Habitat relationships of forest and early successional bird communities in Iowa

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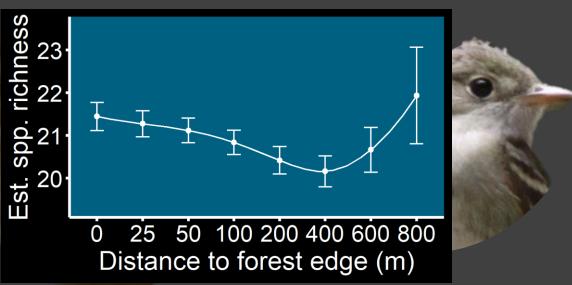
Introduction

With recent declines in avian populations, it is important to identify areas of high avian diversity and to determine habitat characteristics associated with that diversity. Our study focused on breeding bird communities three Bird **Conservation Areas in** south-central lowa; these areas are primarily forested but also contain successional and agricultural habitats. Our primary goals were to 1) compare species richness estimates which do and do not consider imperfect detection probability, and 2) determine relationships between species diversity and habitat metrics.

»We detected
77 breeding bird
species meeting several
criteria needed for
analysis, 24 were listed
as lowa Species of
Greatest Conservation
Need.

» Point-scale
species richness
estimates
accounting
for imperfect detection
(range = 16.7 to 25.7)
were, on average,
7.7 species higher
than species richness
calculated from raw
data.

» Bird species richness was related to habitat at multiple spatial scales; related metrics included landscape-scale forest cover at a 10 km scale (+), tree species richness (+), and leaf litter cover (-). Both forest edges and interior forests had high species richness (see figure below).



» We recommend management practices in this area that preserve interior forest habitat, maintain or increase landscape-scale forest cover, and that any selective harvests maintain tree species richness at a site by not completely eliminating any tree species.

Methods

Distance sampling bird point surveys (493 points, 3,944 surveys) were arranged in grids with 300 m spacing. Surveys had 10 min. duration and 100 m truncation. They occurred May-Aug for four years (2016-2019) with two visits per point per year.

Vegetation surveys were conducted at the bird survey points, from Jul-Aug 2019. Landscape scale forest cover was derived from the National Cropland Data Layer, and forest patch boundaries were digitized from aerial imagery.

Hierarchical distance sampling models in R package 'unmarked' were used to calculated occupancy probability (ψ) at each point for each year. Values of ψ were summed to estimate species richness for each year at a point, and years were averaged. Estimated species richness was related to habitat metrics using linear regression.

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