

THE EFFECTS OF HUMAN RESIDENTIAL DEVELOPMENT ON AVIAN COMMUNITIES ALONG THE SNAKE RIVER RIPARIAN CORRIDOR IN JACKSON HOLE WY ♦ USA

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♦ ABSTRACT

Throughout North America, bird declines may be attributable to loss of habitat on the breeding grounds. Human land uses, especially residential development pressures within western ecosystems, are greatly impacting riparian landscapes and biota, particularly breeding birds. While most studies have focused on eastern birds, it is unclear how western bird assemblages respond to development and its concomitant effects on habitat alteration and fragmentation. I sampled bird community parameters and habitat variables at three spatial scales (microhabitat, macrohabitat, and landscape) along a human development gradient along the Snake River riparian corridor in Jackson Hole, WY, USA. Fifty-six cottonwood forest patches were surveyed during the 2001-02 breeding season. Principal component analysis, canonical correspondence analysis, and multiple linear regression statistical tests were used to determine the effects of housing densities on avian assemblages, guilds, and habitat features. Overall species richness and diversity declined with increasing human development. Neotropical migrant species were most negatively impacted and consistently declined in proportional representation on forested plots as human development densities increased. Short-distance migrants, food generalists, ground gleaners, and avian nest predators all increased with increasing human development. Brood parasites, on the other hand, did not increase with increasing fragmentation and their distribution may reflect the availability of nest host species. Of

the environmental variables measured, landscape features were most affected and metrics consistent with habitat fragmentation were most correlated with human development. These results suggest that residential development within riparian habitats may be exerting a strong negative influence on western bird communities and at high densities may lead to a depauperate avian biota and reproductive sinks.

♦ METHODS

Study Area

Bird, vegetation, and landscape data were collected in riparian forest patches along the Snake River in Jackson Hole, Wyoming, USA. The study area was approximately 40 km long from the southern portion of Grand Teton National Park (GTNP) (4839833N, 526157E, Zone 12T) to the town of Wilson, Wyoming, USA (4815635N, 512057E, Zone 12T) in its northern and southern extents, respectively. Surveys were conducted on private land (n=42) located within Teton County and within the boundaries of GTNP (n=14). Dominant tree species were cottonwood (*Populus spp.*) and spruce (*Picea spp.*) with a variable understory. Grasses, forbs, bare ground, and detritus formed the dominant ground cover classifications. Smith (2002) provides a complete overview of the study area as well as additional information on methods, results, analyses, and implications.

I selected a stratified random sample of plots along a human development density gradient based on a query of a 1999 high-resolution (0.3m² pixel size) digital aerial photograph of Teton County (Horizons, Inc. 1999). ArcView 3.2 (ESRI) was used to digitize the raster data to ensure that all sets met the following criteria: 1) each plot was separated by at least 150m to insure independence of bird data (Reynolds et al. 1980), 2) sites were selected in the major categories of riparian forest, cottonwood and spruce, 3) plots were located within patches with an area >2,500m², and 4) had a negligible slope. Sites were visited to ensure accuracy of site-specific attributes.

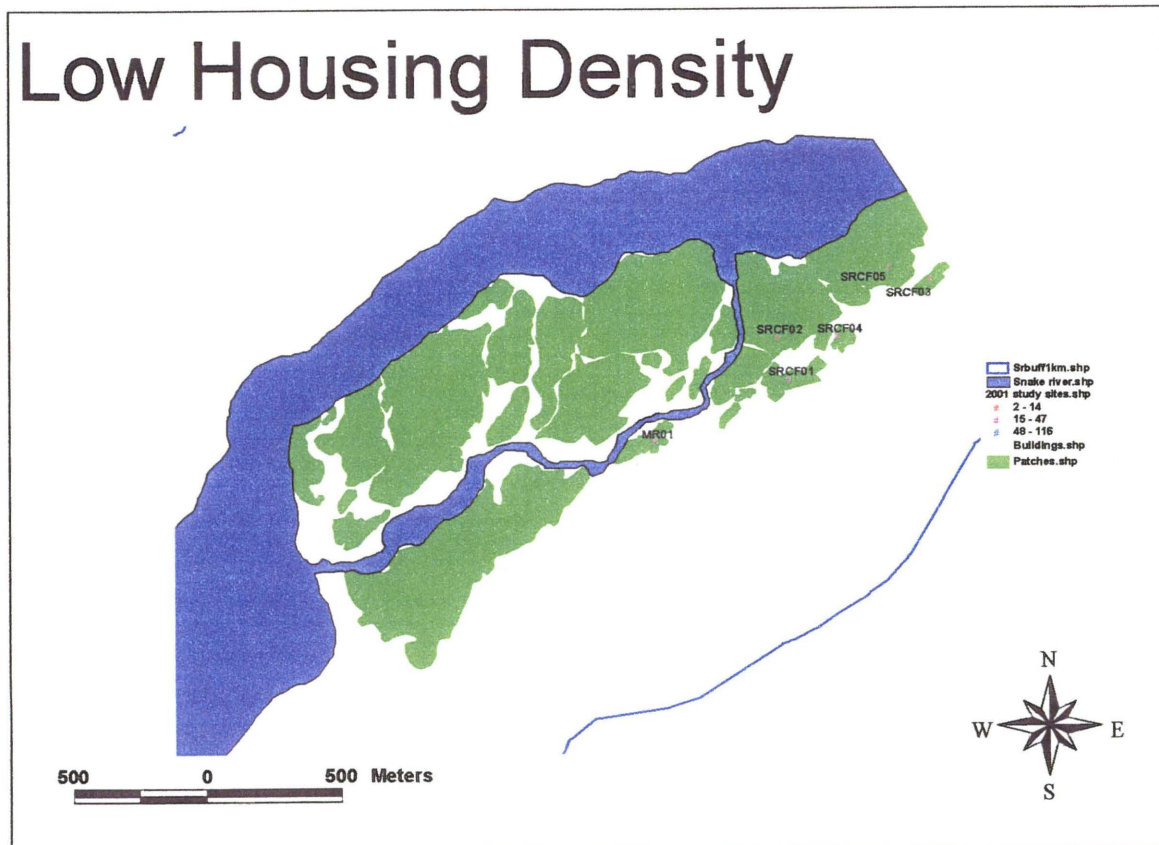
From the pool of potential survey sites, fourteen plots were randomly selected in each of the low- (Figure 1), medium- (Figure 2), and high-density (Figure 3) development areas. Landowner permission was obtained to conduct point counts throughout the breeding season. Fourteen reference plots were established in areas with no human development (reference plots) located in the southern portion of GTNP.

Point Count Protocols

Avian surveys were conducted between May 20 and June 30, 2001 and 2002. Bird abundance and diversity data were collected using the 25m fixed-radius and unlimited-radius point count described by Hutto et al. (1986). Count duration was 15 minutes long. Point counts were conducted between the hours of 0600 and 1000. To control for biases associated with varying avian activity levels throughout the survey period, each site was surveyed in the early, mid, and late portions of the morning (0600-0720, 0720-0840, 0840-1000, respectively). Point counts were not conducted during periods of inclement weather (e.g., strong winds, precipitation).

Bird data generated from the observations included species richness, abundance, diversity, and guild composition. The mean number of detections over the three visits was used to calculate relative abundance values. *A priori* life history guilds, including nesting, migratory, foraging, and food, were classified according to Erlich et al. (1988).

Figure 1: Figure of Low Density Housing



Migratory status was determined from previously published data, including Johnsgard (1986), Erlich et al. (1988), Freemark and Collins (1992). Diversity was calculated according to the Shannon's Diversity Index (Zar 1984 and Krebs 1989).

Scale Perspectives

I selected a nested project design in order to determine the effects of human development on bird habitat use at a variety of spatial scales. Finer resolution of microhabitat provided insight into habitat utilization while coarser metrics (i.e., landscape) placed habitats within a context. Table 1 summarizes all variables measured.

Table 1: Summary Statistics of Environmental Variables

Num.	Variable	Units	Mean	Standard Deviation	Sum	Minimum	Maximum
Microhabitat							
1	Deciduous Trees	#	15.580	9.024	872.500	4.333	49.000
2	Coniferous Trees	#	3.958	4.145	221.667	0.333	20.000
3	Snags	#	1.333	1.378	74.667	0.000	8.000
4	Tree Density	#/ha	514.658	240.585	28820.83	200.000	1325.000
5	Mean Tree dbh	cm	31.018	9.419	1737.025	8.941	64.787
6	Canopy Cover	%	73.884	17.738	4137.497	12.567	93.825
7	Canopy Height	ft.	67.106	10.543	3757.922	44.500	90.750
8	Deciduous Shrubs	#	54.891	49.053	3073.875	4.000	302.000
9	Coniferous Shrubs	#	11.385	16.552	637.542	0.000	82.333
10	Nearest Log Distance	m	7.796	2.158	436.579	2.650	11.300
11	NL Diameter	cm	18.452	9.540	1033.287	0.000	44.000
12	NL Length	m	6.441	3.065	360.700	0.000	15.300
13	Understory Dens. (0-0.3m)	%	71.748	20.085	4017.889	10.000	100.000
14	Understory Dens. (0.3-1m)	%	45.309	21.648	2537.285	0.714	100.000
15	Understory Dens. (1-2m)	%	33.949	19.292	1901.125	0.000	100.000
16	Understory Dens. (2-3m)	%	34.622	19.569	1938.833	0.000	100.000
17	Ground Cover Grass	%	23.360	14.452	1308.137	1.050	61.150
18	GC. Forbs	%	10.149	6.656	568.346	0.050	29.567
19	GC. Bare Ground	%	3.648	5.691	204.308	0.000	30.250
20	GC. Shrubs	%	7.885	6.719	441.558	0.083	29.075
21	GC. Litter	%	53.437	17.915	2992.475	17.600	92.950
Macrohabitat							
22	Patch Area	ha	2.360	2.873	132.160	0.351	14.459
23	Core Area	ha	0.337	1.127	18.880	0.000	5.195
24	Patch Perimeter	m	702.532	421.971	39341.810	241.626	2195.795
25	Perimeter:Area	m/ha	0.042	0.013	2.336	0.013	0.074
Landscape							
26	Number of Houses	#	23.661	28.708	1325.000	0.000	114.000
27	Nearest Neighbor	m	70.659	47.487	3956.910	10.530	221.318
28	Area of Neighboring Patches	ha	2.273	1.619	127.271	0.445	7.910
29	Total Area of Surrounding Patches	ha	42.979	27.539	2406.801	8.928	109.206
30	Forest Coverage	%	0.299	0.154	16.727	0.109	0.732
31	Ground Coverage	%	0.416	0.140	23.284	0.114	0.645
32	Water Coverage	%	0.098	0.070	5.497	0.004	0.286
33	House Coverage	%	0.016	0.017	0.903	0.000	0.071
34	Road Coverage	%	0.045	0.039	2.509	0.000	0.129
35	Sand Coverage	%	0.057	0.050	3.178	0.000	0.183
36	Wetland Shrub Coverage	%	0.070	0.053	3.904	0.000	0.236
Other							
37	Human Disturbance	#	9.679	9.522	542	0.000	29.667

Microhabitat Sampling

Vegetation sampling techniques used were those developed by James and Shugart (1970), modified by Noon (1981). Circular plots (0.04ha) were constructed around the point count location and at random points within the forest patch. Large, medium, and small area plots were sampled using three, two, or one plots, respectively, in order to

standardize sampling effort. Forest structural components measured were % canopy and ground cover, understory density, shrub and ground cover dispersion, dbh of all trees >3cm within the plot, representative tree height (five trees), shrub and stem density of coniferous and deciduous shrubs (stem diameter ≤3cm), number of snags present, and dominant shrub species (Table 1).

Macrohabitat Metrics

In addition to the fine grain analysis of habitat, I employed a macrohabitat-level approach. Using Arc View 3.2 GIS (ESRI), digital orthoquads, and aerial photographs of the study area, I incorporated methods described by McGarigal and Marks (1995) to analyze contextual and structural landscape information. These included measuring patch area, perimeter, perimeter:area ratio, and core area (see also Freemark et al. 1995).

Landscape Metrics

To determine the effects of housing development on landscape-scale features and subsequent alterations within the bird community, several measures were computed. Landscape metrics included number of houses and roads within 500m, mean distance to the five closest patches (i.e., degree of isolation), mean area of five closest patches, and total area of patches within 500m of each patch. Landscape metric values were calculated based on queries and measurements taken from the GIS database.

Landscape coverages were also calculated within 1 km² of each study plot. Digital aerial photographs of each study site were overlaid with a reference grid at a scale of 1:6,000. Estimates were made for the percentage of land cover in each 200m x 200m plots (n=25) and averaged across the 1 km² area. Land coverage classifications included: non-forested habitats (bare ground, gaps in forests, grass, and upland sagebrush habitats), forest coverage, water, wetland shrub (*Salix spp.*), sand, buildings, and roads (Table 1).

◆ ANALYSES

Avian species richness, abundance and diversity were calculated for each site using PC-ORD (McCune and Mefford 1999). Each of the 37 environmental variables measured was tested for normality (i.e., skewness, kurtosis, normal probability plots, and Ryan-Joiner tests for normality)

using MINITAB statistical software (MINITAB 1998). Data that exhibited significant departures from normality were transformed prior to analysis using standard transformation techniques, including square root, logarithmic, and arcsine square root transformations (Neter et al. 1989).

I used principal component analysis (PCA) to condense the 37 environmental variables to composite values using MINITAB statistical software (MINITAB 1998). Significant components were then used in multiple linear regression analysis (MRA) to determine the effects of the dominant environmental gradients on species richness, abundance, diversity, and migratory status.

I employed Canonical Correspondence Analysis (CCA) to determine the effects of the dominant environmental gradients on individual bird, food, foraging, and nesting guild distributions in environmental space using PC-ORD (McCune and Mefford 1999).

★ RESULTS OVERVIEW

Bird Occurrence

During the 2001-02 breeding seasons, over 9,000 individual birds were detected representing 61 species during 336 visits made to the 56 study sites (refer to Smith 2002 for complete species lists). Species richness ranged from 10 to 24 species with a mean of 17 species per plot. Simpson's Diversity for each plot ranged from 2.829 to 1.71 with a mean of 2.429.

Habitat

Habitat variables were collected and analyzed at three spatial scales (microhabitat, macrohabitat, and landscape). Sites were dominated by a cottonwood canopy and had variable understory and ground cover. Patches were small with large amounts of edge habitat and limited core area. All patches were surrounded by upland sagebrush habitats or agricultural lands.

Six principal components for the 37 environmental variables were deemed to be meaningful according to the scree criterion and broken-stick eigenvalues (McCune and Mefford 1999; McGarigal et al. 2000). The component loadings were considered to be significant if >0.450 (McGarigal et al. 2000) and were examined for

relationships among variables. Most components loaded on a set of conceptually related variables and I assigned ecological meaning as appropriate. Of the total variation within the environmental matrix, 67.2% was captured by the six selected components. Table 2 summarizes the PCA loadings for the 37 environmental variables.

Table 2: Principal Component Loadings of 37 Environmental Variables

Principal Component	PC1	PC2	PC3	PC4	PC5	PC6
Variables	Human Development Gradient and Associated Trends	Understory Density and % Wetland Vegetation	% Water, Snags, and Ground Cover Type	Patch Characteristics	Stand Age and size of adjacent patches	# of Conifers
Eigenvalues	7.996	5.770	3.639	3.182	\$2.334	1.939
% Explained	21.600	15.600	9.800	8.600	\$6.300	5.200
Cumulative %	21.600	37.200	47.000	55.600	\$61.900	67.200
# Deciduous Trees	-0.719	-0.149	0.065	0.242	\$0.383	-0.052
# Conifers	0.394	-0.385	-0.071	0.133	\$0.137	-0.495
# SNAGS	0.258	-0.105	-0.427	0.244	\$0.042	-0.371
Tree Density	-0.493	-0.418	-0.014	0.340	\$0.459	-0.335
Tree dbh	0.283	0.506	-0.078	-0.433	\$0.477	0.283
Canopy Cover	-0.305	0.125	0.326	-0.292	\$0.255	0.229
Canopy Height	0.100	0.314	-0.326	-0.513	\$0.176	0.205
# Deciduous Shrubs	-0.202	-0.604	0.271	0.174	\$0.204	0.348
# Coniferous Shrubs	0.615	-0.420	0.226	0.188	\$0.006	-0.203
Nearest Log Distance	-0.362	-0.045	0.554	0.180	\$0.127	0.057
Nearest Log Diameter	0.081	0.321	-0.167	-0.194	\$0.133	0.165
Nearest Log Length	0.287	0.074	-0.476	0.039	\$0.163	-0.234
Understory Dens. (0-0.3m)	-0.083	-0.729	-0.298	-0.030	\$0.033	0.306
Understory Dens. (0.3-1m)	-0.282	-0.746	-0.209	-0.058	\$0.096	0.350
Understory Dens. (1-2m)	-0.255	-0.630	-0.444	0.147	\$0.176	0.145
Understory Dens. (2-3m)	-0.263	-0.565	-0.463	0.182	\$0.159	0.123
Ground Cover Grass	-0.112	-0.275	-0.569	-0.440	\$0.065	-0.308
GC. Forbs	-0.003	-0.578	-0.348	-0.070	\$0.202	0.092
GC. Bare Ground	-0.092	0.075	0.453	0.123	\$0.180	-0.384
GC. Shrubs	-0.108	-0.526	0.502	0.129	\$0.027	0.352
GC. Litter	0.250	0.542	0.353	0.336	\$0.206	0.171
Housing Density	-0.688	-0.202	0.183	-0.408	\$0.067	-0.186
H. Dist	-0.685	-0.233	-0.018	-0.314	\$0.070	0.009
Core Area	0.537	-0.164	0.094	-0.414	\$0.392	0.017
Area	0.632	-0.262	0.103	-0.554	\$0.379	-0.041
Perimeter	0.568	-0.285	0.117	-0.549	\$0.331	-0.023
Perimeter:Area	-0.678	0.180	-0.040	0.480	\$0.301	0.168
Nearest Neighbor Dist.	-0.530	0.373	-0.187	0.221	\$0.081	-0.096
N. N. Area	0.504	-0.297	0.063	0.186	\$0.558	-0.284
Total area	0.704	-0.441	0.295	0.061	\$0.144	0.105
% Forest	0.609	-0.443	0.372	0.115	\$0.236	0.049
% Ground	-0.524	0.142	0.120	-0.260	\$0.224	-0.234
%Water	0.169	0.188	-0.640	0.273	\$0.444	0.169
% Houses	-0.750	-0.075	0.049	-0.385	\$0.031	-0.202
% Roads	-0.830	-0.221	0.038	-0.310	\$0.172	-0.020
%Sand	0.554	0.113	-0.399	0.295	\$0.253	0.216
%Wetland	-0.056	0.747	-0.229	0.085	\$0.318	0.151

The dominant primary component described a decreasing human development gradient. Plots positively associated with this component were larger in size, occurred in closer proximity to neighboring patches, and had significantly fewer roads, houses, deciduous trees, and human disturbance scores. The second component primarily loaded on understory density and % of wetland shrub within the landscape matrix. Plots with positive associations with this component had dense understories and large amount of wetland shrubs within 1 km². The third component described % of water within the landscape matrix and dominant ground cover type. Positively associated plots had more snags, more dispersed downed logs, more ground cover dominated by shrubs and bare ground with less grass cover, while having less water in the landscape matrix. The fourth component captured patch (macrohabitat) characteristics. Patches that were positively associated with this component were

larger with a lower perimeter:area ratio. Tree density, mean dbh, and area covered by adjacent patches loaded the fifth component. The abundance of conifers loaded the final component. Plots associated with the fifth component had more trees with smaller mean dbh and were located next to larger patches. The increased tree density and smaller dbh is indicative of younger stands, therefore it was described as an indicator of stand age (i.e., positively associated plots were younger and located closely to larger forest patches). Plots positively related to the final component were characterized by fewer coniferous trees.

◆ BIRD HABITAT RELATIONSHIPS

Avian Species Richness, Diversity, and Abundance

Based on multiple linear regression analysis (MRA), 2 components were significantly related to trends in species richness ($r^2=0.458$, $F=6.90$, $P=0.0001$). The best predictors for species richness were principal components 1 (human development) ($t=5.89$, $P=0.0001$) and 5 ($t=-2.08$, $P=0.043$) (stand age and proximity to larger patches).

MRA was used to determine bird diversity response to the environmental matrix. Two principal components were significantly related to trends in avian diversity along the development gradient ($r^2=0.497$, $F=8.06$, $P=0.0001$). PC1 (human development) was the most significant predictor ($t=6.13$, $P=0.0001$) and PC2 (understory characteristics and % wetland shrubs in the landscape) was the second best predictor of avian diversity along the development gradient ($t=-2.63$, $P=0.012$). MRA showed that bird abundance was not significantly related to any of the principal components ($r^2=0.13$, $F=1.22$, $P=0.313$).

Individual Species Responses

Canonical Correspondence Analysis (CCA) was used to determine how bird distributions were structured relative to the measured environmental variables. Bird species associated with human development included Black-billed Magpie, European Starling, Brewer's Blackbird, Northern Oriole, and Common Grackle (Figure 4). Species negatively associated with human development included Dusky Flycatcher, Hammond's Flycatcher, Ruffed Grouse, Dark-eyed Junco, and Swainson's Thrush.

Food Guild Responses

CCA was used to determine how food guilds were structured according to the environmental variables. Axis 1 described the human development gradient with human-associated guilds located on the left side of the axis and less human tolerant species on the right side. Generalists were negatively associated with Axis 1 (i.e., more abundant in areas of high development). Axis 2 defined a gradient characterized by positive associations with coniferous trees and shrubs and negatively associated with nearest neighbor distance. Plots and guilds associated with this axis were less isolated from neighboring patches and contained more coniferous trees and shrubs. Seed eaters were positively associated with Axis 2 while nectivores were negatively correlated to the axis. Berry eaters and insectivores did not correlate to any of the first 3 axes, indicating they are not significantly structured by either of these gradients.

Foraging Guild Responses

CCA was also used to determine how foraging guilds were structured according to trends within the environmental variables. Axis I described the human development gradient (i.e., more developed sites occurred along the right side of Axis 1). Environmental variables shown had $r^2>0.10$. Axis 3 described forest structural elements, particularly mean tree dbh and downed log characteristics. Foraging guilds associated with human development (i.e., values on the right side of Axis I) included aerial and ground gleaners. Guilds negatively associated with human development were foliage gleaners and hawkers. The bark gleaning guild was associated with Axis 3 and related to patches with larger mean tree dbh and with longer fallen logs.

Nesting Guild Responses

CCA revealed how nesting guilds were structured according to the dominant environmental variables. Axis 1 described the human development gradient with human dominated sites and human-associated species occurring to the left along Axis 1. Nesting guilds associated with development were sphere and cavity nesters while open cup nesters were negatively associated with this gradient (i.e., occurring on the right side of Axis 1). Axis 2 described trends in the number of coniferous trees and nearest neighbor distances (i.e., less isolated patches with more coniferous trees). Pendant and saucer nesters were positively associated with Axis 2

while brood parasites and cup nesters were not as strongly structured by either axis.

Migratory Guild Response

CCA could not be conducted on migratory guild responses to environmental variables due to high levels of multicollinearity. Therefore MRA was employed to determine if migratory strategy was related to the principal components generated from the environmental matrix.

Neotropical migrants were significantly related to PC1 and PC4 ($t^2=0.539$, $F=9.55$, $P=0.0001$). PC1 (human development gradient) was the strongest predictor ($t=5.85$, $P=0.0001$) and PC4 (patch characteristics) was the second best predictor ($t=4.53$, $P=0.0001$). These scores indicate that Neotropical migrants were more abundant on larger patches with fewer anthropogenic elements in the landscape. Short-distance migrants were negatively related to PC1 (human development gradient; $r^2=0.337$, $F=4.15$, $P=0.002$; $t= -4.67$, $P=0.0001$). Short-distance

Figure 4: CCA biplot of Individual Species and Environmental Gradients



Axis 1= Fragmentation and Human Development

migrants were more abundant on disturbed sites than on areas with little or no human disturbance. Resident species were not significantly related to any of the principal components ($r^2=0.157$, $F=1.52$, $P=0.191$).

◆ DISCUSSION AND CONSERVATION IMPLICATIONS

As development and habitat fragmentation increased, local bird communities were significantly impacted. The primary effects of residential development on bird community structure occurred at the landscape scale and included fragmentation effects such as increased patch isolation, smaller forest patches, as well as alterations within the habitat matrix (i.e., forests were surrounded by more roads, houses, and had higher human disturbance scores). Saab (1999) found similar results along the Snake River in Idaho. Shifts within the bird community included decreased species richness and diversity, a trend documented by others (Clergeau et al. 1998; Fernández-Juricic 2000; Jokimäki 1999). Bird abundance was not affected, however this is often the case. Aldrich and Coffin (1980) and Blair (1996) both reported increases in avian abundances as housing densities increased, primarily due to the addition of edge-tolerant and edge-specialist species. I discovered a similar trend along the development gradient in Jackson Hole. The decrease in avian diversity showed that while abundances remained comparable in high-density development, the most dominant species were locally common generalists or ground gleaners, which represent a drastically different community (i.e., lower species richness, diversity, and guild dominance) than those found in undeveloped areas.

I found increases in generalists, short-distance migrants, ground gleaners, aerial foragers, and cavity and sphere nesters as development increased. Guilds negatively associated with development included foliage gleaners, hawkers, open-cupped nesters, and Neotropical migrants. Given the decline of Neotropical migrant species (Robbins et al. 1989), increasing human development could have serious repercussions on breeding bird habitat conservation in the western United States.

The trends in abundances of brood parasites and avian predators reveal interesting patterns within western bird assemblages along the development gradient. As human densities increased, bird predators, particularly Black-billed Magpies, dramatically increased.

This phenomenon has been documented by several other researchers (Tomialojć 1978; Andrén 1992; Nilon et al. 1995; Kluza et al. 2000). While I did not directly measure the impact Black-billed Magpies had on bird nesting success, fledging rates, or overall survivorship, it is likely the presence of these important nest predators exerts a negative pressure on heterospecifics. I am currently investigating the effects of development and associated increases in avian nest predators on nest productivity through an artificial nest study.

Brood parasites such as the Brown-headed Cowbird did not increase with increasing housing densities or habitat fragmentation, which contrasts with results of similar studies in the eastern United States (Robinson et al. 1995). Similar findings on the decline of cowbird parasitism associated with increasing fragmentation have been found in Montana. Simple linear regression analysis indicated a significant relationship between the abundances of cowbirds and Yellow Warblers ($P=.0091$, $r^2=.119$), the primary nest host species in this study area (Erlich et al. 1986). CCA analysis of individual species distribution in environmental space also indicated a close relationship between Yellow Warbler and cowbird distributions (Fig. 4). This trend in cowbird distributions could be due to fragmentation effects operating in opposite directions within western ecosystems. As fragmentation levels increase, cowbird host species (particularly open cup nesting Neotropical migrants) become less represented within the bird communities. Fragmentation and edge availability is a common feature of western riparian corridors (Knight 1994) and it is likely that cowbirds are able to parasitize a large proportion of undeveloped forest patches. Within the west, brood parasitism is not likely to heavily affect bird populations as human development increases and cowbird distributions may be structured by availability of host species rather than increasing fragmentation.

Conservation efforts in the west should therefore focus on maintaining adequate areas of undeveloped riparian forest tracts. Retaining landscape and macrohabitat (i.e., patch) level characteristics such as large areas of forest, high contagion, and core area could help retain local bird populations. Furthermore, reducing both primary (i.e., disturbance levels) and secondary human impacts (e.g., road and housing densities) may also contribute to conservation efforts. It is clear from the results of my study that increasing housing densities may support more generalist species and avian predators, particularly Black-billed Magpies. It may

be that the presence and increased abundance of these species may be even more impactful than the results of this study indicate. Hansen and Rotella (2000) have suggested that cottonwood patches immersed within a human-dominated landscape may be acting as reproductive sink areas and may thus be exerting considerably more negative influence on the larger metapopulation than indicated by simple declines in species richness, diversity, and underrepresentation of certain assemblage guilds. Despite the attractiveness of the areas for residential development (Hansen and Rotella 1999), it will be vital to limit development in riparian forests to low to moderate densities to ensure the persistence of breeding Neotropical migrant species.

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