

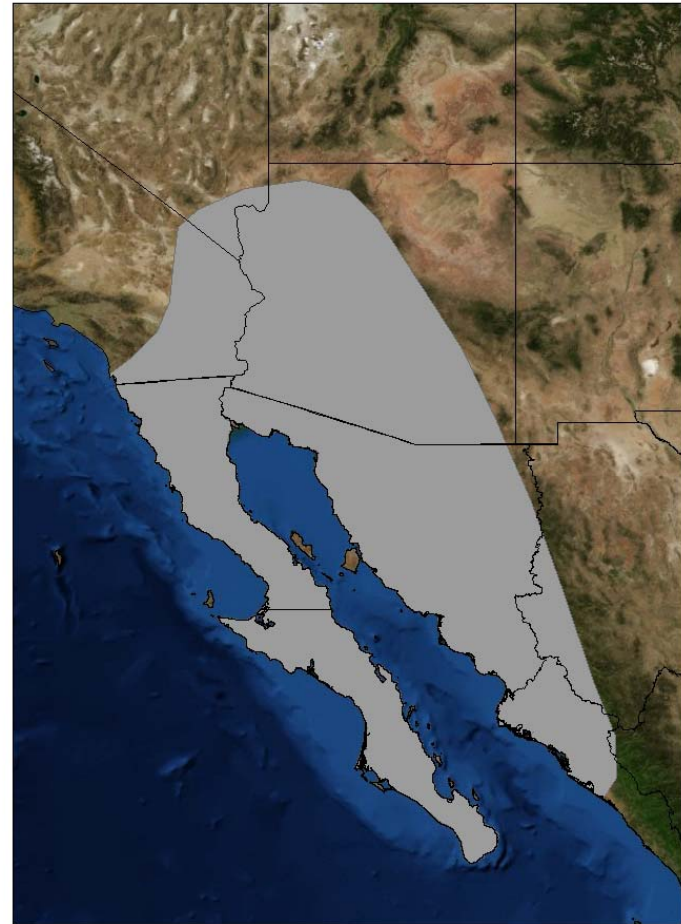
Determining space use and  
population demographics of the  
California leaf-nosed bat  
(*Macrotus californicus*): a  
population genetics approach

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# Background

- MSCP evaluation species
- Non-migratory, colonial, cave roosting bat
- SW Deserts



# Background

- Maternity colonies
- Males maintain harem groups of females
- Single young born in mid summer
- Forage on insects within 10 km of their roost

# Stop being difficult, bat.

- Not readily detected using acoustic surveys
  - Low intensity calls
  - Similar to other species
- Low capture rate in mist netting
  - Makes gathering demographic data difficult

# A different approach

- Natural history via molecular genetics
- Microsatellites
  - Identifying population structure
  - Gene flow
  - *M. waterhousii*
  - Relatively inexpensive

# Purpose

- Develop a conservation strategy for the LCR population



# Goals

1. Demographic data along LCR
  - Population structuring
  - Genetic diversity
  - $N_e$
2. Space use and foraging habits
  - Document movement patterns and site preferences
  - Identify differences between sexes in roosting, foraging, and gene flow

# Sampling

- 15 ind/cave along LCR
- Sample during post restoration monitoring
- “Outgroups” from Mexico and off LCR





# Sequencing

- Control region of mtDNA
  - Maternal
  - Moderately variable
- Microsatellites
  - Biparental
  - Highly variable

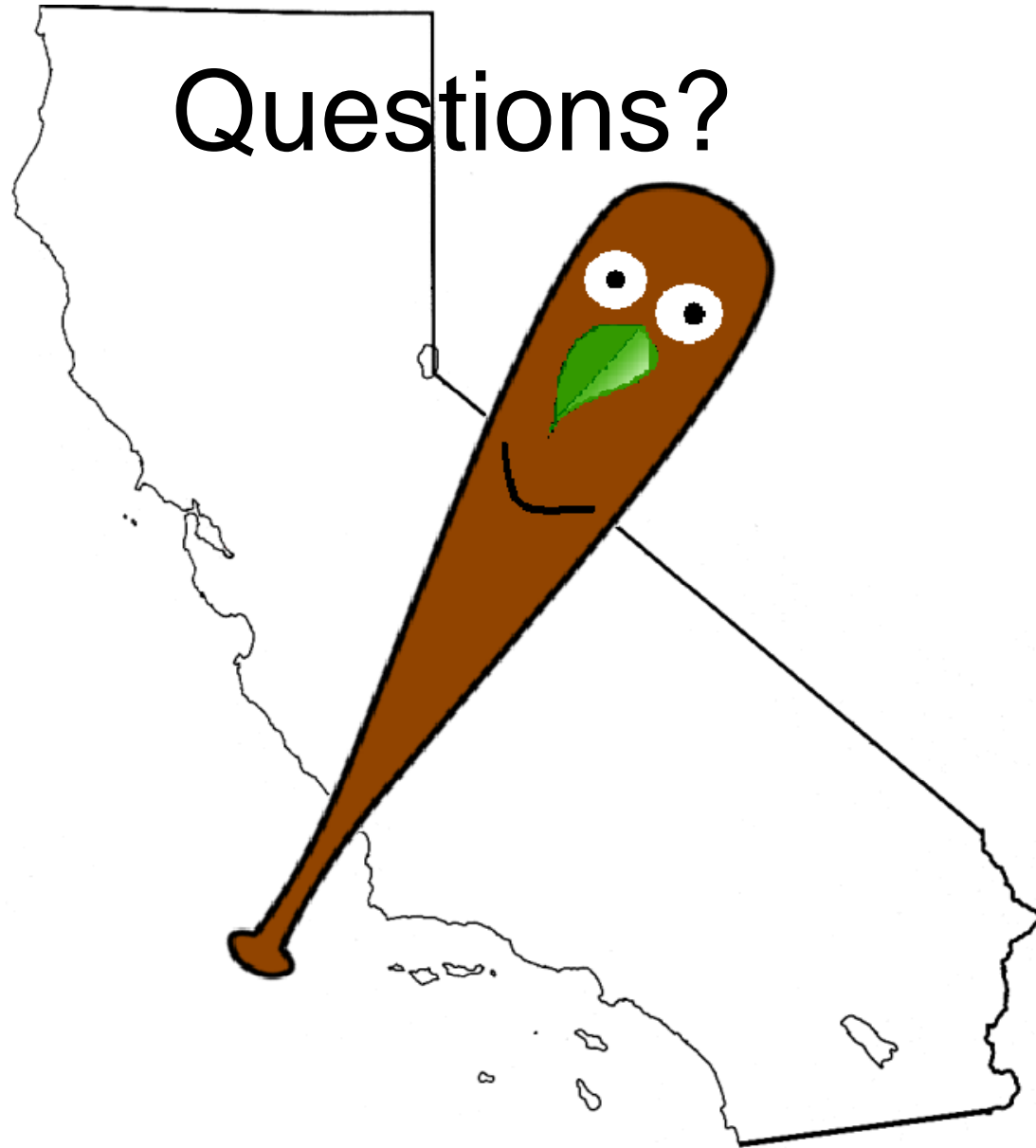
# Analyses

- Demographic data along LCR (Goal 1)
  - Population structuring among caves
  - Phylogenetic networks
  - $F_{st}$  and  $F_{is}$
  - AMOVA
  - $N_e$
  - Bayesian Skyline plot

# Analyses

- Space use and Foraging Habits (Goal 2)
  - Assignment of foraging bats to caves
    - Program Structure
  - Document movement patterns and site preferences
    - Between roosts
    - From roost to habitat
  - Identify differences between sexes
    - Compare mtDNA to Microsat

Questions?



*Macrotus californicus*