

A conservation genetic assessment of the Florida snail kite

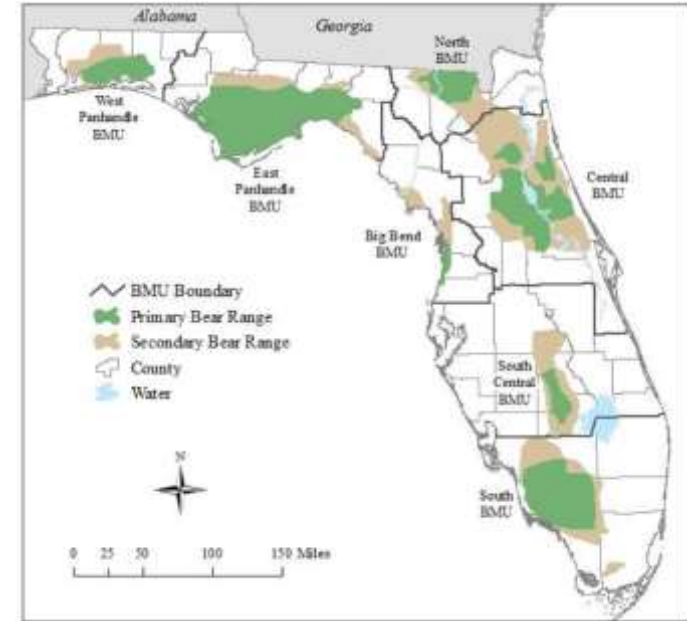


Ellen Robertson
Rob Fletcher
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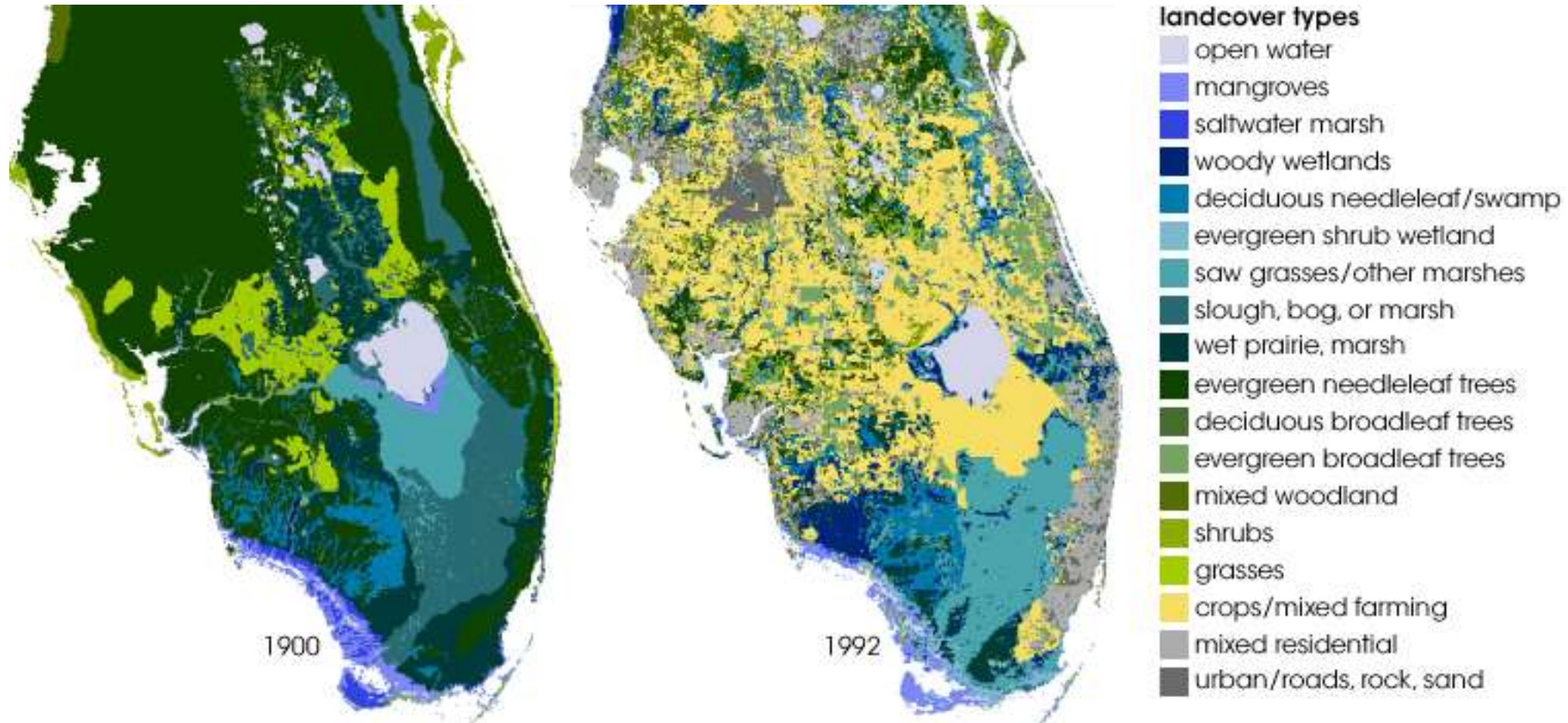


Conservation genetics

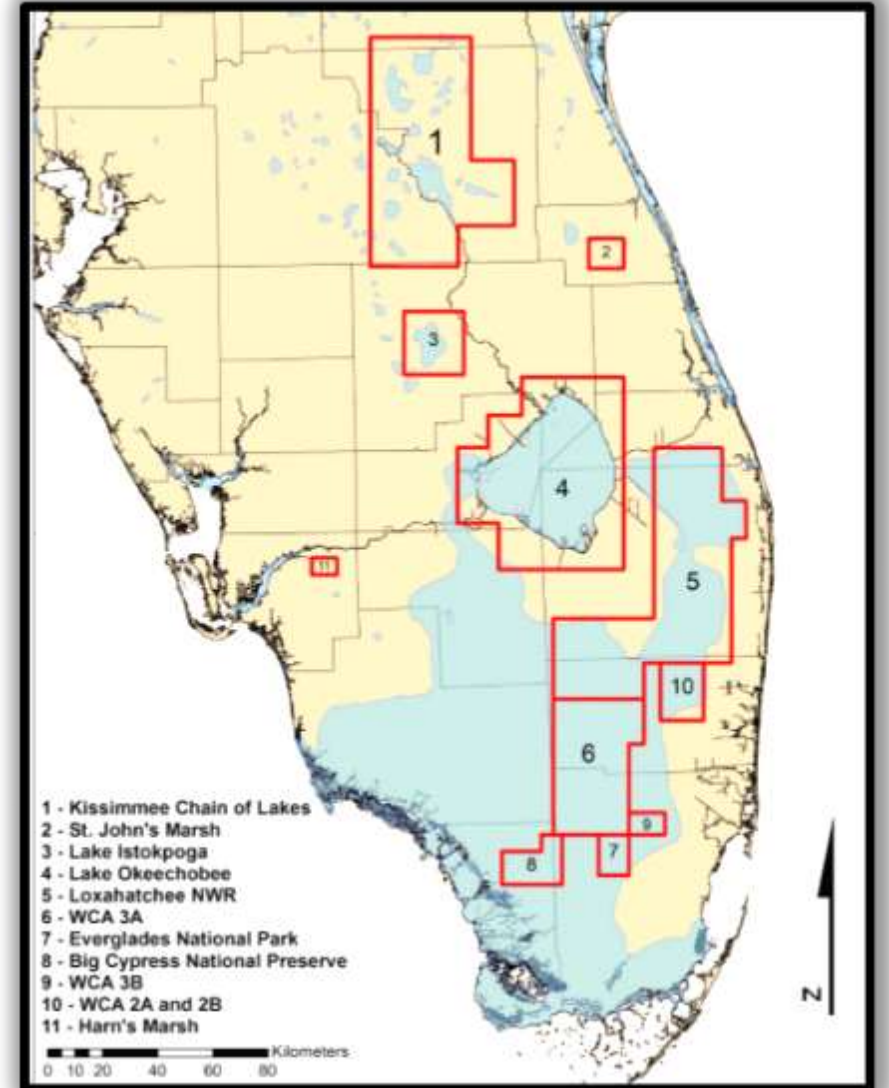
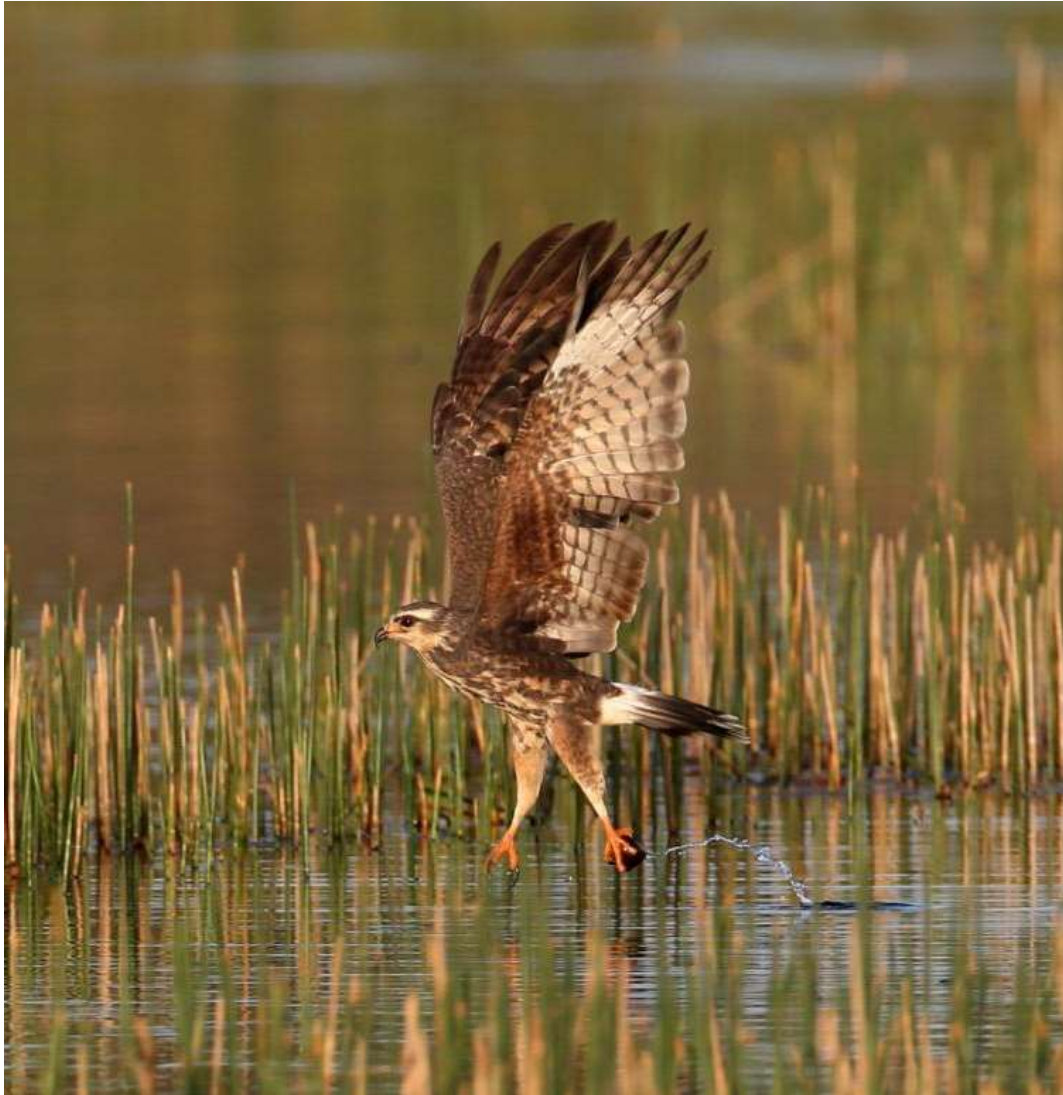
- Genetic structure
 - delineating management units
- Genetic diversity
 - indicator for potential genetic problems
 - inbreeding
 - reduced evolutionary potential



Florida: potential for genetic changes:

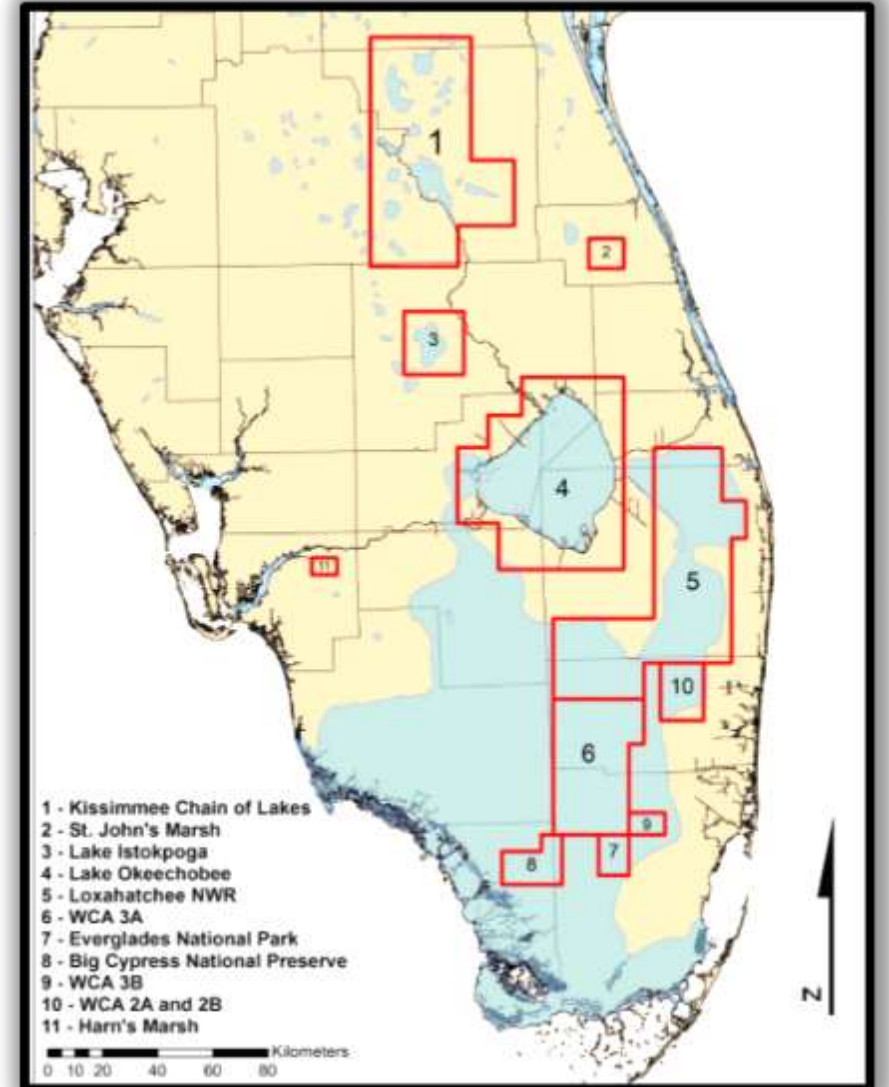


Snail kites (*Rostrhamus sociabilis plumbeus*)

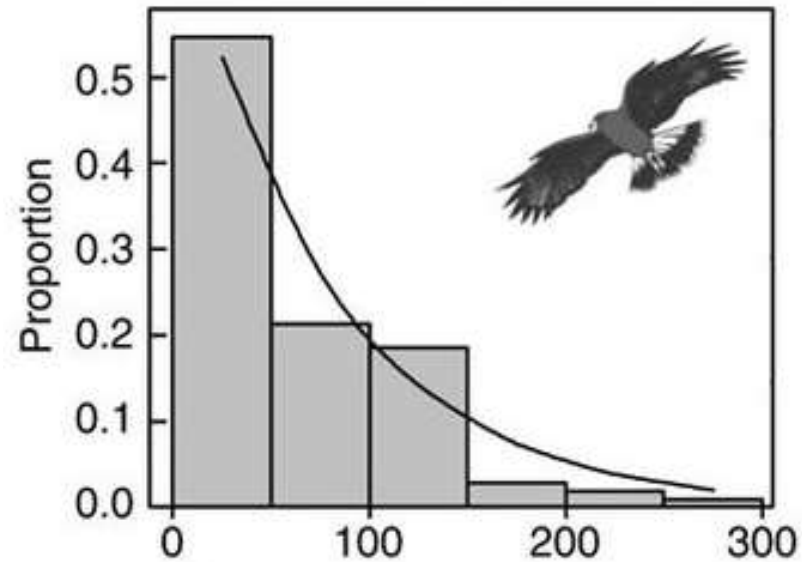


The monitoring program

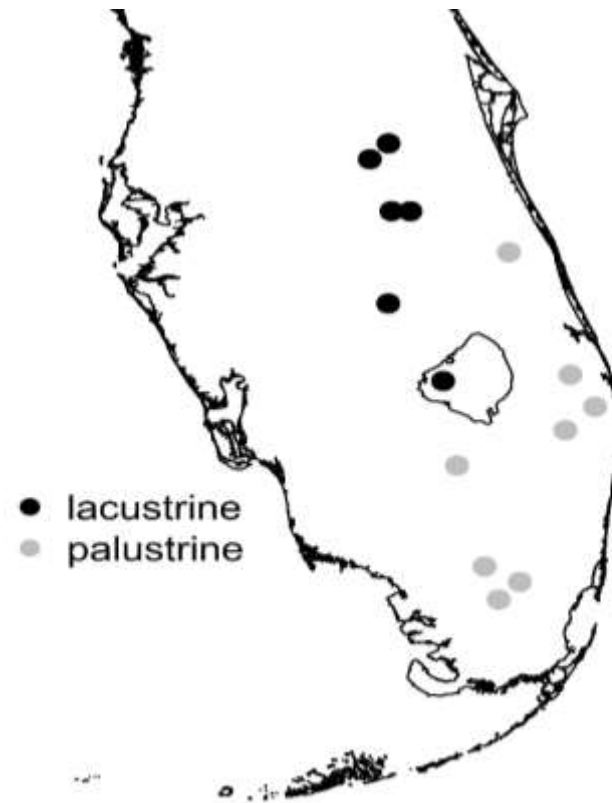
Standardized mark-resight surveys and nest finding across the geographic range



Movements are fairly frequent and widespread but still potentially structured



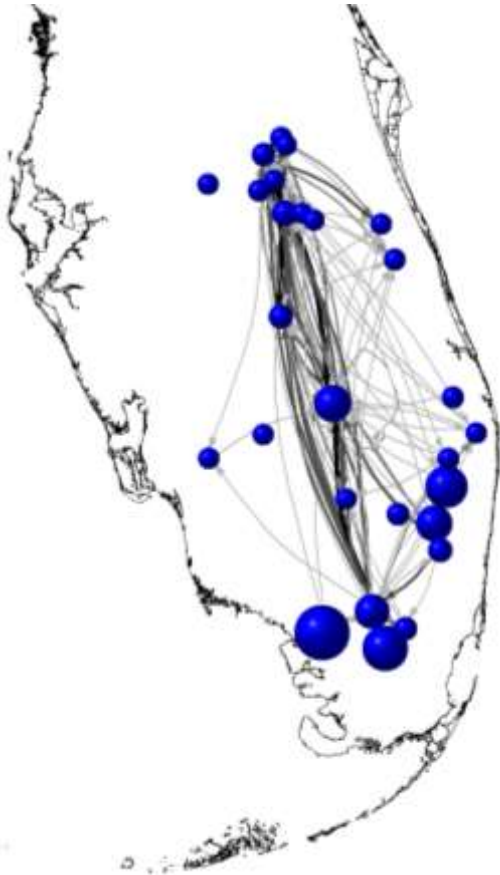
Histogram of movements
Fletcher et al. 2013



Fletcher et al. 2015

Movements are fairly frequent and widespread but still potentially structured

Movement



dispersal +

successful reproduction



Effective dispersal:

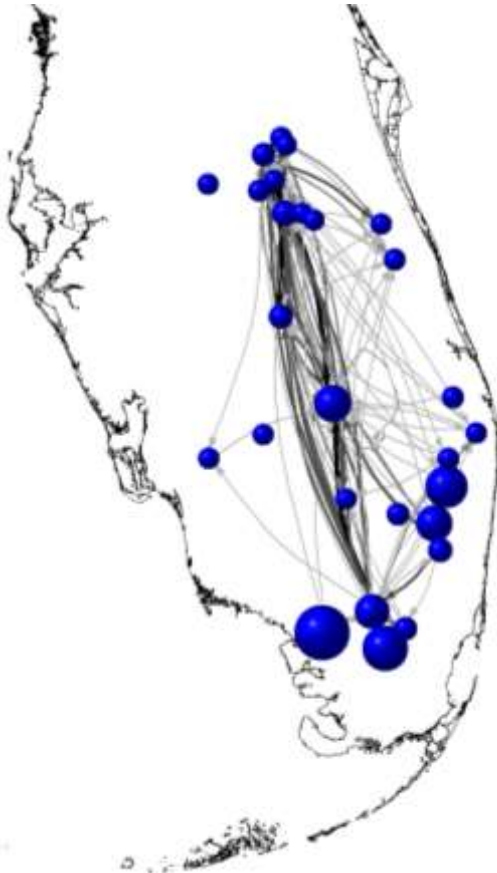
dispersal +

successful reproduction

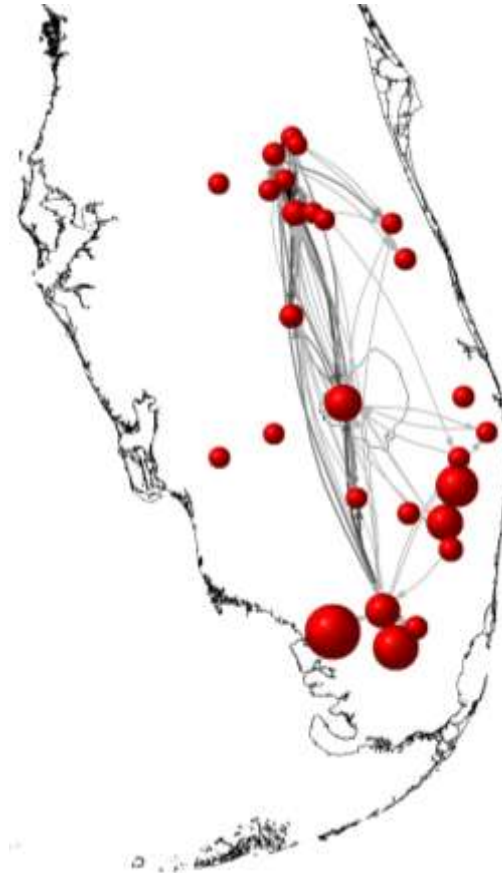


Reproduction following movement is less frequent and less widespread

Movement

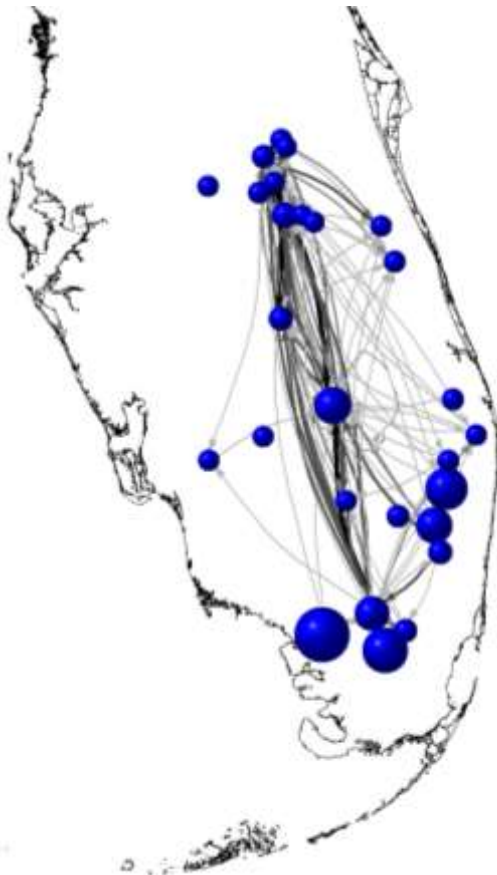


*Movement + Nesting Attempt
(dispersal)*

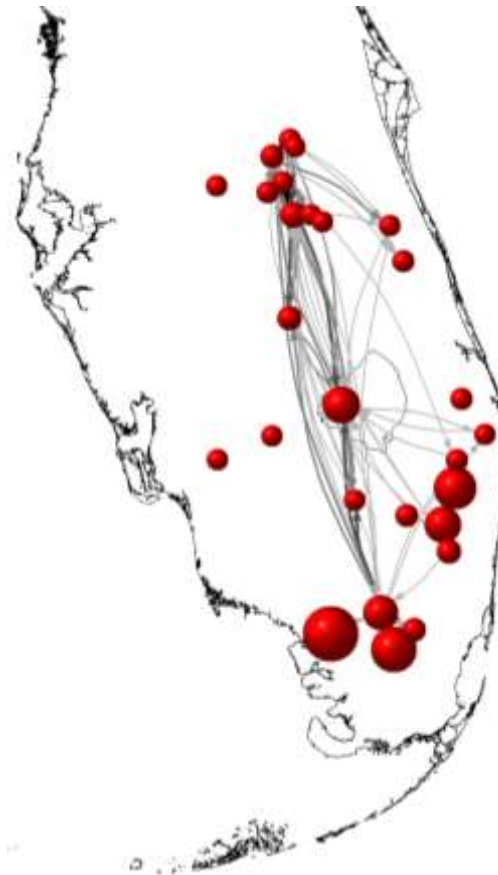


Reproduction following movement is less frequent and less widespread

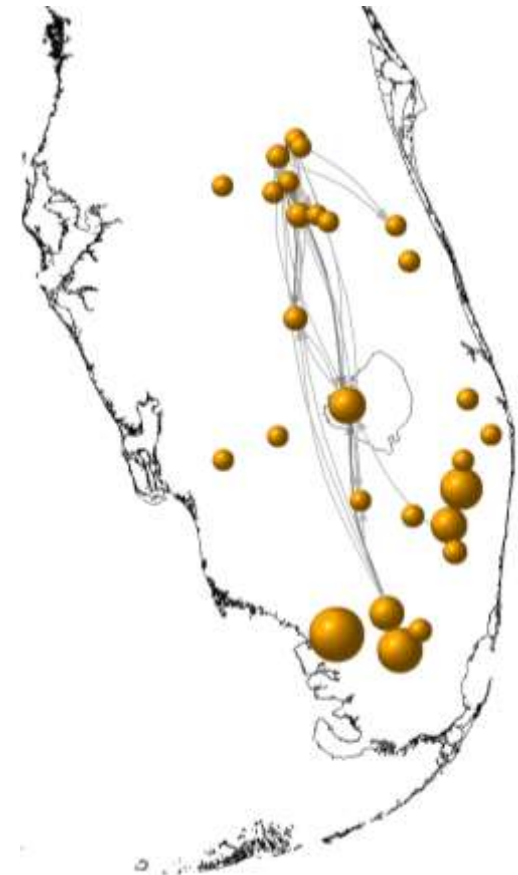
Movement



*Movement + Nesting Attempt
(dispersal)*



*Movement + Successful Nesting
(effective dispersal)*



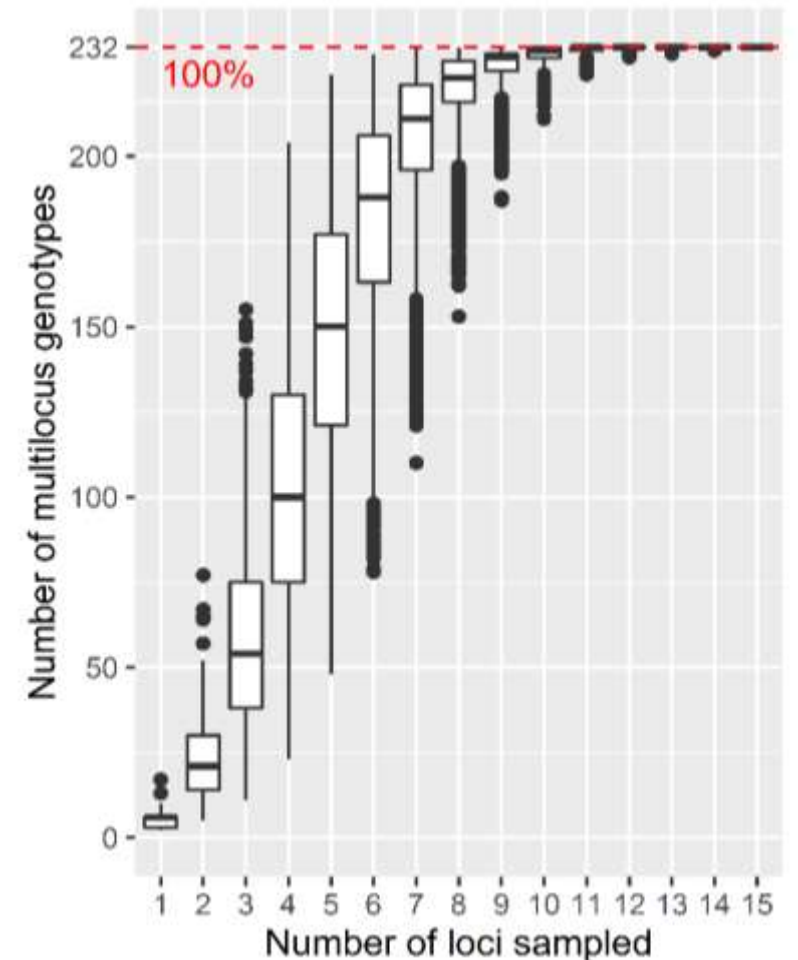
Is there genetic structure in Florida snail kites?

- Developed microsatellite markers:
 - 235 feather samples (from nestlings)
 - 2013 (N = 114) and 2014 (N = 121)
 - 14 wetlands across the breeding range



15 microsatellite markers

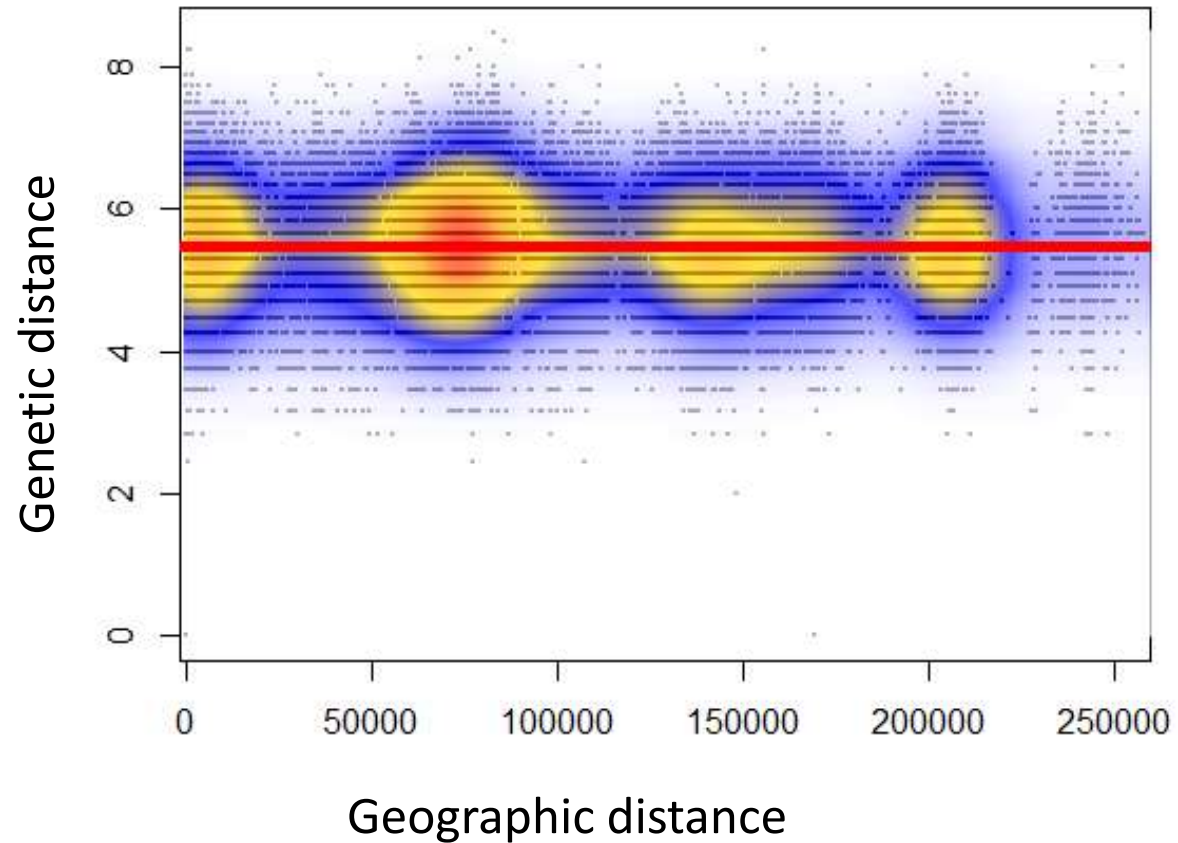
- HWWE
- Linkage equilibrium
- Checked for null alleles, allelic dropout
- Genotyping error
 - Repeated PCR and genotyping on 19% of samples



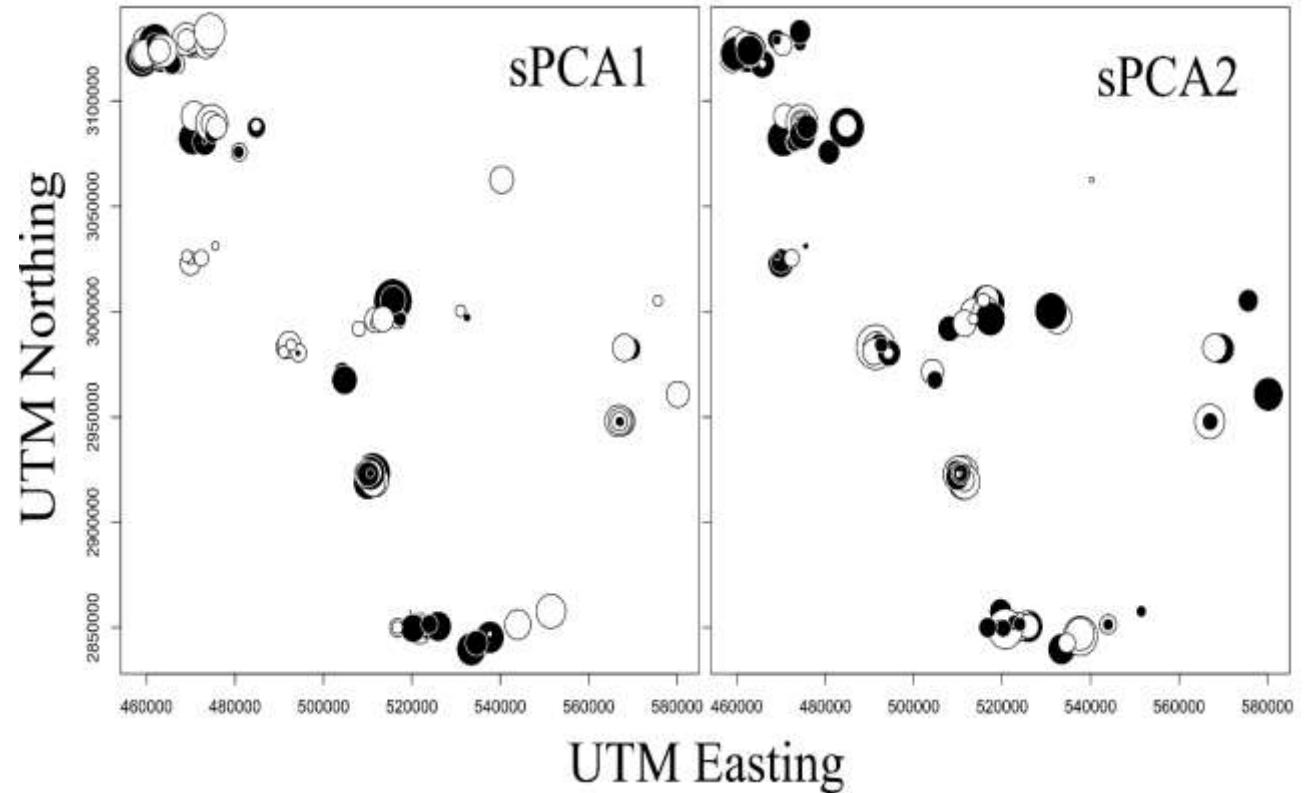
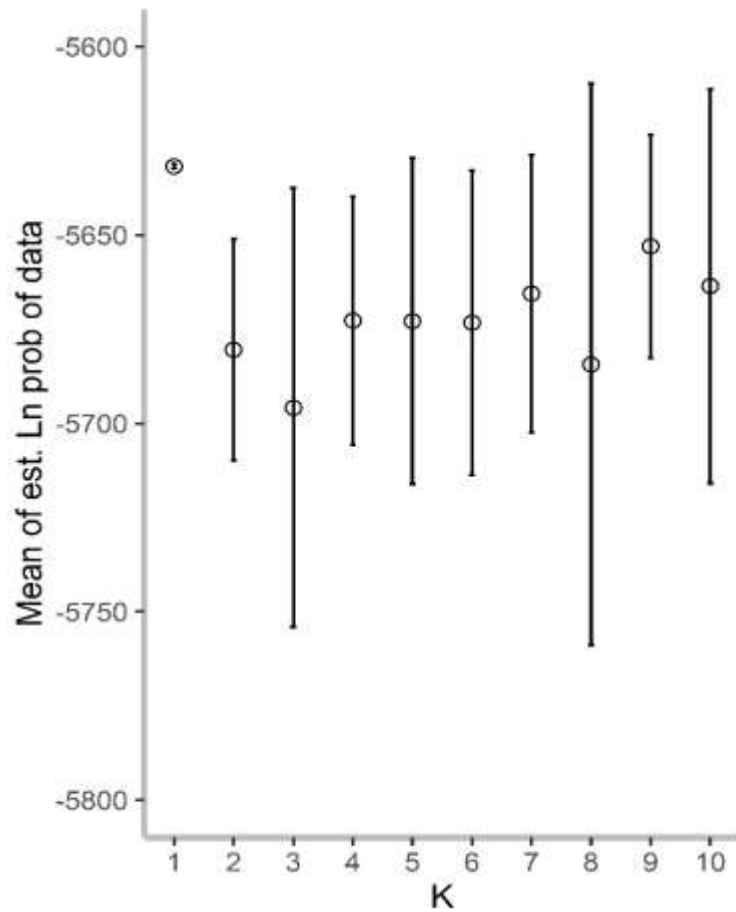
Spatial analyses:

- Isolation by distance
 - Mantel test
- Structure
 - Program Structure
 - Geneland
 - spatial coordinates as a prior
 - Ordination
 - Spatial Principal Component Analysis (sPCA)
 - Memgene
 - Spatial autocorrelation

No evidence for genetic isolation by distance



No evidence for genetic structure



Low genetic diversity relative to other rare raptors

Species	IUCN Status	A	Ar	N (for Ar)	He
snail kites	least concern	2.5	2.28-2.61	8-60	0.37
northern goshawk	least concern	10.4	NA	NA	0.81
bearded vulture	near threatened	4.3-6.0	NA	NA	0.54-0.68
common kestrel	least concern	4.8-7.8	4.5-5.0	14-28	0.63-0.71
peregrine falcon	least concern	4.25	NA	NA	0.51
Spanish imperial	vulnerable	4.9	4.9	10-20	0.55
Egyptian vultures	endangered	NA	2.4-3.0	143-242	0.44-0.56
golden eagle	least concern	4.62	NA	NA	0.49
mountain hawk-eagle	least concern	3.67	NA	NA	0.56


Conclusions and future research

- Snail kites are genetically panmictic
 - Should monitor and manage as a single population
 - Management in one location can have implications across the range





Conclusions and future research

- Snail kites are genetically panmictic
 - Should monitor and manage as a single population
 - Management in one location can have implications across the range
 - Snail kites have relatively low genetic diversity
 - USFWS Species-level Recovery Action
 - Fitness consequences?
- 



Acknowledgments

Monitoring design and implementation: Wiley Kitchens, Rob Bennetts, Vicky Dreitz, Don DeAngelis, James Nichols, Bill Kendall, Jim Hines, FL Coop Unit

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Why the disconnect between dispersal and genetics?

- These are different metrics:
 - can be time lags (ecological vs evolutionary time scales)
 - only a small amount of dispersal can result in low genetic differentiation
- Genetic structure is also affected by effective population size (N_e) through genetic drift
- Complicated system that is likely not at equilibrium
 - a possible bottleneck in mid-1900s
 - recent range expansion (with exotic snail)

Potential drivers of reduced effective dispersal?

