# **Recovery Criteria**

Reclassify to Threatened:

1. Two viable populations of at least 240 individuals (adults and subadults) maintained for 12 years

2. Sufficient habitat ... to support these populations is protected for the long-term

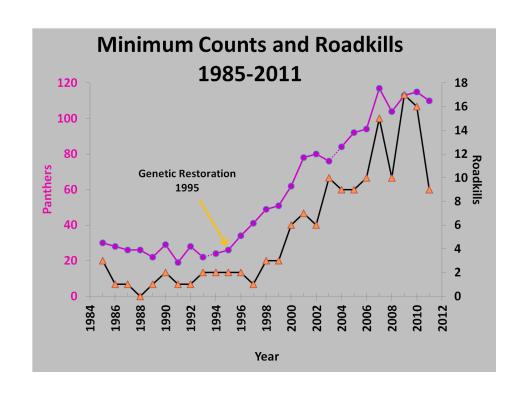
Delist (no longer Endangered or Threatened):

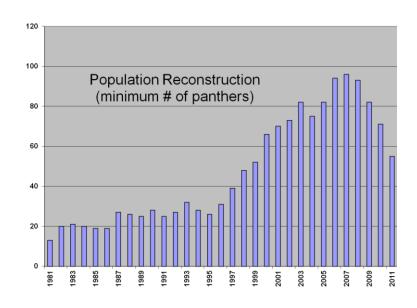
1. Three viable populations of at least 240 individuals (adults and subadults) maintained for 12 years

2. Sufficient habitat ... to support these populations is protected for the long-term

# Population size and trend

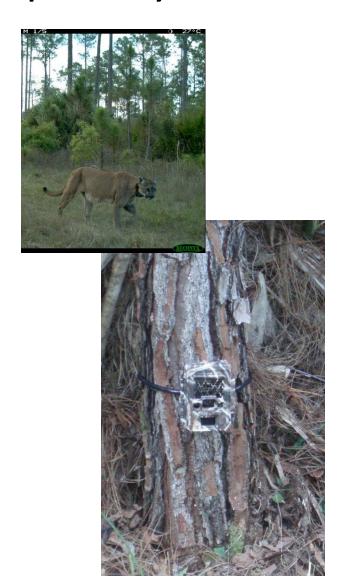
 Develop inventory and monitoring techniques that produce reliable population estimates





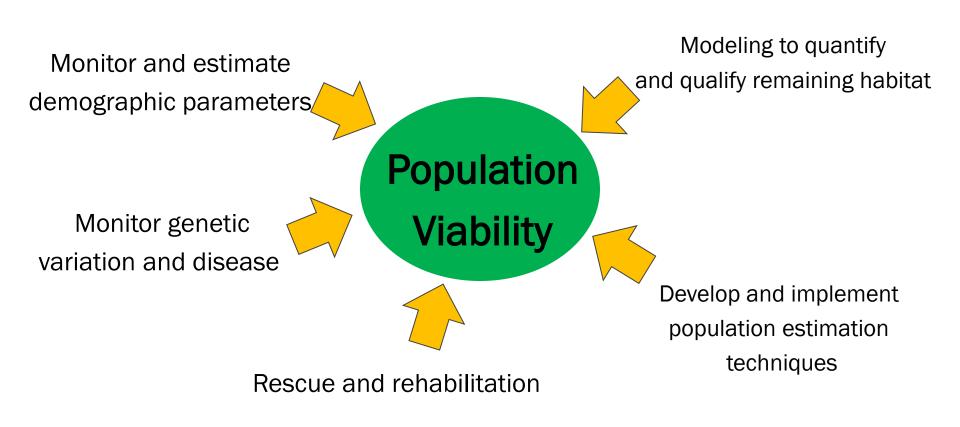
## FWC Panther Camera Trap Study

- Objective
- Phased approach
- Addition Lands of BCNP
- 162 km<sup>2</sup> study area
- 4 month sampling period
- Collect data on
  - Panthers
  - Deer
  - Other wildlife



## **Current Research Priorities**

# Assessing Progress Towards Recovery



# Genetic Management



### Florida Panther Taxonomic Status

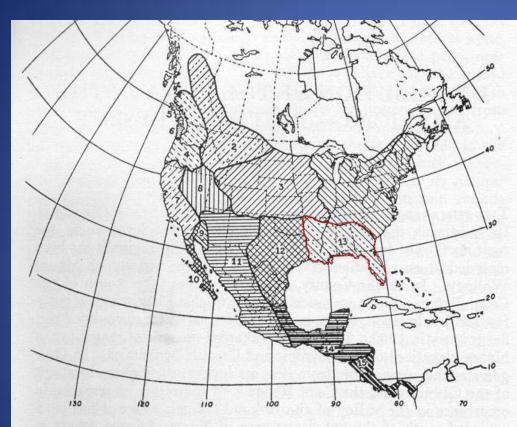


Figure 1. Distribution of subspecies of Felis concolor in North and Middle America

1. Felis concolor couguar

2. F. c. missoulensis

3. F. c. hippolestes

4. F. c. oregonensis

5. F. c. vancouverensis

6. F. c. olympus

7. F. c. californica

8. F. c. kaibabensis

9. F. c. browni

10. F. c. improcera

11. F. c. axteca

12. F. c. stanleyana

13. F. c. coryi

14. F. c. mayensis

15. F. c. costaricensis

15 Subspecies of Puma concolor

Young and Goldman 1946 Hall 1981 J Hered. 2000 May-Jun;91(3):186-97.

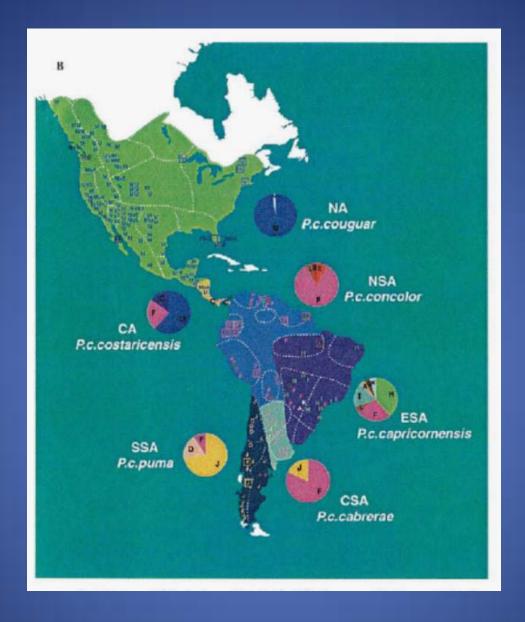
#### Genomic ancestry of the American puma (Puma concolor).

Culver M, Johnson WE, Pecon-Slattery J, O'Brien SJ.

Laboratory of Genomic Diversity, National Cancer Institute, Frederick Cancer Research and Development Center, MD 21702-1201, USA.

#### Abstract

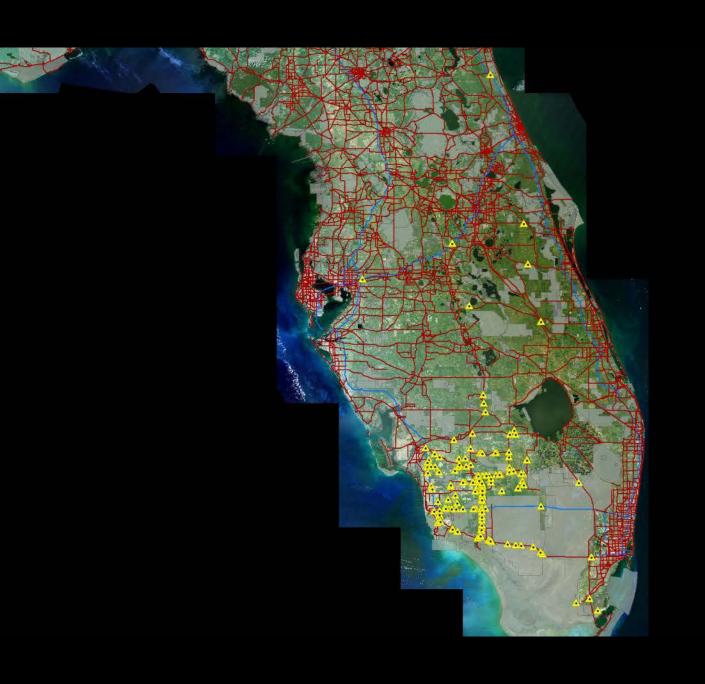
Puma concolor, a large American cat species, occupies the most extensive range of any New World terrestrial mammal, spanning 110 degrees of latitude from the Canadian Yukon to the Straits of Magellan. Until the recent Holocene, pumas coexisted with a diverse array of carnivores including the American lion (Panthera atrox), the North American cheetah (Miracynonyx trumani), and the saber toothed tiger (Smilodon fatalis). Genomic DNA specimens from 315 pumas of specified geographic origin (261 contemporary and 54 museum specimens) were collected for molecular genetic and phylogenetic analyses of three mitochondrial gene sequences (16S rRNA, ATPase-8, and NADH-5) plus composite microsatellite genotypes (10 feline loci). Six phylogeographic groupings or subspecies were resolved, and the entire North American population (186 individuals from 15 previously named subspecies) was genetically homogeneous in overall variation relative to central and South American populations. The marked uniformity of mtDNA and a reduction in microsatellite allele size expansion indicates that North American pumas derive from a recent (late Pleistocene circa 10,000 years ago) replacement and recolonization by a small number of founders who themselves originated from a centrum of puma genetic diversity in eastern South America 200,000-300,000 years ago. The recolonization of North American pumas was coincident with a massive late Pleistocene extinction event that eliminated 80% of large vertebrates in North America and may have extirpated pumas from that continent as well.



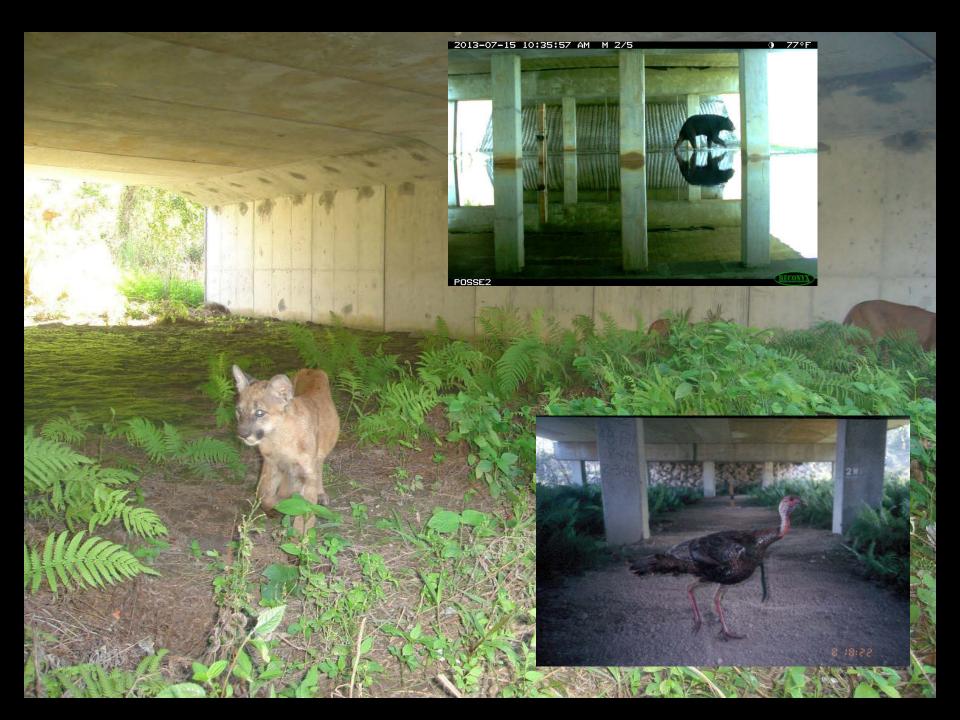
North American puma = P. concolor couguar

# Transportation Subteam





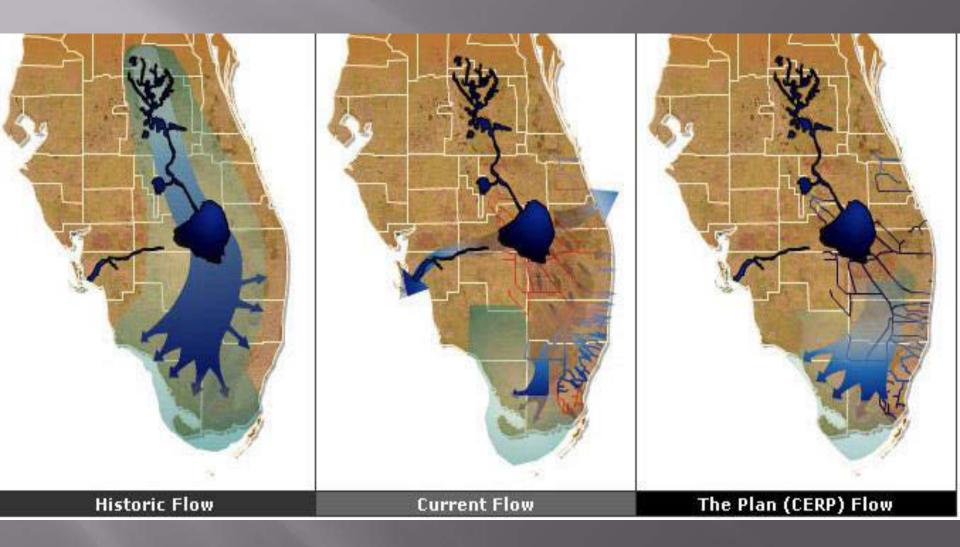




# HABITAT RESTORATION AND MANAGEMENT

Florida Panther Recovery
Implementation Team Meeting
May 22, 2014

## Hydrologic Restoration





## Control invasive plants



## Accomplishments

- Recognized the need for a sub-team for this topic
- Combine hydrological, botanical, and fire ecology expertise
- Relationship between panther and prey populations
- Guidance where restoration efforts could occur

# Importance of Private Lands

