

# ECOLOGICAL DRIVERS OF MAMMALIAN TREE ISLAND USE

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## Tree Islands as Habitat

- Wildlife relies heavily on tree island habitat
  - Forage, dry refugia, forested cover, breeding/nesting/rearing sites
- Mammals especially reliant because of terrestrial affinity
- Quality mediated by tree island characteristics (e.g., size, elevation, plants)
  1. Diversity
  2. Composition
  3. Distributions (Zarnetske et al., 2017; Hamer et al., 2021; Ferreira Neto et al., 2021)
- Limited mammal research in Everglades
  - Multidecadal decline

# Methods

## ■ Site Selection

- Areas of interest (distinct hydrologically)

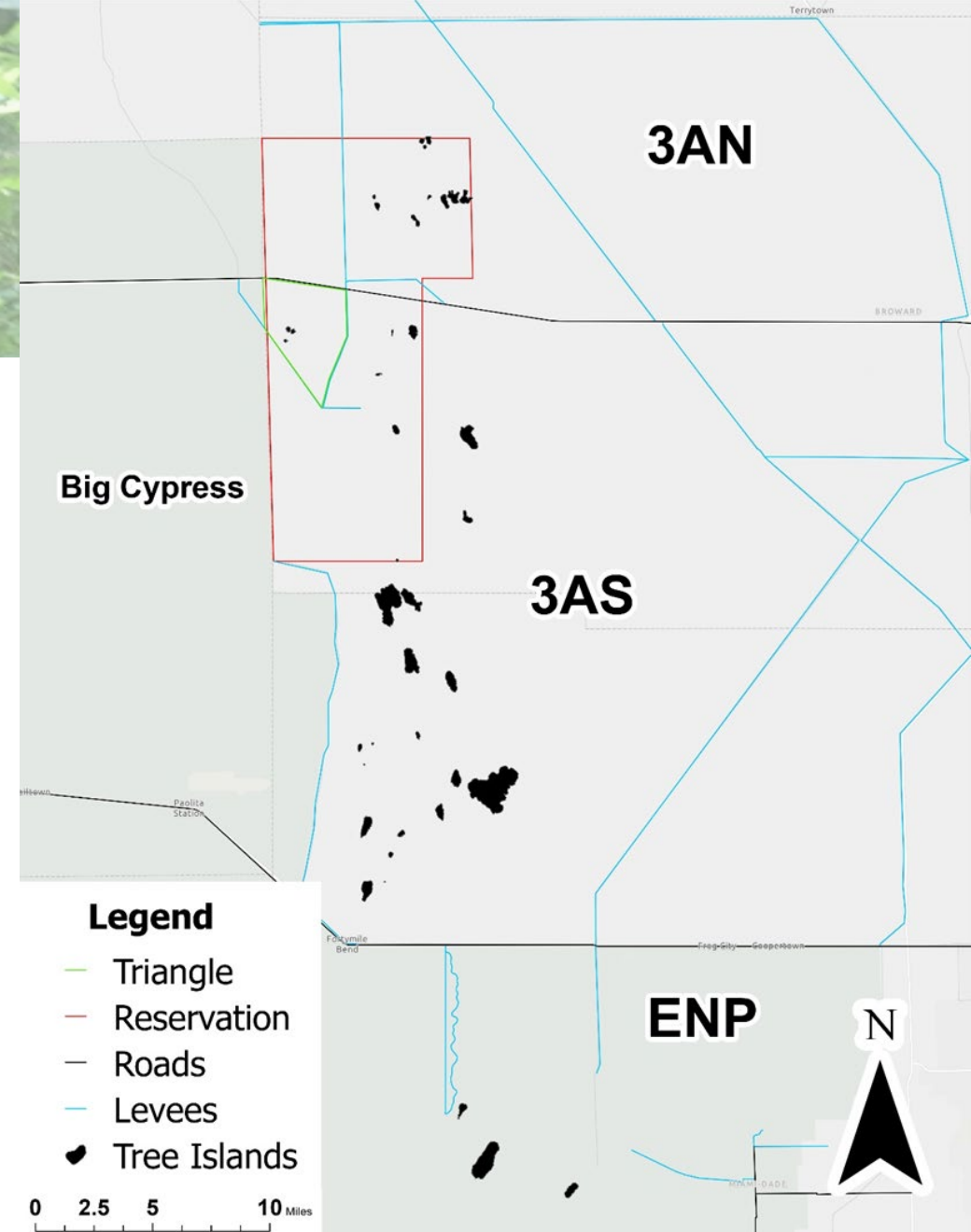
1. **3AN** (10 TIs)

2. **TRI** (3 TIs)

3. **3AS** (24 TIs)

4. **ENP** (3 TIs)

- 34 islands randomly along depth transect
- TRI and ENP added to ensure full range of combinations



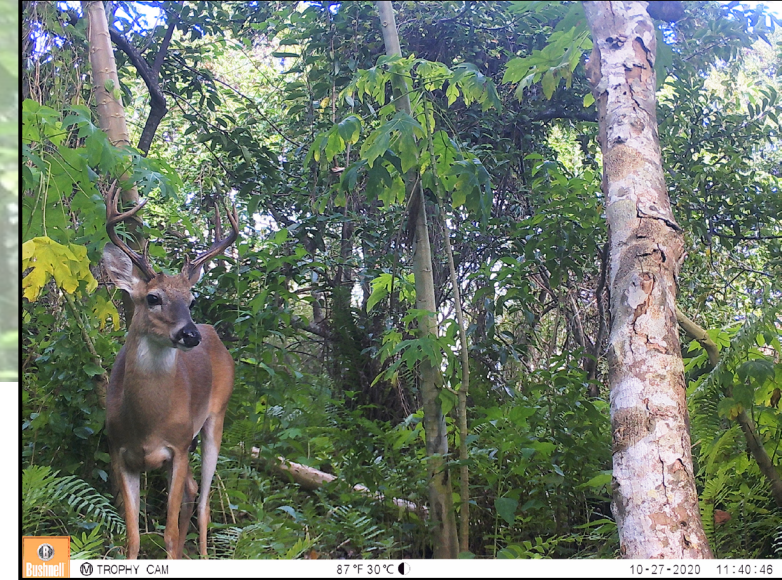
\* Polygons are 5x the scale of their true areal extent



# Methods

## ■ Camera Trapping

- Deployed using consistent protocol (i.e., height, angle, settings)
- Cams on “head” of TIs
  - More diverse than downstream areas (All species use head, not subset)
  - Last place to flood. During high-water, wildlife congregate here
  - Methodological consistency. ‘Head’ is not ‘habitat’
- Cameras positioned to optimize data collection
- Hourly occurrences → Relative abundance index (RAI)  
$$RAI_{spp1} = (\text{Number of occurrences}) / (\text{Number of trap days})$$





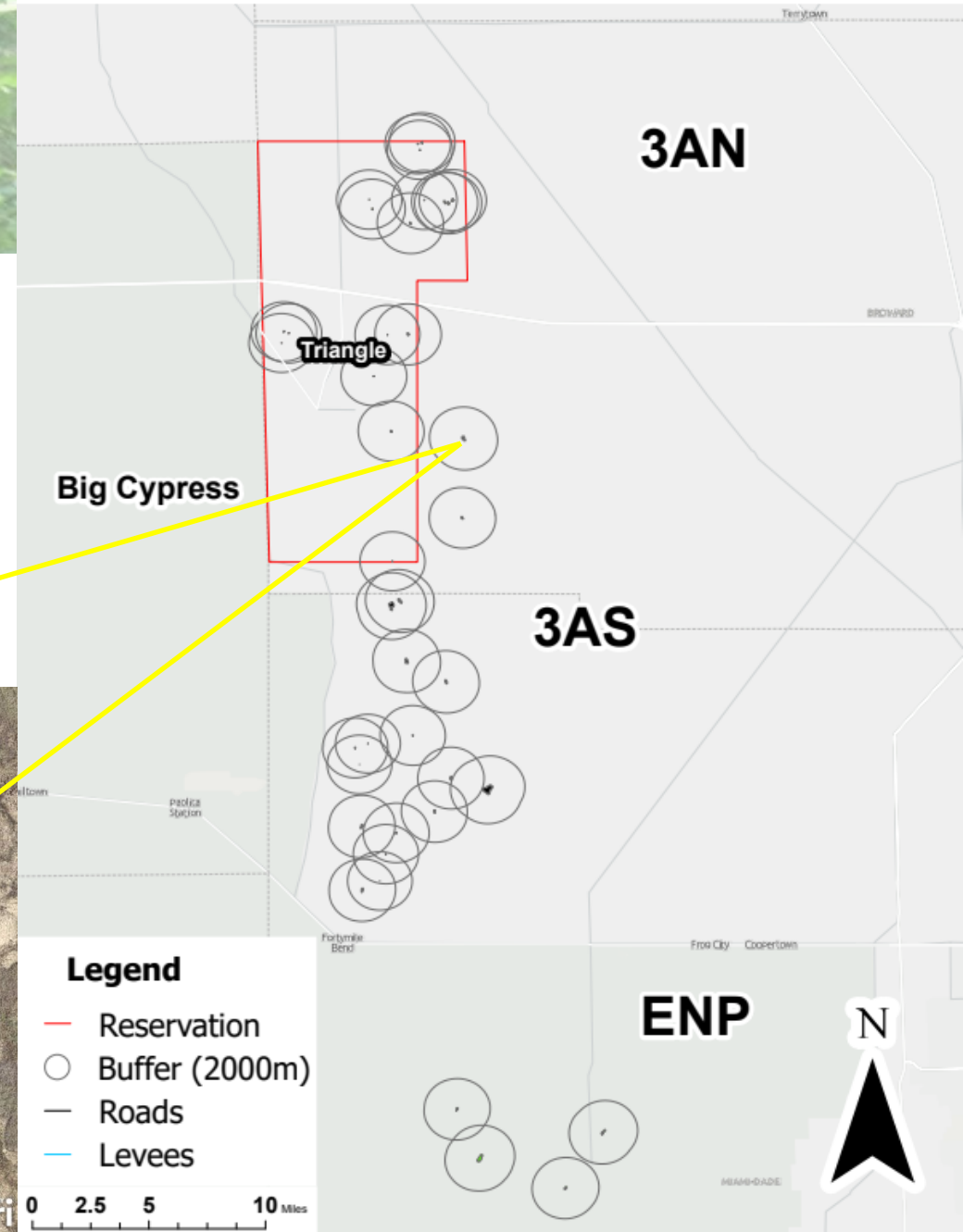
# Methods

## ■ Landscape Characteristics

- ArcGIS Pro to digitize heads
- Measure head area
- Spatial Rings
  - 100/250/500/1000/2000m
  - # of Neighbors
  - Area of Neighbors



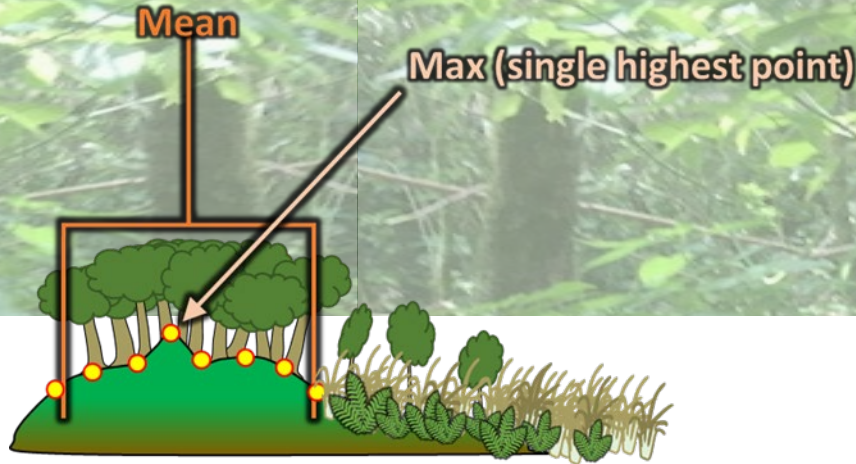
## Study Tree Islands





# Methods

## ▪ Elev./Hydrology



- Head elevations surveyed (post-Eta in Nov. 20)
  - Transect -10m (marsh), through head (not tail)
  - Measured 3x every 5 – 10 m; Hammock vs Bayhead/swamp, respectively
    - 3AS Triangle 3AN
    - ENP (SOFTTEL)



- $(\text{EDEN water level}) - (\text{Survey Depth}) = (\text{Plot Elevation})$ 
  - Derive tree island & marsh (rings) hydrologic variables  
Max. elev., Relative water, Hydroperiod, etc.





# Methods

## ■ Vegetation

- Same transect as the elevation survey

### 4x4 m square plot

- Ground cover/Vines/Saplings/Trees
  - DBH
  - Crown cover
  - Canopy height
  - Canopy structure

### 2x2 m subplot

- Herbs/Vines
  - Stem density
  - % Crown cover

Table 2. Stratification and defining criteria of plants in vegetation surveys.

<i>Strata</i>	<i>Criteria</i>
Herbs	Plants, including vines and seedlings, with a maximum height <1 m.
Shrubs	Plants, excluding vines, with a maximum height >1 m; woody species must have a DBH <1 cm.
Vines	Vines with a maximum height >1 m.
Saplings	Woody species with a maximum height >1 m and DBH of 1-5 cm.
Trees	Woody species with a maximum height >1 m and DBH ≥5 cm.

(Sah, 2004)





# Species Diversity

## ■ Coverage

- Occurrence data used to calculate **coverage**  
(How completely sample represents population)

Mean **coverage** = 99% (min. = 86%)

- Standardize to **asymptotic estimate of coverage** (100%)
  - Minimal extrapolation  
(only 6 increased) (Roswell et al., 2021)

$$C = 1 - \frac{f_1}{n} \left[ \frac{(n-1)f_1}{(n-1)f_1 + 2f_2} \right]$$

## Asymptotic Estimate

	<u>Richness</u>	<u>Hill-Shannon</u>
<b>Gamma</b>	12	3.2
<b>Alpha</b>	1 – 12	1 – 6
<b>Mean alpha</b>	5	2.4
<b>Beta</b>	2.4	1.3

Common Name	Scientific Name
Black bear	<i>Ursus americanus floridanus</i>
Bobcat	<i>Lynx rufus</i>
Coyote	<i>Canis latrans</i>
Eastern gray squirrel	<i>Sciurus carolinensis</i>
Feral hog	<i>Sus scrofa</i>
Florida panther	<i>Felis concolor coryi</i>
Marsh rabbit	<i>Sylvilagus palustris</i>
Raccoon	<i>Procyon lotor</i>
River otter	<i>Lutra canadensis</i>
Virginia opossum	<i>Didelphis virginiana</i>
Whitetail deer	<i>Odocoileus virginianus seminolus</i>
Rat	<div><div>Cotton mouse</div><div>Marsh rice rat</div><div>Hispid cotton rat</div><div>Norway rat</div><div>Roof rat</div></div> <div><i>Peromyscus gossypinus</i> <i>Oryzomys palustris</i> <i>Sigmodon hispidus</i> <i>Rattus norvegicus</i> <i>Rattus rattus</i></div>



# Species Diversity

## ■ Generalized Linear Mixed Models (GLMMs)

- GLMMs modelled variables as fixed effects
  - Site as random effect
  - AIC identified parsimonious models

❖ Ave. marsh depth best explained

**Species Richness** ( $R^2_{\text{marg}} = 0.34$ )

**Hill-Shannon** ( $R^2_{\text{marg}} = 0.49$ )

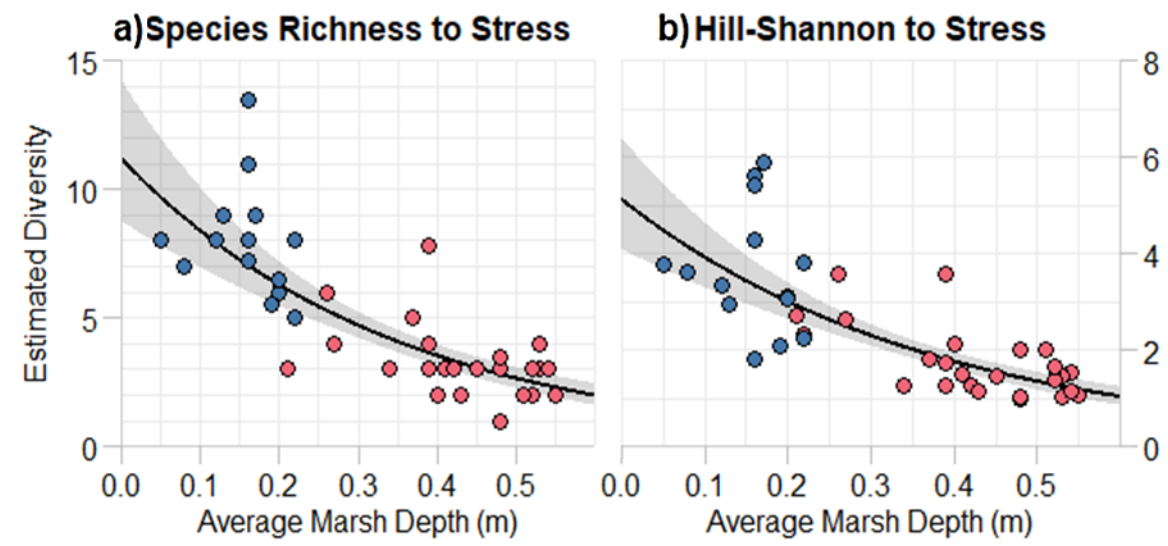


Figure 5. GLMMs of the effect of matrix-derived hydrologic stress on tree island site's estimated alpha species diversity. Species richness (a) and Hill-Shannon diversity (b). Cluster 1 is blue; Cluster 2 is pink. Gray band depicts the 95% confidence interval.



# Metacommunity

## EMS Framework

- Reciprocal Averaging ordination of RAI
- Coherence =  $\oplus$  ; Turnover =  $\ominus$  ; Clumping =  $\oplus$

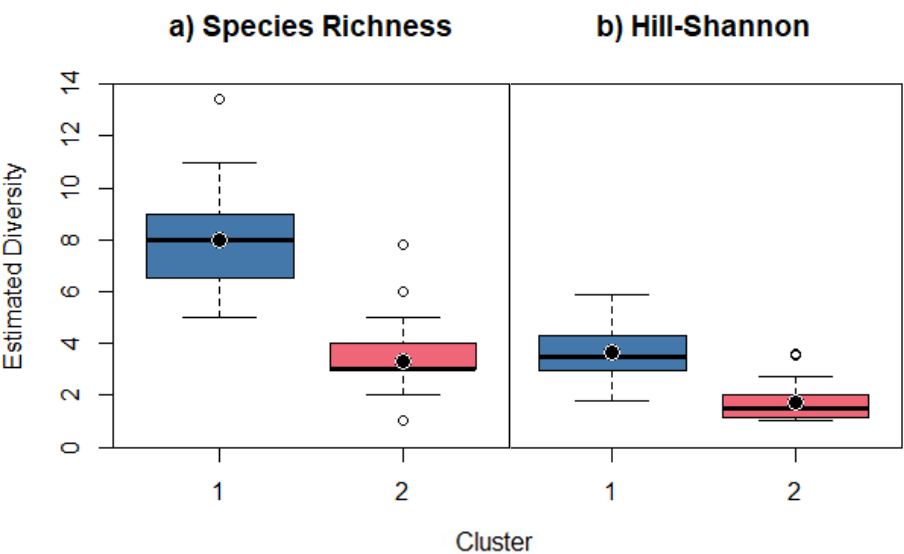
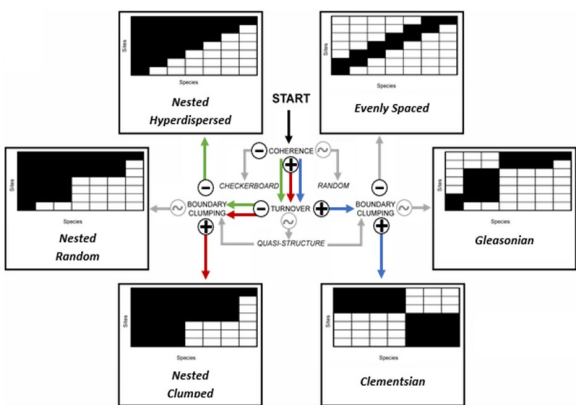


Figure 6. Boxplots of clusters' estimated alpha species diversities. Species richness (a) and Hill-Shannon diversity (b) were estimated from coverage-based asymptotes.



Modified from Eden et al. (2022)

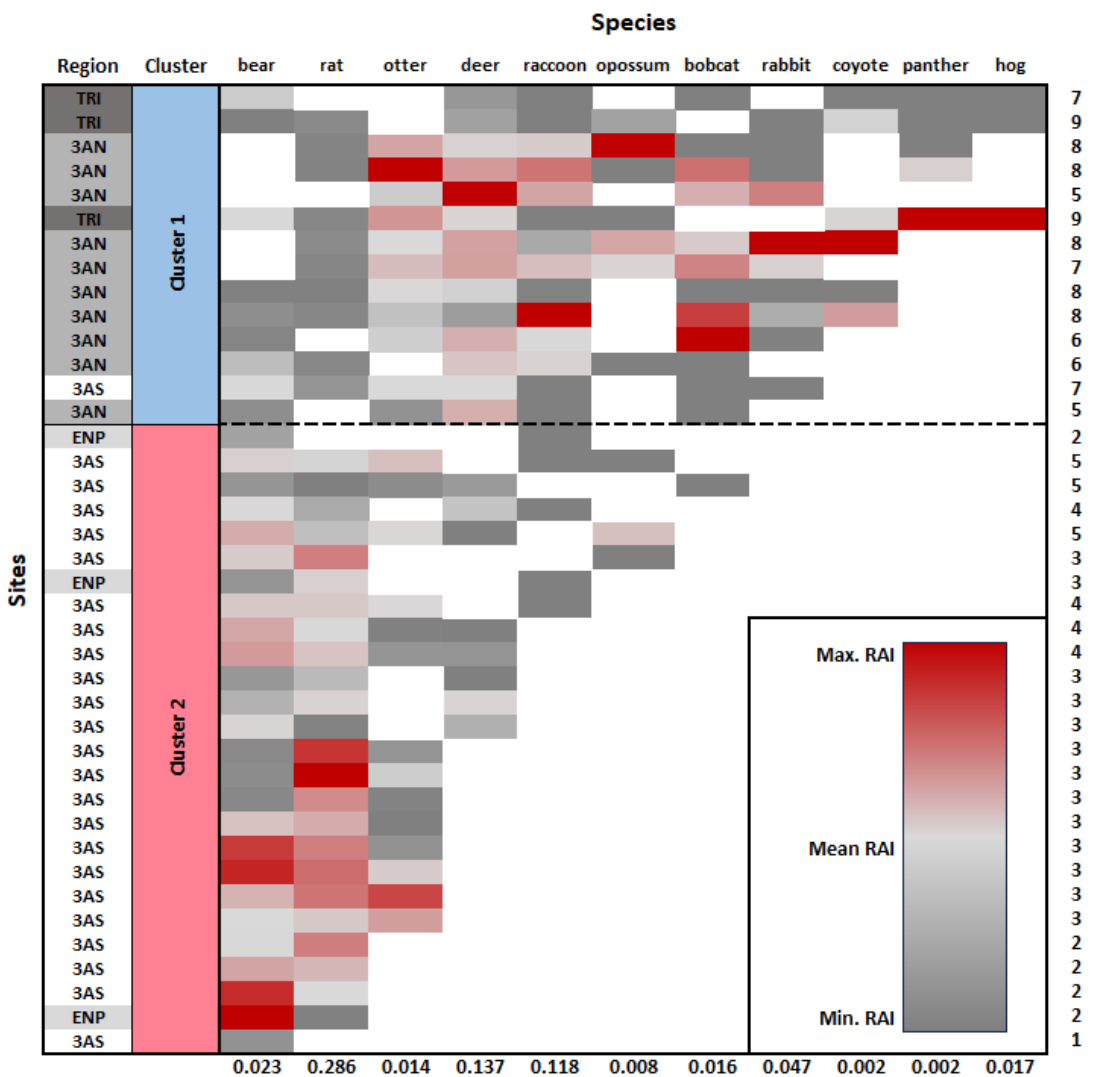


Figure 3. Ordinated sites-species matrix of all tree island sites and mammals detected during camera trapping. Sites and species' relative abundances (RAI) were organized according to their reciprocal averaging Axis-1 score. The right y-axis displays row counts (i.e., number of species at each site) and the bottom x-axis displays column means (i.e., the mean RAI for each species across).



# Metacommunity

## ■ Distance-Based Redundancy Analysis

(1) ordinate BC dissimilarity (2) Multiple regression

- Species in Euclidean space (Angle = Correlation)  
(spp. ~ spp.) (var. ~ var.) (spp. ~ var.)

- Parsimonious dbRDA reduced to 4 variables
  - Explained 44% total variation

❖ DEEP = (high-water depth) = 23% of var. (x6)

❖ MRSH = (marsh amplitude)

❖ AREA = (log area of head)

❖ NBR% = (area of neighbors in 1000 m)

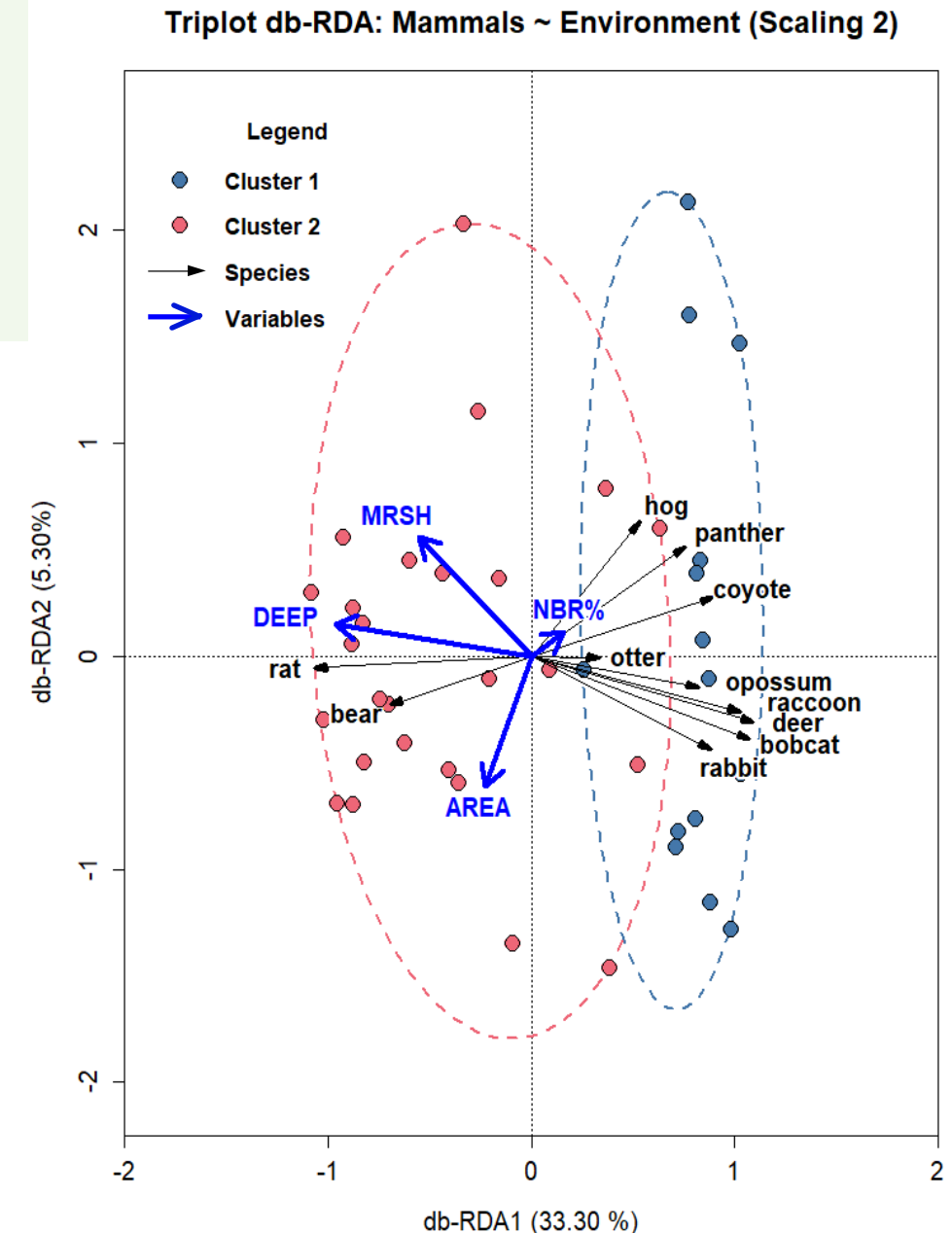


Figure 7. Distance-based redundancy analysis correlation triplot of constraining variables, Hellinger transformed species, and sites scores fitted as orthogonal linear combinations of constraining variables (*i.e.*, linear combination (lc) scores).

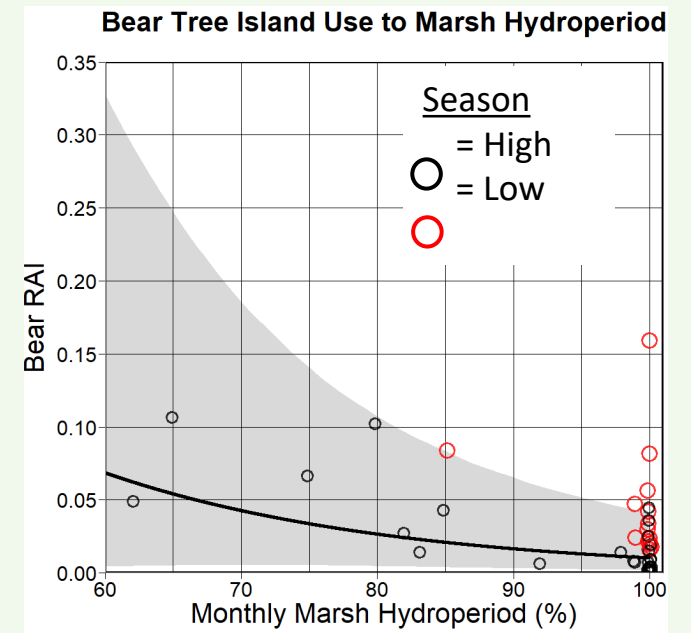
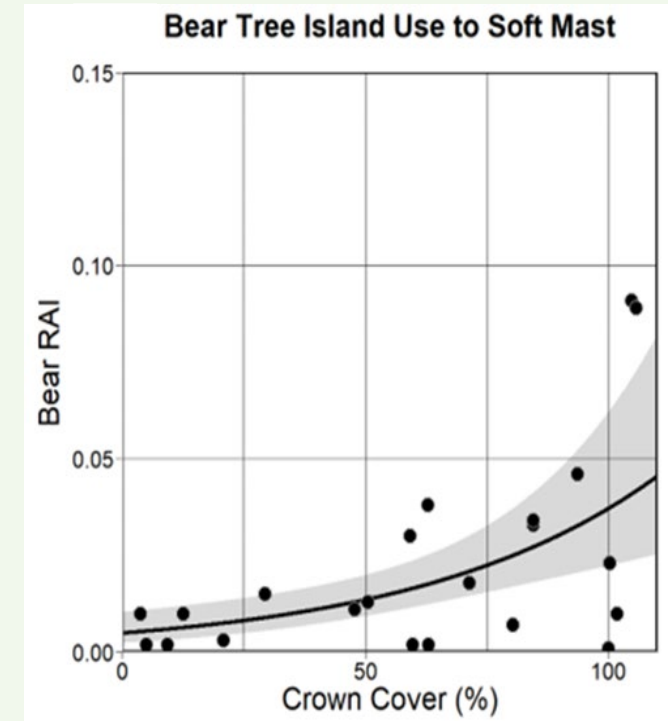


## Distance-Based Redundancy Analysis

- Multiple regions & species omits temporal & blurs species-specific trends

### Ex. Bears in WCA 3A

- Bears on all 24 TIs
  - RAI +10x higher on most used than least used
  - Availability of fruit mast is vital
    - Pond apple, cocoplum, strangler fig
  - **+10%** hydroperiod → **-50%** likelihood of use





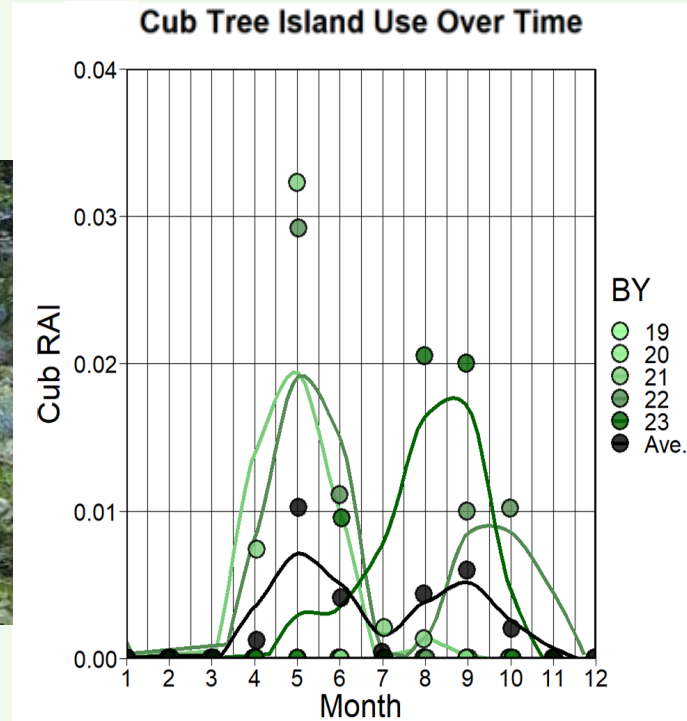
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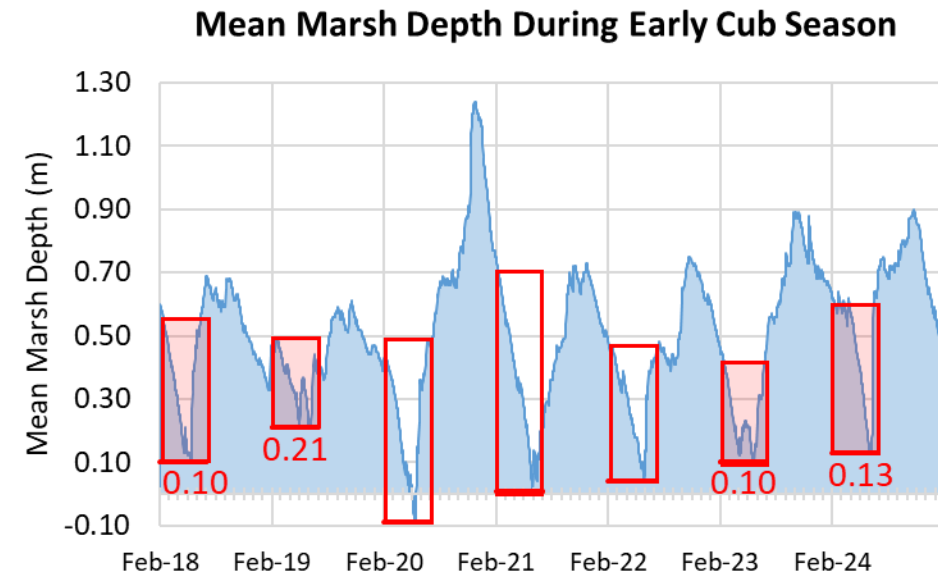
- RAI +10x higher on most used TI
- Availability of fruit mast is vital
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- **Cubs** on 9 TIs

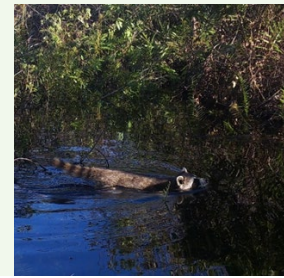
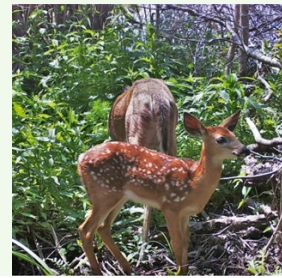
- 8x more likely on TI with hammock
- Born in February, emerge April, disperse June
- Need drydown (<4" = 0.1m) for recruitment
  - Very sensitive to pre- & post-denning conditions

(Elowe & Dodge, 1989; Garrison, 2004)



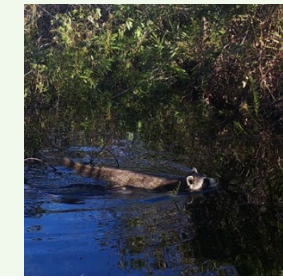
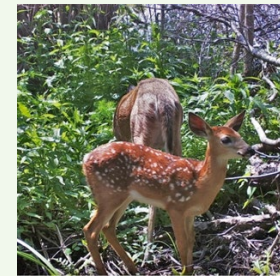


# Implications



- **Depth** (wet-season) drive diversity, distributions, habitat use, abundance
  - Patch size and connectivity important but secondary to hydrology
- Multicollinear = islands with  $>0.4$  m are same with regular, prolonged flooding of hammocks
  - Flooding hammocks → loss of forage, habitat, reproductive failure, mortality
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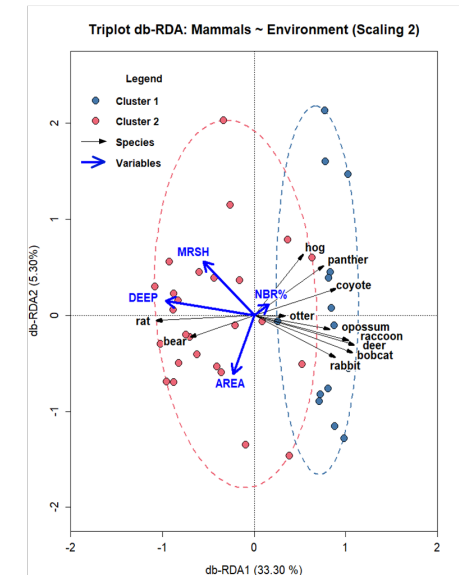
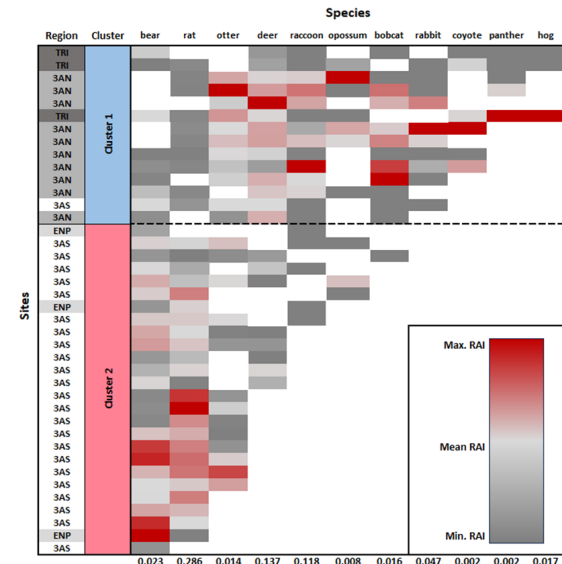
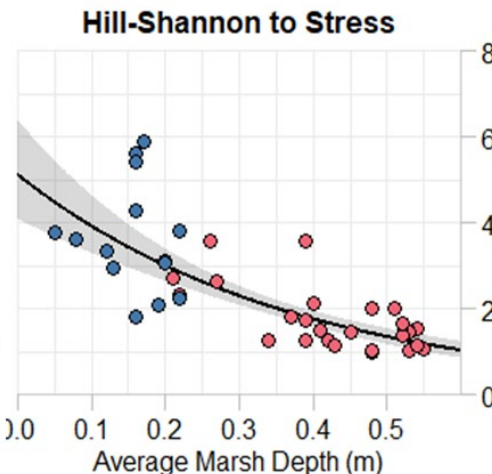
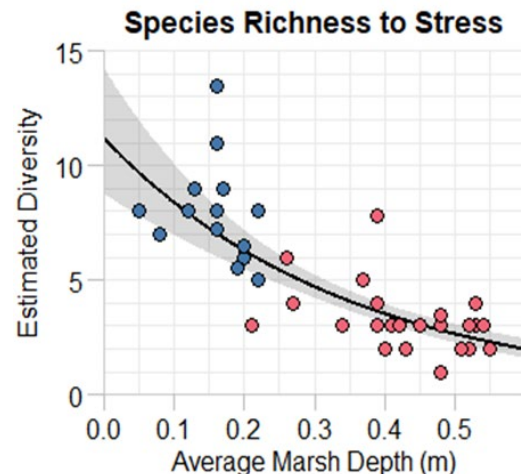
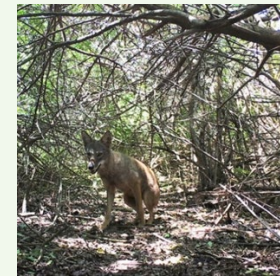
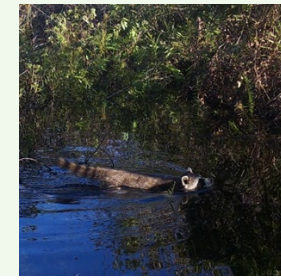
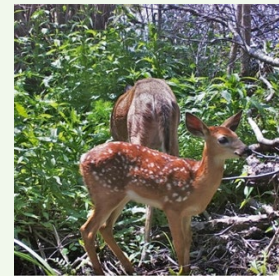


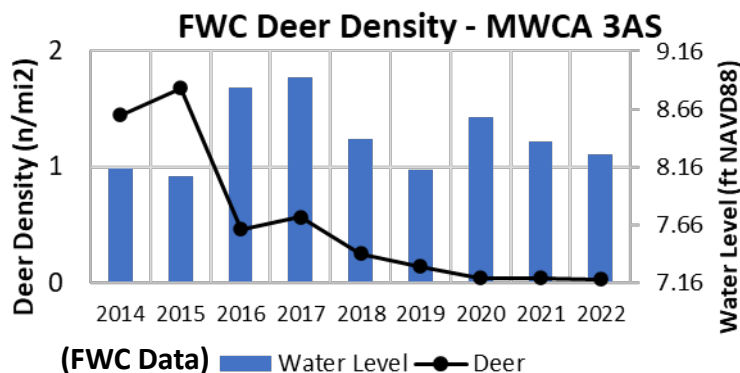
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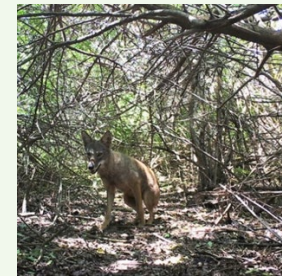
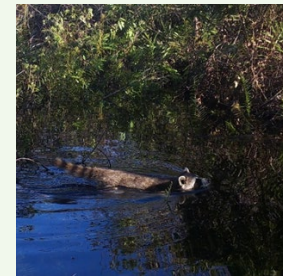
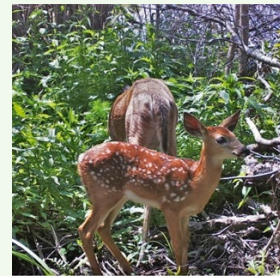
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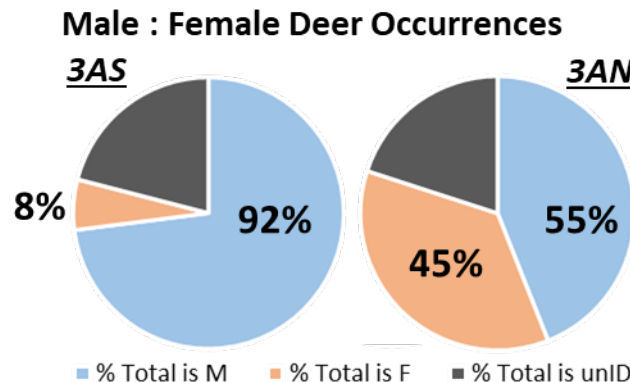
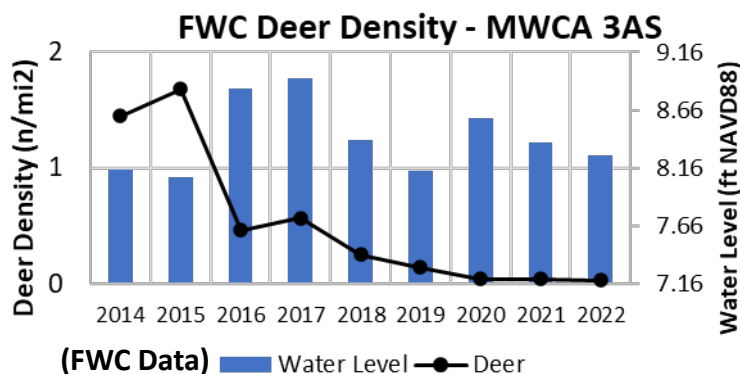
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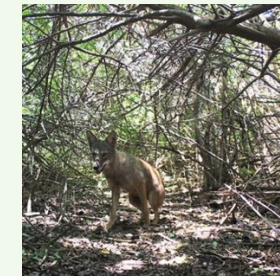
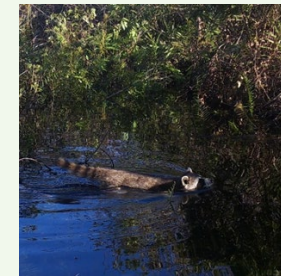
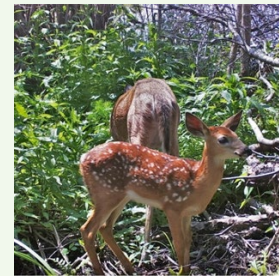


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  - c) Mass mortality ( $\sim 50\%$ ) during high-water; demographics (MacDonald-Beyers, & Labisky, 2005)  
(Cherry et al., 2019)

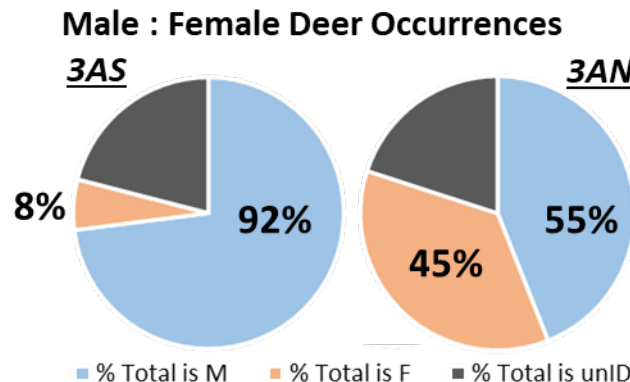
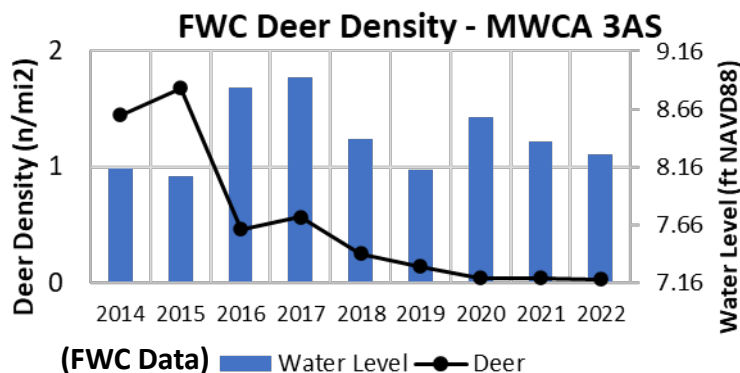




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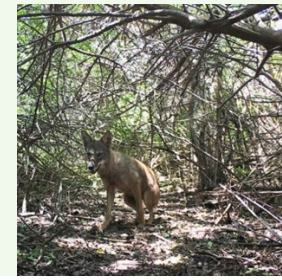
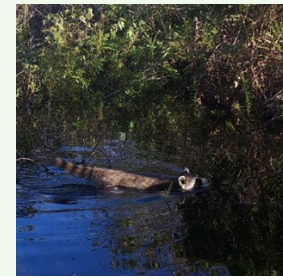
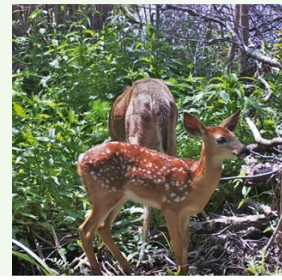


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  - Mass mortality ( $\sim 50\%$ ) during high-water; demographics (MacDonald-Beyers, & Labisky, 2005)
  - Diversity & occurrences higher where drier (Cherry et al., 2019)



RAI of 3AS to	3AN	TRI
■ Coyote	✓	✓
■ Panther	✓	✓
■ Bobcat	27x	11x
■ Deer	13x	15x
■ Opossum	2x	7x
■ Marsh rabbit	76x	2x

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  - Multiple lines of evidence suggest high-water may be leading driver, at least regionally (3AS)
    - \* Need direct species interaction data (e.g., python predation or density)
    - \* Need species-specific studies to refine understandings (e.g., raccoons sensitive to 500m scale)
- **Restore** Quantity suitable habitat (TI Area and Number)  
Quality (Forage + Hydrologic stress)



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