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Abstract: Whether considering the cumulative impact of chronic, smallsize oil discharges or accidents associated with marine traffic and offshore oil and gas development, seabirds face a variety of threats in the marine environment. Assessing the vulnerabilities of seabirds to maritime hazards requires an understanding of their species distribution, and a means for combining this information across groups. Using at-sea survey data gathered as part of a regional monitoring program, an efficient framework for integrating multi-species data was developed. Density estimates, derived from counts of species known to be particularly vulnerable to oil pollution, were used to construct multiple species distribution models (SDMs). The structural difficulties associated with sampling sparsely distributed individuals that can occur in large, localized concentrations led to the use of three modelling techniques potentially well suited for this type of data: negative binomial, "hurdle", and random forest methods. Predicted abundances were combined to produce an ensemble forecast, and multi-species potential vulnerability maps developed to identify core areas. The maps confirmed the general importance of the shelf break and sea banks, but distribution was highly seasonally specific. A pattern of shifting presence emerged, with some species disappearing from the region (e.g., storm-petrels and shearwaters in winter) and others increasing at the same time (e.g., Black-legged Kittiwakes, Rissa tridactyla). Across all species, spring (March - April) stood out as a period of peak importance, though seasonal variation in the usage of the offshore marine environment may render the seabird community vulnerable to different threats throughout the year. This study illustrates how multi-species distribution modelling can predict seasonal usage patterns, and contribute to an awareness of the collective vulnerability of groups of seabirds to hazards distributed over wide geographic areas.

*Cover Letter

August 7, 2013

Dr. David Lieske Department of Geography and Environment Mount Allison University 144 Main Street Sackville, New Brunswick, Canada E4L 3G1

To the Editor(s),

On behalf of my colleagues and myself, please find attached the manuscript entitled "Maps, Models, and Marine Vulnerability: Assessing the Community Distribution of Seabirds at-Sea" for consideration in *Biological Conservation*.

This paper is an original application of ensemble modelling to assess the distribution of a community of seabirds that occur in Maritime Canada. As this is an area subject to significant marine traffic, chronic small-scale oil pollution, and ongoing oil and gas extraction and exploration, this group of seabirds (Black-legged Kittiwakes, Dovekie, murres, Northern Gannet and storm-petrels) constitute a vulnerable community. By combining the predictions from three modelling methods (negative binomial GLM, hurdle GLM, and random forest machine learning) to produce an ensemble prediction, we assessed the predictive accuracy of these techniques and demonstrate how multi-species vulnerability maps can be generated to identify core areas of conservation concern.

This work builds nicely on papers recently published in *Biological Conservation*, e.g., Oppel et al. (2012): Comparison of five modelling techniques to predict the spatial distribution and abundance of seabirds, Biol. Cons. 156, 91-104, and feel that the results will be of wide interest to its readership.

The manuscript is slightly less than 7500 words in length and has not been submitted for publication elsewhere.

Should you have any questions, please do not hesitate to contact me at the following address. I look forward to hearing from you in due course.

Sincerely,

David breske

David Lieske, Ph.D. Department of Geography and Environment, Mount Allison University, 144 Main Street, Sackville, New Brunswick, E4L 1A7, Canadaph: (506) 364-2315; fax: (506) 364-2625 dlieske@mta.ca

1 Highlights

- 2
- 3 Distribution models were constructed for a vulnerable group of seabirds.
- 4 Generalized linear, hurdle and random forest modelling methods were used.
- 5 Predicted seabird abundances were combined to form ensemble estimates.
- 6 Relative community importance for each regional $1^0 \ge 1^0$ unit was calculated.
- 7 Community vulnerability maps showed strong seasonal and spatial patterns.

Article Type: Research Article

Maps, Models, and Marine Vulnerability: Assessing the Community Distribution of Seabirds at-Sea

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Keywords: Species distribution modelling; multi-species distributions; ensemble prediction; seabirds; negative binomial GLM; hurdle, random forest; Maritime Canada

ABSTRACT

Whether considering the cumulative impact of chronic, small-size oil discharges or accidents associated with marine traffic and offshore oil and gas development, seabirds face a variety of threats in the marine environment. Assessing the vulnerabilities of seabirds to maritime hazards requires an understanding of their species distribution, and a means for combining this information across groups. Using at-sea survey data gathered as part of a regional monitoring program, an efficient framework for integrating multi-species data was developed. Density estimates, derived from counts of species known to be particularly vulnerable to oil pollution, were used to construct multiple species distribution models (SDMs). The structural difficulties associated with sampling sparsely distributed individuals that can occur in large, localized concentrations led to the use of three modelling techniques potentially well suited for this type of data: negative binomial, "hurdle", and random forest methods. Predicted abundances were combined to produce an ensemble forecast, and multi-species potential vulnerability maps developed to identify core areas. The maps confirmed the general importance of the shelf break and sea banks, but distribution was highly seasonally specific. A pattern of shifting presence emerged, with some species disappearing from the region (e.g., storm-petrels and shearwaters in winter) and others increasing at the same time (e.g., Black-legged Kittiwakes, Rissa tridactyla). Across all species, spring (March - April) stood out as a period of peak importance, though seasonal variation in the usage of the offshore marine environment may render the seabird community vulnerable to different threats throughout the year. This study illustrates how multi-species distribution modelling can predict seasonal usage patterns, and contribute to an awareness of the collective vulnerability of groups of seabirds to hazards distributed over wide geographic areas.

1. Introduction

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Successful conservation planning hinges on an ability to identify hazards, as well as an awareness of species vulnerability. In marine contexts, threats originate from a number of factors, including: small-scale, chronic oil discharges associated with maritime traffic (Wiese and Ryan, 2003); wide-scale accidental releases of oil (Henkel et al., 2012); fisheries bycatch (Tasker et al., 2000; Hamel et al., 2009); collisions with and loss of habitat associated with offshore wind farms (Exo et al. 2003); and negative interactions with offshore oil drilling platforms (Wiese et al., 2001). Heavier-bodied, diving seabirds are at greater risk of exposure to these hazards by virtue of the amount of time they spend interacting with the sea-surface-air interface (Camphuysen, 1998) but they are also vulnerable as a result of low fecundity (Votier et al., 2005). Mitigating anthropogenic risks first requires information about organisms' usage of the marine environment, in both time and space, so that priority areas can be identified and effective management strategies developed. Unfortunately, the very large area of ocean habitat potentially utilized by seabirds makes it logistically difficult to enumerate these species (Brown, 1980; McKinnon et al., 2009). Pelagic surveys rely heavily on moving platforms ('ships of opportunity') which may or may not be primarily tasked with gathering seabird information. Large gaps in knowledge inevitably occur, especially during the non-breeding season when, no longer constrained by central place foraging within range of their colonies, species are free to roam over much larger distances (Huettmann and Diamond, 2001; Oppel et al., 2012).

In terrestrial contexts, there are many successful applications of species distribution modelling (SDM, Guisan and Zimmermann, 2000; Franklin, 2009) to predict habitat usage in unsurveyed areas (Austin, 2002; Shriner et al., 2002). SDMs work by combining empirical patterns of occurrence with Geographic Information System (GIS)-derived information about environmental conditions, and are a valuable tool for conducting conservation planning (Jones-Farrand, 2011). Applying this methodology in marine contexts can be quite difficult, however. Marine environments are highly dynamic (Haney, 1989), and difficult to represent within a standard GIS. For instance, fronts, widely recognized as important areas of nutrient entrapment and increased biological productivity (Bakun, 1996), can shift position over a period of hours (Durazo et al., 1998). Species may respond by rapidly shifting their distributions to benefit from these changing patterns of food abundance. Additionally, seabirds simultaneously display sparse distributions over wide areas, as well as a tendency to form large localized aggregations in productive foraging areas (Briggs et al., 1987; Hunt and Schneider, 1987; Fauchald et al., 2002; Clarke et al., 2003). This produces data distributions with both a large number of non-observations (the so-called 'zero-inflation' problem) and occasionally large concentrations of highly variable abundance (the so-called 'overdispersion' problem). Take together, zero inflation and overdispersion pose serious problems for SDMs based on traditional techniques (Hilbe, 2008).

Over the past 20-years, there has been a notable proliferation of algorithms for modelling animal occurrence and abundance (Hegel et al., 2010). Such methods range from presence-only techniques (e.g., maximum entropy or MAXENT, Phillips et al., 2006; ecological niche factor analysis or ENFA, Hirzel et al., 2002) to presence-absence (logistic generalized linear or logistic generalized additive models), to abundance (Poisson or negative binomial regression). Machine learning methods, such as random forest (Breiman, 2001), can be applied in any of these cases. The very large number of available methods renders it difficult to choose a technique (Jones-Farrand et al., 2011), so is a decision often based on investigator preference or familiarity (Araújo and New, 2007; Jones-Farrand et al., 2011). Current research, however, points to the advantages of combining the predictions of different modelling methods, a process referred to as ensemble modelling (Araújo and New, 2007; Oppel et al., 2012) or decision fusion (Das et al., 2008). This approach allows models to reinforce each other when concensus exists, while dampening divergent predictions when they are discrepant. The implicit assumption is while the true underlying process is unknown, different forecasting models are able to capture different components of the underlying signal (Clemen, 1989). The advantages are expected to be maximized when different, but complementary, modelling algorithms are employed.

Marine SDMs are rare (Robinson et al., 2011), and SDMs based on shipboard surveys are even rarer (Oppel et al., 2012). Due to four years of pelagic surveys conducted as part of the east coast Eastern Canada Seabirds at Sea monitoring program, this study to was able to extend the approach developed by Oppel et al. (2012) and examine the atsea distribution of a range of seabirds that occur on or near the Scotian Shelf of Nova Scotia. Density data were corrected for variation in detectability, and three different approaches selected for their theoretical suitability to deal with zero-inflated, overdispersed data: negative binomial (NB) generalized linear modelling, Hurdle modelling, and random forest (RF) machine learning. Results from each of these methods were combined to produce an ensemble prediction, the accuracy of which were carefully assessed and compared. In addition to providing a case-example of how multi-method, multi-species models can be informatively combined, seasonally-specific community vulnerability maps were generated to assist regional conservation planning.

2. Methods

2.1. Data collection

The focal area was the Scotian Shelf with a maximum depth of about 200 m, extending 150-250 km offshore of the province of Nova Scotia, Canada. Also included were portions of the Bay of Fundy as well as the Cabot Strait (Figure 1). At the shelf break sea depth increases rapidly to abyssal levels exceeding 3000 m. The topography of the shelf,

combined with tidal forces, wind, and fluctuations of offshore currents associated with the Gulf Stream, result in upwelling and partial gyres capable of retaining nutrients and concentrating marine biomass (Smith and Petrie, 1982; Hannah et al., 2001; NSMN, 2010). A number of banks occur throughout the shelf, some of which (e.g., Browns Bank, Sable Island Bank, Georges Bank) experience tidally-produced gyres of sufficient strength to also retain nutrients and concentrate prey (Hannah et al., 2001)

2.2. At-sea survey data

The focal species of this study were: (1) Black-legged Kittiwake (*Rissa tridactyla*); (2) Dovekie (*Alle alle*); (3) murres (Thick-billed, *Uria lomvia*; Common Murres, *U. aalge;* unidentified *Uria* sp.); (4) Northern Gannet (*Morus bassanus*); (5) storm-petrels (Wilson's Storm-Petrel, *Oceanites oceanicus*; Leach's Storm-Petrel, *Oceanodroma leucorhoa*; unidentified sp.) and (6) shearwaters (Greater Shearwater, *Puffinus gravis*; Sooty Shearwater, *P. griseus*; Manx Shearwater, *P. puffinus*; Cory's Shearwater *Calonectris diomedea*; Audubon's Shearwater, *P. lherminieri*; unidentified sp.), also grouped to enhance sample size and maximize precision of the density estimates (see Section 2.3). As a group, these species exhibit high vulnerability to sea-surface oil pollution (Brown, 1980; Camphuysen, 1998), are a significant presence on the Scotian Shelf, and encompass a cross section of ecological niches: e.g., surface feeders (Black-legged Kittiwake, Leach's Storm-Petrel), deep-water pursuit divers (murres, Dovekie), plunge divers (Northern Gannet), and surface pursuit divers (shearwaters).

Observations from 76 surveys were gathered from March 1, 2006 to October 31, 2009 as part of the Eastern Canada Seabirds at Sea (ECSAS) program (Fifield et al., 2009). The ECSAS survey protocol (Gjerdrum et al., 2012) incorporates the recommendations of Tasker et al. (1984), as well as modern distance sampling techniques (Buckland et al., 2001), and relies on ships of opportunity to conduct the survey.

All birds were counted in a 90[°] arc to one side of the vessel, out to a maximum distance of 300 m. Birds on the water were counted and assigned to one of four distance classes (0-50 m, 51-100 m, 101-200 m, 201-300 m) on the basis of perpendicular distance from the vessel path, confirmed (when necessary) using a distance gauge. Flying birds were sampled using a "snapshot" methodology such that all flocks present within the 90[°] arc 300 m to one side and 300 m ahead of the ship were recorded. The application of the "snapshot" methodology was necessary to offset the tendency to over estimate the abundance of flying birds, which enter and re-enter the field of view (Tasker et al., 1984). Due to ongoing development of the survey protocol, there was some variation in the way in which distance was estimated to flying birds. From October to December, 2007, distances were not measured for flying birds. Between January and July, 2008, flying birds were assigned to distance classes based on perpendicular distance to the vessel path. After July, 2008,

distances to flying birds were measured radially from the vessel. As a result of this variation in sampling, there were really three different protocols applied to flying birds, necessitating three separate density estimates be calculated in program Distance 6 Release 2 (Thomas et al., 2010) and combined at the end of the analysis (Fifield et al., 2009).

Observations were only conducted when the ship was travelling at a minimum speed of 4 knots (7.4 km/h) and a maximum of 19 knots (35.2 km/h). When visibility was poor (due to rain or fog) the actual width of the visible transect (e.g., 200 m) was recorded.

2.3. Density estimation

Distance 6 Release 2 (Thomas et al., 2010) was used to estimate seabird densities, stratified by season and $1^0 \ge 1^0$ grid cell. We started with basic key functions (half-normal), and optional series expansion terms chosen from one of three families (cosine, hermite, or polynomial). Visual inspection and the χ^2 goodness-of-fit test were used to assess detection function model fit, and then an attempt was made to improve fit by either choosing a different key function (hazard rate or uniform with optional series expansion terms) or by including explanatory covariates (such as wind speed, sea state, wave height and/or observer) using the multi-covariate distance sampling engine (Marques et al., 2007). Analyses were conducted separately for each of the sampling regimes (birds on water vs. flying birds under three different distance measurement protocols, see Section 2.2), and yielded separate estimates of detection probability (\hat{p}), mean flock size (\bar{s}), seabird densities were corrected for the proportion of birds present but not observed and densities estimated (by program Distance) as: $\hat{D} = \hat{D}_s * \bar{s}$. Final densities were computed as the weighted average of the three different sampling regimes, based on kilometers surveyed during each survey method.

2.4. Environmental data

A wide range of oceanographic and biological variables have been implicated to explain seabird distribution (Table 1). Broadly speaking, previous studies have either focused on modelling distributional patterns or relating occurrence to oceanographic and bathymetric features or processes. For instance, Certain et al. (2007) used GAM models to produce trend surfaces of seabird occurrence purely as a function of latitude and longitude (although predictions were further refined using residual kriging) but no oceanographic information informed the resulting prediction surface. In contrast, Yen et al. (2004a) integrated factors such as sea-surface temperature (averaged over a 5-month period) and an index of herring spawning to produce predictive maps.

Given the prominence of the Scotian Shelf in the study region, as well as previous work documenting the influence of shelf-break and sea bank processes on seabird distribution in this area (e.g., Brown, 1988a; Brown 1988b; Huettmann and Diamond, 2001), regional predictive models were constructed with a focus on bathymetric properties (Table 2). Data from the ETOPO2v2c data set of 2-minute resolution distributed by the National Geophysical Data Centre (NGDC 2009) were integrated within a GIS and used to produce the variable DEPTH. The shelf break was identified using the 300 m isobath, and distances to this linear feature measured using the "Euclidean Distance" tool in ArcGIS 9.3 (ESRI 2008) as variable SHELFDIST. In keeping with earlier findings of the potential importance of sea floor "roughness" (e.g., Yen et al., 2004b) the coefficient of variation of sea depth (DEPTH CV) was also calculated. As with the shelf break, sea banks are important sources of nutrient upwelling and are well documented as primary spawning areas for many species of commercial fish (Ashmole, 1971; Brander and Hurley, 1992). To model the potential influence of sea banks on seabirds, 100 m isobath polygons were identified within the GIS, corresponding to the banks labeled in Figure 1. The proportion of each $1^{0} \times 1^{0}$ prediction cell intersecting with sea bank polygons was then calculated using the "Zonal Statistics" tool in ArcGIS 9.3 (ESRI, 2008), resulting in variable PBANK. Finally, preliminary analysis, as well as earlier atlas work (Lock et al., 1994), indicated that broad-scale distributional patterns were highly seasonal. To capture seasonal variation in abundances, variable SEASON was defined using the following categories: "spring" (March-April), "summer" (May-August), "fall" (September-October) and "winter" (November-February). In the case of negative binomial and hurdle generalized linear models (Section 2.5), fall densities served as the reference category for assessing seasonal effects.

2.5. Model construction

Species distribution models (SDMs) were constructed using the base library of the *R* Statistical Package (R Development Core Team 2009) as well as the MASS library of Venables and Ripley (2002). The glm.nb function (Venables and Ripley, 2002) was used to construct negative binomial, generalized linear models (NB GLMs). This approach uses a combination of maximum likelihood to estimate the overdispersion parameter (θ) and iterativelyreweighted least squares to estimate the model coefficients. Because of the overdispersion parameter, NB GLMs are more flexible than Poisson GLMs in dealing with extra count variance (Hilbe, 2008).

Hurdle NB models (referred to henceforth as 'Hurdle' models) were implemented using the *pscl* library of Zeileis et al. (2007). These two-component models apply a binomial GLM to the 'zero' (absence) vs. non-zero (y > 0) observations, and a NB GLM to the non-zero (y > 0) count data. In effect, an observation must be non-zero in order to pass over the "hurdle" (Zuur et al., 2009) and be modelled as a count-generating process. Predictions at any given location were

generated as the expected number of seabirds (via the NB GLM) times the fraction of the probability of a non-zero count binomial to a non-zero count negative binomial (Zuur et al., 2009). Final Hurdle models, therefore, involve estimation of two sets of coefficients: one for the binomial (the presence-absence part), and one for the NB GLM (the abundance part).

No variable selection algorithms were used to determine the covariates to retain/discard in the final NB GLM and Hurdle models. Instead, predictive models were constructed using an information-theoretic approach similar to that of Gray et al. (2010) and Lieske et al. (2012). An all-combinations algorithm, implemented in the *R* Statistical Package by D. Lieske (available upon request), computed predictive models for all possible-combinations and sorted them (lowest to highest) on the basis of Akaike Information Criterion (AIC) values. Only the top 95% of models were retained, and the weighted average of each coefficient was computed using the AIC weight, W_i (Burnham and Anderson, 2002). In the case of the hurdle algorithm, model averaged coefficients were computed for both the binomial and NB GLM components.

Random forests (RF, Breiman, 2001) constitutes a machine-learning alternative to NB and Hurdle methods, and supports its own internal ensemble learning algorithm to construct multiple tree-structured classifiers (Liaw and Wiener, 2002). Using the RF approach, novel predictions are obtained by "dropping" new inputs into the RF classifiers and then combining the "votes" to determine the most popular class or value (Breiman, 2001). The *randomForest* library of Liaw and Wiener (2002) was used to construct RF models. Library *randomForest* also provides information on variable importance, which is determined by how much prediction error increases when testing data for that variable is permuted while all others are left unchanged (Liaw and Wiener, 2002).

Ensemble predictions were generated by taking the arithmetic mean of the predictions from the NB GLM, Hurdle GLM, and RF models (the so-called "committee averaging" method, Araújo and New, 2007; Das et al., 2008). A recent comparative analysis performed by Marmion et al. (2009) indicated that other measures of calculating consensus (e.g., the use of median values) performed about the same or worse, which is consistent with the general findings reported in the ensemble forecasting literature (Clemen, 1989).

2.6. Model evaluation

Ten sets of randomly chosen training and testing data (90% and 10% of original data, respectively) were used to iteratively construct and test the predictions from NB, Hurdle, and RF models. Prediction error was assessed using root mean square error (RMSE):

$$RMSE = \sqrt{\frac{\sum_{i=1}^{n} (y_{obs} - y_{pred})^2}{n}}$$
(1)

2.7. Assessment of potential seabird vulnerability

The predictions from each of the species distribution models were combined into a single, seasonally-specific vulnerability map. Potential vulnerability (V) was quantified for each location in the prediction surface as the sum of the relative importances (R) for each of the k species:

$$V = \sum_{i=1}^{k} R_i \tag{2}$$

Relative importance was derived from quintile classification of the SDM abundance predictions, but with the break points determined using data pooled across all seasons. Adopting this approach ensured that seasonal shifts in seabird community usage patterns could be readily visualized and detected. It should be noted that while specific hazards were not introduced into equation 2 (e.g., offshore wind farms), they could easily be incorporated by applying species-specific weighting to reflect differing species sensitivity. Furthermore, when regional maps of the distribution and intensity of particular hazards are available, the vulnerability score could be multiplied by the hazard risk to produce a composite risk map.

3. Results

3.1. Species observations

Distance-sampling methods allowed for the estimation of seabird densities, by species group and season, at the level of $1^{0} \times 1^{0}$ survey blocks. Preliminary inspection of the resulting density distributions suggest significant offshore aggregations of Black-legged Kittiwake in the winter and spring seasons (Figure 2a), concentrations of Dovekie on the shelf and shelf break during the winter-spring period (Figure 2b), significant near-shore abundances in murres (Figure 2c) and Northern Gannet (Figure 2d) during the spring season, and noteworthy aggregations of storm-petrels (Figure 2e) and shearwaters (Figure 2f) throughout the region during the summer (see also Fifield et al., 2009). Storm-petrels are entirely absent from the study area during winter.

3.2. Model construction and evaluation

Species distribution models were constructed, for each species, using NB, Hurdle, and RF algorithms. In the case of NB and Hurdle models, an all-combinations approach was applied and standardized coefficients obtained using AIC-weighted model averaging (see Section 2.5). This process was repeated for 10 sets of training-testing data in order to generate 95% confidence intervals around the model-averaged coefficients obtained using the complete data set. Physical variables were then sorted from largest amplitude, negative-value to largest amplitude, positive-value coefficients, for both the NB and Hurdle algorithms (Figures 3 and 4, respectively). For RF models, variable importance was assessed using the importance values generated by the *randomForest* software (Figure 5).

Season was a key covariate for all species, regardless of modelling algorithm, though it varied in importance. Relative to fall occurrences, Black-legged Kittiwake, Dovekie, and murres showed strong winter responses under NB models (Figure 3). The same was true for Hurdle models (Figure 4), though uncertainty in the coefficients for the count-component tended to be higher. Storm-petrels and shearwaters showed strong summer responses under NB and Hurdle models, while densities of the Northern Gannet peaked in the fall. Depth (DEPTH) and distance to shelf break (SHELFDIST) were influential for some species, e.g., Black-legged Kittiwake, Dovekie, and Northern Gannet. DEPTH and season tended to be the most influential covariates in RF models (Figure 5), with SHELFDIST most important for Northern Gannet. Variation in seadepth (DEPTH_CV) also played an important role in storm-petrel and shearwater RF models.

No one algorithm (NB, Hurdle, RF) exhibited consistently superior performance over all species and iterations, but occasionally, NB and Hurdle algorithms failed to compute (Figure 6). RF models appeared better able to cope with sparse data and, in some cases, to also generate lower prediction error (e.g., Northern Gannet, Figure 6). More often, however, variation in performance meant that methods alternated in predictive power, working better in some iterations than others. Simple averaging of the NB, Hurdle and RF results produced ensemble predictions that either tracked the consensus or constituted a compromise value.

3.3. Assessment of species vulnerability

Vulnerability maps (Figure 7) represented the influence of season and bathymetric factors on the relative distribution of all species combined. Potential vulnerability for this group of species were highest in the spring, when large numbers of

seabirds were widely distributed throughout the Nova Scotian shelf and shelf break. This contrasts with the fall season,
when vulnerability was distributed along a narrower front concentrated primarily over the shelf break. There was a nearshore shift in vulnerability in the summer, presumably reflecting the central place foraging limitations imposed on
species such as the Northern Gannet. But it is important to note that this was counteracted by the presence of large
numbers of non-breeding species such as shearwaters, which would not be similarly range restricted. Potential
vulnerability was lowest during the winter season (Figures 7 and 8) when patchier distributions combined with the
absence of storm-petrels, for example, lowered the vulnerability scores.

4. Discussion

Conservation planning is a spatially-explicit exercise (Magness et al., 2011). The effective management of human activity in marine environments, e.g., through the designation of marine protected areas (MPAs), requires a synthesis of all available information (Araujo and New, 2007; Ronconi et al., 2012), and SDMs offer a powerful way to combine biological surveys with environmental information to better understand habitat usage through space and time. This study demonstrates an efficient framework for integrating multiple species data from an ongoing monitoring program across several implementation steps: GIS-based data management, state-of-the-art modelling, and production of a key spatial planning product. Each aspect of this framework will be discussed in turn.

Through a regular, grid-based approach ("Eulerian" data, see Tremblay et al., 2000) data gathered through opportunistic surveys was aggregated at 1^o x 1^o survey blocks to allow accurate density estimation. Factors which impact seabird detection, e.g., sea condition, or whether a bird was in the air or on the water, were statistically accounted for as covariates using distance sampling (Buckland et al., 2001; Marques et al., 2007). While unable to account for all the variability in distribution, season, sea depth and distance to shelf breaks were important and influential predictors. Proximity to the shelf break, for instance, stood out as an important factor influencing Black-legged Kittiwake numbers, while seafloor 'ruggedness' (DEPTH_CV) was associated with shearwaters and Northern Gannet. Adopting an aggregated, grid-based framework meant that seabird counts had to be pooled over the entire study period, but as fine-scale marine processes were not available contemporaneous for the entire study area, this was a sensible approach that permitted a focus on stable marine features (e.g., bathymetry). This approach did allow for seasonal summaries of abundance, which was important given that season was influential regardless of modelling method.

Despite an extensive history in other disciplines (e.g., macroeconomics and psychology, see Clemen, 1989), ensemble prediction/forecasting has only recently been applied in SDM (Oppel et al., 2012; Araujo and New, 2007). The focus of most SDM papers published in the first decade of the 21st century focused on comparing relative performance (e.g., Elith et al., 2006). As a consequence, the SDM literature is replete with an enormous array of single-method modelling

approaches where no attempt is made to gather the "consensus opinion" presented by a diversity of algorithms. While individual methods might be well calibrated and offer high predictive accuracy for the data at hand, there is no way to assess how realistic the predictions will be in unsampled locations, or under changing environmental conditions (Marmion et al., 2009). Jones-Farrand et al. (2011) argue that basing conservation decisions on the basis of predictions from a single model is "risky", in that problems are formulated under a specific set of objectives, constraints and assumptions that may not apply more generally. Ensemble approaches have an excellent track record for producing robust predictions, helping to reduce uncertainty and, in the process, increasing confidence in the decisions derived from them. In this study, no single algorithm (NB, Hurdle, RF) exhibited consistently superior performance over all species and iterations, though occasionally, NB and Hurdle algorithms failed altogether. RF models appeared better able to cope with sparse data and, in some cases, to also generate lower prediction error (e.g., Northern Gannet, Figure 6). In general, variation in performance mean that methods alternated in predictive power, working better in some iterations than others. Simple averaging of the NB, Hurdle and RF results produced ensemble predictions that either tracked the consensus or constituted a compromise value.

Deriving inferences from model predictions requires useful spatial products. In this study, predicted abundance based on the ensemble model was used to define the relative importance of n = 1243 locations in the study area for each species independently. Quintile classes were assigned, based on the entire range of predicted abundances across all seasons, in order to rank the relative usage of each of these locations. Potential vulnerability to anthropogenic disturbance was then assessed as a function of the sum of the relative usage across species, enabling the visualization of changing patterns of usage across the entire community. The composite picture that emerges is one of shifting presence, with some species disappearing from the region (e.g., petrels and shearwaters in winter) and others increasing at the same time (e.g., Black-legged Kittiwake). Spring (March - April) stood out as a period of peak usage by the entire community, which corresponded with a wide geographic distribution throughout the shelf and shelf break. As this is an area exposed to extensive shipping traffic and ongoing petroleum extraction and exploration, there remains an ongoing potential impact by acute (e.g., accidental) and chronic (e.g., illegal) discharges of oil pollution. A previous study indicated that nearshore areas, particularly in the vicinity of harbours, tend to suffer the highest pollution loading but elevated levels also occurred within and beyond the shelf break (Lieske et al., 2011).

The value of at-sea surveys are clear when one considers how difficult it can be to assess the potential vulnerability of seabirds outside the breeding season, when species are widely distributed throughout remote locations (Votier et al., 2005) and others may be present as transient non-breeders. Future work should pursue greater integration of models from different data sets, which would allow broad and fine-scale patterns to be explored and important patterns to be detected. For instance, combining telemetry tracking information with broad-scale distributional information would allow for analysis of seabird activity at multiple scales, widening the range of hypotheses that could be tested.

Furthermore, seabird research programs would greatly benefit from the coordinated acquisition of marine hazard data (e.g., fishing activity, oil discharges) at regional scales, which would support geographic analysis of the potential impact of specific risks.

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Figures

Figure 1. Coastline (dark grey shading), bathymetry (indicated by 100m isobath lines), and the location of key seabanks (100m depth, light grey shading) within the Canadian Maritimes.

Figure 2a. Density of Black-legged Kittiwake (per km^2) for $1^0 \times 1^0$ survey blocks, for spring (a, Mar - Apr), summer (b, May - Aug), fall (c, Sep - Oct), and winter (d, Nov - Feb).

Figure 2b. Density of Dovekie (per km^2) for $1^0 \times 1^0$ survey blocks, for spring (a, Mar - Apr), summer (b, May - Aug), fall (c, Sep - Oct), and winter (d, Nov - Feb).

Figure 2c. Density of murres (per km^2) for $1^0 \times 1^0$ survey blocks, for spring (a, Mar - Apr), summer (b, May - Aug), fall (c, Sep - Oct), and winter (d, Nov - Feb

Figure 2d. Density of Northern Gannet (per km²) for $1^0 \times 1^0$ survey blocks, for spring (a, Mar - Apr), summer (b, May - Aug), fall (c, Sep - Oct), and winter (d, Nov - Feb

Figure 2e. Density of storm-petrels (per km²) for $1^0 \times 1^0$ survey blocks, for spring (a, Mar - Apr), summer (b, May - Aug), fall (c, Sep - Oct), and winter (d, Nov - Feb).

Figure 2f. Density of shearwaters (per km²) for $1^0 \times 1^0$ survey blocks, for spring (a, Mar - Apr), summer (b, May - Aug), fall (c, Sep - Oct), and winter (d, Nov - Feb Figure 3. Variable importance for species distribution models developed using the negative binomial (NB) generalized linear model as indicated by regression coefficients for standardized covariates. Also shown are the estimated confidence intervals (95%) derived from testing and training data.

Figure 4. Variable importance for species distribution models developed using the hurdle model, as indicated by the regression coefficients for standardized covariates. Results are presented for both the presence component (a, binomial distribution) and the count component (b, negative binomial distribution). Also shown are the estimated confidence intervals (95%) derived from testing and training data.

Figure 5. Variable importance for species distribution models developed using random forest machine learning. "Importance" was assessed as the percentage change in prediction error when each variable is permuted out of the set of covariates used by the algorithm classifier.

Figure 6. Variation in prediction error, as measured by root mean squared error (RMSE), across 10 iterations of randomly selected training and testing data modelled using: negative binomial (NB) glm, Hurdle, Random Forest (RF), and ensemble.

Figure 7. Seasonally-specific potential vulnerability maps, based on the sum of the relative densities derived from ensemble model predictions for each of six species (Black-legged Kittiwake, Dovekie, murres, Northern Gannet, storm-petrels, and shearwaters). Seasons were defined as: spring (Mar - Apr), summer (May - Aug), fall (Sep - Oct), and winter (Nov - Feb).

Figure 8. Density distribution of the cell-specific vulnerability scores derived from the seasonally-specific vulnerability maps (Figure 7).

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)	50100	200	300	400
				Kilometers

















Figure6





summer

winter



fall



Symbol Range 10 - 12 12 - 15 15 - 18 18 - 21 21 - 25

160 240

320

80



Vulnerability (total)

				Oceanographic &		
Species	Scale	Period	Location	Biological Factors ¹	Source	Model?
Storm Petrels (Hydrobatidae)	~ 3.7-km x 0.40-km	1980, 1985, 1986	Scotian shelf, Cabot strait	Shelf break (shelf slope), sea banks (e.g., Georges Bank)	Brown (1988a)	No
Pelagic birds	Mega: >3000 km Macro:1000-3000 km Meso: 100-1000 km Coarse: 1-100 km Fine: 1-100 m	1987	Global	 Mega: major current systems and water masses; margins of oceans Macro: gyres, zonal bands, fronts Meso: rings, eddies, jets, broad upwelling events, freshwater plumes, fronts Coarse: island wakes, ice edge, sea mounts, shelf edges Fine: wind streaks 	Hunt & Schneider (1987)	No (review)
Marbled Murrelet (Brachyramphos marmoratus)	10 km shore lengths	Densities: 1972-1993 Counts: 1922- 1989	British Columbia, Canada	Average sea-surface temperature (SST), Apr. to Aug.; herring spawn index; estuary locations; distribution of sand and fine gravel substrate; proximity to glaciers	Yen et al. (2004a)	Yes (Poisson GLM ² , CART ³ , Tree ⁴ , MARS ⁵ ANN ⁶)

Table 1. Summary of the results of studies with the expressed goal of associating seabird distribution with oceanographic and biological factors.

Cassin's Auklet (<i>Ptychoramphus</i> <i>aleuticus</i>); Common Murre (<i>Uria</i> <i>aalge</i>); Sooty Shearwater (<i>Puffinus griseus</i>); Phalaropes (<i>Phalaropus</i> spp.)	~ 1 km ²	1996-1997, 2001-2002	California	Median depth; contour index (benthic "roughness"); distance to continental shelf break (200-m isobath); distance to banks, canyons and/or archipelago	Yen et al. (2004b)	Yes (ordinal logistic regression)
Western Gulls (<i>Larus</i> occidentalis); Common Guillemot (<i>Uria</i> aalge); Waves Albatrosses (<i>Phoebastria irrorata</i>)	15-min. survey time period x strip width of 300-600 m	1985-1994, continuously for gulls and guillemots; discontinuously for albatrosses	Farallon Is., California; Galapagos	Latitude; longitude; distance to mainland; distance to breeding colony; depth (model selection results not provided)	Clarke et al. (2003)	Yes (GAM ⁷)
Guillemots (Uria spp.)	20 km x 20 km grid	1986-1994	Barents Sea, Norway	Sea-surface temperature (SST); Sea-surface salinity (SSS); variance in SST; variance in SSS	Fauchald et al. (2002)	Yes (linear regression)
Northern Gannets (<i>Morus bassanus</i>); Large gulls; Auks; Kittiwakes	10 km x 10 km grid	2001-2002	Bay of Biscay, France	Latitude and longitude	Certain et al. (2007)	Yes (Two- stage modelling: GAM ⁷ + kriging of residuals)
Northern Gannets; Black- legged Kittiwake (<i>Rissa</i> <i>tridactyla</i>); Northern Fulmar (<i>Fulmaris</i>	1 ⁰ x 1 ⁰ grid	1966-1992	Atlantic Canada	Sea surface temperature (SST); sea surface salinity (SSS); water temperature at 30m depth; salinity at 30m	Huettmann and Diamond (2001)	Yes (Two- stage modelling: logistic

glacialis); Atlantic Puffin (Fratercula arctica); Thick-billed Murre (Uria lomvia); Common Murre				depth; air pressure at sea level (asl); standard deviation (STD) of atmospheric pressure; windspeed; difference between air and sea temperature; distance to coastline; shelf aspect; shelf edge; sea mount; sea depth; slope; STD of seafloor; aspect of sea floor		GLM + CART ³)
Black-footed Albatross (<i>Phoebastria nigripes</i>); Northern Fulmar; Sooty Shearwater; Fork-tailed Storm-Petrel (<i>Oceanodroma furcata</i>); Leach's Storm Petrel (<i>O. leucorhoa</i>); phalaropes; Heermann's Gull (<i>Larus heermanni</i>); California Gull (<i>L. californicus</i>); Glaucous-Winged Gull; Sabines' Gull (<i>L. sabini</i>); Black-legged Kittiwakes; Arctic Tern; Common Murre; Cassin's Auklet (<i>Ptychoramphus aleuticus</i>); Rhinoceros Auklet (<i>Cerorhinca</i>	4 km x 4 km grid	1996-2004	Southeastern Gulf of Alaska	SST (from AVHRR sensor); SST gradient (using edge detection algorithm of Etnoyer et al. 2004); Chlorophyll a (from SeaWiFS sensor)	O'Hara et al. (2006)	Yes (logistic GLM)

monocerata); Tufted Puffin (*Fratercula*

cirrhata)

Cory's Shearwater (<i>Calonectris diomedea</i>)	Continuous sampling along transect (sampled in 15 minute increments)	1983	South Atlantic Bight off northeast Florida and southeast Georgia	Distance from Gulf Stream front; change in SST; change in fluorescence	Haney and McGillivray (1985)	Yes (Pearson's r)
Thick-billed Murre	50 m, 100 m, 250 m, 500 m, 1000 m and 2000 m transect segments	1982-1983	Nuvuk Is., Hudson Bay	Depth; depth gradient (maxdepth – mindepth); depth x depth gradient (interaction); nekton density (via acoustic sampling)	Cairns and Schneider (1990)	Yes (Pearson`s r)
Dovekie (Alle alle)	Continuous sampling along transect (sampled in 10 minute increments)	1969-1983	Atlantic Canada	Distance to shelf break	Brown (1988b)	Yes (Chi- square)
Red-tailed Tropicbird (<i>Phaethon rubricauda</i>); White-tailed Tropicbird (<i>P. lepturus</i>); Red-billed Tropicbird (<i>P. aethereus</i>)	2 [°] x 2 [°] grid	1980-1995	Pacific Ocean	Latitude; longitude; shortest distance to mainland; ocean depth; distance to primary breeding colony	Spear and Ainley (2005)	Yes (GAM ⁷)
Manx shearwater (<i>Puffinus puffinus</i>); Guillemot; Razorbill (<i>Alca torda</i>); Kittiwake; Fulmar	~ 2 km resolution (transect sampled in 10-minute intervals)	1990	Irish Sea	SST; change in SST; SSS	Durazo et al. (1998)	Yes (Kruskal- Wallis ANOVA)
Northern Gannet	1 km x 1 km grid	2003	Western North Sea	Distance to colony; coastal water mass; front	Skov et al. (2008)	Yes (ENFA ⁸)

Northern Fulmar, Dovekie	Continuous sampling along 6- km length transects	1986-1990	Skagerrak, North Sea	Shelf break front	Skov and Durinck (1998)	Yes (ANOVA)
Balearic Shearwater	4 km x 4 km	2004-2009	Southwest Iberia	Distance to coast; mean depth; depth gradient; SST; Chl-a; sea surface height (SSH); density of ocean fronts; mean distance to nearest ocean front; 3-month SST, Chl-a, SSH; change in SST; change in Chl-a	Oppel et al. (2012)	Yes (GLM, GAM, BRT ⁹ , Maxent, random forest, ensemble)
Northern Fulmar	3 km x 3 km grid	1975-2009	Bering Sea and Aleutian Islands	Depth; slope; bathymetric features (e.g., plains, ridges); distance to land; colony effect; SST; primary productivity; fish catch	Renner et al. (2013)	Yes (GLMM, GAM, MARS, random forest, ensemble)
Black-legged Kittiwake	1 km x 1 km grid	2009-2010	Lambay and Raithlin Is., Republic of Ireland	Distance to colony; distance to land; depth; slope; Chl-a; SST; July fronts	Chivers et al. (2013)	Yes (Maxent)

¹ Factors which were either implicated by qualitative association, or were selected as significant predictors through formal model selection ² GLM = Generalized Linear Model.
³ CART = Classification and Regression Tree.
⁴ Tree (SPLUS).
⁵ MARS = Multivariate and Adaptive Regression Splines.

⁶ ANN = Artificial Neural Network.
⁷ GAM = Generalized Additive Model.
⁸ ENFA = Ecological Niche Factor Analysis.
⁹ BRT = Boosted Regression Tree.

Abbreviation	Variable	Unit	Data Provider	Website Address
DEPTH	Mean seadepth	m	National Geophysical	http://www.ngdc.noa
			Data Center (NOAA)	a.gov/mgg/global/eto
				po2.html
DEPTH_CV	Coefficient of	Standard-	GIS-derived product	NA
	variation in	ized	of DEPTH data	
	seadepth	variation		
PBANK	Percentage of	%	GIS-derived from	NA
	area with		100 m seadepth	
	seabank		contours	
SEASON	Categorical	dummy	NA	NA
	variable	indicator		
	(fall,spring,	variable		
	summer, winter)			
SHELFDIST	Mean distance	m	GIS-derived product	NA
	to shelfbreak		-	

Table 2. Environmental variables used to construct seabird species distribution models.