



Applying hierarchical distance sampling in 'unmarked' to Iowa forest birds

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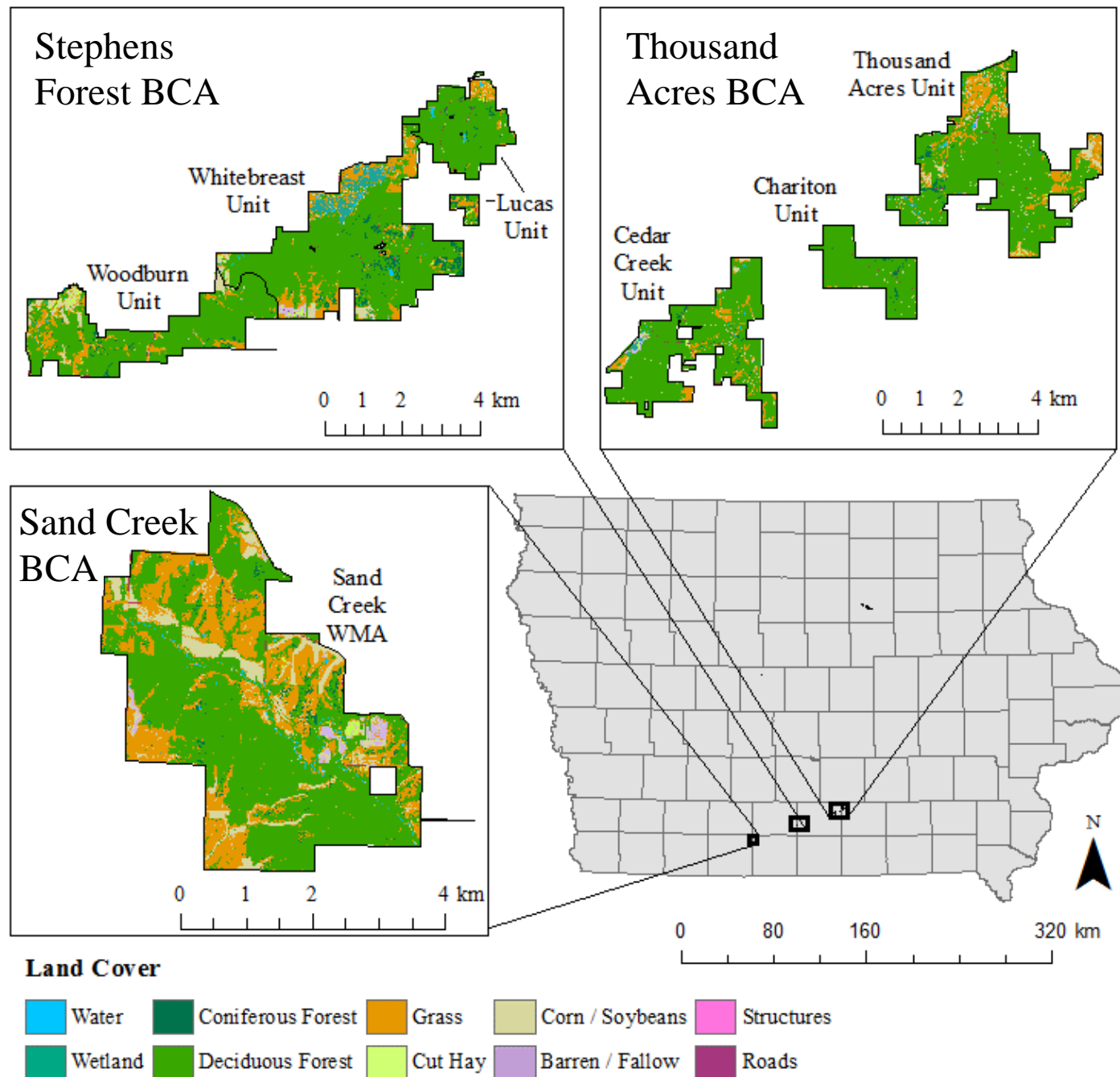
Study Objectives

Acadian Flycatcher

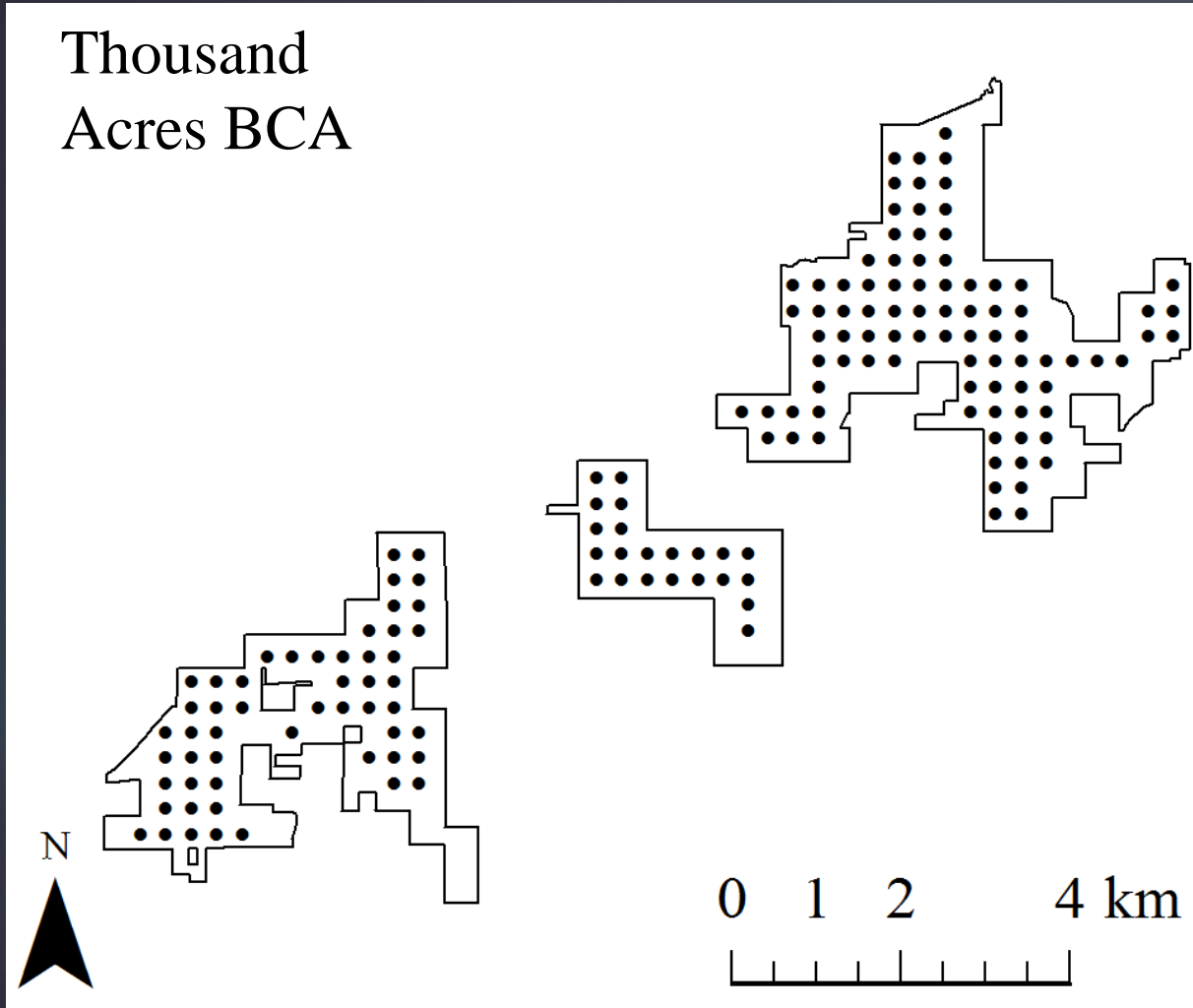


- Estimate densities of bird species of conservation concern in south-central Iowa forests
- Determine relationships between habitat characteristics and bird densities to potentially inform management

Study Area

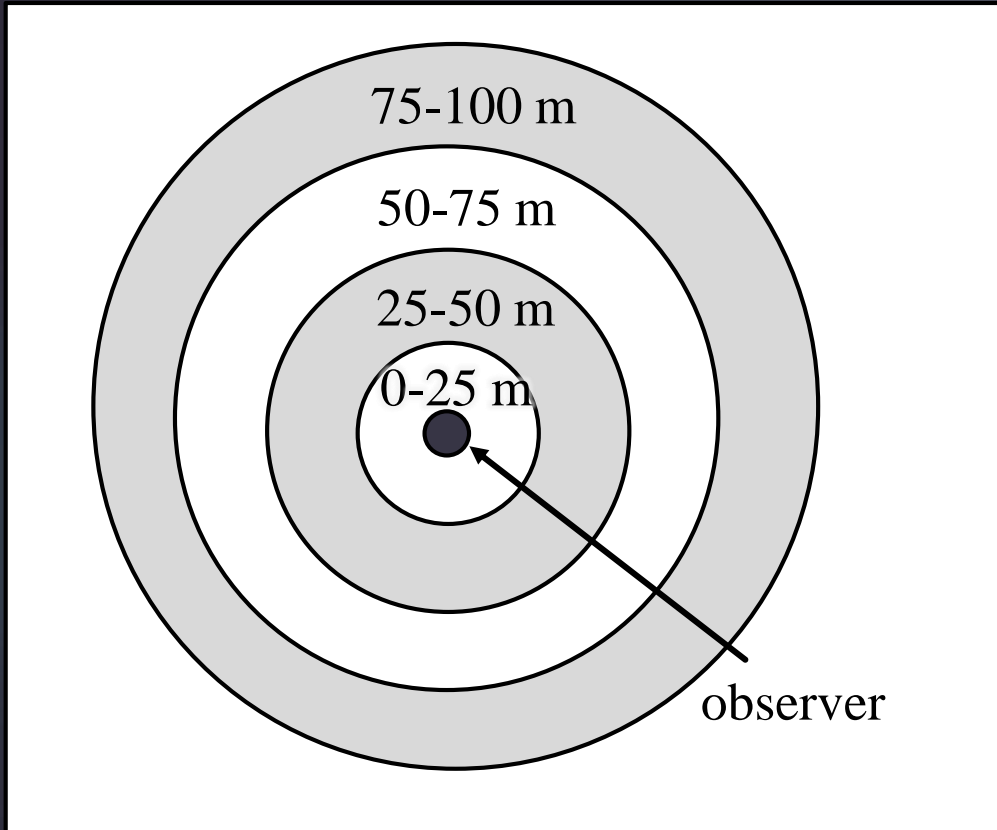


Bird Point Counts



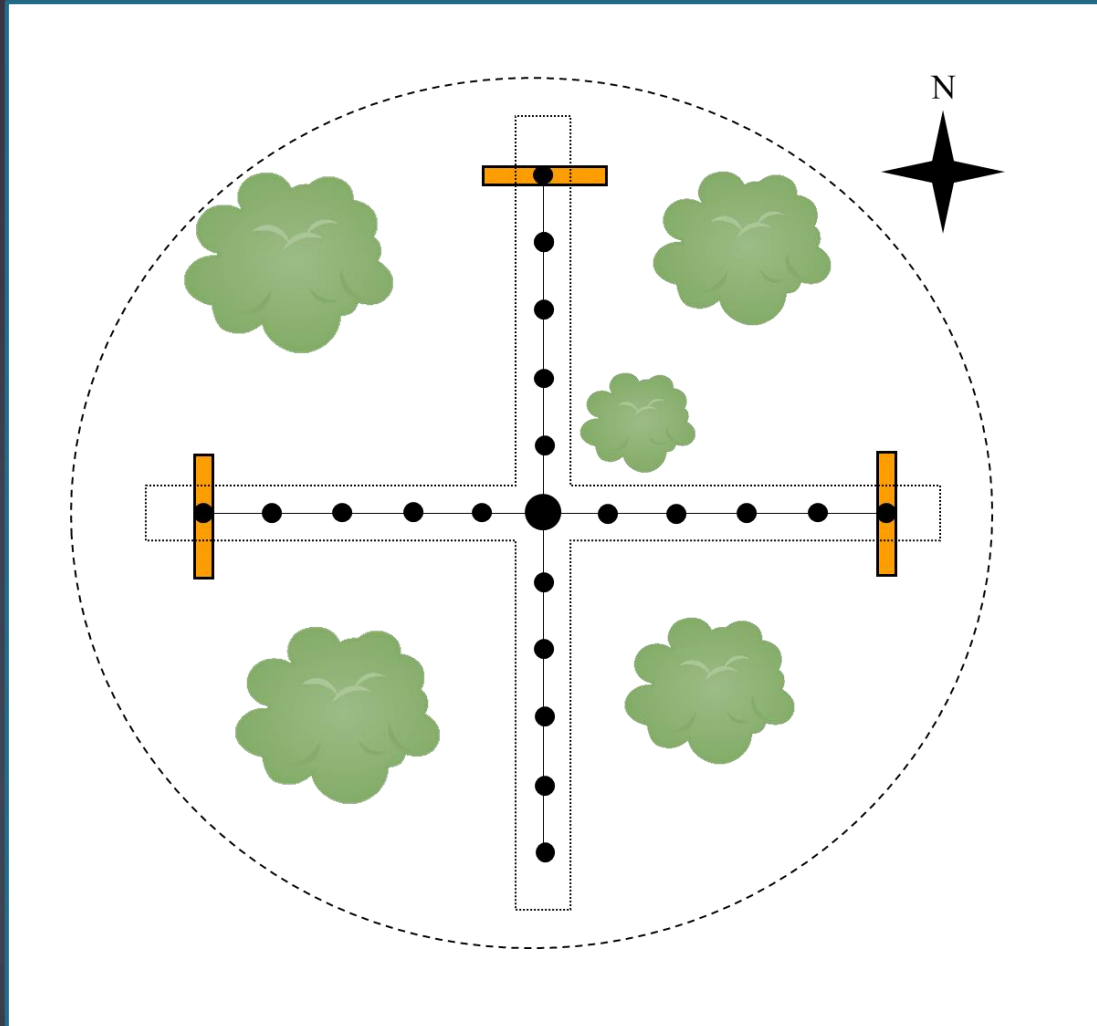
- 493 points
- 2016-2019
- Two visits/year during breeding season

Bird Point Counts



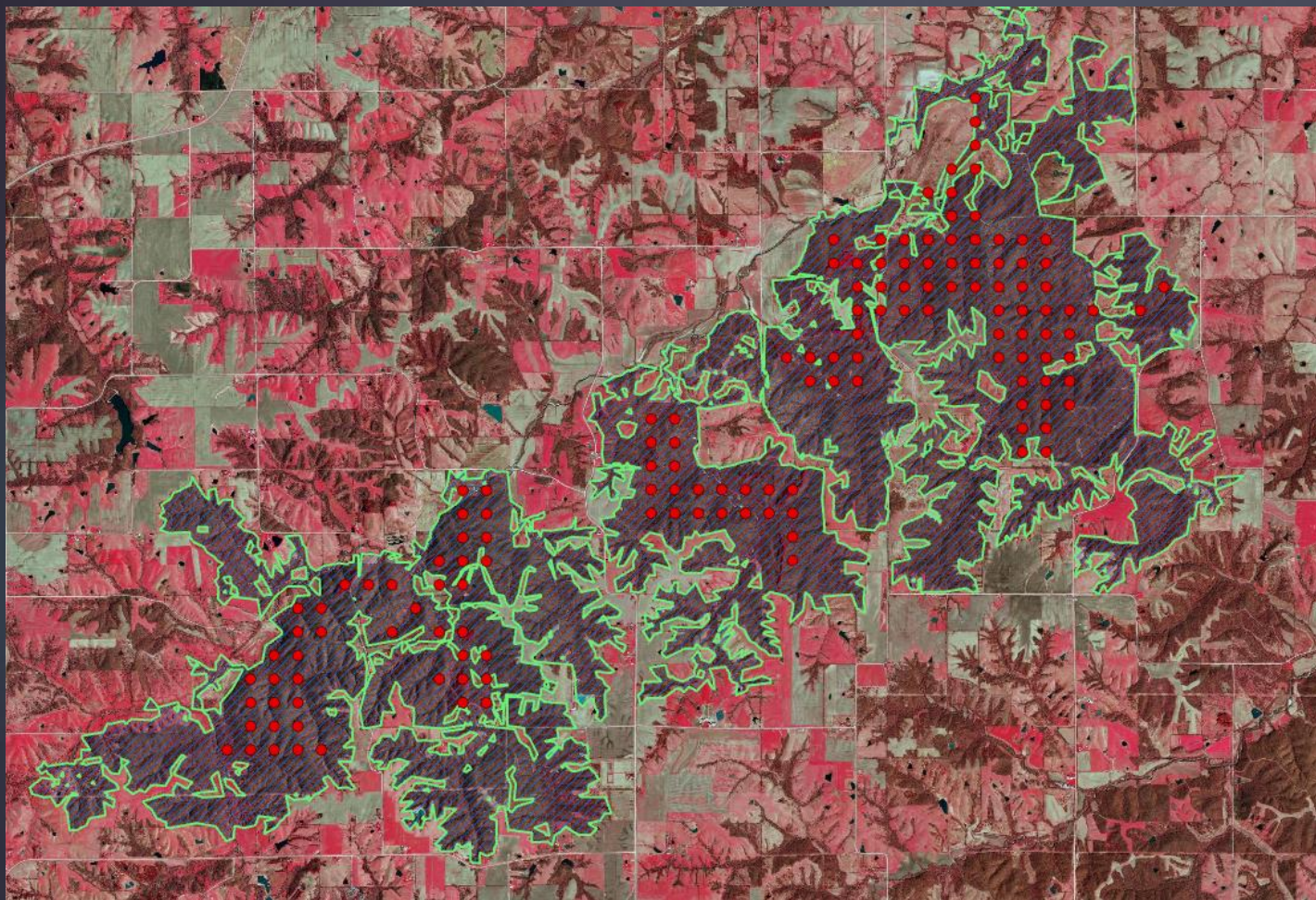
- Distance sampling
 - 4 bins
 - 100 m truncation
 - 10 minutes
 - No visual only or known female observations

Vegetation Surveys



- 493 points (same as bird surveys)
- July – August 2019
- Forestry prism sample, ground cover, shrub stems

Landscape Scale Data



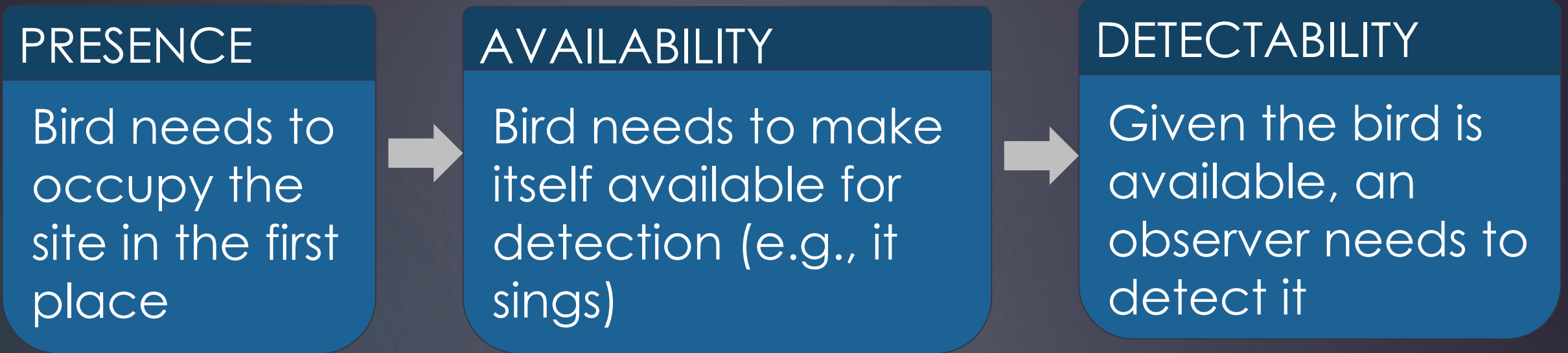
- Cropscape data and ArcGIS used to estimate % forest cover within 1 and 10 km of point
- Digitized 2016-2018 Iowa Spring Color Infrared Orthophotos to get forest patch boundaries (Cropscape as an initial template)

Data Analysis Conundrum



- It is virtually impossible to detect all of the birds at a site
- Need fancy(ish) statistics to get good estimates of densities, and even fancier statistics to related density to habitat

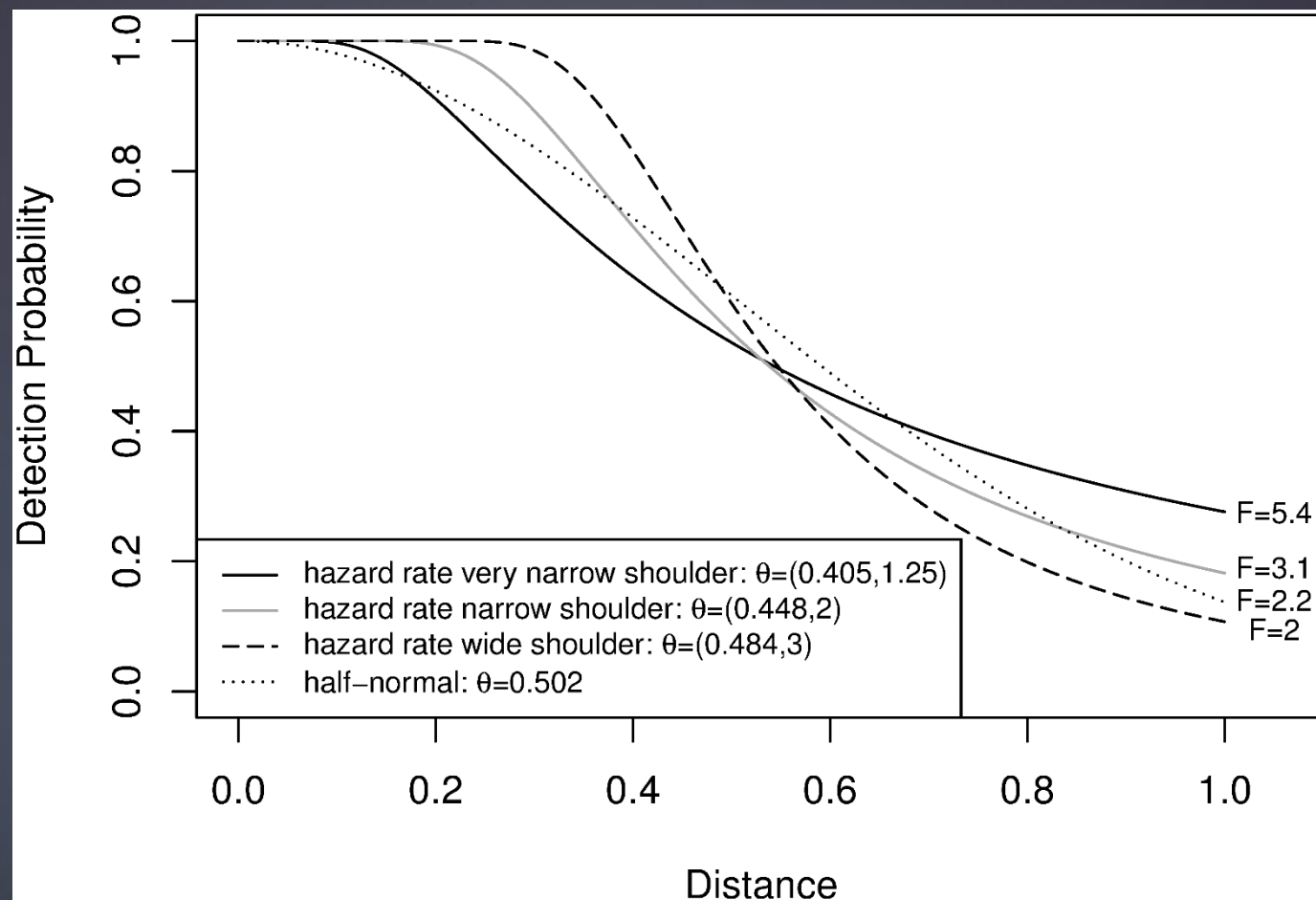
Detection Process



Modeling detectability

- Detectability can be modeled using distance sampling
 - Most commonly used for birds and marine mammals, but can be applied to many other taxa (e.g., herps)
 - Detection probability (p) declines with distance
 - Different functions can be used to describe the decline in detection probability
 - Can include covariates (e.g., observer, wind speed)
 - R packages: 'unmarked', 'detect', 'Distance'

Detection functions




Clark, R.G., 2016. Statistical efficiency in distance sampling. *PloS One*, 11(3).



Selecting detection functions and
detection covariates for distance
sampling in 'unmarked'

Modeling availability

- At least two different sampling schemes
 - Multiple visits: N-mixture models or hierarchical distance sampling incorporate this ('unmarked' can do both)
 - Removal sampling
 - Chunk a point count into multiple time bins, once you hear a bird don't count it in later time bins (thus "removing" it)
 - Implemented for point counts in R package 'detect;' can be combined with distance sampling
- Covariates can include things like time of year and time of day
- Combining multiple visits AND removal sampling with distance in a non-Bayesian framework is difficult (no good way to do this in R)
- Reference for removal sampling in 'detect': Sólymos, P., Matsuoka, S.M., Bayne, E.M., Lele, S.R., Fontaine, P., Cumming, S.G., Stralberg, D., Schmiegelow, F.K. and Song, S.J., 2013. Calibrating indices of avian density from non-standardized survey data: Making the most of a messy situation. *Methods in Ecology and Evolution*, 4(11), pp.1047-1058.



Incorporating availability
covariates into hierarchical
distance sampling in 'unmarked'

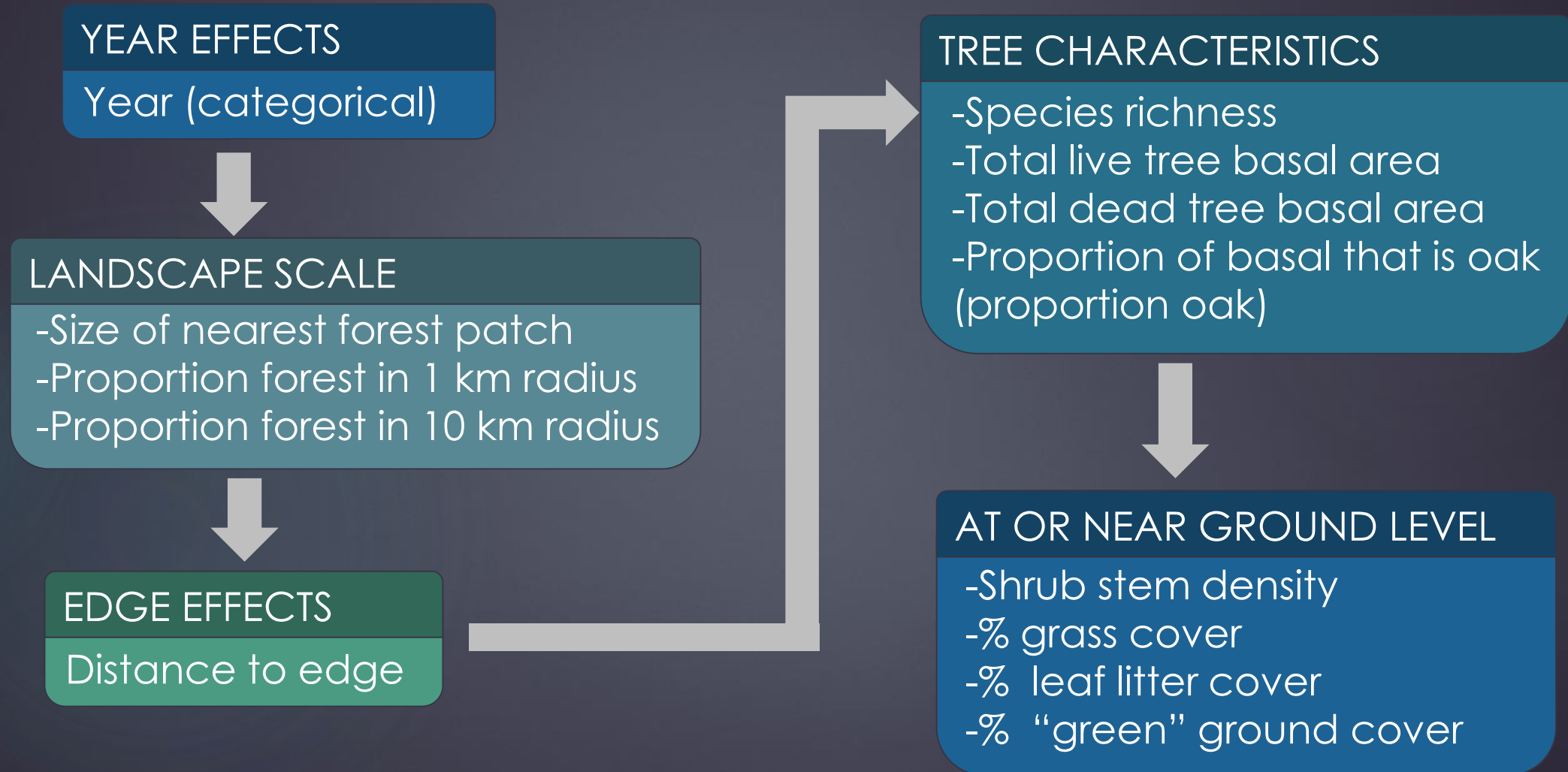
Modeling abundance (or density)

- I will focus on hierarchical distance sampling here
- Given $p = P(\text{detectable})$, $\phi = P(\text{available})$, a = area sampled, and d = number detected, overall abundance in its simplest form can be calculated as:

$$d/(\phi * p) * a$$

- With hierarchical distance sampling, abundance as a point scale is treated as a random variable and can accommodate covariates
- Reference for hierarchical distance sampling: Chandler, R.B., Royle, J.A. and King, D.L., 2011. Inference about density and temporary emigration in unmarked populations. *Ecology*, 92(7), pp.1429-1435.

Abundance covariate hierarchy for my study



Acadian Flycatcher Predictions



- ▶ Forest cover at 1 km and 10 km (+)
- ▶ Patch size (+)
- ▶ Distance to edge (+)
- ▶ Live tree basal area (+)

Incorporating abundance covariates into hierarchical distance sampling in 'unmarked'

- Covariate key (all numeric covariates center scaled)
 - year = survey year (factor)
 - nearest_patch_size = area of closest forest patch
 - fprop1km and fprop10km = proportion of land area covered by forest in 1 km and 10 km, respectively
 - dist_edge = distance to forest edge (before center-scaling, 0 if not in forest)
 - spp_rich = tree species richness in 1m factor prism sample
 - live_basal = basal area of all live trees in 1m factor prism sample
 - dead_basal = basal area of all dead trees in 1m factor prism sample
 - oak_prop = proportion of basal area comprised by oaks (0 if no trees)
 - grass_prop = proportion of ground covered by graminoids (e.g., grass)
 - litter_prop = proportion of ground covered by leaf litter
 - green_prop = proportion of ground covered by herbs or green leaves of small woody plants
 - shrub_dens = density of shrub stems

Acadian Flycatcher Results

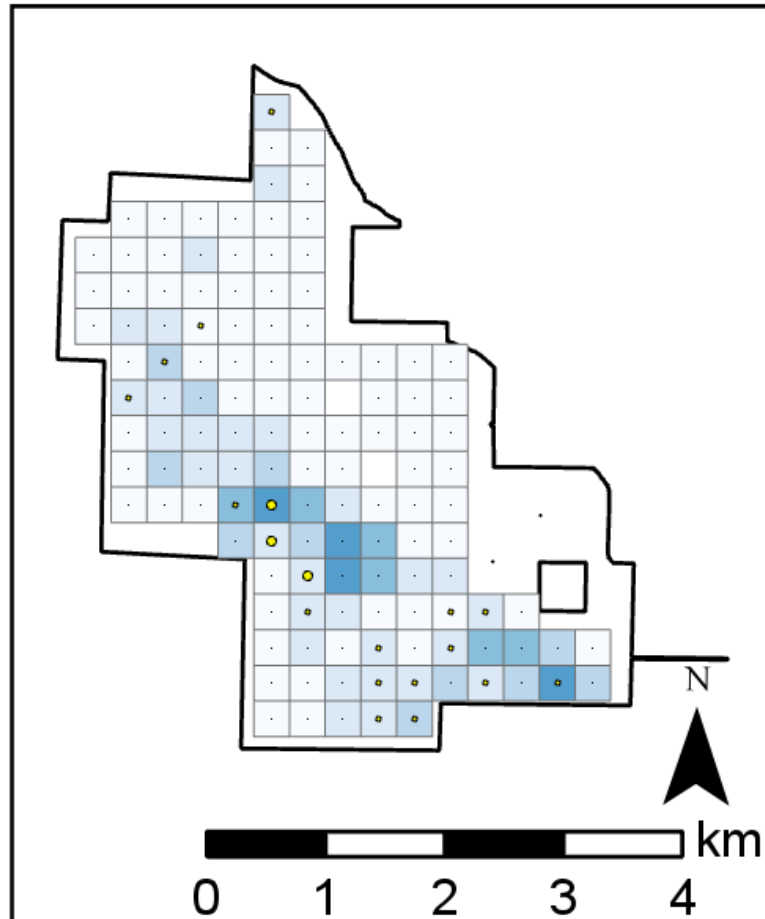
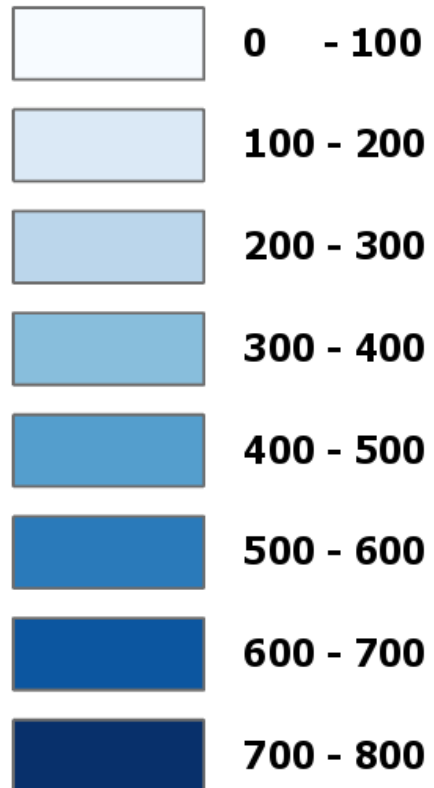


- ▶ Forest cover at 10 km (+)
- ▶ Patch size (-)
- ▶ Distance to edge (+)
- ▶ Live tree basal area, spp. richness, proportion oak(+)
- ▶ Shrub density and grass cover (-)
- ▶ Litter cover (+)

Displaying results in ArcGIS

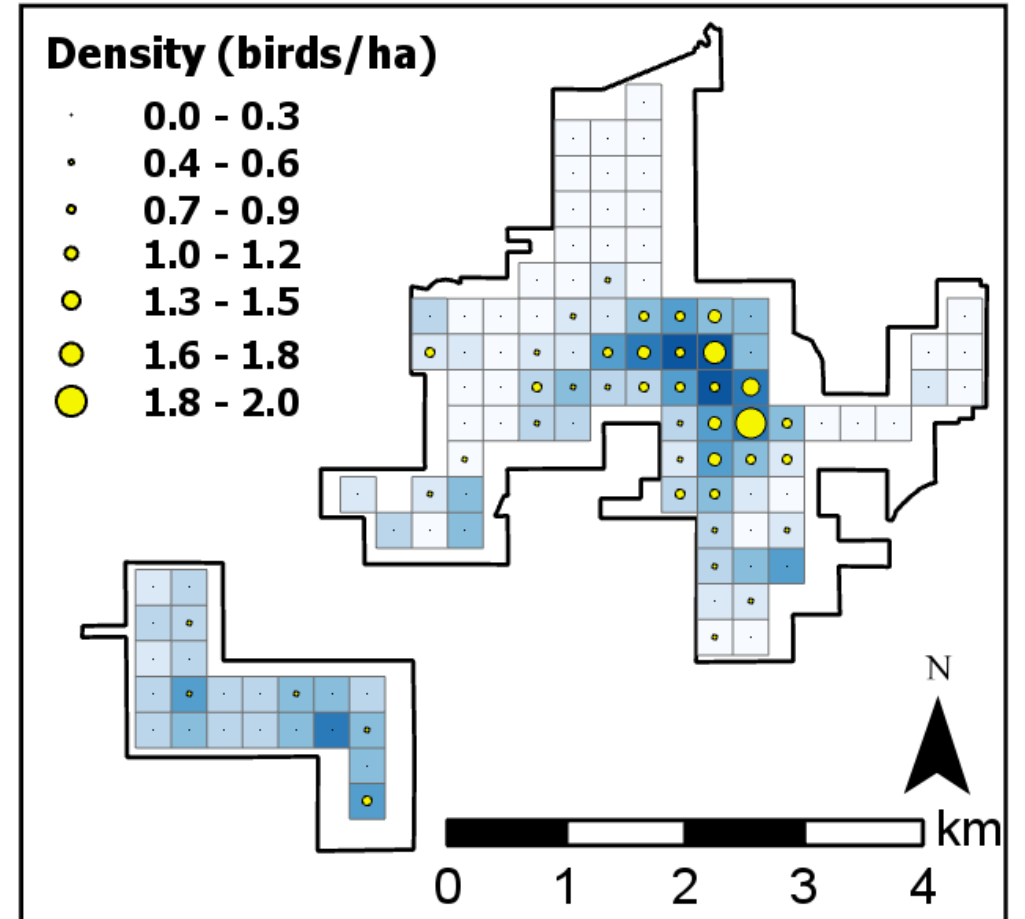
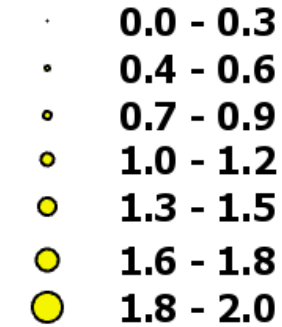
Sand Creek WMA

Distance to Edge (m)



**Stephens SF -
Chariton and Thousand Acres Units**

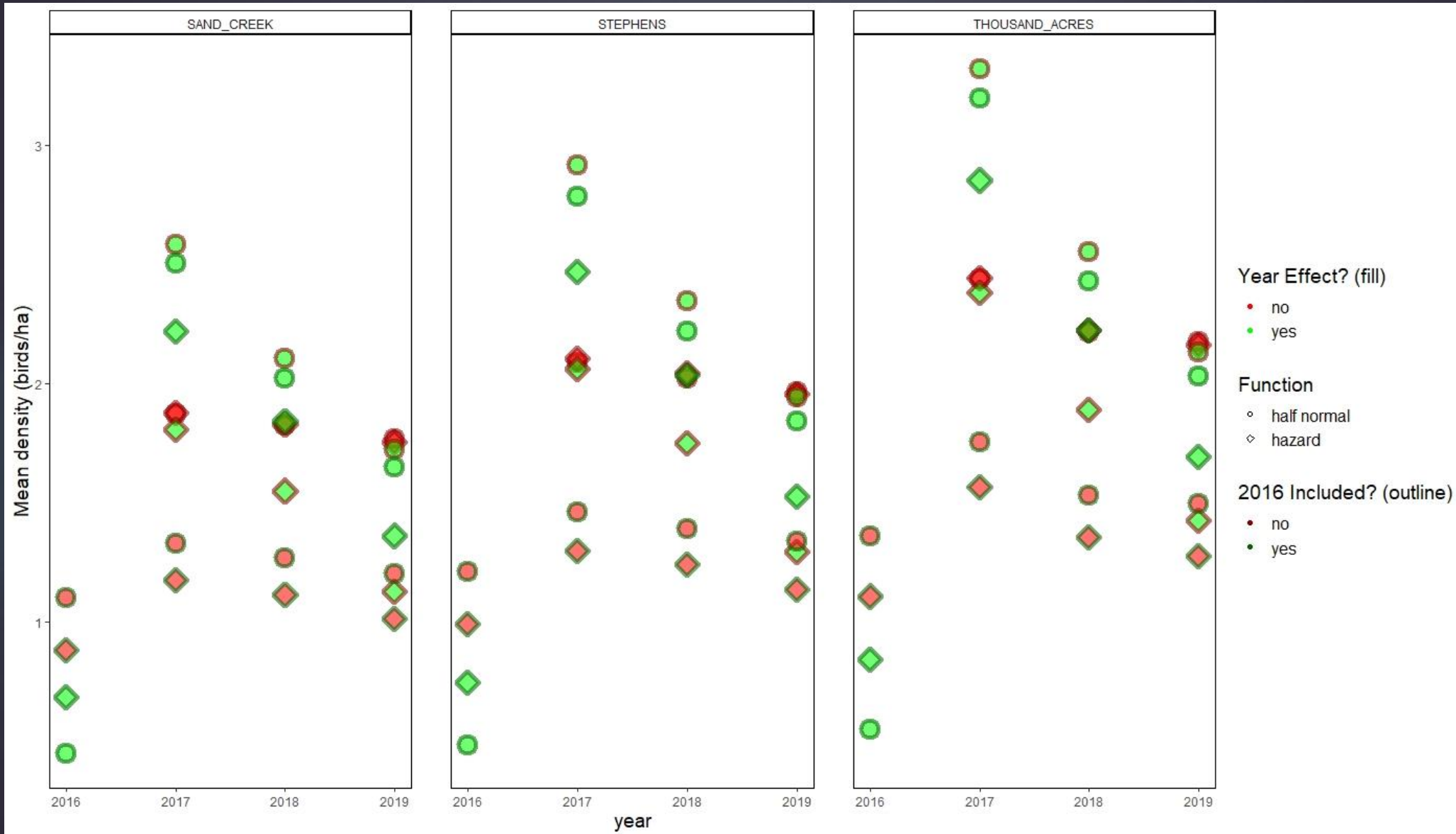
Density (birds/ha)



Issues

- Four years worth of data; 2016 had two observers that didn't observe during any other year; makes models with observer and year difficult
 - Hazard function: singular Hessian (no variance/covariance matrix, so no precision estimate)
 - Half normal: doesn't always fit as well as hazard, and sometimes I don't get estimates for one of my observers
- Some species are too loud for distance sampling (detectability doesn't decline with distance); going to try N-mixture and see how that goes
- Getting precision on density estimates for year-area combinations (i.e., subset points in a single year)
 - Data aren't quite distributed Poisson (lower variance at a point scale)
 - Might try bootstrapping

Looking at confounded observer-year effects



Future Directions

- Adding in a midstory foliage density component to my model hierarchy
- Using point-scale density estimates from a variety of species to 1st) estimate diversity at a point-scale 2nd) use those diversity estimates to determine relationships between diversity and habitat

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