

1 **1. Introduction**

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3 Many agencies and programs encourage prairie landowners to implement practices to
4 maintain or increase biodiversity (Gottschalk et al., 2009). Often there is inadequate monitoring
5 to measure progress toward biodiversity-related objectives (Mac Nally and Fleishman, 2002),
6 since it is either forgotten or too complicated, expensive, or time-consuming. However,
7 indicators can be used to partially overcome these constraints, in order to measure progress
8 towards a program's objectives (Kati et al., 2004; Mac Nally and Fleishman, 2002).

9 Indicators are species or groups of species “whose population trends reflect the average
10 behavior of the constituent species, but also cast light on trends in attributes of other taxa and act
11 as a surrogate for ecosystem health” (Gregory et al., 2005, p. 271). In short, indicators can
12 provide valuable information on complex issues in a relatively accessible way (Niemeijer and de
13 Groot, 2008). Bioindicators should have biological, methodological, and societal relevance
14 (Burger, 2006), and require three kinds of validation (Bockstaller and Girardin, 2003): (1) for
15 *design validation*, indicators should be scientifically founded, representative, meaningful,
16 credible, reliable, based on accessible data, and derived with consistent methods (Gregory et al.,
17 2005); (2) for *output validation*, indicators should be timely, immediate, responsive, and applied
18 at the appropriate spatial and temporal scales (North American Bird Conservation Initiative,
19 2004; Niemeijer and de Groot, 2008; Gottschalk et al., 2009); and (3) for *end use validation*,
20 indicators should be understandable, relevant to landowners and decision-makers, simple, cost-
21 effective, consistent, and provide a broader context (Bildstein, 2001; Burger, 2006; Turnhout et
22 al., 2007; Niemeijer and de Groot, 2008).

1 Among many other considerations, a biodiversity indicator would rate highly if its
2 taxonomy is stable, natural history is well known, populations are readily surveyed and occupy a
3 breadth of habitats, populations are relatively sensitive to habitat change, patterns are reflected in
4 other related and unrelated taxa, and presence has economic importance (Pearson, 1994;
5 Papazoglou and Phillips, 2003; Burger, 2006; Öster et al., 2008). For many of these reasons,
6 birds are useful indicators of biodiversity and environmental quality (Bildstein, 2001; Trulio,
7 2004; Padoa-Schioppa et al., 2006). Birds have been used as indicators of environmental and
8 biodiversity change by many organizations (Bildstein, 2001; Gregory et al., 2005; Weber et al.,
9 2008), even though there are some limitations (Furness et al., 1993; Canterbury et al., 2000;
10 Gregory et al., 2005).

11 Monitoring to evaluate a program's success should be conducted at an appropriate scale
12 (Peterjohn, 2003). Many programs promote biodiversity-friendly practices at the farm scale, but
13 most biodiversity monitoring programs evaluate trends at ecosystem, provincial, or national
14 scales. Thus, the goal of this paper is to examine validation aspects of using birds as indicators of
15 biodiversity at the farm scale in rural Canada. First, this paper will evaluate two variations of
16 bird monitoring with point counts (50 m radius versus unlimited radius). Second, this paper will
17 examine the consistency of four biodiversity measures (species richness, abundance, Shannon
18 Index, and Inverse Simpson Index) across different farmland habitats.

19

20 **2. Methods**

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22 The study area was located in the prairie and aspen parkland ecoregions of east-central

1 Alberta, Canada. This area was chosen because it had significantly reduced natural habitats
2 (Bjorge et al., 2004), declining trends for some bird groups (Canadian Wildlife Service 2008),
3 and existing programs to promote biodiversity-friendly practices on farms. Landowners within
4 2.5 hours of Camrose were chosen through informal networks, ensuring a wide variety of
5 habitats, farming practices, and levels of farming intensities.

6 Using a stratified sampling design at 22 farms, field staff chose 1-2 sites, wherever
7 possible, to represent each of six habitat types: 1) upland forest: overstory dominated by
8 trembling aspen (*Populus tremuloides*); 2) tame pasture: perennial non-native grass cover for
9 cattle grazing; 3) wetland/riparian: primarily cattail (*Typha latifolia*) marshes, sedge (*Carex* spp.)
10 meadows, and creek corridors; 4) cultivated cropland: producing barley, wheat, and canola; 4)
11 native prairie: dominated by native grasses and a few shrubs; and 6) homestead habitats: areas
12 around landowners' farm buildings with attendant gardens and hedges (both native and non-
13 native species).

14 All counts took place during peak vocal bird activity (5 to 10 am), from May 30 to July 8,
15 2005 in conditions with no rain and wind speed less than 20 km/h. Individual counts took place
16 during a single morning to minimize travel costs and survey time. Sites were located near the
17 center of each habitat block, and were separated by at least 250 m. Field staff used 10-minute
18 point counts (Ralph et al., 1993; 1995), to record all birds seen and heard, with two variations in
19 the counting method, a 50 m radius count circle and an unlimited radius count circle (Ralph et
20 al., 1993). For each method, species richness was calculated by totaling the number of species
21 recorded during all counts on a single farm. Abundance was calculated by totalling the number
22 of individual birds recorded during all counts on a single farm.

1 For the other two diversity measures, only 50 m radius count data were used, using
2 species counts weighted according to the area of each habitat. The Shannon index formula was
3 $-\sum p_i \log_{10} p_i$ and the Inverse Simpson index formula was $1/\sum(p_i^2)$, where p_i is the proportion of
4 birds for species i (Feinsinger, 2001). The Shannon and Inverse Simpson indices are measures of
5 evenness by species within a population. The species richness and the Shannon measures are
6 biased toward rare species, while the abundance and Inverse Simpson measures are biased
7 toward dominant species (Stiling, 2002). The Shannon index was represented by its absolute
8 value multiplied by ten to produce values of the same magnitude as the other indices. Each
9 measure was calculated per farm and per site.

10 These measures were compared among habitat types. Statistical analyses used one-way
11 analyses of variance (post-hoc multiple comparisons used Tukey's b test) and Pearson
12 correlation tests (one-tailed). Furthermore, the species richness from each farm was compared
13 with the 5-year average of total species from the nearest Breeding Bird Survey (BBS) route (if
14 within 50 km of the farm). These one-day counts are conducted during the same time of year, but
15 involve 50 stops of 3 minutes each, using a 400 m fixed-radius point count (Downes and Collins,
16 2003). BBS data provide long-term continent scale data for tracking breeding bird populations
17 (Peterjohn, 2003).

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19 **3. Results**

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21 The mean farm size was 862 ha (mode = 413; range = 28-5400). Field staff sampled 178
22 sites, with an average of eight per farm (range = 3-12). Of these sites, 21% were classified as

1 cultivated cropland, 20% as tame pasture, 18% as upland forest, 16% as wetland/riparian, 15%
2 as native prairie, and 11% as homestead habitat types. Field staff recorded 4,350 individual birds,
3 representing 90 species.

4 Using the unlimited radius method, the mean species richness per farm was 35.2 and the
5 mean bird abundance per farm was 197.7 (Table 1). Using the 50 m radius method, the mean
6 species richness per farm was 25.8 and the mean bird abundance per farm was 99.0. For all
7 farms, the mean percentage of unlimited radius species richness accounted for by the 50 m radius
8 data was 52%. Similarly, the mean percentage of unlimited radius abundance accounted for by
9 the 50 m radius data was 50%. For comparison, the mean species richness of the nearest BBS
10 route within 50 km was 17 per site (versus 7.0 in this study) and 53 per survey (versus 25.8 per
11 farm in this study).

12 Average bird diversity measures for each habitat type are shown in Fig. 1. The 50 m
13 radius data showed differences among 3-5 habitat types, whereas the unlimited radius data
14 showed differences between only two habitat types for each diversity measure (Table 2).
15 Because the 50 m radius method showed more differentiation than the unlimited radius method,
16 subsequent analyses used only 50 m radius data.

17 Using any diversity measure, the wetland/riparian habitat scored highest, followed by the
18 homestead, upland forest, native prairie, tame pasture, and cultivated cropland habitats. By site,
19 all diversity indices were strongly correlated. By farm, species richness was correlated with
20 abundance ($r = 0.87$, $p < 0.001$) and the Shannon index ($r = 0.48$, $p = 0.011$), but not with the
21 Inverse Simpson index. The Shannon and Inverse Simpson indices were also correlated ($r = 0.91$,
22 $p = 0.001$). Similar results were found when the farms were ranked from highest to lowest by

1 each diversity measure.

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3 **4. Discussion**

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5 Species richness and bird abundance measures using the unlimited radius method were
6 about twice as large as those using the 50 m radius method. This was expected as the unlimited
7 radius counts had a greater area from which to record birds than the 50 m radius counts. Bird
8 diversity measures varied consistently across habitat types, except for the unlimited radius
9 abundance (which had higher values for the latter three habitats; Fig. 1). When comparing habitat
10 types, the 50 m radius data were more discriminating than the unlimited radius data. This is due
11 to the potential for variation in habitat beyond 50 m (e.g., an upland forest that is small or narrow
12 may transition to a different habitat within 50 m). Thus, bird monitoring programs generally use
13 count circles with some distance limits (Ralph et al., 1995).

14 These results help evaluate design, output, and end use validation for indicators
15 (Bockstaller and Girardin, 2003). For design validation, the bird diversity measures used in this
16 study are scientifically founded, are representative of bird populations, and are consistent across
17 different habitat types. In terms of output validation, these measures were assessed at the farm
18 scale, whereas many monitoring programs (e.g., BBS) use measures for larger regional or
19 ecosystem scales. Since many programs to encourage landowners to adopt biodiversity-friendly
20 practices are delivered at the farm scale, it is appropriate that these programs also be evaluated
21 through monitoring at the farm scale. Finally, for end use validation, diversity measures should
22 be understandable and relevant to users. Among the diversity measures used, species richness is

1 readily understood and most likely to be seen as a unit of practical management for legislators,
2 policy-makers, and landowners (Gaston and Spicer, 2004). Given that diversity measures are
3 consistent across habitats and are highly correlated among themselves (Gaston, 1996), it makes
4 sense to use the simplest measure.

5 Based on the measures used, wetland/riparian habitats had the highest species diversity,
6 followed by homestead, upland forest, native prairie, tame pasture, and cultivated cropland
7 habitats. This is consistent with other studies (e.g., Mosley et al., 2006; Ribic et al., 2009), and is
8 explained by higher productivity in riparian/wetland habitats and intensive agricultural
9 management (Herzon et al., 2008) during the breeding season in managed habitats such as
10 cultivated land and tame pasture. Homestead habitats had greater species diversity than expected
11 because of the potential for habitat diversity and supplemental feeding.

12 Overall, these indicators can be used, for example, to identify priority habitats for nature
13 conservation objectives, demonstrate differences among farms or habitat types in the common
14 ecoregions of east-central Alberta, or to show anthropogenic effects on target species. Future
15 research should explore how bird trends at the farm scale correlate with trends at different spatial
16 scales (e.g., provincial ecosystem types) and with different survey methods (e.g., BBS). As well,
17 research should explore how well bird diversity measures represent the diversity of other
18 taxonomic groups and with various natural processes (Moonen and Barberi, 2008; van Strien et
19 al., 2009). Last, research should evaluate the value of incorporating other environmental
20 parameters (eg. slope, aspect, habitat area, and current and historical land use), into a monitoring
21 program.

22

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