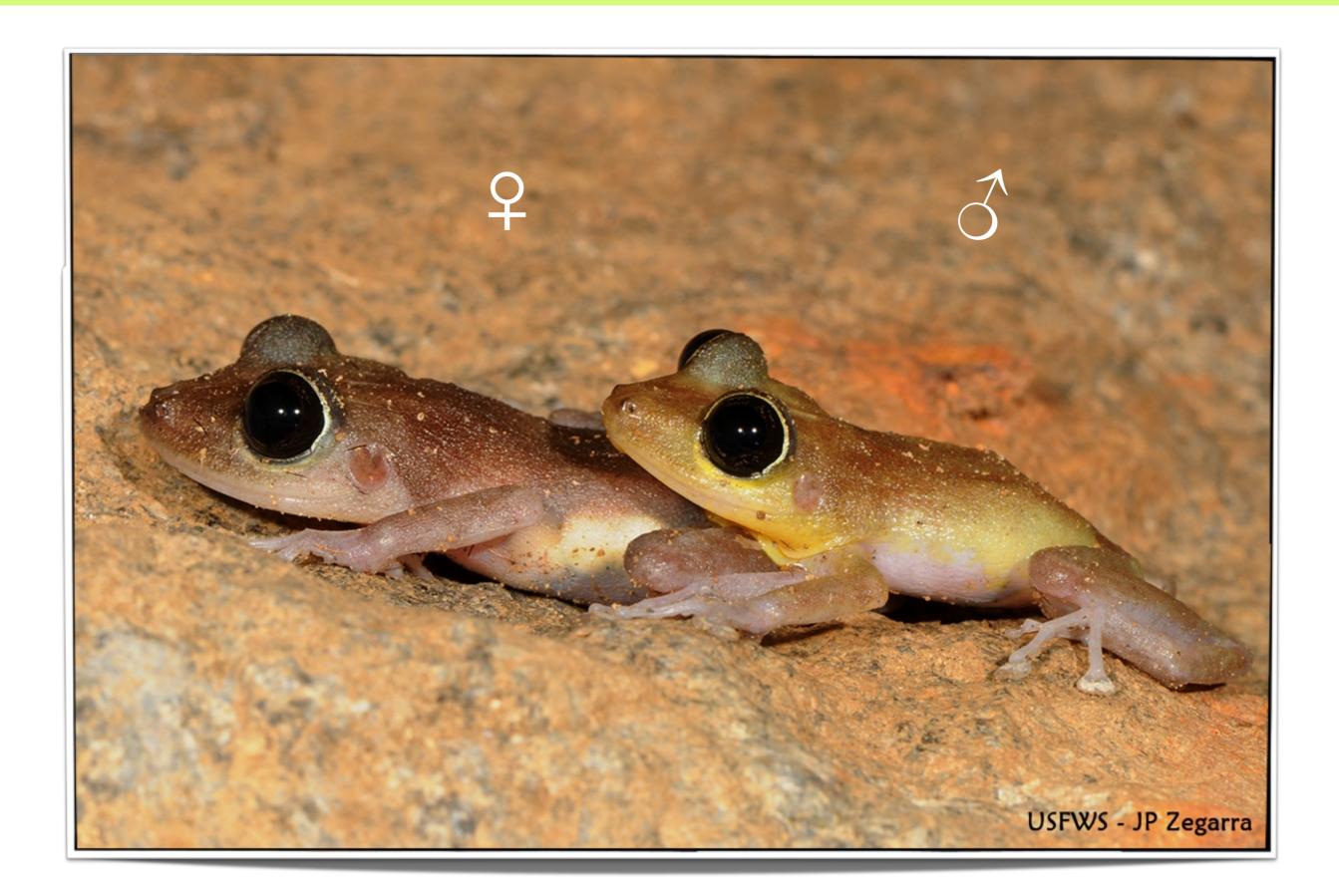




Special thanks to all the landowners who gave us permission to work in their lands!



#### Granodiorite is essential for Guajón's reproduction

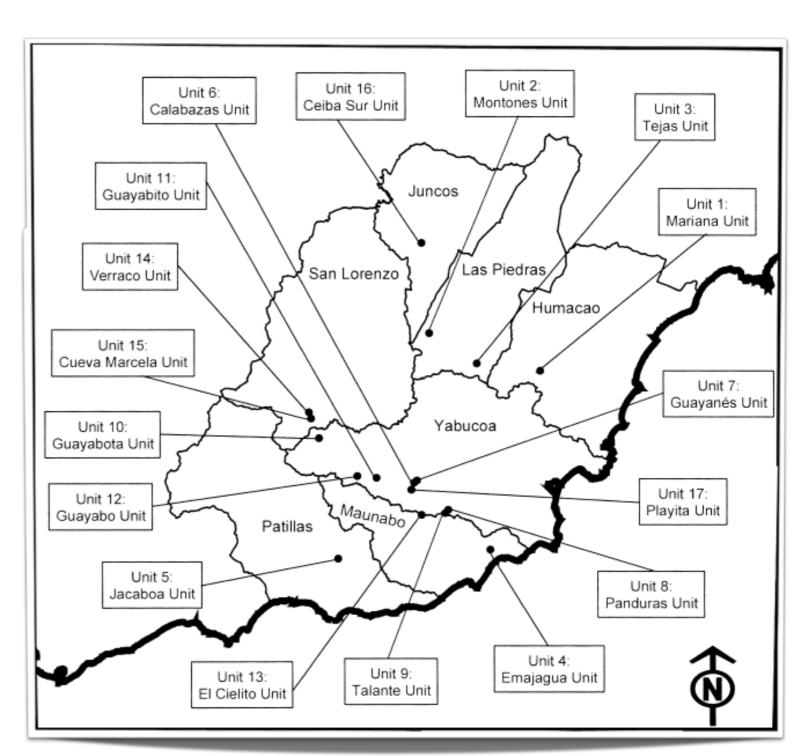






#### Coquí Guajón (Eleutherodactylus cooki)

- Listed as threatened in 1997 by USFWS.
  - Vulnerable: DNR
- Recovery plan released in 2004.
- 17 critical habitats declared in 2007.
  - Restricted to private lands

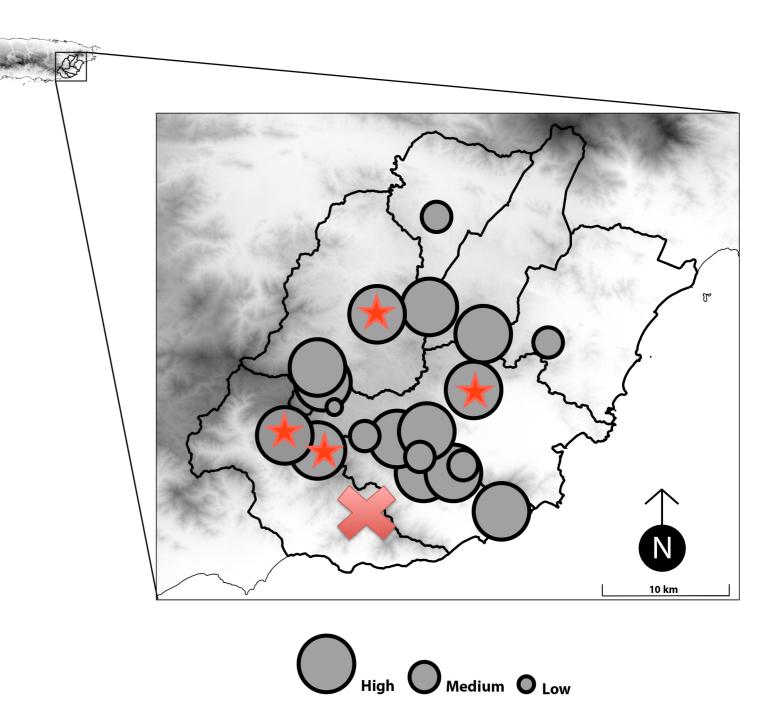


#### Aims of the cooperative agreement

- I. Monitor the population status by performing demographic studies.
- 2. Determine the impacts of the fungal pathogen Batrachochytrium dendrobatidis (Bd) and parasitic tick Ornithodoros talaje (OT).
- 3. Assess the genetic diversity and patterns of gene flow (connectivity) between localities.
- 4. Develop an outreach program to promote species awareness and to improve habitat quality.

#### Monitor the population status of Coquí Guajón.

- 16 habitats (+4 additional localities ★) visited twice:
   November 2012 & 2013 (wet); and March 2013 & 2014 (dry).
- Audio & visual-encounter surveys.
- Collected tissues: swabs, toe clips for genetic analyses, and ticks.
- No individuals found in Jacaboa.



#### Monitor the population status of Coquí Guajón.

Locality	Trash	Invasive Predator	Parasites	Bd	Habitat Modification	Agriculture	Erosion	Abundance Score	Threat Level Score	Abundance Category	Threat Level Category
Claudio Hernández	0	1	0	1	0	2	3	1	8	High	Moderate
Casas de la Selva	0	1	0	1	0	0	0	1	3	High	Low
Car. 181 Patillas	2	0	0	2	0	0	3	1	8	High	Moderate
Fuentes	2	1	1	1	0	2	0	1	8	High	Moderate
Unit_01_Mariana	0	0	0	2	3	0	0	2	7	Medium	Moderate
Unit_02_Montones	0	1	1	2	0	2	0	1	7	High	Moderate
Unit_03_Tejas	2	0	0	2	3	2	3	1	13	High	High
Unit_04_Emajagua	2	1	1	2	3	0	0	1	10	High	Moderate
Unit_05_Jacaboa*	2	1	0	2	3	2	3	3	16	Low	High
Unit_07_Guayanes	0	1	1	1	0	0	0	1	4	High	Low
Unit_08_Pandura	2	1	1	1	0	0	0	2	7	Medium	Moderate
Unit_09_Talante	2	0	1	1	0	0	3	1	8	High	Moderate
Unit_10_Guayabota	2	1	1	1	3	2	3	3	16	Low	High
Jnit_11_Guayabito	2	0	1	1	0	0	0	1	5	High	Low
Jnit_12_Guayabo	0	0	1	1	0	2	0	2	6	Medium	Moderate
Jnit_13_Cielito	0	1	1	1	0	0	0	1	4	High	Low
Jnit_14_Verraco	0	1	1	1	0	0	0	1	4	High	Low
Jnit_15_Marcela	0	1	1	1	0	2	3	1	9	High	Moderate
Jnit_16_Ceiba Sur	2	1	0	1	3	0	3	2	12	Medium	High
Jnit_17_Playita	0	1	0	1	3	2	0	2	9	Medium	Moderate
Number of locations with threat	10	14	12	20	7	9	8		Higher numbers ind	icate higher threat le	vel

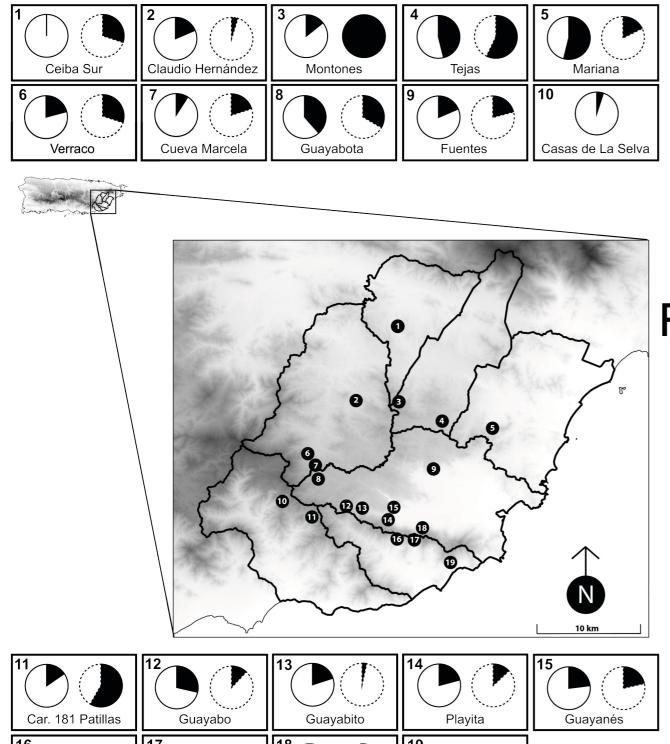
<sup>\*</sup> No E cooki were found & Bd Prevalence on Common Coqui 77%

Threat Level	Score
High	11 16
Moderate	6 10
Low	1 5

Threats	Values*	
Agriculture	2	
Erosion	3	
Habitat Modification	3	
Trash	2	
Bd prevalence < 50%	1	
Bd prevalence > 50%	2	
Invasive Predators	1	
Parasites	1	
Low Abundance	3	
Medium Abundance	2	
High Abundance	1	
	Agriculture Erosion Habitat Modification Trash Bd prevalence < 50% Bd prevalence > 50% Invasive Predators Parasites Low Abundance Medium Abundance	

<sup>\*</sup>The higher the number the larger the threat

## Pathogen prevalence and load significantly increased during the cool-dry season.



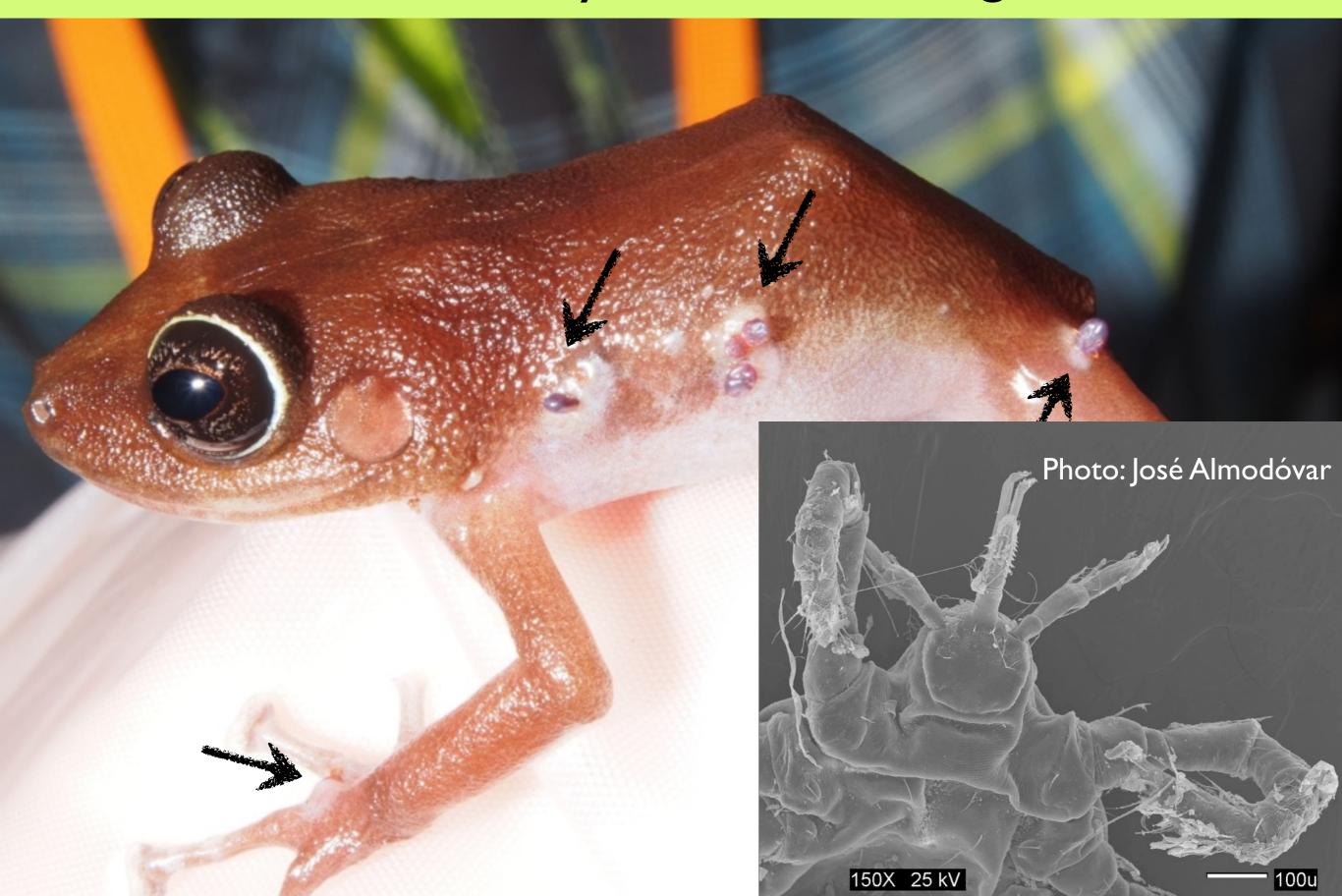
100% of populations are infected:

 Prevalence > 50% at 6 localities (30% of total)

Patterns are not associated to:

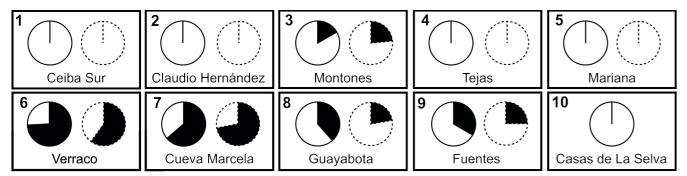
- Elevation
- Lat/Long
- Habitat type (caves vs streams)

## Ticks mostly infect adult frogs.

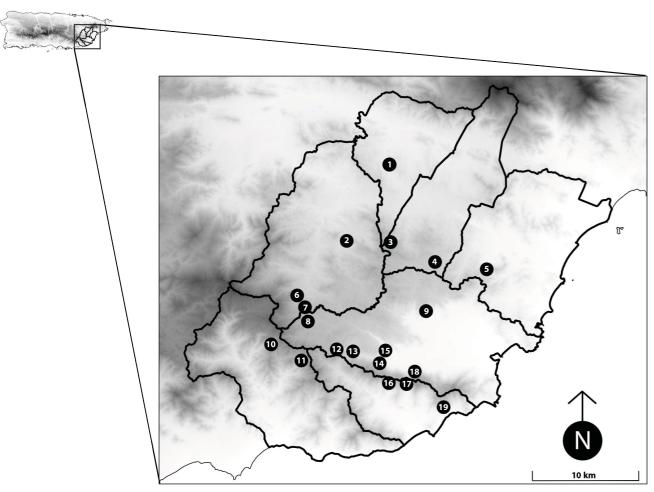




#### Parasite prevalence significantly increased during the cool-dry season.



Ticks also showed seasonal patterns:

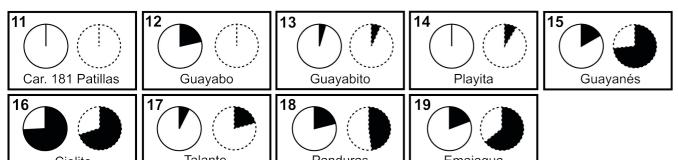


Avg. of 4 (I-30 ticks).

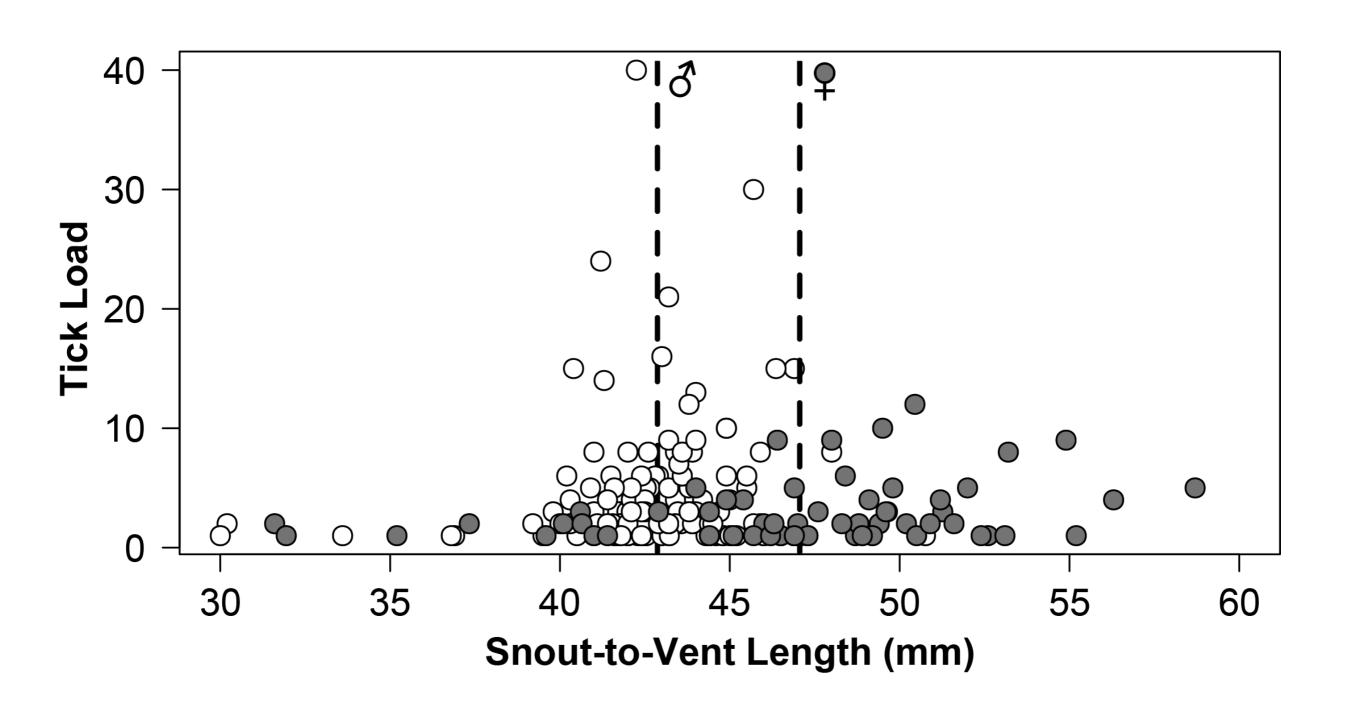
Some localities were "tick-free"

68% of populations were parasitised by ticks.

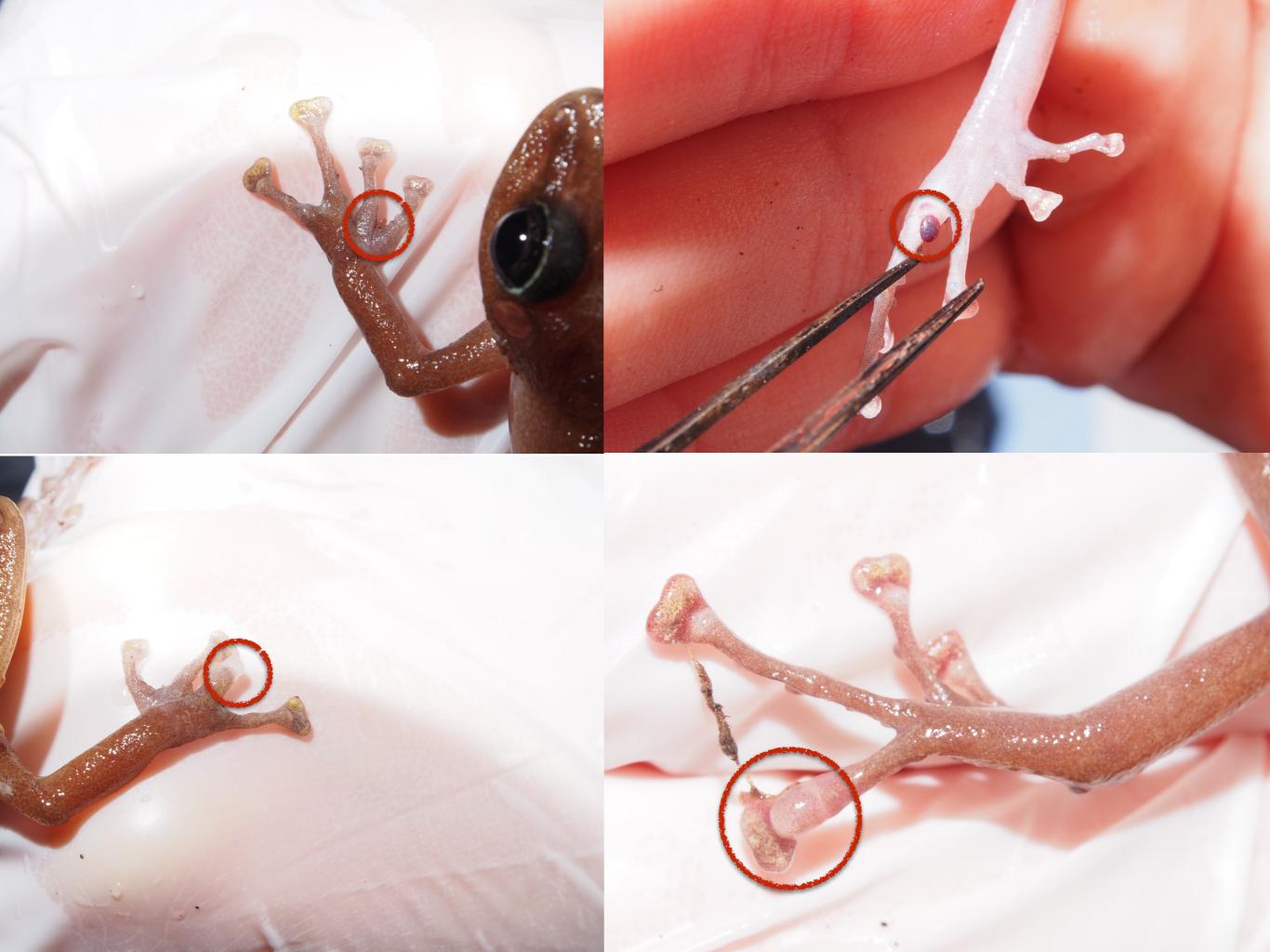
Skin discoloration marks



#### Ticks mostly infect male frogs.



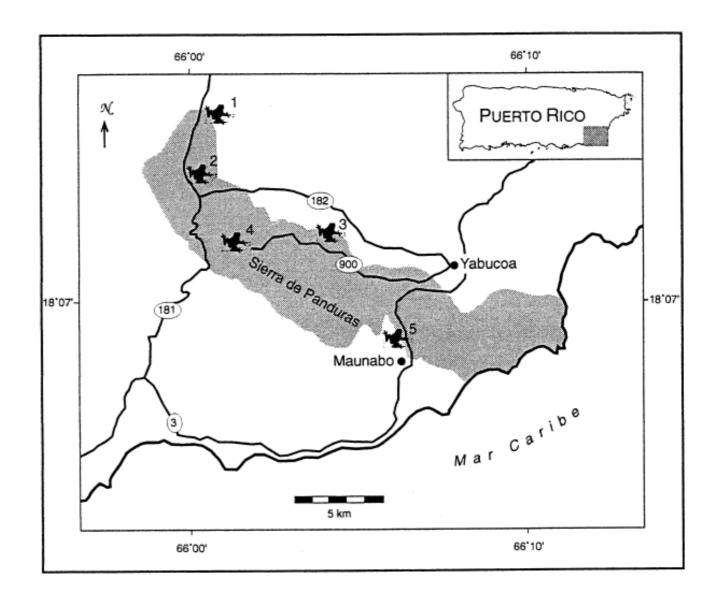




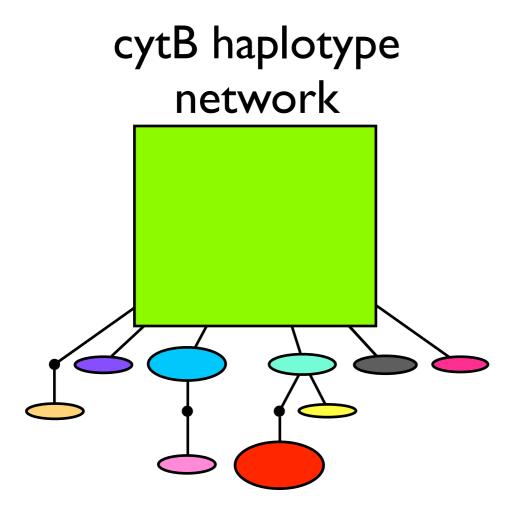
#### Cave-dwelling species should show genetic structure

- Burrowes & Joglar (1999) did not find genetic differentiation among 5 populations in Sierra Pandura.
- Guajones move approximately 30m to forage (Vega-Castillo, 2001).





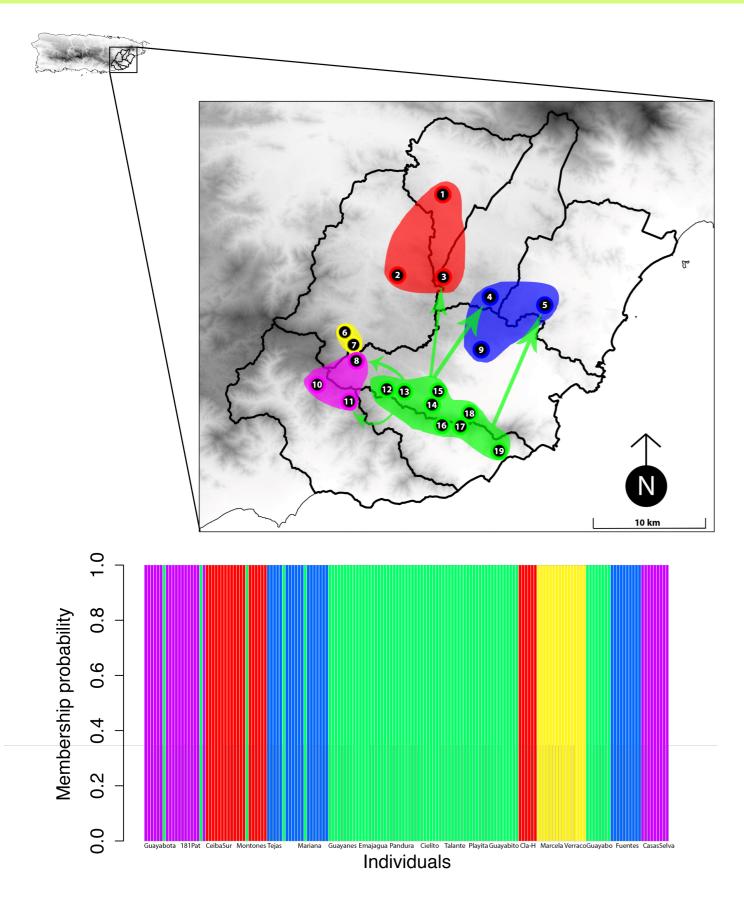
#### Mitochondrial DNA: only 10 haplotypes!



- 10 haplotypes in 96 individuals.
  - 49 haplotypes in 77 individuals (E. coqui-Velo-Antón et al. 2007)
- 17% divergence from E. coqui.
- Typical pattern of population expansion followed by bottlenecks.

Very little genetic variation with no apparent population structure.

#### High population structure among critical habitats

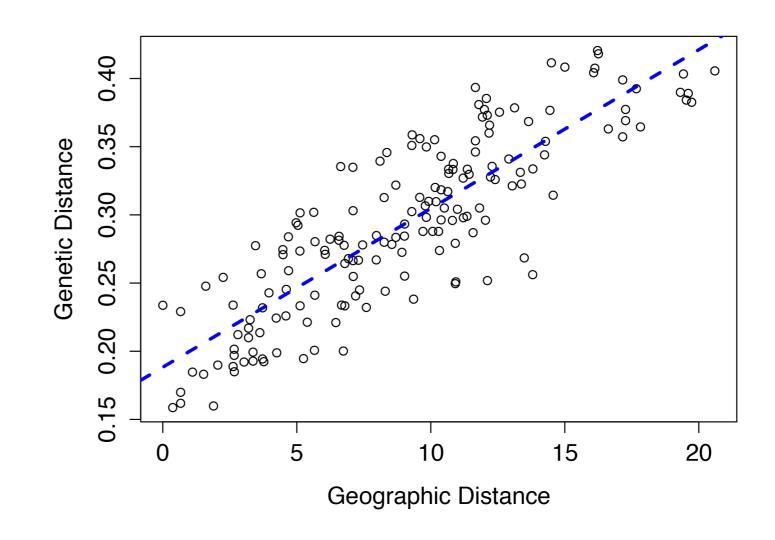


We used genotyping-by-sequencing (GBS) to genotype 175 E. cooki individuals at 3125 SNP loci throughout their entire genome. We then performed multivariate analyses to identify genetic clusters using the variation in allele frequencies across all 3125 SNP loci. Discriminant analyses revealed that the 19 populations of E. cooki are divided into five genetic clusters.

#### Isolation by distance only in North-eastern habitats

We found that gene flow was limited to very short distances, which led to increased genetic differentiation as the geographic distance became greater.

The largest geographic distance is between Mariana and Casas de la Selva (20.6 km), and it corresponds with individuals showing the highest genetic distances.



#### Increase awareness among landowners and students

# Conoces al Coqui Guajón? www.coquiguajon.org











## Increase awareness among landowners and students









#### Next steps

- I. Genotype more individuals around cueva marcela and verraco.
- 2. Examine patterns of genetic differentiation from males vs. females.
- 3. Link phenotypes with genotypes to find resistant individuals.

### Questions?

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