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Comparison of breeding bird and vegetation communities in primary and secondary forests of Great Smoky Mountains National Park

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ABSTRACT

We compared breeding bird communities and vegetation characteristics at paired point locations in primary (undisturbed) and mature secondary forest (70–100 years old) sites in Great Smoky Mountains National Park, USA to understand how sites logged prior to creation of the park compare to undisturbed sites following 70 years of protection from human disturbance. We found that bird and vegetation communities are currently similar, but retain some differences in species composition. Rank abundance curves for primary and secondary forest bird communities showed very similar patterns of species dominance. Species composition was also similar on the two sites which shared 24 of the 25 most frequently recorded species. Nonetheless, comparisons of density estimates derived from distance sampling showed three bird species were more abundant on primary forest sites and that one bird species was significantly more abundant on secondary forest sites. Notably, comparisons based on raw counts (unadjusted for potential differences in detectability) produced somewhat different results. Analyses of vegetation samples for the paired sites also showed relative similarity, but with some differences between primary and secondary forests. Primary forest sites had more large trees (trees greater than 50 cm diameter at breast height) and late successional species. Primary forest sites had a denser tall shrub layer while secondary forest sites had a denser canopy layer. Nonetheless, tree species richness, basal area of live trees and number of standing snags did not differ between primary and secondary forest sites. Results indicate that breeding bird communities on sites within the park that were logged commercially 70 years ago are currently quite similar to bird communities on sites with no history of human disturbance. Similarities between the bird communities on previously disturbed and undisturbed sites in Great Smoky Mountains National Park may exceed those on more fragmented landscapes because large patches of primary forest, adjacent to commercially logged sites, remained in the park when it was established in 1935. These patches of primary forest may have served as source areas for commercially logged sites.

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1. Introduction

Many studies have documented differences in animal species community structure and composition on old-growth and second-growth forest sites (e.g., Haney, 1999; Jung et al., 1999; Okland, 1996; Thompson et al., 1999; Lomolino and Perault, 2000; Martikainen et al., 2000; Blake and Loiselle, 2001; Hyde and Simons, 2001). Similarly, many studies have investigated animal community responses to silvicultural practices (e.g., Morrison, 1992; Herbeck and Larsen, 1999; Sallabanks et al., 2000). Most comparative studies have focused on differences between early second-growth and old-growth, or early and late second-growth forests (e.g., Diaz et al., 2005). Comparative studies are often constrained by a lack of replication and the difficulty of pairing small and widely scattered remnants of old forest with comparable sites that have regenerated from previous clearing. Few studies have examined animal community differences between undisturbed primary forest and mature second-growth forests (but see Herbeck and Larsen, 1999; Hyde and Simons, 2001). Such comparisons are an important step in identifying the point at which wildlife species no longer discriminate habitat differences associated with historic land use.

Animal community differences along successional gradients are generally attributed to changes in vegetation structure and composition (Thompson et al., 1999, 2003). Comparisons of old-growth and mature second-growth temperate forests have shown that forests regenerated over the last century have developed many of the structural characteristics of old-growth (Schuler and Gillespie, 2000; Ziegler, 2000). Despite measurable differences between old-growth and second-growth forests, there is no general agreement on a definition of old-growth (Leverett, 1996; White and White, 1996). Assigning an age-based definition is problematic because different forest types are dominated by species with differing life history parameters. Furthermore, researchers do not agree on whether or not forests that have undergone widespread natural disturbance, but have not been altered by human disturbance, should be defined as old-growth (Leverett, 1996; White and White, 1996). White and White (1996) argue that forests that are relatively young due to natural disturbance should be considered old-growth because they contain undisturbed soils and characteristic amounts of coarse woody debris, and because such forests are important and natural elements of old-growth forest mosaics. Moreover, Runkle (2000) showed that while undisturbed old-growth forests in the eastern United States change very slowly over time, they are always in a state of flux.

Despite these issues, most researchers agree on the general characteristics of old-growth forests, such as old trees, large trees, woody debris, high density of snags, canopy gaps, pits and mounds, identifiable growth layers, late successional species, and lack of human disturbance (Leverett, 1996). These features provide unique structural habitat characteristics that have been identified in several studies as offering optimum conditions for North American bird species such as Red-breasted Nuthatch (*Sitta canadensis*), Brown Creeper (*Certhia americana*), Winter Wren (*Troglodytes troglodytes*), and Golden-crowned Kinglet (*Regulus satrapa*) (Haney et al., 2001), as well as the Rufus Treecreeper (*Climacteris*

rufa) in western Australia (Luck, 2002), and the Eurasian Treecreeper (*Certhia familiaris*) in central Finland (Suorsa et al., 2005).

Great Smoky Mountains National Park, USA includes more than 45,000 ha of primary forest that lack evidence of direct human disturbance from timber harvest, settlement, or agriculture (Pyle, 1985; Davis, 1996). Although these forests are considered old-growth, most areas have been subject to large scale natural disturbances at some point in the past. Natural disturbances have included extensive fire, windstorms, ice and snow storms, landslides, and insect outbreaks (SAMAB, 1996; Greenberg and McNab, 1998). Furthermore, exotic pests and the loss of predators have influenced forest structure and composition. Invasions of chestnut blight (*Endothia parasitica*) and balsam woolly adelgid (*Adelges piceae*) have led to the eradication of American chestnut (*Castanea dentata*) and Fraser fir (*Abies fraseri*) as canopy trees in most forests of the southeast (Liebhold et al., 1995). The loss of large predators and subsequent increases in deer and feral hog populations have altered understory communities (SAMAB, 1996). Nonetheless, these forests retain essential qualities of primary forest (Busing, 1998) and are characterized by diverse plant and animal populations. Great Smoky Mountains National Park, like much of the southern Appalachians region, also contains forests regenerating from the extensive industrial logging that took place in the early part of the twentieth century (Pyle, 1985; SAMAB, 1996; Yarnell, 1998).

Hyde and Simons (2001) showed that salamander populations are more abundant and salamander communities are more diverse on undisturbed sites compared to mature second-growth sites in Great Smoky Mountains National Park. Furthermore, several salamander species show strong positive associations with undisturbed sites. While no similar study has been conducted on breeding birds in the southern Appalachians, Haney (1999) documented increased abundance of several bird species in old-growth forests of the Allegheny Plateau in Pennsylvania, USA compared with the surrounding landscape.

In this paper we report on breeding bird community structure and composition in primary forest (undisturbed old-growth) and mature secondary forest (70–100 years post logging) in Great Smoky Mountains National Park. We compare the two forest types to investigate whether bird communities on mature second-growth sites are indistinguishable from those on undisturbed sites, or if community differences continue to persist after 70–100 years of regeneration. Similarly, we evaluate vegetation composition and structure to identify factors potentially associated with differences in bird communities. We also discuss the importance of adjusting for differential detectability when point count surveys are used to compare breeding bird communities in different habitats. Although increasing attention has been focused on the need to use sampling and analytic methods that account for habitat or species-specific differences in detection probabilities (Buckland et al., 1993; Lancia et al., 1994; Thompson et al., 1998; Yoccoz et al., 2001; Rosenstock et al., 2002), relatively few studies (e.g., Boulinier et al., 2001) incorporate such methods, and none have demonstrated habitat specific differences in detection probabilities.

2. Methods

2.1. Study area

The southern Appalachians region of the southeastern United States is an area of nearly 15 million ha extending from northern West Virginia and Virginia southwest to northern Alabama and Georgia (SAMAB, 1996). Much of the region was cleared as a result of industrial logging in the early twentieth century (Yarnell, 1998). The area is now 70% forested as a result of reforestation over the last century. Current forest composition in the southern Appalachians reflects a history of logging, fire suppression, settlement, and invasions of chestnut blight and balsam woolly adelgid.

We sampled vegetation and bird communities in Great Smoky Mountains National Park (35°37'N, 83°53'W) which serves as the nucleus of a group of publicly owned protected areas in the southern Appalachians, including more than 2 million ha of National Forests, federally designated wilderness areas, state lands, Tennessee Valley Authority reservoirs, and National Parks. The park, located along the border between North Carolina and Tennessee (Simons et al., 2000), is an International Biosphere Reserve and World Heritage Site. It is the largest National Park, and the largest federal wilderness area in the eastern United States (Johnson et al., 2000).

When established in 1935, over three quarters of Great Smoky Mountains National Park had been cleared by humans for timber harvest and settlement (Pyle, 1985) (Fig. 1). Highly mechanized industrial logging from 1900 to 1930 created extensive clear cuts that resulted in significant soil erosion (Pyle, 1985). The 23% of the park that has no history of logging or clearing by humans represents the largest contiguous tract of primary forest in the eastern US (Davis, 1996). Logged sites have regenerated since the park was established and today more than 95% of the approximately 205,000 ha park is forested.

Great Smoky Mountains National Park is characterized by wide elevational gradients (575–1830 m) and complex topography which support a rich diversity of habitat and vegetation types. More than half the park is deciduous mesic hardwoods, with spruce-fir and northern hardwoods found at the highest elevations, and xeric deciduous hardwoods, evergreen pines, and areas of mixed hardwoods and pines found at lower elevations.

2.2. Bird surveys

We conducted 7535 variable circular plot point transects (Reynolds et al., 1980) at 4157 point locations from mid-May to the end of June during 1996–1999 for a park-wide inventory of breeding birds (Shriner, 2001; Shriner et al., 2002). We established points approximately 250 m apart throughout the park. Most (>95%) points were established on low-use hiking trails, although some were located on low-traffic roads or off-trail transects. We recorded a location for each point using a differentially corrected global positioning system (GPS) (GeoExplorer II; Trimble Navigation 1996). Surveys followed protocols recommended by Ralph et al. (1995), recording all birds seen or heard during 10 minute counts conducted between sunrise and 10:15 AM in favorable weather. We recorded the horizontal distance from the observer to each bird detected. We used a laser range finder (Bushnell, Yardage Pro 400™) to aid in distance estimation by sighting a 50-m radius circle prior to each survey. At each point we also recorded the time of day and scored levels of background noise (generally a function of stream noise) using a 5-category index.

2.3. Vegetation surveys

We sampled vegetation in 10 m radius plots at each of the 4157 bird point count locations. For bird points along trails or roads, we shifted vegetation plots 10 m away from point

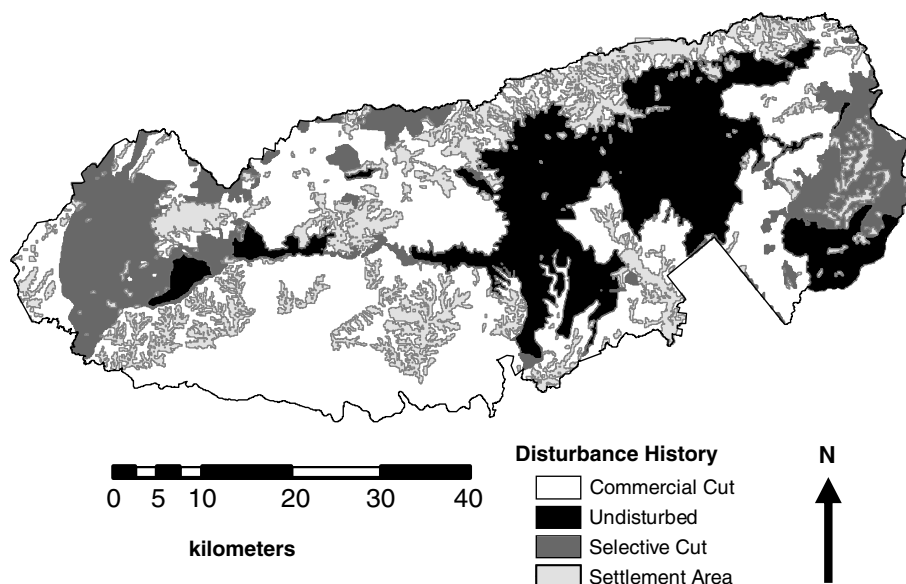


Fig. 1 – Disturbance history of Great Smoky Mountains National Park (Pyle, 1985). Comparisons of primary and secondary forest communities were restricted to survey locations in either undisturbed primary forest (black) or secondary forests that have re-grown following commercial logging (white).

centers, tossing a coin to determine which side of the road or trail to sample; off-trail points were sampled at bird point count centers. We visually estimated percent cover (defined as percent foliage blocking sunlight to layers below) for five vegetation layers (tree canopy, subcanopy, tall shrub, low shrub/seedling, and herbaceous) and assigned each layer to one of the following percent cover classes: <0.1, 0.1–1, 1–2, 2–5, 5–10, 10–25, 25–50, 50–75, 75–95, >95. We estimated height ranges for each layer, using laser range finders to determine the height of the tallest tree in canopy and subcanopy layers. We determined species composition of canopy, subcanopy, and tall shrub layers by identifying species within each layer, and visually estimated species-specific percent contribution to the layer. We also used a wedge prism (basal area factor 20, Avery and Burkhardt, 1983) at the center of each vegetation plot to sample trees. Wedge prism samples are plotless so we included all individuals detected in the wedge prism regardless of distance. We identified all trees observed in the wedge prism sample, and visually assigned each tree to one of six dbh (diameter at breast height) classes: 0–10, 11–25, 26–50, 51–75, 76–100 and >100 cm.

2.4. Paired point selection

We limited our analyses to survey points located in either undisturbed primary forest or commercially logged secondary forest (Fig. 1). In an effort to eliminate possible effects of site covariates on community comparisons, we paired primary and secondary forest survey points according to the following selection criteria. We first identified a pool of secondary forest points that matched primary forest points based on vegetation community type, census date (within 14 days), time (within 2 h), and background noise (within 1 noise level). From the pool of points that met these criteria we paired primary and secondary forest points by minimizing elevational differences. Although we allowed a maximum difference of 150 m, the paired points had an average elevation difference less than 30 m. We limited pair selection to vegetation types that were represented by at least 30 pairs of points (cove hardwood, mixed mesic hardwood, and northern hardwood), identifying 247 sample pairs that met these selection criteria.

2.5. Bird communities

We compared bird communities in primary and secondary forests using measures of relative abundance and density. We tested for differences using two-tailed paired t-tests (Zar, 1999). We compared the number of detections per point for individual species from primary forest locations with those from secondary forest locations. We also plotted rank abundance curves (Magurran, 1988) for the two forest samples using the percent of the total sample for each species as its index of abundance.

Because we were concerned about differential detectability of birds in primary forest sites compared to secondary forest sites, we estimated effective detection radii (EDR) for each bird species in each habitat type using Program DISTANCE (Thomas et al., 1998). Program DISTANCE is appropriate for use with distance sampling data in which observations are accompanied by an estimate of distance from the observer.

The EDR is approximately the distance at which the probability of detection declines to one half, depending on the function used to model detectability as a function of distance (Laake et al., 1993). We compared EDR estimates for each species to identify differences in detectability between primary and secondary forest sites. Program DISTANCE uses a maximum likelihood based function of detection and distance to estimate a sample density. We compared density estimates using z-tests, and we limited our analyses to species for which we had a minimum of 75 detections on both primary and secondary forest sites.

2.6. Vegetation communities

We compared per plot coverage estimates for three vegetation layers (canopy, subcanopy, and tall shrub) on primary and secondary sites using paired t-tests (233 paired plots, a subset of the 247 paired bird points for which we had complete data). We used the midpoint of the field estimated cover class at each point as our cover estimate. We evaluated species-specific differences between the two communities (for canopy, subcanopy, and tall shrub layers) by comparing per plot importance values for individual species using paired t-tests. We calculated importance values by multiplying the percent contribution of each species by the midpoint of the cover class. We assessed tree size distribution with wedge prism data for 191 paired plots by summing the number of stems counted in each dbh class for the primary and secondary forest samples. We also tested for differences in the number of snags by comparing the per plot number of standing snags counted in the wedge prism samples for primary and secondary forest plots.

3. Results

3.1. Bird communities

We detected 4757 individual birds of 68 species on the 247 paired point transects (494 total points). We observed 2406 individuals of 56 species on primary forest sites and 2351 individuals of 66 species on secondary forest sites (Appendix). Twelve species were limited to secondary forest sites, two species were detected only on primary forest sites, and 54 species were observed on both forest types. Rank-abundance plots for primary and secondary forest sites indicate that the bird community structure was very similar at the two sites (Fig. 2).

Paired comparisons of unadjusted counts showed Winter Wren and Black-throated Blue Warbler (*Dendroica caerulescens*) were recorded significantly more frequently on primary forest sites compared to secondary forest sites. Black-throated Green Warbler (*Dendroica virens*) was recorded significantly more frequently on secondary forest sites (Fig. 3). Effective detection radii (EDR) varied between primary and secondary forest habitats for individual bird species with significant differences for three species (Fig. 3). Dark-eyed Junco (*Junco hyemalis*), Winter Wren, and Blue-headed Vireo (*Vireo solitarius*) had significantly lower EDR on primary forest sites than on secondary forest. In nine of the 10 species with a minimum of 75 detections in both primary and secondary forests,

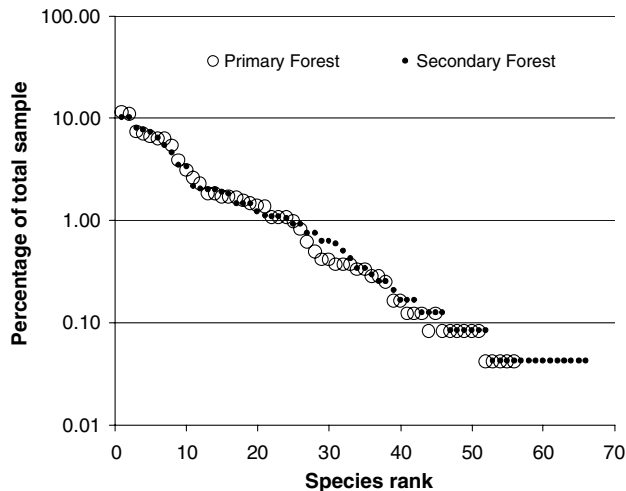


Fig. 2 – Rank abundance plot for unadjusted counts of breeding bird species detected in 247 pairs of point counts. For each pair of points one count was conducted in primary forests (open circles) and one count was conducted in secondary forests (solid circles).

estimated EDRs were lower in primary forests than in secondary forests.

Species specific comparisons using density estimates derived from Program DISTANCE showed three bird species (Winter Wren, Black-throated Blue Warbler, and Dark-eyed Junco) were significantly more abundant in primary forests compared with secondary forests and one bird species (Scarlet Tanager, *Piranga olivacea*) was significantly more abundant in secondary forests compared with primary forests (Fig. 3).

3.2. Vegetation communities

Vegetation structure, as measured by the number of trees in different size classes, showed differences between primary and secondary forests (Fig. 4a). While trees with a diameter at breast height (dbh) ≤ 25 cm were similar for the two forest types, medium trees (26–50 cm dbh) were significantly more abundant on secondary forest sites and large trees (>50 cm dbh) were significantly more abundant on primary forest sites.

Comparisons of per plot cover in the two communities revealed significantly higher canopy cover in secondary forests ($p < 0.005$), no difference in the subcanopy layer ($p > 0.05$), and significantly higher tall shrub cover in primary forests ($p < 0.005$). Standing snag densities were not significantly different between the two forest sites (155 standing snags on primary forest sites and 153 standing snags on secondary forest sites). However, the number of standing snags >50 cm dbh was significantly greater in the primary forest sample compared to the secondary forest sample ($p = 0.005$).

Tree species composition exhibited some significant differences between sites (Fig. 4(b)). We observed 50 tree species in the canopy on 233 paired vegetation plots. Ten tree species were limited to primary forest sites, 10 species were limited to secondary forest sites, and 30 species were common to

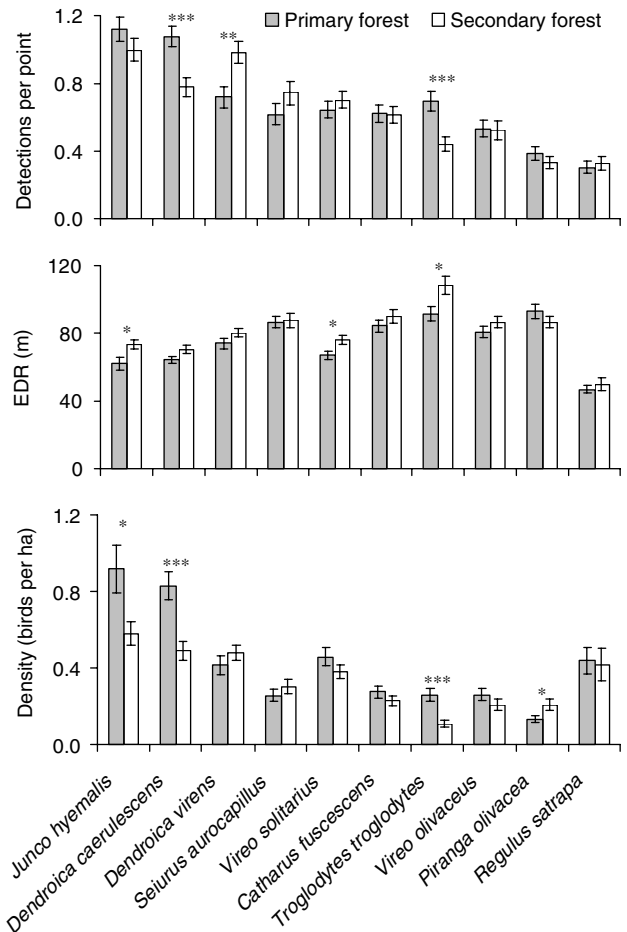


Fig. 3 – Comparisons of breeding birds recorded for 247 paired survey locations in primary and secondary forests. (Top) Mean relative abundance; counts are not adjusted for differences in detectability. (Middle) Effective detection radii (EDR) estimates (calculated using Program DISTANCE) for breeding bird species in primary and secondary forests. (Bottom) Estimated density of breeding birds in primary and secondary forests (calculated using Program DISTANCE). Error bars represent standard errors. Significant differences are denoted by asterisks (* $p < 0.05$, ** $p < 0.01$, * $p < 0.001$).**

both forest types. Four canopy species were significantly more abundant on primary forest sites (eastern hemlock [*Tsuga canadensis*], red oak [*Quercus rubra*], white oak [*Quercus alba*], and white pine [*Pinus strobus*]) and five species were significantly more common on secondary forest sites (yellow birch [*Betula allaghaniensis*], black birch [*Betula lenta*], American basswood [*Tilia heterophylla*], sugar maple [*Acer saccharum*], and pin cherry [*Prunus pensylvanica*]). Species composition patterns in the subcanopy and tall shrub layers were similar to the canopy, but with fewer significant differences. The subcanopy reflected differences in the canopy with eastern hemlock more abundant on primary forest sites and yellow birch more abundant on secondary forest sites. In the tall shrub layer, red oak, *Rhododendron* spp., and American chestnut (*C. dentata*) were significantly more abundant on primary forest sites while yellow buckeye (*Aesculus octandra*) was more common on secondary forest sites.

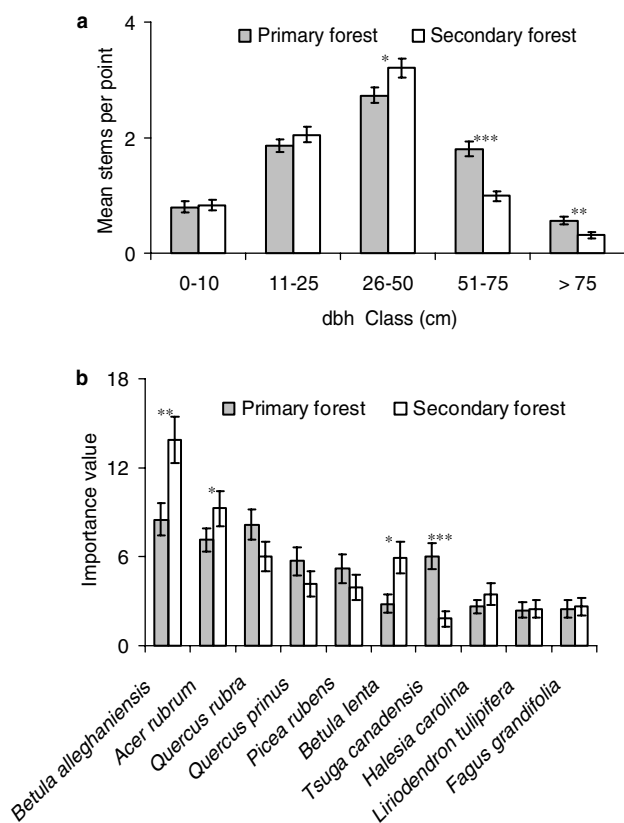


Fig. 4 – (a) Number of trees counted for five dbh classes. Data are from 191 paired survey plots in primary and secondary forests. (b) Mean importance values for tree species observed in 233 paired survey plots in primary and secondary forests. Significant differences are denoted by asterisks (* $p < 0.05$, ** $p < 0.01$, * $p < 0.001$).**

4. Discussion

Breeding bird and vegetation community structure were largely similar on primary forest and mature secondary forest 70 years after establishment of Great Smoky Mountains National Park. Primary and secondary forests shared 24 of the 25 most common bird species, as measured by number of detections (Appendix). Nonetheless, three breeding bird species (Winter Wren, Black-throated Blue Warbler, and Dark-eyed Junco) showed significantly higher densities on primary forest sites than on closely paired secondary forest sites. Scarlet Tanagers were more abundant on secondary forest sites. The paucity of extensive tracts of similar primary forests in the American Southeast limits our knowledge of historic primary forest bird communities in this region. Differences in breeding bird density probably reflect habitat preferences linked to differences in vegetation characteristics on primary and secondary forest sites. The greater abundance of eastern hemlock on primary forest sites likely contributes to increased densities of Winter Wren at those sites because of the wren’s preference for coniferous species (Hansen et al., 1995; Haney and Lydic, 1999). Similarly, the higher abundance of *Rhododendron* spp. in the tall shrub layer may have contributed to increased Black-throated Blue Warbler density on pri-

mary forest sites due to its use of *Rhododendron* spp. as a foraging and nesting substrate (Holmes, 1994). Higher vegetation density in the tall shrub layer of primary forest sites may benefit those two species, as well as Dark-eyed Juncos, that prefer to nest and forage in lower vegetation strata. Likewise, higher canopy cover on secondary forest sites may explain higher Scarlet Tanager densities on those sites because the species prefers to nest and forage in closed-canopy forest (Mowbray, 1999).

Primary forest sites, as expected due to the age difference in the two forest types, had significantly more trees in large diameter at breast height (dbh) classes. Nonetheless, stem densities for smaller dbh classes were similar for primary and secondary forest sites, and we did not find a significant difference in the number of standing snags. Cover estimates for canopy, subcanopy, and tall shrub layers showed significantly higher canopy cover in secondary forests, no difference in the subcanopy layer, and significantly higher tall shrub cover in primary forests. Although we do not have tree fall gap data, these results may indicate more canopy openings in primary forests which could, in turn, result in higher cover estimates in the tall shrub layer of the forest. In general, these results are similar to those outlined in Ziegler’s (2000) comparison of old-growth and mature second-growth hemlock-hardwood forests with the exception that we did not find a difference in snag density between the two sites. Primary forest sites had significantly higher importance values for late successional canopy species (eastern hemlock, red oak, white oak, and white pine) and significantly lower values for canopy species associated with intermediate successional forests (yellow birch, black birch, American basswood, and sugar maple). These compositional differences are similar to those identified by Latty et al. (2004) for old-growth and mature second-growth forests and also mirror Ziegler’s (2000) finding of significantly higher densities of eastern hemlock on old-growth sites.

Stronger differences between undisturbed and disturbed habitats were identified in a study of salamander communities in Great Smoky Mountains National Park (Hyde and Simons, 2001). Two factors may explain the larger differences in salamander populations on these two forest types. First, differences in the size and vagility of salamanders and birds suggests they may respond to habitat features at different spatial scales. Consequently, vegetation differences between undisturbed and disturbed forests in the park may occur at a scale relevant to salamanders, but below the resolution at which many breeding bird species select habitat. Second, salamanders and breeding birds respond to very different habitat features. Soil conditions and leaf litter characteristics are vital habitat features for salamanders, while vegetation structure is a key habitat feature for most breeding forest birds. Although we lack soil and leaf litter data for our study site, Latty et al. (2004) showed that differences in soil characteristics persist nearly 100 years after selective logging in northern hardwood forests.

The differing results for bird and salamander communities in primary and secondary forests in Great Smoky Mountains National Park have important management implications for resource managers who rely on indicator taxa to draw inferences about other species or communities. While breeding birds are more readily sampled than salamanders, breeding

birds may not be adequate indicators of forest condition for species that are potentially more sensitive to fine scale habitat features. Similarly, vegetation data should be accompanied by information about soil characteristics in discriminating between primary and secondary forests. These results add strength to the recommendations of Rolstad et al. (2002) who stress the importance of identifying appropriate structural features in the identification of old-growth and suggest that dispersal-limited species may be better indicators of old-growth forests than more vagile species.

The differences in our bird community results for unadjusted point count data and density estimates which accounted for differences in detection probabilities underscore the importance of using methods that estimate detection probabilities in different sampling units. Had we not adjusted our point transect data to account for differential probabilities of detection in primary and secondary forests, we would have missed the significant difference in densities for Dark-eyed Junco and erroneously reported a significant difference for Black-throated Green Warbler. Both of these species exhibited lower effective detection radii (EDR) in primary forests compared with secondary forests, effectively increasing the community differences for Dark-eyed Juncos and decreasing the differences for Black-throated Green Warblers. Failure to account for such differences in detection probabilities in different sampling units, such as primary and secondary forests, weakens inferences from comparative studies (Yoccoz et al., 2001; Pollock et al., 2002).

Our results indicate that measures of bird abundance in southern Appalachian forests should be adjusted for differences in detection probability associated with both bird species and habitat types. Nine of the 10 most common species in this study had lower EDRs in primary forests, suggesting reduced detection probabilities in primary forest for common forest bird species. Differences in detection probability may reflect differences in vegetation structure. For example, the dense tall shrub forest layer may have reduced observers' ability to hear distant birds on primary forest sites. Likewise the dense canopy layer on secondary forest sites may explain why EDRs for Scarlet Tanagers were lower on secondary forest sites. Fixed radius plots, scaled to reduce differences in detection probability among species or habitats, can reduce sampling bias, but often at the cost of reduced sample size and lower statistical power. Alternatively, detection probabilities can also vary with bird density (Bart and Schoultz, 1984).

Although bird and vegetation communities on primary and mature secondary forest in Great Smoky Mountains National Park are currently similar, some breeding bird species still differentiate between the two forest types, and some differences in vegetation structure and composition remain. These results suggest that forests logged 70–100 years ago have substantially recovered as habitat for most forest songbirds, but several caveats must be considered. First, there is no way to determine if the remaining primary forest sites are representative of historic primary forest conditions, because most of the primary forest remaining in the eastern United States survives on steep, high elevation slopes that were inaccessible to loggers. Second, results for the bird community cannot be generalized to other species, such as salamanders, that respond to different habitat features and which may respond at different spatial and temporal scales. Third, the bird communities on undisturbed and previously logged sites in Great Smoky Mountains National Park may be more similar today than those on more fragmented landscapes because many secondary forest sites in the park are contiguous with large patches of undisturbed forest. The proximity of logged sites to extensive tracts of primary forest may have buffered the effects of disturbance, if the primary forest sites served as population sources for regenerating habitats. Coarse-scale island biogeographic (MacArthur and Wilson, 1967; Newton, 1995) and population source-sink (Robinson et al., 1995) relationships are well established for bird populations. At a finer scale, the putative role source areas for birds (Simons et al., 2000; Brotons et al., 2003) and plants (Duffy and Meier, 1992; Matlack, 1994) play in the colonization of regenerating habitats, has proven difficult to verify. Finally, because our findings are based on abundance measures, our comparisons ignore potentially important differences in the survival and fecundity of populations (Van Horne, 1983).

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Appendix - Number of detections of breeding bird species observed in paired primary and secondary forest sites (N = 247 pairs). Species are in descending order of commonness, as measured by number of detections (unadjusted for detectability).

Common name	Scientific name	Primary forest	Secondary forest
Dark-eyed Junco	<i>Junco hyemalis</i>	277	247
Black-throated Blue Warbler	<i>Dendroica caerulescens</i>	266	192
Black-throated Green Warbler	<i>Dendroica virens</i>	178	243
Ovenbird	<i>Seiurus aurocapillus</i>	153	184
Blue-headed Vireo	<i>Vireo solitarius</i>	159	174
Veery	<i>Catharus fuscescens</i>	154	152

Appendix – continued

Common name	Scientific name	Primary forest	Secondary forest
Winter Wren	<i>Troglodytes troglodytes</i>	172	109
Red-eyed Vireo	<i>Vireo olivaceus</i>	132	129
Scarlet Tanager	<i>Piranga olivacea</i>	95	82
Golden-crowned Kinglet	<i>Regulus satrapa</i>	75	81
Red-breasted Nuthatch	<i>Sitta canadensis</i>	64	47
Canada Warbler	<i>Wilsonia canadensis</i>	56	52
Black-capped Chickadee	<i>Poecile atricapilla</i>	44	49
Black-and-white Warbler	<i>Mniotilta varia</i>	44	45
Brown Creeper	<i>Certhia americana</i>	36	48
Hooded Warbler	<i>Wilsonia citrina</i>	38	43
Northern Parula	<i>Parula americana</i>	41	35
Chestnut-sided Warbler	<i>Dendroica pensylvanica</i>	42	27
Eastern Towhee	<i>Pipilo erythrophthalmus</i>	33	35
Pileated Woodpecker	<i>Dryocopus pileatus</i>	26	35
Chimney Swift	<i>Chaetura pelagica</i>	42	18
Blackburnian Warbler	<i>Dendroica fusca</i>	34	22
American Crow	<i>Corvus brachyrhynchos</i>	26	29
Blue Jay	<i>Cyanocitta cristata</i>	26	25
Tufted Titmouse	<i>Baeolophus bicolor</i>	24	26
Hairy Woodpecker	<i>Picoides villosus</i>	15	26
Rose-breasted Grosbeak	<i>Pheucticus ludovicianus</i>	20	10
Indigo Bunting	<i>Passerina cyanea</i>	8	22
Wood Thrush	<i>Hylocichla mustelina</i>	9	18
Acadian Flycatcher	<i>Empidonax virescens</i>	12	14
White-Breasted Nuthatch	<i>Sitta carolinensis</i>	10	15
Worm-eating Warbler	<i>Helmitheros vermivorus</i>	6	15
Common Raven	<i>Corvus corax</i>	10	5
Eastern Wood-Peevee	<i>Contopus virens</i>	9	8
Downy Woodpecker	<i>Picoides pubescens</i>	9	4
Carolina Chickadee	<i>Poecile carolinensis</i>	8	8
Carolina Wren	<i>Thyrothorus ludovicianus</i>	4	12
American Robin	<i>Turdus migratorius</i>	7	4
Pine Siskin	<i>Carduelis pinus</i>	7	3
Cedar Waxwing	<i>Bombycilla cedrorum</i>	4	6
Gray Catbird	<i>Dumetella carolinensis</i>	3	7
American Goldfinch	<i>Carduelis tristis</i>	2	6
Yellow-billed Cuckoo	<i>Coccyzus americanus</i>	3	4
Brown Thrasher	<i>Toxostoma rufum</i>	3	2
Broad-winged Hawk	<i>Buteo platypterus</i>	3	1
Ruby-throated Hummingbird	<i>Archilochus colubris</i>	2	2
Red Crossbill	<i>Loxia curvirostra</i>	2	2
Ruffed Grouse	<i>Bonasa umbellus</i>	2	2
Northern Flicker	<i>Colaptes auratus</i>	1	3
Wild Turkey	<i>Meleagris gallopavo</i>	2	1
Yellow-throated Warbler	<i>Dendroica dominica</i>	2	1
Northern Cardinal	<i>Cardinalis cardinalis</i>	0	3
Purple Finch	<i>Carpodacus purpureus</i>	0	3
Red-bellied Woodpecker	<i>Melanerpes carolinus</i>	2	0
American Redstart	<i>Setophaga ruticilla</i>	1	1
Eastern Phoebe	<i>Sayornis phoebe</i>	1	1
Yellow-throated Vireo	<i>Vireo flavifrons</i>	1	1
Louisiana Waterthrush	<i>Seiurus motacilla</i>	0	2
Blue-gray Gnatcatcher	<i>Poliophtila caerulea</i>	0	2
Red-tailed Hawk	<i>Buteo jamaicensis</i>	1	0
Brown-headed Cowbird	<i>Molothrus ater</i>	0	1

(continued on next page)

Appendix – continued

Common name	Scientific name	Primary forest	Secondary forest
Great Crested Flycatcher	<i>Myiarchus crinitus</i>	0	1
Barred Owl	<i>Strix varia</i>	0	1
Mourning Dove	<i>Zenaida macroura</i>	0	1
Swainson's Warbler	<i>Limnothlypis swainsonii</i>	0	1
Eastern Screech-owl	<i>Otus asio</i>	0	1
Summer Tanager	<i>Piranga rubra</i>	0	1
Saw-whet Owl	<i>Aegolius funereus</i>	0	1

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