*Multi-seasonal species distribution models better facilitate habitat conservation for a migratory bird*

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**Abstract**

Species distribution models have issues with cross-seasonal transferability when data collected during a single season do not reflect habitat relationships across other seasons. This issue can be addressed using spatial decision support systems, which allow users to incorporate multiple season-specific distribution models into a single tool to prioritize habitat management. We demonstrated a potential application of this framework through an analysis of multi-season habitat use by American woodcock in Pennsylvania, USA. We modeled woodcock breeding and migratory habitat distributions in Pennsylvania using random forest classifiers and integrated the predictions of both models into a single decision support system using a Shiny application. The Shiny application accepts user input through breeding and migratory season weights, allowing the user to customize the tool based on area-specific management priorities. Multi-season distribution models like this one allow us to overcome lack of cross-seasonal transferability by incorporating multiple season-specific species distribution models into a single management prioritization framework. This framework is best suited for taxa with distinctly different habitat associations throughout the full annual cycle, such as migratory birds.

**1 Introduction**

Species distribution models are frequently used to assist conservation decision-making (Miller, 2010), however, they are known to have issues with transferability. For example, models developed in one area may not be reflective of animal distributions in other parts of their range (spatial transferability; Randin et al., 2006), or may fail to project species distributions into the future due to changing conditions (temporal transferability; Dobrowski et al., 2011). We posit that an additional subcategory of spatiotemporal transferability exists, called cross-seasonal transferability, for situations where species habitat associations differ among seasons or life stages, resulting in the species using fundamentally different space throughout the year. For example, wildlife science has a long history of bias towards studying animals during the breeding season, which may neglect the value of non-breeding habitat for survival and ignore carry-over effects into the breeding season (Norris and Marra, 2007). By building species distribution models which focus solely on occurrence data collected during breeding, we may disregard portions of a species’ distribution that are essential to persistence.

Spatial decision support systems (SDSS; Hopkins and Armstrong, 1985)) may provide a useful mechanism to circumvent issues of cross-seasonal transferability by combining distribution models from multiple seasons of the full annual cycle during the decision-making process. SDSS utilize user-friendly, interactive toolsets to guide users through making a set of spatial prioritization decisions (Sugumaran and Degroote, 2010). SDSS frequently come as extensions of existing geographic information systems (McConnell and Burger, 2011), but the learning curve and costs associated with professional geographic information systems can often be an impediment to reaching the intended user base (Harper, 2006). The widespread adoption of interactive online mapping tools, such as leaflet (Agafonkin, 2022) and ArcGIS Online (ESRI, 2023), has greatly expanded the capacity to custom build SDSS that are accessible via a web browser and can be easily used by decision makers with little additional training (Sugumaran and Sugumaran, 2007). SDSS provide an interface which allows users to interact with multiple spatial data layers, such as species distribution models. In circumstances where species distribution models have low cross-seasonal transferability, SDSS can also be used to compensate by incorporating multiple season-specific species distribution models into the decision-making process.

Migratory birds are one group of organisms that are clearly sensitive to issues of cross-seasonal transferability through use different geographic areas throughout their annual cycle that can be coarsely divided into breeding, wintering, and migratory seasons (Marra et al., 2015). Resource requirements frequently differ among these three seasons, often resulting in bird use of fundamentally different habitat types (Allen et al., 2020; Rice et al., 1980; Stanley et al., 2021). However, there are circumstances in which breeding, wintering, and migratory habitat may occur in the same region, especially for short-distance migrants which travel less than 2000 km during their migration (cite tk). Short-distance migrants include wintering waterfowl, such as sea duck species tk, where more northern populations overwinter in the same regions that more southern populations breed, and nomadic finch species such as tk, which migrate, breed, and overwinter in the same regions despite differential resource requirements among seasons (cite tk). Managing for these species, which occur in the same areas but have issues of cross-seasonal transferability in their models, could be done more effectively using an SDSS approach which combines species distribution models from different seasons into a single predictive layer using user-specified weights. For example, a user might want to primarily conserve breeding habitat for a species and conserve migratory habitat as a secondary objective. Using an SDSS, the user could weight a breeding season species distribution model to 70% and a migratory season species distribution model to 30% and output a single predictive habitat layer that prioritized management areas according to the user’s seasonal management objectives.

We demonstrate a SDSS framework to spatially-prioritize habitat management while accommodating the cross-seasonal transferability necessary to capture multiple seasons of a migratory bird’s full annual cycle. Our case study is focused on American woodcock (*Scolopax minor*;hereinafter woodcock) in the state of Pennsylvania, USA. Woodcock are short distance migrants with considerable overlap among migratory, breeding, and wintering ranges (Myatt and Krementz, 2007; Fig. 1), but fundamentally different habitat requirements among seasons (Allen et al., 2020). Our goal was to develop a SSDS tool to aid conservation practitioners considering trade-offs between managing for breeding and migratory season habitat in Pennsylvania. We used a multi-season modeling framework to predict the distribution of migratory and breeding habitats, which we combine in a SSDS to facilitate habitat prioritization.



Figure 1. Seasonal ranges and hypothetical migration routes of American woodcock in eastern North America. Seasonal ranges were delineated by eBird’s Status and Trends project (Fink et al., 2022) using citizen science data. Migration routes illustrate potential connections among eastern (dashed line), central (solid line), and western (dotted line) population segments. Migration routes were originally proposed by Glasgow (1958) and later reproduced by Moore et al. (2019). Inset illustrates multiple migration routes intersecting with the breeding range in the state of Pennsylvania.

**2 Methods**

*2.1 Study area*

We modeled woodcock habitat suitability throughout the state of Pennsylvania, which provides breeding habitat for an estimated 2.3% of the global woodcock population (52,400 birds), and potentially provides stopover habitat for nearly 1/3rd of the global woodcock population breeding throughout the northeastern United States and eastern Canada (30.5% of global woodcock, 684,500 birds; Kelley et al., 2008). Pennsylvania also contains some wintering habitat for woodcock in the southwestern portion of the state, but the amount of wintering habitat in Pennsylvania is negligible compared to breeding and stopover habitat (Fig. 1). Pennsylvania is composed of 11 EPA level 3 ecoregions (Omernik and Griffith, 2014), which reflect the topological gradient from Pennsylvania’s ridge-and-valleys in the central portion of the state to the coastal lowlands along the edge of Lake Erie. Lowland areas in Pennsylvania tend to be heavily agricultural, with development primarily concentrated around the two largest cities, Philadelphia (pop. Tk) and Pittsburgh (pop. Tk). Mountainous areas, such as the North Central Appalachians ecoregion, remain in mostly contiguous forest cover (percent agricultural tk).

Woodcock in Pennsylvania are managed by the Pennsylvania Game Commission, which manages hunting and provides habitat for wildlife throughout the state. The Pennsylvania Game Commission owns more than 600,000 hectares of land in Pennsylvania, referred to as state gamelands, which are managed primarily for wildlife and to provide hunting and trapping opportunities for the public (cite tk). Managing woodcock habitat for both breeding and migratory seasons have been identified as priorities by the Pennsylvania Game Commission, and the commission has expressed interest in tools allowing for ranked prioritization of woodcock seasonal habitat on state gamelands. We designed the SDSS tool to aid the Pennsylvania Game Commission with woodcock management prioritization of state gamelands and allow consideration of habitat needs across multiple seasons during the commission’s management activities.

*2.2 Breeding and migratory season data*

To model woodcock habitat suitability during the breeding and migratory seasons, we used separate data sources that described woodcock occupancy during each of those time periods. For the breeding season (March–May), we used survey data collected as part of the federally-coordinated American Woodcock Singing-Ground Survey (Seamans and Rau, 2020) and additional state-level monitoring conducted by the Pennsylvania Game Commission. Both state and federal surveys consisted of 5.76 km routes with 10 evenly spaced points, where observers listened for woodcock calls during their crepuscular breeding display. Observers recorded counts of all males calling during 2-minute intervals shortly after dusk. We used survey data collected from 2016–2020, and distilled records to presence or likely absence at each point based on detection of at least one male during the 5-year period.

“Male woodcock are often assumed to display near female nesting habitat, and so we felt good about using male presence as an indicator of male and female presence at the scale of application (cite tk)”

We delineated woodcock occurrence during the migratory season using GPS-tracking data from the Eastern Woodcock Migration Research Cooperative, a collaboration of 42 federal, state, provincial, non-profit, and university partners throughout the United States and Canada (www.woodcockmigration.org). We captured woodcock at 34 sites in Quebec, Ontario, Nova Scotia, Maine, Vermont, New York, Rhode Island, Pennsylvania, Maryland, West Virginia, Virginia, North Carolina, South Carolina, Georgia, Alabama, and Florida using mist nets during mornings and evening flights (Sheldon, 1960), or using spotlights and dip nets at night (McAuley et al., 1993; Rieffenberger and Kletzly, 1966). We attached 4g, 5g, or 6.3g PinPoint GPS Argos transmitters (Lotek Wireless Inc., Newmarket, Ontario, CA) to captured woodcock. Transmitters recorded GPS locations at 12–60m accuracy and were programmed to record diurnal locations every 1–3 days. Transmitters, bands, and attachment materials never exceeded 4% of a bird’s body weight, and all capture and handling were conducted with methods approved by the University of Maine Institutional Animal Care and Use Committee (Protocol # A2020-07-01).

We used woodcock location data to identify stopover locations, defined as any place where a migrant bird can land and survive until the next migratory flight (Mehlman et al., 2005). We consider woodcock to be migrating after they have made their first >16.1 km movement in fall or spring, and to complete their migration after they have made their last >16.1 km movement in the respective season. The >16.1 km threshold is based on tk (cite tk). Because woodcock migrate at night, we considered all diurnal locations between migratory initiation and termination to be stopovers. We grouped successive locations within 3 km into a single stopover, based on our observations that movements <3 km tended to be recursive rather than directional, and removed all but one location from each stopover from the analysis to reduce pseudoreplication and spatial autocorrelation of closely clustered locations. We also generated 10,000 locations randomly distributed throughout Pennsylvania, which we considered pseudoabsence locations.

*2.3 Species distribution modeling*

We constructed separate species distribution models for migratory and breeding seasons to accommodate differences in habitat associations and data sources. Each model used explanatory variables with presumed relevance to woodcock habitat associations (McAuley et al., 2020), with suites of variables including land use/land cover, forest successional class, topography, region, and soil moisture (Table 1). We additionally calculated landscape metrics from the landscapemetrics package (Hesselbarth et al., 2019) in program R (R Core Team, 2022), which represented landscape composition and configuration. To generate composition metrics, we resampled the National Land Cover Dataset to a 90m resolution, and then calculated the percent of each cover type within 500m, 1km, 5km, and 10km radii for each pixel. For configuration metrics, we used the National Land Cover Dataset to create a binary forest/non-forest layer, which we resampled to 90m resolution, and then calculated each configuration metric within 500m, 1km, 5km, and 10km radii of each pixel.

We conducted a pilot evaluation of several potential modeling techniques fit to a subset of woodcock occurrence data, including using MaxEnt (Phillips et al., 2006), random forest classification (Breiman, 2001), classification and regression trees (Breiman et al., 1984), and neural networks (Hopfield, 1982). All models were fit using the R package SDMtune (Vignali et al., 2020). We compared model outputs using area-under-the-curve (AUC), a common metric of predictive accuracy for classification models (Fielding and Bell, 1997). The random forest classifier had the highest AUC among modeling approaches, and we therefore used random forest techniques for all subsequent models.

For the breeding season model, we used a random forest classifier designed for clustered data (Wang and Chen, 2016), and applied survey route as a clustering variable to compensate for spatial autocorrelation among points on the same survey route. As federal survey routes were randomly distributed (Clark, 1970) and state surveys were distributed opportunistically, we included survey type (state vs federal) as an explanatory variable to accommodate differences in route distribution. For the migratory season, we used a traditional random forest classification model, written using the randomForest package in R (Liaw and Wiener, 2002). We assessed the accuracy of all models using a k-fold cross validation approach, where separate training and testing datasets were randomly sampled for each fold. We sampled folds at a survey route level for the breeding season model to accommodate autocorrelation within survey routes and prevent data leakage between the training and testing datasets. We used 10 folds for the breeding season model (90% training, 10% testing), but only 5 folds for the migratory season model (80% training, 20% testing) to accommodate the smaller sample size of the stopover dataset. We averaged AUCs for each of the folds to produce a mean AUC for each model and created predictive layers at 90m resolution that averaged predictions across folds.

To avoid overwhelming final predictive models with highly correlated or uninformative variables, we used the R package VSURF (Genuer et al., 2022) to implement a three-step backwards variable-selection approach, where each step produced a more parsimonious model. The first step eliminated irrelevant variables with lower variable importance than a defined threshold value (determined based on guidelines in Genuer et al., 2015). The second step retained only variables with the smallest out-of-bag error rates when training the model, effectively eliminating variables with some relevance but not critical for prediction. The third step used a stepwise process to test whether each included variable led to an out-of-bag error decrease that was larger than a defined threshold value, effectively removing redundant variables from consideration (Genuer et al., 2015). We compared predictive accuracy of models created from each step using AUC, and retained the model from the step that produced the highest AUC to create a final predictive layer for each season.

Table 1. Explanatory variables used to model woodcock distributions in Pennsylvania, USA. Suites indicate conceptual grouping of variables into classes relevant to woodcock occurrence.

|  |  |  |
| --- | --- | --- |
| **Suite** | **Covariate** | **Source** |
| Land use/land cover | Land use/land cover | National Land Cover Dataset (Jin et al., 2019) |
| Forest successional class | Forest successional class | LANDFIRE (USGS and USDA, 2020) |
| Topography | Elevation | USGS (2000) |
| Slope | Derived from elevation |
| Region | EPA level 3 ecoregions | Omernik and Griffith (2014) |
| Soil moisture | Soil drainage | Web soil survey (NRCS, 2021) |
| Topographic wetness index | Derived from elevation (Fink, 2013) |
| Landscape composition  (0.5, 1, 5, and 10km scales) | % Forest | Derived from National Land Cover Dataset using landscapemetrics (Hesselbarth et al., 2019) |
| % Agricultural |
| % Developed |
| Landscape configuration  (0.5, 1, 5, and 10km scales) | Aggregation index | Derived from National Land Cover Dataset using landscapemetrics (Hesselbarth et al., 2019) |
| Cohesion |
| Edge density |

*2.4 Analysis of covariate relationships and spatial distribution of seasonal habitat*

Random forest techniques do not provide easily interpretable covariate relationships, leaving the user to determine how covariates might influence the outcomes of the model (Breiman, 2001). While we were not interested in exploring woodcock-habitat relationships per se, we sought to understand how environmental variables contributed to model predictions. We also sought to highlight regional differences in the distribution of breeding and migratory habitat among ecoregions. To depict these differences, we sampled covariate values, ecoregion type, and model-predicted suitability at 10,000 randomly distributed points throughout Pennsylvania. We used hex plots to visualize trends between covariates and predictions for each season, and visualized variation among each EPA level 3 ecoregion in Pennsylvania using box-and-whisker plots.

*2.5 Spatial Decision Support System*

We created a SDSS in the Shiny ecosystem (Chang et al., 2022), named the Woodcock Priority Area Siting Tool (W-PAST), to facilitate user choice in the importance of breeding and migratory season habitat to local woodcock management planning. The SDSS allowed users to assign weights to each seasonal habitat suitability layer in 10% increments (ex. 20% migratory and 80% breeding season), which combined seasonal predictions into a single multi-season layer (Fig. 2). The weighting was conducted on a pixel-by-pixel basis as a simple weighted average where *pw* indicates the value of the weighted pixel value, *wm* the weight of importance for migratory habitat, *wb* the breeding season weight, *pm* the migratory pixel value, and *pb* the breeding season pixel value.

Practitioners often benefit from SSDS features customized to their management applications. In the case of the Pennsylvania Game Commission, a state wildlife management agency, a primary goal was to increase availability of woodcock habitat on state-managed gamelands, requiring functionality within the tool to compare habitat suitability among gamelands. We built four comparison metrics into the SDSS that were calculated using the weighted averages of the breeding and migratory season predictive layers: average pixel value, total habitat, % high quality, and % medium quality. Average pixel value was the arithmetic mean of all pixels within a state gameland, which tended to favor small gamelands predominantly composed of woodcock habitat and was intended to demonstrate where a small amount of habitat management could increase local woodcock populations. Total habitat was average pixel value multiplied by the acreage of the gameland, which favored larger gamelands that contained relatively large amounts of woodcock habitat in aggregate by virtue of their size. Total habitat could be used to determine which gamelands would provide the most habitat in aggregate if they were managed for woodcock. Percent high quality habitat was the percentage of cells within a gameland that had values greater than the 33rd percentile of all pixel values in the state, and percent medium quality was the percentage of cells falling between the 66th and 33rd percentile. These percentile-based metrics allowed users to quantify the proportion of a gameland which might be suitable for woodcock management. By multiplying the percent high or medium quality by the gameland acreage (also provided in the tool), the user could also derive the acreage in each gameland that could be managed for woodcock effectively. Further instructions for using these metrics in management are included in a user manual, publicly available with the SDSS at www.woodcock.shinyapps.io/W-PAST.



Figure 2. Conceptual diagram of user decision options in the Woodcock Priority Area Siting Tool (W-PAST). Users can choose the weighting of migratory and breeding season habitat based on management priorities. The resulting weights are used to generate a statewide predictive layer and gameland prioritization metrics, which allow the user to compare the suitability of gamelands for woodcock management.

**3 Results**

We collected data from tk migrant woodcock marked with GPS transmitters from fall 2017 to spring 2021 throughout the eastern portion of the woodcock’s range. Eighty-two individuals (tk%) recorded GPS locations at 113 stopovers in Pennsylvania. Breeding season survey data were available for 770 locations along 77 federal American Woodcock Singing-Ground Survey routes and 800 locations along 80 Pennsylvania Game Commission state survey routes. The most predictive breeding season model (AUC = 0.83) was the result of the second variable selection step, in which all variables with low predictive capacity were removed. This model was heavily informed by landscape variables at 5 and 10 km scales (Table 2), and no variables at the finest spatial scale (0.5km) or in the suite of soil moisture characteristics were included in the most informative model. Graphs of habitat suitability for each covariate showed strong, non-linear relationships (Fig. 3). Suitability was highest for landscapes with 0–25% developed land area, 0–50% agricultural land area, and aggregation index values of 80–100, all at the 10km scale. At the 5km scale, the breeding season model also showed high suitability in landscapes with 30–100% forest cover (Fig. 3).

The most predictive migratory model (AUC=0.78) was the full model, including all landscape, land cover, geographic, and soil moisture covariates (Table 2). Likely due to the wide array of covariates influencing the model, individual covariate graphs do not show clear visual patterns between migratory habitat suitability and any one covariate. However, the migratory model illustrated greater tolerance of migrant woodcock for developed and dis-aggregated landscapes at a 10km scale than the breeding season model (Fig. 3). The two models were also distinguished by the scale at which covariates influenced habitat suitability; the most informative breeding season model was not influenced by any landscape covariates at the 500m scale, and only 1 landscape covariate at the 1km scale, whereas the most informative migratory model included all available small-scale landscape covariates. Because of the influence of covariates at 500m and 1km scales, the migratory model predicted much more spatial variation in habitat distribution than the breeding season model, despite identical pixel resolutions (Fig. 4).

Breeding season habitat was not evenly distributed among ecoregions (Fig. 5A), with mean habitat suitability values ranging from 22.9–86.0%. Migratory habitat was more evenly distributed among ecoregions, with mean habitat suitability values ranging from 46.5–87.5%. Most of the difference between the distribution of migratory and breeding season habitat was in the Northern Piedmont, Middle Atlantic Coastal Plain, and Central Appalachians ecoregions, which had mean breeding season habitat suitability values of < 30% and mean migratory season habitat suitability values of >60% (Fig. 5B).

Table 2. Variables selected via backwards variable selection using the R package VSURF (Genuer et al., 2022) for the migratory and breeding season models. The migratory model employs the full set of variables, while the breeding season model uses a subset of variables inclined towards coarse resolution landscape variables.

|  |  |  |
| --- | --- | --- |
| Suite | Migratory | Breeding |
| Landscape (500m) | Aggregation Index, Cohesion, Edge Density, % Forest, % Agricultural, % Developed |  |
| Landscape (1km) | Aggregation Index, Cohesion, Edge Density, % Forest, % Agricultural, % Developed | % Agricultural |
| Landscape (5km) | Aggregation Index, Cohesion, Edge Density, % Forest, % Agricultural, % Developed | Cohesion, % Forest, % Agricultural, % Developed |
| Landscape (10km) | Aggregation Index, Cohesion, Edge Density, % Forest, % Agricultural, % Developed | Aggregation Index, Cohesion, % Agricultural, % Developed |
| Land Cover | Forest, Successional Class |  |
| Geography | Elevation, Slope, Ecoregions | Elevation, Ecoregions |
| Soil Moisture | Drainage, Topographic Wetness Index |  |

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Figure 3. Comparison of relationships between landscape variables and habitat suitability for breeding and migratory season models of American woodcock in Pennsylvania, USA. During the breeding season, woodcock habitat suitability is highest in highly aggregated landscapes with ~75% forest and ~25% agricultural cover. During the migratory season, however, woodcock become far more tolerant of landscapes that are unsuitable during the breeding season, including landscapes with higher proportions of developed cover. Habitat suitability is displayed on a percentile scale, indicating whether a certain pixel was more suitable for woodcock occupancy than the corresponding percentage of other pixels in the state.

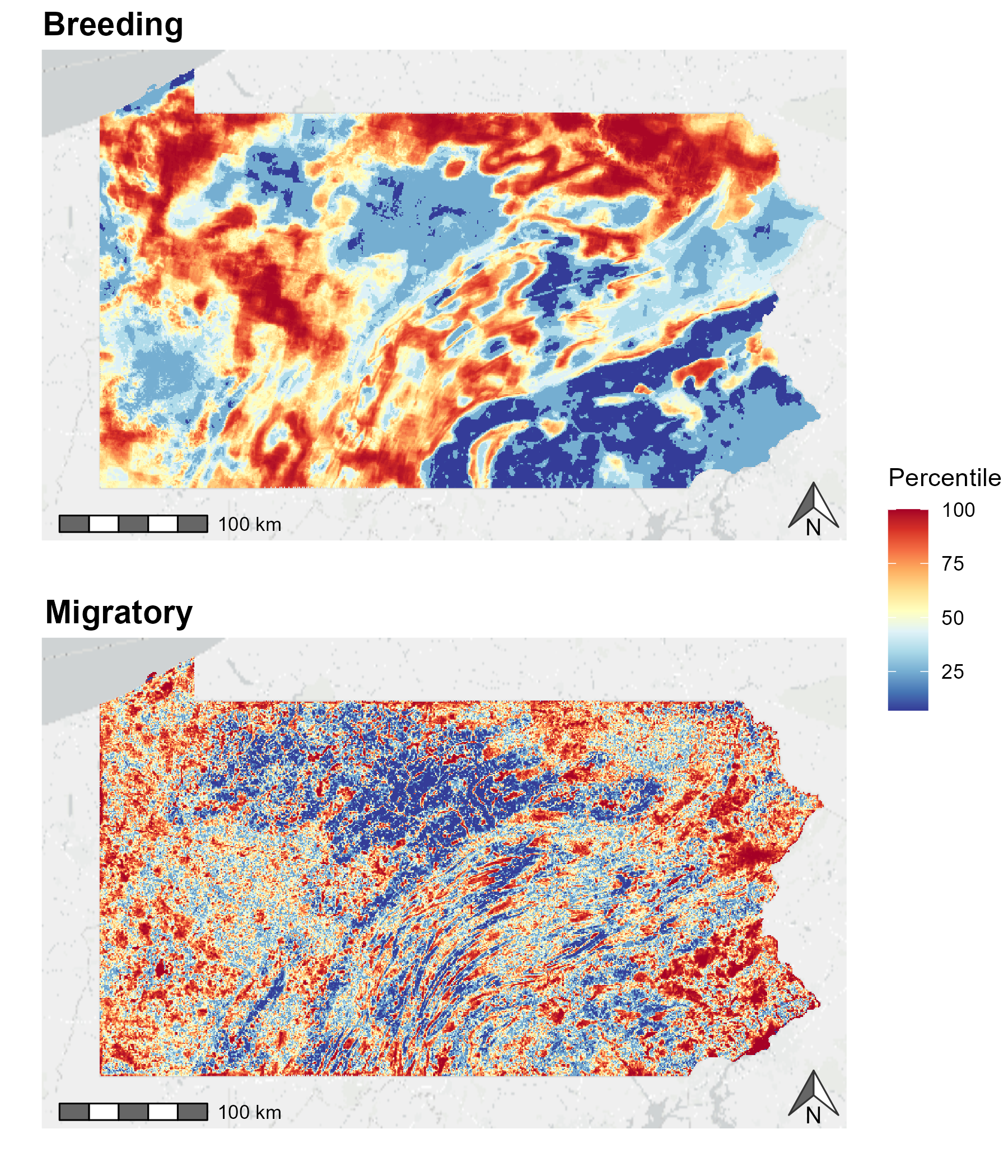


Figure 4. Breeding and migratory predictive habitat suitability layers suggest that woodcock select habitat at different scales in different seasons. Certain areas which are not appropriate for breeding season habitat management, such as southeastern Pennsylvania, may be appropriate for migratory habitat management. Percentile indicates whether a certain pixel was more suitable for woodcock occupancy than the corresponding percentage of other pixels in the state.



Figure 5. Breeding and migratory season habitat suitability for woodcock (5A) by EPA level 3 ecoregion (5B) in Pennsylvania (Omernik and Griffith, 2014). Three ecoregions, Northern Piedmont, Middle Atlantic Coastal Plain, and Eastern Great Lakes Lowlands, had mean breeding season habitat suitability values of <30% and mean migratory season habitat suitability values of >60%. Habitat suitability is calculated based on randomly sampled locations within each ecoregion and uses a percentile scale, indicating whether a certain pixel was more suitable for woodcock occupancy than the corresponding percentage of other pixels in the state. Box plots, arranged in the same order as the figure legend, indicate the median and interquartile range while whiskers extend to the largest/smallest value within 1.5 times the interquartile range.

**4 Discussion**

We demonstrated an integration of multi-season species distribution models into a single SDSS tool. The SDSS emphasizes the importance of practitioner input in management prioritization by allowing user choice in the weighting of breeding and migratory habitat to meet local management objectives. This approach allows users to overcome issues with low cross-seasonal transferability in species distribution models and prioritize management in a way that conserves habitat across stages of the life cycle.

We also demonstrate that, due to low cross-seasonal transferability, a single-season modeling approach would not adequately conserve woodcock habitat during multiple seasons. The correlation value between breeding and migratory habitat in the model was TKTKTK, and several Pennsylvania ecoregions have abundant migratory habitat but little breeding habitat. The five gamelands with the most abundant breeding habitat ranked tk, tk, tk, tk, and tk for migratory habitat among all tk gamelands. This indicates that an approach that manages solely for woodcock breeding habitat is unlikely to result in substantial conservation of woodcock migratory habitat. The use of a SDSS allows users to consider how their management strategy influences birds throughout multiple stages of their life cycle and encourages informed decision-making regarding the importance of habitat during those stages.

Regional differences between the breeding and migratory models underscore the importance of multi-season distribution models in delineating regional priorities for migratory bird management. For the woodcock model, we found relatively low breeding season suitability within the Northern Piedmont, Middle Atlantic Coastal Plain, and the Eastern Great Lakes Lowlands ecoregions, despite high migratory suitability. User-weighted prioritization of seasonal habitat might allow managers in regions in which woodcock breeding habitat is scarce to instead prioritize migratory habitat management. On the other hand, a manager of an area that provides breeding habitat in a region where breeding habitat is scarce might decide that their most effective decision would be to prioritize breeding habitat as much as possible. We posit that there might be several effective management strategies based on the information provided in multi-season distribution models, and the incorporation of user-specified weights empowers users to consider multiple possible management decisions and customize the information provided to inform their management strategy.

We showed that American woodcock occurred in distinctly different habitat during the breeding and migratory seasons in Pennsylvania and were associated with different spatial scales between seasons. During the breeding season, woodcock habitat suitability was dependent primarily on covariates at 5 and 10 km scales, while during the migratory season habitat suitability was additionally dependent on covariates at 500m and 1 km scales. This pattern supports past observations that migratory birds select habitat at a finer scale during the migratory season (Stanley et al., 2021). Due to these differences in the scale, managers may need to adjust management to match the scale of the season of interest. For example, woodcock management for breeding season habitat in Pennsylvania might focus on conserving broad swaths of habitat on large public lands, such as Pennsylvania state gamelands. As the predictive layer is fairly uniform across even large state gamelands, performing habitat management at that scale would likely be effective. However, the migratory model had a much finer spatial resolution, and was much more prone to having small pockets of habitat in areas not traditionally targeted by wildlife management agencies, such as urban areas. Differences in the spatial scale of habitat associations between seasons demonstrate the necessity of modeling occupancy for each season separately, to ensure that management supports the habitat requirements of animals throughout the full annual cycle.

Multi-season distribution modeling may also highlight areas of potential for conservation that are not traditionally managed for wildlife habitat due to a lack of breeding season occupancy. Woodcock were more tolerant of developed land cover during the migratory season than the breeding season, and the migratory season model predicted use of highly developed areas such as suburban Philadelphia and Pittsburg. This corresponds with findings of Buler and Dawson (2014), who found that migratory birds heavily used urban greenspaces during stopover, presumably due to attraction to high levels of artificial light at night (McLaren et al., 2018) and lack of other stopover options. One implication is that, in addition to management for woodcock at smaller spatial scales, practitioners may need to consider management of urban greenspaces for migratory birds. Opportunities for urban habitat conservation might come through partnerships with public and private landowners, such as park authorities and utility companies, to conserve migratory habitat in urban greenspaces. Another opportunity for urban habitat conservation might be the Urban National Wildlife Refuge program (USFWS, 2023), which has dual roles in preserving wildlife habitat and expanding access to natural areas for historically excluded communities. Pennsylvania is host to one Urban National Wildlife Refuge, John Heinz National Wildlife Refuge at Tinicum, located in the Philadelphia suburbs. Our model predicted high migratory habitat suitability for woodcock within this refuge, demonstrating how urban wildlife refuges may provide crucial stopover habitat in heavily urbanized areas.

A multi-season distribution model framework is particularly well suited to migratory bird management due to its flexibility in application of multiple data sources, which is particularly useful for species that are studied using separate techniques and surveys during each season. While there are several surveys for examining bird distribution during the breeding and wintering seasons (e.g. Bonter and Greig, 2021; Robbins et al., 1986), examining bird habitat use during the migratory period continues to be a challenge. Individually-marked birds with GPS transmitters are the gold standard for this type of analysis, as stopover locations can be separated from breeding and wintering locations for each tagged bird. However, GPS transmitters are still too large to attach to many small migratory birds, and the low number of stopovers attained per individual (mean = 1.4, sd = 0.6 in this study) combined with the considerable price of these transmitters may make attaining a large sample size a financial difficulty for most study species. The use of citizen science data collected during migration, such as the eBird data collection platform (Sullivan et al., 2009), may provide a more generalizable way to collect stopover location data, but certain assumptions must be made to distinguish true migratory locations from early breeding/wintering season arrivals. Decisions on seasonal management priorities can also be informed by other data sources and models, such as multi-season survival models to determine whether breeding or migratory habitat has a greater role in limiting survival or migratory corridor models to identify high densities of migrants (Cohen et al., 2022). SDSS provide a framework for blending these multi-season datasets and models to improve management and conservation decision making for migratory birds.

**CRediT authorship contribution statement**

LA Berigan, AM Roth, LM Williams, KR Duren, S Bearer, K Wenner, P Kasper, and EJ Blomberg contributed to the conceptualization of this study and design of the methodology. KR Duren acquired funding for the creation of W-PAST, while EJ Blomberg and LM Williams acquired funding for the expansion of the EWMRC into Pennsylvania. LA Berigan and AC Fish curated the GPS data and LM Williams curated the survey data used in this study. LA Berigan conducted the formal analysis and built the W-PAST tool. LA Berigan, AM Roth, and EJ Blomberg wrote the original draft of the manuscript, and all authors contributed to revisions and editing.

**Declaration of competing interest**

The authors have no conflicts of interest to declare.

**Data availability**

The code used in this analysis is publicly available at github.com/EWMRC/AMWO-seasonal-weighted-SDM. Data are available from the corresponding author upon reasonable request.

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