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## **Volume 7B Proposed Development (Offshore) Appendices**

Appendix 6-1 Offshore Ornithology Baseline Characterisation Report Annex 16 MRSea Modelling Report

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# **Volume 7B Appendix 6-1 Annex 16 MRSea Modelling Report**



*This document contains the following report: 'Caledonia Offshore Windfarm. MRSea modelling of key seabird species' as prepared by Black Bawks Data Science Ltd in February 2024. For the purpose of Consent Application, the document has been retitled to: 'Volume 7B, Appendix 6-1, Annex 16: MRSea Modelling Report', alongside the addition of a new front cover.*





# Caledonia Offshore Windfarm

MRSea modelling of key seabird species







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- List of Abbreviations
- ANOVA Analysis of Variance
- BIC Bayesian Information Criterion
- CReSS Complex Regional Spatial Smoother
- CV Coefficient of Variation
- GAM Generalised Additive Model
- GEBCO General Bathymetric Chart of the Oceans
- GIS Geographic Information System
- GLM Generalised Linear Model
- JNCC Joint Nature Conservation Committee
- MRSea Marine Renewables Strategic Environmental Assessment
- OWF Offshore Wind Farm
- QL Quasi-likelihood
- RSE Relational Standard Error
- SALSA Spatially Adaptive Local Smoothing Algorithm
- SST Sea Surface Temperature
- UTM Universal Transverse Mercator
- VIF Variance Inflation Factor

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#### 1 Introduction

<span id="page-16-0"></span>As part of the environmental impact assessment framework for the proposed Caledonia offshore wind farm, an assessment of the distribution and abundance of key species was contracted to be undertaken using the Marine Renewables Strategic Environmental Assessment (MRSea Windows package) in R (Scott-Hayward, 2013). The goal of this work was to first quantify the abundance and distribution of key seabird species while considering environmental covariates that could impact their ecology. Further to this, we developed and utilized a spatial apportioning approach for observations unable to be identified to species level which allowed for modelling and thus appropriate accounting for variability in the data. We also applied the random forests algorithm to create predictions of the distribution of birds based on several wind farm scenarios. The use of random forests also allowed for us to make some assessment of displacement based on the partial dependence plots. Finally, using the MRSea outputs, we performed a hot-spot analysis to determine if there were any parts of the proposed development area that had persistent hot-spots for the key species of the analysis.

#### <span id="page-16-1"></span>2 Methods statement

Seabird observation and effort data from 24 digital aerial surveys conducted by APEM between May 2021 – April 2023 were used for the spatial modelling of species' monthly (or yearly, if too few observations) distribution and abundance. Aerial surveys were flown over the proposed Caledonia Offshore Wind Farm (OWF) development, wherein transects included the site and a 4km buffer. Data for eight species were processed: kittiwake (Rissa tridactyla), razorbill (Alca torda), quillemot (Cepphus grylle), puffin (Fratercula arctica), gannet (Morus bassanus), fulmar (Fulmarus glacialis), great black-backed gull (Larus marinus), and herring gull (Larus argentatus). All species were modeled with data for "all birds" (both flying and sitting behaviours). For kittiwake, gannet, and fulmar, there were sufficient observations to create additional models specifically for birds observed while flying. For all species except auks (puffin, razorbill, guillemot), only detections identified to species level were included in the analysis. For the auks, observations identified to species level were supplemented with counts apportioned using a proportional technique based on species groupings provided by APEM. Proportional no-id apportioning was made at the transect level to ensure a more accurate proportional representation of non-identified species (see section [2.5.1](#page-20-0) for more detail).

To model survey-specific bird distribution and abundance, we used the Complex Regional Spatial Smoother (CReSS) spatial modelling method using Spatially Adaptive Local Smoothing Algorithm (SALSA) based model selection (Scott-Hayward, 2013). The models effectively fit the relationship between the observations (count response variable) and the environment (candidate covariates, listed in table 1) at each location which was then be used to estimate the animal density over the area of interest. For each survey, individual counts for each species were assigned to the midpoint of the respective aerial image footprint to produce the input data for each species-specific model. This generated a count variable (the dependent variable) for each footprint, and the footprint area thus became the offset for the model to ensure predicted outputs represented density. Covariate values were then assigned to the midpoint of each segment such that the resulting model input data frame included survey-specific species counts and covariate values for each transect segment.



Count data from aerial transect surveys were correlated to consecutive measurements through space and time. Furthermore, due to environmental and prey conditions, the number of animals in any given area is more likely to be similar for points closer temporally, than those more distant in time. Models fitted to (relative) abundance data attempt to explain animal abundance at any location, but the information (covariate data) that describes why animals are found in high/low numbers at specific locations is frequently missing from the model, leaving patterns in the model's noise component (model residuals). These patterns are also expected to be similar along the track lines. This (positive) correlation in model residuals along the track lines violates a critical assumption for standard statistical models that require an independent set of residuals (such as Generalised Linear Models (GLMs) / Generalised Additive Models (GAMs)). Ignoring this violation can invalidate all model-based precision estimates (e.g., standard errors, confidence intervals, and p-values), resulting in overly complex models that can suggest irrelevant environmental covariates are statistically significant. Transect data are frequently prone to such spatiotemporal autocorrelation, which violates a core assumption of GLMs/GAMs. Thus, transect ID was included as a blocking factor in the analysis to control for autocorrelation in the model. This informed the model that correlation within a transect is accepted and that transect independence is assumed.

#### 2.1 Model Inference

<span id="page-17-0"></span>For all models we assumed that the data followed a quasi-poisson distribution, which is typical when working with overdispersed count data (overdispersion occurs when the variance of the data is greater than the mean). This distribution assumes: 1) observations can be counted; 2) the rate at which observations occur can be calculated; 3) No two observations can occur at the same time and/or in the same place; 4) the variance is a linear function of the mean; and 5) that all observations are independent (Ver Hoef and Boveng, 2007). However, count data from aerial transect surveys are likely to be correlated (i.e., dependent) to consecutive measurements through space and time. Furthermore, due to environmental and prey conditions, the number of animals in any given area is more likely to be similar for points closer temporally, than those more distant in time. Models fitted to (relative) abundance data attempt to explain animal abundance at any location, but the variables that determine why animals are found in high/low numbers at specific locations are frequently missing from the model, leaving patterns in the model's noise component (model residuals). These patterns are also expected to be similar along the track lines. This, temporal and spatial autocorrelation in model residuals along the track lines violates a critical assumption for standard statistical models which require an independent set of residuals (such as Generalised Linear Models (GLMs) / Generalised Additive Models (GAMs)). Ignoring this violation can invalidate all model-based precision estimates (e.g., standard errors, confidence intervals, and p-values), resulting in overly complex models that can result in a Type 1 statistical error whereby irrelevant environmental covariates are statistically significant. Transect data are frequently prone to such spatiotemporal autocorrelation, which violates a core assumption of GLMs/GAMs. To address this issue, transect ID was included as a blocking factor in the analysis to control for autocorrelation in the model. Including transect ID mitigates autocorrelation issues by including it as a factorial variable that describes the distribution of the individuals, accounting for the correlation within the transect. This is standard practice when accounting for autocorrelation in transect data. A one-way Analysis of Variance (ANOVA) was performed to determine the statistical significance of covariates in the predictive model. Partial dependence plots were used to investigate covariates that have significant relationships with the data in the model. Further model inference was made by examining the cumulative residual plots output by the models.



#### 2.2 Selection of Model Covariates

<span id="page-18-0"></span>First, a full model with all appropriate terms (e.g., as identified from [Table 1\)](#page-18-1) was fit for each species without a smooth term for the spatial component. This allowed the potential relationships between covariates and species observations to be initially unhindered by spatial information. Variance Inflation Factors (VIF) were then used to select covariate terms from the initial model fitting process that should be removed due to collinearity based on a VIF threshold value of 2. There is not a standardised approach to choosing this value, but in general, lower threshold values constitute a more conservative approach to eliminating issues with multicollinearity. The flexibility of the smoother-related term for each model term was then chosen, followed by the model selection for the two-dimensional smoother term for the spatial component. Segment area was then incorporated into the model as an offset term as the transects' division may have resulted in slightly different dimensions. Each model was permitted to retain the covariates as a smooth or linear term (or omitted completely). SALSA was used to fit a smooth function for each covariate. Model selection for both the covariates and spatially based smoothers was conducted by using an objective fit measure (Bayesian Information Criterion (BIC) for quasi-likelihood (QL) models). Models that permit over-dispersion for Poisson-style counts are QL based, necessitating QL-based fit scores.

<span id="page-18-1"></span>





#### 2.3 Knot Placement and Basis Function Details

<span id="page-19-0"></span>The number of "knots" used for the model and the effective range of each knot (the spatial extent to which each knot influences the fitted surface) are both key factors in determining the model flexibility for the spatial surfaces in this setting. The candidate models were chosen from a range of models that varied in the number of knots provided and the effective range (R-value) of each knot because the optimal choices for both values are always unknown. The starting knot positions on the spatial surface were chosen to maximize coverage across the spatial area (via a space filling algorithm; (John et al., 1995)), and these positions were allowed to move according to the SALSA (Walker et al., 2011) model selection technique. We used the local exponential basis function, defined as:

#### $((exp(-d/r2)))$

where d is the Euclidian distance, allowing for varied R-values over the surface. A variable number of knots (2-40 depending on data sparsity; the number is denoted by the degrees of freedom in the model) were used for the candidate models, and an objective fit criterion was employed to select the best model(s). In effect, the position of the knot placement, and to a lesser extent the number of knots, reflect the complexities of the spatial relationship between bird abundance and the covariates chosen for the study. Knot locations were identified separately for each survey to accommodate differences in survey effort and bird distributions across surveys.

#### 2.4 Geo-Referenced Results

<span id="page-19-1"></span>The species-specific fitted surfaces were generated by making predictions within a grid using the final model at a 1km x 1km resolution. The grid is a series of regular points spaced at 1km resolution across the surface of the area of interest. These regular points are associated with the same environmental covariates as those used in the modelling process. This allows the trained MRSea model to make predictions of animal density on each of those points. Those data can then be visualized or interpolated to create surfaces. These grids were projected as the Universal Transverse Mercator (UTM; Zone 30) projection. To measure uncertainty spatially and in the population estimates, the model was bootstrapped 1,000 times (wherein random subsets of the modelled coefficients are drawn from a multivariate normal distribution and predictions are made for each grid cell, 1,000 times). From this we calculated the mean predicted density, the upper and lower 95% confidence limits, and the coefficient of variation (CV; as defined by the ratio of the standard deviation to the mean). These measures of uncertainty were visualized and are presented in the results section.

#### 2.5 Abundance Estimates from MRSea Density Surfaces

<span id="page-19-2"></span>Abundance estimates were calculated by summing the grid cells across the prediction surface at the temporal scale specified by the exploratory analysis. To calculate abundance estimates within the survey area, we summed grid cells that fall within the boundary or touch the edges, ensuring that grid cells at the boundary are clipped to the boundary footprint and adjusted for the new area. The upper and lower confidence limits of the population estimate were calculated by determining the 95% confidence limits of the 1000 bootstrapped surfaces. The bootstrap outputs were not normally distributed, and the mean was more sensitive to outliers therefore the median was chosen as a more representative measure of central tendency when presenting abundance estimates. We



note that in previous work in the offshore wind industry in the UK, the term "coefficient of variation" is often applied to population estimates derived from bootstrap exercises. However, this is a misnomer because the standard deviation of a bootstrapped mean is the standard error. The calculation of the ratio of the standard error to the mean is called the relational standard error (RSE), and not the CV as has been previously used. For the sake of accuracy in the vocabulary used and to distinguish this measure of uncertainty from the spatial CV, we present the uncertainty as the RSE, but note that this is equivalent to what has been referred to as the CV in past work.

#### 2.5.1 Species Apportioning

<span id="page-20-0"></span>To account for auks that have not been identified to species level, we applied a spatial proportional apportioning technique based on the species groupings defined by the digital aerial survey provider. For example, if there is a species grouping called "large auks" which is made up of possible guillemots or razorbill, then the proportional species composition will be applied within spatial bins that will be used for modelling. In other words, if 80 guillemot and 20 razorbill were identified in a survey, and a spatial bin/cluster had 10 unidentified large auks, then 8 guillemot and 2 razorbills would be apportioned into that bin. Spatial proportional no-id apportioning was first attempted at a transect level (which ensures a more accurate proportional representation of non-identified species). In cases when that was unable to be performed, then it was performed at the survey level. In the case that a proportional approach leads to fractional values (for example, if the split was 80/20 for guillemot to razorbill, and there were 7 unidentified large auks in a cluster, this would equate to 5.6 guillemot and 1.4 razorbill. This would be rounded to 6 guillemot and 1 razorbill). The new observations apportioned from the non-identified birds were then used for MRSea modelling.

#### 2.6 Hotspot Analysis

<span id="page-20-1"></span>A hotspot analysis was performed upon completion of all species models. The analysis was informed by the "all birds" survey-level MRSea models. Model outputs from all species and surveylevel outputs were normalized on a scale of 0 –1 and then averaged. The cell-by-cell coefficient of variation was calculated by dividing the standard deviation by the mean. This provided a single magnitude (I.e., mean normalized prediction layer) and persistence (I.e., variability as defined by the CV) layer. The upper and lower 95<sup>th</sup> percentiles of all values in the magnitude and persistence layers were computed to use as thresholds for determining if a grid cell is persistent hot or cold spot. This is categorized as per table 2.

<span id="page-20-2"></span>

| $ny \cup y$   |                                     |                      |
|---|-------------------------------------|----------------------|
| Persistence   | Magnitude                           | Classification       |
| $CV > 95th$ percentile  | Mean $> 95th$ percentile            | Persistent hot spot  |
| $CV > 95$ <sup>th</sup> percentile  | Mean $<$ 5 <sup>th</sup> percentile | Persistent cold spot |
| $CV < 5th$ percentile   | Mean $> 95th$ percentile            | Volatile hot spot    |
| $CV < 5th$ percentile   | Mean $<$ 5 <sup>th</sup> percentile | Volatile cold spot   |
| $CV > 5th$ percentile, < 95 <sup>th</sup>   | Mean $> 95th$ percentile            | Transient hot spot   |
| percentile  |                                     |                      |
| $CV > 5$ <sup>th</sup> percentile, < 95 <sup>th</sup> Mean < 5 <sup>th</sup> percentile |                                     | Transient cold spot  |
| percentile  |                                     |                      |

Table 2. Classifications of hot spots as defined by percentiles from magnitude and persistence layers

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#### 2.7 Assessment of Displacement from MRSea

<span id="page-21-0"></span>We made use of the precise Moray East offshore wind farm operational turbine locations and timing of installation to build a series of "distance to turbine" predictor layers that were used in the modelling process. We constructed monthly "distance to turbine" layers that were representative of the active turbines in the water at the time of each survey (akin to monthly SST layers). We then examined the partial dependence plots of the distance to turbine layers to make assessments of displacement because inflection points in those figures could indicate if displacement is occurring, and at what spatial scales.

#### 2.8 Random forests

<span id="page-21-1"></span>An additional analysis was conducted which performed spatial displacement scenarios based on proposed turbine locations. The distance to turbine layers from the operational Moray East offshore wind farm were used to train a spatial model that employs a machine learning algorithm called random forests. The random forest algorithm is a supervised machine learning algorithm used widely for regression and classification problems in machine learning (Breiman, 2001). A random forest is a classifier that includes many decision trees on various subsets of a given dataset. The classifier takes the average decision of that subset to improve the predictive performance. It is built on the idea of ensemble learning, where multiple classifiers are integrated to solve complex problems and improve model performance. In the same way that a forest with many trees is more robust, a random forest algorithm with more decision trees will have greater accuracy and higher predictive ability.

Random Forests for regression represent a powerful ensemble learning approach, particularly adept at handling complex relationships in data (Hastie et al., 2008). The foundation of Random Forests lies in the construction of decision trees. In the context of regression, each tree is essentially a sequence of binary decisions or splits that recursively partition the input space. What sets Random Forests apart is the introduction of randomness in the process. During the creation of each tree, a random subset of the training data, known as a bootstrap sample, is drawn with replacement. Moreover, at each node of the tree, only a random subset of predictors is considered for determining the best split. This deliberate injection of variability helps prevent individual trees from fitting noise in the data and encourages diversity among the constituent trees.

The aggregation of these independently grown and varied trees defines the strength of Random Forests. For regression, the final prediction is an average (or weighted average) of the predictions from all the trees, providing a robust and accurate estimate of the target variable. Importantly, Random Forests are equipped to handle both numerical and categorical predictors, making them



versatile across a range of real-world datasets. The algorithm's adaptability, resilience to overfitting, and ability to capture non-linear relationships make it particularly well-suited for regression tasks where the underlying data structure is complex and multifaceted.

The calculation of regression trees within the Random Forest framework involves recursive binary splits based on predictor variables. At each node, the algorithm identifies the predictor and split point that minimizes the sum of squared differences between the observed and predicted values. The splitting process continues until a predefined stopping criterion is met, such as a minimum node size or a maximum tree depth. This recursive partitioning results in a tree structure where each terminal node, or leaf, represents a distinct subset of the data. During prediction, an input observation traverses the tree, and its target value is determined by the average of the training observations in the terminal node to which it belongs. This process is repeated for each tree in the ensemble, and the final prediction for regression is the aggregated result of all individual tree predictions. The collective strength of Random Forests lies in their ability to capture intricate relationships in the data while minimizing the risk of overfitting (Breiman 2001; Cutler et al., 2012; Hastie et al., 2008). Furthermore, its non-parametric nature means that many of the typical frequentist statistical assumptions do not have to be adhered to, thus increasing the likelihood of generating statistically appropriate estimations.

In a spatial context, this is applied in an almost identical fashion to MRSea, where observations or counts are associated with environmental covariates which allows for a model to be trained. This trained model is then applied to a grid of regular points to generate predictions in space. We used the same environmental covariates as MRSea, however, the spatial component was replaced by spatial autocorrelation terms, which helped capture the nature of flocking behaviour by marine birds. To generate the predicted scenarios of displacement, Ocean Winds was consulted to create three plausible wind turbine configurations, and each of those scenarios was used to create distance to wind turbine layers, which was used to predict displacement for Guillemot, Razorbill, Kittiwake, Fulmar, Puffin and Gannet. This algorithm was used as opposed to MRSea because MRSea requires the spatial knots as calculated using the raw observations which have been captured prior to any turbines being installed. The aim of this process was to create predictive scenarios of turbines being installed in various spatial configurations without being constrained by the locations of birds as they existed prior to the installation.

#### 2.9 Displacement simulation

<span id="page-22-0"></span>To assess the influence of the distance to turbine predictor layer in the model, we built a random forests model using the caret (Kuhn 2022) and ranger (Wright and Ziegler 2017) packages in R version 4.3.1. Using a grid-based tuning approach, we tested values of mtry (i.e., the number of randomly selected variables at each split), from 3 to 10, and the number of minimum observations in each terminal node between 80 and 100. As a metric for model selection, we used root mean squared error (RMSE) with five-fold cross validation. The model with the lowest RMSE was selected as the best model.

To assess the effect of turbines on the distribution of key species, the partial dependence plots of the distance to turbine covariate were generated and examined. A partial dependence plot is a graphical method used in machine learning to visualize the marginal effect of a feature on the predicted outcome of a model while marginalizing over the values of all other features. It helps to understand the relationship between a feature and the target variable in isolation, holding other



features constant or averaging over their values. However, in terms of understanding displacement, we note that we are incorporating observations of birds throughout the entire survey area (i.e., outside of the likely distance of immediate impact of turbines on distribution as they would be outside of visual range). Furthermore, relationships noted (e.g., higher densities of birds predicted further away from turbines) could simply be due to other factors (e.g., presence of more favourable foraging conditions at a site that happens to fall at some distance away from the turbine blades), Thus these interpretations must be taken cautiously. Visual acuity varies between species depending on their foraging ecology, but based on data collected for Northern fulmar, the smallest low-contrast object that can be seen near sea-level would have to be 13m in diameter at 11 km distance (Mitkus et al., 2016). Thus, a conservative estimate of 20 km was applied as the maximum distance along the x-axis for interpretation of the partial dependence plots.

Next, we used plausible scenarios of turbine distributions to generate spatial predictions, under the assumption that the distribution of birds in the region was representative of the post-construction phase. For the purposes of this analysis, we use all flying and sitting birds combined under the assumption that the impact will be the same for both behaviour types.

Four scenarios were modeled: Scenario 1; where no new turbines were constructed - this represents the baseline scenario as per the digital aerial surveys. Scenario 2; where turbines were constructed only in the northern part of the proposed wind farm. Scenario 3; where turbines were constructed only in the southern part of the proposed wind farm. And Scenario 4; a combination of scenarios 2 and 3 (Figure 1).



Figure 1: Turbine scenarios for examining distributional responses post-construction



Grid cells in the predicted outputs were adjusted for unequal areas around the borders of the survey region and predicted counts were summed to generate population estimates for each survey and turbine scenario.

#### 2.10 Interpretation of Random Forests outputs

<span id="page-24-0"></span>Although random forests offers a number of advantages as a predictive algorithm, interpretation of the outputs must be done carefully, particularly when attempting to understand mechanistic relationships. Firstly, Random forests are comprised of multiple decision trees, where each tree is trained on a subset of the data and with a subset of features. This ensemble approach often leads to improved performance compared to individual decision trees. However, interpreting the collective decision-making process of many trees can be complex, especially when there are many trees in the forest. Also, the building blocks of random forests are capable of learning complex non-linear relationships in the data. As a result, random forests can model intricate decision boundaries that may not be easily understandable or interpretable, especially in high-dimensional spaces. Further to this, even though decision trees are simpler models compared to some other machine learning algorithms, individual trees in a random forest can still be quite complex, especially if the dataset contains many features or if the trees are allowed to grow deep. Interpreting the decisions made by these trees can be challenging, particularly when trying to understand how they collectively contribute to the final prediction.

#### <span id="page-24-1"></span>3 Results

Models were generated for each species using observations of that species for all behaviours, these will be referred to as the "all-birds" models. Additional models were generated for kittiwake, gannet, fulmar, and great black-backed gulls using only observations of flying birds, these models will be referred to as "flying-birds" models. Mean density surfaces for each survey from MRSea outputs mapped to the Caledonia OWF site are provided in figures 2 - 17 (all kittiwake), figures 21-24 (flying kittiwake), figures 40-41 (all gannet), figures 51-52 (flying gannet), figures 62-66 (all fulmar), figures 85-89 (flying fulmar), figures 108-112 (guillemot), figures 130-133 (razorbill), figures 148-150 (puffin), figure 162 (all great black-backed gull), figure 169 (flying great blackbacked gull), figure 176 (herring gull).

#### 3.1 Kittiwake

#### <span id="page-24-2"></span>3.1.1 All Birds Model

<span id="page-24-3"></span>Table 3. Candidate and final covariates for all kittiwakes model.

<span id="page-24-4"></span>



Distribution maps generated using MRSea (Figure 2-17) suggest that kittiwake are widely distributed throughout the area during the breeding season (15 April – 31 August), with higher concentrations in the south. In the non-breeding season, distribution declines, but is again generally higher in the south. The highest densities were observed in the southern third of the study area during July 2021.

Model fit was poor with a marginal R squared value of 0.04 and root mean squared error of 13.13. Cumulative residuals in the model showed that there was a moderate relationship between predicted and observed values across the lower range of predicted values, though the model tends to under-predict for values between  $\sim$  1.8 and  $\sim$  5 birds/ km<sup>2</sup> (bottom row, Figure 19).





<span id="page-26-0"></span>Figure 2. Median density of all kittiwake in the survey area for months with sufficient observations between May and September 2021





<span id="page-27-0"></span>Figure 3. Median density of all kittiwake in the survey area for months with sufficient observations between October 2021 and March 2022





<span id="page-28-0"></span>Figure 4. Median density of all kittiwake in the survey area for months with sufficient observations between April and August 2022





<span id="page-29-0"></span>Figure 5. Median density of all kittiwake in the survey area for months with sufficient observations between September 2022 and April 2023





<span id="page-30-0"></span>Figure 6. Lower confidence limit of density of all kittiwake in the survey area for months with sufficient observations between May and September 2021





<span id="page-31-0"></span>Figure 7. Lower confidence limit of density of all kittiwake in the survey area for months with sufficient observations between October 2021 and March 2022





<span id="page-32-0"></span>Figure 8. Lower confidence limit of density of all kittiwake in the survey area for months with sufficient observations between April and August 2022





<span id="page-33-0"></span>Figure 9. Lower confidence limit of density of all kittiwake in the survey area for months with sufficient observations between September 2022 and April 2023





<span id="page-34-0"></span>Figure 10. Upper confidence limit of density of all kittiwake in the survey area for months with sufficient observations between May and September 2021





<span id="page-35-0"></span>Figure 11. Upper confidence limit of density of all kittiwake in the survey area for months with sufficient observations between October 2021 and March 2022




Figure 12. Upper confidence limit of density of all kittiwake in the survey area for months with sufficient observations between April and August 2022

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Figure 13. Upper confidence limit of density of all kittiwake in the survey area for months with sufficient observations between September 2022 and April 2023





Figure 14. Spatial coefficient of variation of predicted densities of all kittiwake from MRSea across the survey area for months with sufficient observations between May and September 2021





Figure 15. Spatial coefficient of variation of predicted densities of all kittiwake from MRSea across the survey area for months with sufficient observations between October 2021 and March 2022





Figure 16. Spatial coefficient of variation of predicted densities of all kittiwake from MRSea across the survey area for months with sufficient observations between April and August 2022





Figure 17. Spatial coefficient of variation of predicted densities of all kittiwake from MRSea across the survey area for months with sufficient observations between September 2022 and April 2023





Figure 18. Autocorrelation test for kittiwake density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)





Figure 19. Fitted (MRSea predictions) versus observed counts of all kittiwake

Table 4. ANOVA results from the best MRSea model for all kittiwake as selected by cross-validation

| Variable           | Degrees of Freedom | Chi-square | P-value |
|--------------------|--------------------|------------|---------|
| Sandeel Density    |                    | 31.23      | <<0.001 |
| Distance to Colony | ∽                  | 7.63       | 0.003   |
| X/Y (location)     |                    | .85<br>41  | <<0.001 |

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Figure 20. Partial dependence plots for significant variables for all kittiwake from MRSea. Note that distance to turbine was not a significant variable but was included to demonstrate the relationship. (Clockwise from top left: sandeel density, distance to colony, distance to turbine)

## 3.1.2 Flying Birds Model

Table 5. Candidate and final covariates for flying kittiwake model



Distribution maps generated using MRSea (Figure 21-36) suggest elevated densities of flying kittiwake during the breeding season (15 April – 31 August) of both years, but were higher in



2022. In the non-breeding season, there is a notable drop in kittiwake density. The highest densities were observed in the south of the study area during July 2022.

Model fit was poor with a marginal R squared value of 0.06 and root mean squared error of 3.94. Cumulative residuals in the model showed that there was a poor relationship between predicted and observed values particularly when predicted counts were above  $\sim$  2.5 birds/km<sup>2</sup> (Figure 38).





Figure 21. Median density of flying kittiwakes in the survey area for months with sufficient observations between May and September 2021





Figure 22. Median density of flying kittiwakes in the survey area for months with sufficient observations between October 2021 and April 2022





Figure 23. Median density of flying kittiwakes in the survey area for months with sufficient observations between May and September 2022





Figure 24. Median density of flying kittiwakes in the survey area for months with sufficient observations between October 2022 and April 2023





Figure 25. Lower confidence limit of density of flying kittiwakes in the survey area for months with sufficient observations between May and September 2021





Figure 26. Lower confidence limit of density of flying kittiwakes in the survey area for months with sufficient observations between October 2021 and April 2022





Figure 27. Lower confidence limit of density of flying kittiwakes in the survey area for months with sufficient observations between May and September 2022





Figure 28. Lower confidence limit of density of flying kittiwakes in the survey area for months with sufficient observations between October 2022 and April 2023





Figure 29. Upper confidence limit of density of flying kittiwakes in the survey area for months with sufficient observations between May and September 2021





Figure 30. Upper confidence limit of density of flying kittiwakes in the survey area for months with sufficient observations between October 2021 and April 2022





Figure 31. Upper confidence limit of density of flying kittiwakes in the survey area for months with sufficient observations between October 2021 and April 2022





Figure 32. Upper confidence limit of density of flying kittiwakes in the survey area for months with sufficient observations between October 2022 and April 2023





Figure 33. Spatial coefficient of variation of predicted densities of flying kittiwakes from MRSea across the survey area for months with sufficient observations between May and September 2021





Figure 34. Spatial coefficient of variation of predicted densities of flying kittiwakes from MRSea across the survey area for months with sufficient observations between October 2021 and April 2022





Figure 35. Spatial coefficient of variation of predicted densities of flying kittiwakes from MRSea across the survey area for months with sufficient observations between May and September 2022





Figure 36. Spatial coefficient of variation of predicted densities of flying kittiwakes from MRSea across the survey area for months with sufficient observations between October and April 2023





Figure 37. Autocorrelation test for kittiwake density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)







Figure 38. Fitted (MRSea predictions) versus observed counts of flying kittiwake







Figure 39. Partial dependence plots for significant variables for flying kittiwakes from MRSea. (Clockwise from top left: bathymetry, distance to colony, distance to turbine)

## 3.2 Gannet

3.2.1 All Birds Model

| Starting model covariates after VIF-based collinearity removal | Final model covariates after removal by SALSA |  |
|--|---|--|
| Aspect   | Aspect  |  |
| Slope  | Slope   |  |
| Bathymetry   | Bathymetry                                    |  |
| Sandeel Presence   | Sandeel Presence                              |  |
| Distance to colony   | Distance to colony                            |  |
| Distance to turbine  | Distance to turbine                           |  |
| Standard deviation of Sea Surface Temperature                  | Standard deviation of Sea Surface Temperature |  |

Table 7. Candidate and final covariates for all gannets model

Distribution maps generated using MRSea (Figure 39-40) suggest that gannets are distributed mostly across the southern half of the study area during the breeding season (15 March – 30 September). In the non-breeding season, gannets were largely absent from the study area. The highest densities were observed along the south east border of the study area in July 2022.



Model fit was poor with a marginal R squared value of 0.04 and root mean squared error of 0.87. Cumulative residuals in the model showed that there was a poor relationship between predicted and observed values across most of the range of predicted values, with the model generally over predicting for values under 0.4 birds/km<sup>2</sup>, and over predicting for values above. (Figure 48).





Figure 40. Median density of all gannets in the survey area for months with sufficient observations between June and October 2021





Figure 41. Median density of all gannets in the survey area for months with sufficient observations between May and October 2022





Figure 42. Lower confidence limit of density of all gannets in the survey area for months with sufficient observations between June and October 2021





Figure 43. Lower confidence limit of density of all gannets in the survey area for months with sufficient observations between may and October 2022





Figure 44. Upper confidence limit of density of all gannets in the survey area for months with sufficient observations between June and October 2021





Figure 45. Upper confidence limit of density of all gannets in the survey area for months with sufficient observations between May and October 2022




Figure 46. Spatial coefficient of variation of predicted densities of all gannets from MRSea across the survey area for months with sufficient observations between June and October 2021





Figure 47. Spatial coefficient of variation of predicted densities of all gannets from MRSea across the survey area for months with sufficient observations between May and October 2022





Figure 48. Autocorrelation test for gannet density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)





Figure 49. Fitted (MRSea predictions) versus observed counts of all gannets

Table 8. ANOVA results from the best MRSea model for all gannets as selected by cross-validation

| Variable                     | Degrees of Freedom | Chi-square | P-value |
|------------------------------|--------------------|------------|---------|
| Aspect                       | 5                  | 2.60       | 0.762   |
| Slope                        |                    | 4.07       | 0.539   |
| Bathymetry                   | 5                  | 2.83       | 0.726   |
| Sandeel Presence             | 5                  | 3.79       | 0.58    |
| Distance to Colony           | 5                  | 17.66      | 0.003   |
| Turbine Distance             | 5                  | 6.03       | 0.303   |
| Sea Surface Temperature (SD) | 5                  | 23.91      | <<0.001 |
| X/Y (location)               | 10                 | 84.62      | <<0.001 |



Figure 50. Partial dependence plots for significant variables for all gannets from MRSea. Note that distance to turbine was not a significant variable but was included to demonstrate the relationship. (Clockwise from top left: distance to colony, daily standard deviation of sea surface temperature, distance to turbine)

## 3.2.2 Flying Birds Model





Distribution maps generated using MRSea (Figure 50-51) suggest that flying gannets are distributed mostly across the southern half of the study area during the breeding season (15 March – 30 September). In the non-breeding season, gannets were largely absent from the study area. The highest densities were observed in the southern third of the study area in June 2022.



Model fit was poor with a marginal R squared value of 0.01 and root mean squared error of 0.60. Cumulative residuals in the model showed that there was a moderate relationship between predicted and observed values across most of the range of predicted values, except between ~1.3 and 0.2 birds/km<sup>2</sup>, where the model tended to underpredict density (Figure 59).





Figure 51. Median density of flying gannets in the survey area for months with sufficient observations between June 2021 and June 2022





Figure 52. Median density of flying gannets in the survey area for months with sufficient observations between July and October 2022





Figure 53. Lower confidence limit of density of flying gannets in the survey area for months with sufficient observations between June 2021 and June 2022





Figure 54. Lower confidence limit of density of flying gannets in the survey area for months with sufficient observations between July and October 2022





Figure 55. Upper confidence limit of density of flying gannets in the survey area for months with sufficient observations between June 2021 and June 2022





Figure 56. Upper confidence limit of density of flying gannets in the survey area for months with sufficient observations between July and October 2022





Figure 57. Spatial coefficient of variation of predicted densities of flying gannets from MRSea across the survey area for months with sufficient observations between June 2021 and June 2022





Figure 58. Spatial coefficient of variation of predicted densities of flying gannets from MRSea across the survey area for months with sufficient observations between July and October 2022





Figure 59. Autocorrelation test for gannet density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)







Figure 60. Fitted (MRSea predictions) versus observed counts of flying gannets

Table 10. ANOVA results from the best MRSea model for flying gannets as selected by crossvalidation

| Variable                | Degrees of Freedom | Chi-square | P-value |
|-------------------------|--------------------|------------|---------|
| Turbine Distance I      |                    | 86.        | 0.173   |
| Sea Surface Temperature |                    | 0.66       | 0.416   |
| $X/Y$ (location)        |                    | 52.64      | <<0.001 |





Figure 61. Partial dependence plots for significant variables for flying gannets from MRSea. Note that distance to turbine was not a significant variable, but was included to demonstrate the relationship. (left: distance to turbine; right: daily mean of sea surface temperature)

## 3.3 Fulmar

## 3.3.1 All Birds Model

Table 11. Candidate and final covariate terms for all fulmar model.



Distribution maps generated using MRSea (Figure 61-65) suggest that all fulmar are distributed in higher densities in the southern and western areas of the study area during the breeding season (01 April - 15 September). In the non-breeding season, fulmar tend to be distributed in higher densities along the south edge of the study area. The highest densities were observed around the southern tip of the study area during July 2022.

Model fit was moderate with a marginal R squared value of 0.14 and root mean squared error of 5.20. Cumulative residuals in the model showed that there was a poor relationship between predicted and observed values across most of the range of predicted values, and tended to underpredict bird density (Figure 82).





Figure 62. Median density of all fulmar in the survey area for months with sufficient observations between May and September 2021





Figure 63. Median density of all fulmar in the survey area for months with sufficient observations between November 2021 and March 2022





Figure 64. Median density of all fulmar in the survey area for months with sufficient observations between April and August 2022





Figure 65. Median density of all fulmar in the survey area for months with sufficient observations between September 2022 and February 2023





Figure 66. Median density of all fulmar in the survey area for months with sufficient observations between March and April 2023





Figure 67. Lower confidence limit of density of all fulmar in the survey area for months with sufficient observations between May and September 2021

94





Figure 68. Lower confidence limit of density of all fulmar in the survey area for months with sufficient observations between November 2021 and March 2022





Figure 69. Lower confidence limit of density of all fulmar in the survey area for months with sufficient observations between April and August 2022





Figure 70. Lower confidence limit of density of all fulmar in the survey area for months with sufficient observations between September 2022 and February 2023





Figure 71. Lower confidence limit of density of all fulmar in the survey area for months with sufficient observations between March and April 2023





Figure 72. Upper confidence limit of density of all fulmar in the survey area for months with sufficient observations between May and September 2021





Figure 73. Upper confidence limit of density of all fulmar in the survey area for months with sufficient observations between November 2021 and March 2022





Figure 74. Upper confidence limit of density of all fulmar in the survey area for months with sufficient observations between April and August 2022





Figure 75. Upper confidence limit of density of all fulmar in the survey area for months with sufficient observations between September 2022 and February 2023





Figure 76. Upper confidence limit of density of all fulmar in the survey area for months with sufficient observations between March and April 2023





Figure 77. Spatial coefficient of variation of predicted densities of all fulmar from MRSea across the survey area for months with sufficient observations between May and September 2021





Figure 78. Spatial coefficient of variation of predicted densities of all fulmar from MRSea across the survey area for months with sufficient observations between November 2021 and March 2022





Figure 79. Spatial coefficient of variation of predicted densities of all fulmar from MRSea across the survey area for months with sufficient observations between April and August 2022




Figure 80. Spatial coefficient of variation of predicted densities of all fulmar from MRSea across the survey area for months with sufficient observations between September 2022 and February 2023





Figure 81. Spatial coefficient of variation of predicted densities of all fulmar from MRSea across the survey area for months with sufficient observations between March and April 2023





Figure 82. Autocorrelation test for fulmar density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)





Figure 83. Fitted (MRSea predictions) versus observed counts of all fulmar

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 $10000$ 

Index

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10 \\
\end{array}$  Predicted

 $\frac{1}{15}$ 

 $\overline{20}$ 

Table 12. ANOVA results from the best MRSea model for All fulmar as selected by cross-validation

| Variable            | Degrees of Freedom | Chi-square | P-value    |
|---------------------|--------------------|------------|------------|
| Bathymetry          |                    | 13.73      | <<0.001    |
| Distance to Colony  |                    | 14.84      | 0.002      |
| Distance to Turbine |                    | 62.19      | <<0.001    |
| X/Y (location)      | ゚「へ                | 57.36      | $<<$ N NO1 |



Figure 84. Partial dependence plots for significant variables for all fulmar from MRSea. (Clockwise from top left, bathymetry, distance to colony, distance to turbine)

## 3.3.2 Flying Birds Model

| Starting model covariates after VIF-based collinearity removal | Final model covariates after removal by SALSA |  |  |
|--|---|--|--|
| Aspect   | Aspect  |  |  |
| Slope  | Slope   |  |  |
| Bathymetry   | Distance to turbine                           |  |  |
| Sandeel Density  | Daily Mean of Sea Surface Temperature         |  |  |
| Distance to colony   |   |  |  |
| Distance to turbine  |   |  |  |
| Daily Mean of Sea Surface Temperature                          |   |  |  |
|  |   |  |  |

Table 13. Candidate and final covariates for flying fulmar model.

Distribution maps generated using MRSea (Figure 84-88) suggest that flying fulmar are distributed in higher densities during the breeding season (01 April - 15 September), in particular in July and August. In the non-breeding season, density declines considerably, however there are some interesting hotspots in the north and east corners in November 2021. The highest densities were observed consistently throughout the majority of the study area in July 2022, with slightly lower densities occurring along the western edge.



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Model fit was moderate with a marginal R squared value of 0.12 and root mean squared error of 0.21. Cumulative residuals in the model showed that there was a relatively poor relationship between predicted and observed values across most of the range of predicted values, but were typically bounded around 0 across the whole (Figure 105).





Figure 85. Median density of flying fulmar in the survey area for months with sufficient observations between May and September 2021





Figure 86. Median density of flying fulmar in the survey area for months with sufficient observations between November 2021 and March 2022





Figure 87. Median density of flying fulmar in the survey area for months with sufficient observations between April and August 2022





Figure 88. Median density of flying fulmar in the survey area for months with sufficient observations between November 2022 and March 2023





Figure 89. Median density of flying fulmar in the survey area for months with sufficient observations in April 2023





Figure 90. Lower confidence limit of density of flying fulmar in the survey area for months with sufficient observations between May and September 2021





Figure 91. Lower confidence limit of density of flying fulmar in the survey area for months with sufficient observations between November 2021 and March 2022





Figure 92 .Lower confidence limit of density of flying fulmar in the survey area for months with sufficient observations between April and August 2022

120





Figure 93. Lower confidence limit of density of flying fulmar in the survey area for months with sufficient observations between November 2022 and March 2023





Figure 94. Lower confidence limit of density of flying fulmar in the survey area in April 2023





Figure 95. Upper confidence limit of density of flying fulmar in the survey area for months with sufficient observations between May and September 2021

123





Figure 96. Upper confidence limit of density of flying fulmar in the survey area for months with sufficient observations between November 2021 and March 2022





Figure 97. Upper confidence limit of density of flying fulmar in the survey area for months with sufficient observations between April and August 2022

125





Figure 98. Upper confidence limit of density of flying fulmar in the survey area for months with sufficient observations between November 2022 and March 2023





Figure 99. Upper confidence limit of density of flying fulmar in the survey area in April 2023





Figure 100. Spatial coefficient of variation of predicted densities of flying fulmar from MRSea across the survey area for months with sufficient observations between May and September 2021





Figure 101. Spatial coefficient of variation of predicted densities of flying fulmar from MRSea across the survey area for months with sufficient observations between November 2021 and March 2022





Figure 102. Spatial coefficient of variation of predicted densities of flying fulmar from MRSea across the survey area for months with sufficient observations between April and August 2022





Figure 103. Spatial coefficient of variation of predicted densities of flying fulmar from MRSea across the survey area for months with sufficient observations between November 2022 and March 2023





Figure 104. Spatial coefficient of variation of predicted densities of flying fulmar from MRSea across the survey area in April 2023





Figure 105. Autocorrelation test for fulmar density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)







Figure 106. Fitted (MRSea predictions) versus observed counts of flying fulmar









Figure 107. Partial dependence plots for significant variables (distance to turbine) for flying fulmar from MRSea

## 3.4 Guillemot

Table 15. Candidate and final covatiates for guillemot MRSea model

| Starting model covariates after VIF-based collinearity removal | Final model covariates after removal by SALSA |  |
|--|---|--|
| Bathymetry   | Bathymetry                                    |  |
| Sandeel Density  | Sandeel Density                               |  |
| Distance to colony   | Distance to colony                            |  |
| Distance to turbine  | Distance to turbine                           |  |
| Daily mean of Sea Surface Temperature                          | Daily mean of Sea Surface Temperature         |  |

Distribution maps generated using MRSea (Figure 107-111) suggest that guillemot are widely distributed throughout the study area during the breeding season (01 April – 15 August). In the non-breeding season, distribution is much patchier and density declines. The highest densities were observed along the eastern corner of the study area in May 2022.

Broadly, model fit was better for guillemot than for most species, with a marginal R squared value of 0.18 and root mean squared error of 37.33. Cumulative residuals in the model showed that there was a poor relationship between predicted and observed values across most of the range of predicted values, but typically bounded around 0 across the whole (Figure 128).





Figure 108. Median density of all guillemot in the survey area for months with sufficient observations between May and September 2021





Figure 109. Median density of all guillemot in the survey area for months with sufficient observations between October 2021 and February 2022





Figure 110. Median density of all guillemot in the survey area for months with sufficient observations between March and July 2022





Figure 111. Median density of all guillemot in the survey area for months with sufficient observations between August and December 2022





Figure 112. Median density of all guillemot in the survey area for months with sufficient observations between January and April 2023





Figure 113. Lower confidence limit of density of all guillemot in the survey area for months with sufficient observations between May and September 2021




Figure 114. Lower confidence limit of density of all guillemot in the survey area for months with sufficient observations between October 2021 and February 2022





Figure 115. Lower confidence limit of density of all guillemot in the survey area for months with sufficient observations between March and July 2022





Figure 116. Lower confidence limit of density of all guillemot in the survey area for months with sufficient observations between August and December 2022





Figure 117. Lower confidence limit of density of all guillemot in the survey area for months with sufficient observations between January and April 2023





Figure 118. Upper confidence limit of density of all guillemot in the survey area for months with sufficient observations between May and September 2021





Figure 119. Upper confidence limit of density of all guillemot in the survey area for months with sufficient observations between October 2021 and February 2022





Figure 120. Upper confidence limit of density of all guillemot in the survey area for months with sufficient observations between March and July 2022





Figure 121. Upper confidence limit of density of all guillemot in the survey area for months with sufficient observations between August and December 2022





Figure 122. Upper confidence limit of density of all guillemot in the survey area for months with sufficient observations between January and April 2023





Figure 123. Spatial coefficient of variation of predicted densities of all guillemot from MRSea across the survey area for months with sufficient observations between May and September 2021





Figure 124. Spatial coefficient of variation of predicted densities of all guillemot from MRSea across the survey area for months with sufficient observations between October 2021 and February 2022





Figure 125. Spatial coefficient of variation of predicted densities of all guillemot from MRSea across the survey area for months with sufficient observations between March and July 2022





Figure 126. Spatial coefficient of variation of predicted densities of all guillemot from MRSea across the survey area for months with sufficient observations between August and December 2022





Figure 127. Spatial coefficient of variation of predicted densities of all guillemot from MRSea across the survey area for months with sufficient observations between January and April 2023





Figure 128. Autocorrelation test for guillemot density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)







Figure 129. Fitted (MRSea predictions) versus observed counts of all guillemot







Figure 130. Partial dependence plots for significant variables for all guillemot from MRSea, note that distance to turbine was not a significant covariate in the model but was included for reference. (Clockwise from top left: distance to colony, distance to turbine, daily mean of sea surface temperature.)

## 3.5 Razorbill

Table 17. Candidate and final covariates for razorbill MRSea model.

| Starting model covariates after VIF-based collinearity removal | Final model covariates after removal by SALSA |  |
|--|---|--|
| Bathymetry   | Bathymetry                                    |  |
| Sandeel Density  | Sandeel Density                               |  |
| Distance to colony   | Distance to colony                            |  |
| Distance to turbine  | Distance to turbine                           |  |
| Daily mean of Sea Surface Temperature                          | Daily mean of Sea Surface Temperature         |  |

Distribution maps generated using MRSea (Figure 130 - 133) suggest that the highest densities of all razorbill tend to be distributed throughout the southern region of the study area during the breeding season (01 April - 15 August). The highest densities occur during the non-breeding season, in September 2022, where birds are widely distributed across the entire region.

Model fit was moderate with a marginal R squared value of 0.07 and root mean squared error of 0.85. Cumulative residuals in the model showed that there was a moderate relationship between predicted and observed values across most of the range of predicted values, but bounded around 0 across the whole (Figure 147).





Figure 131. Median density of all razorbill in the survey area for months with sufficient observations between May and September 2021





Figure 132. Median density of all razorbill in the survey area for months with sufficient observations between February and July 2022





Figure 133. Median density of all razorbill in the survey area for months with sufficient observations between August 2022 and February 2023





Figure 134. Median density of all razorbill in the survey area for months with sufficient observations between March and April 2023





Figure 135. Lower confidence limit of density of all razorbill in the survey area for months with sufficient observations between May and September 2021





Figure 136. Lower confidence limit of density of all razorbill in the survey area for months with sufficient observations between February and July 2022





Figure 137. Lower confidence limit of density of all razorbill in the survey area for months with sufficient observations between August 2022 and February 2023





Figure 138. Lower confidence limit of density of all razorbill in the survey area for months with sufficient observations between March and April 2023





Figure 139. Upper confidence limit of density of all razorbill in the survey area for months with sufficient observations between May and September 2021





Figure 140. Upper confidence limit of density of all razorbill in the survey area for months with sufficient observations between February and July 2022





Figure 141. Upper confidence limit of density of all razorbill in the survey area for months with sufficient observations between August 2022 and February 2023





Figure 142. Upper confidence limit of density of all razorbill in the survey area for months with sufficient observations between March and April 2023





Figure 143. Spatial coefficient of variation of predicted densities of all razorbill from MRSea across the survey area for months with sufficient observations between May and September 2021





Figure 144. Spatial coefficient of variation of predicted densities of all razorbill from MRSea across the survey area for months with sufficient observations between February and July 2022





Figure 145. Spatial coefficient of variation of predicted densities of all razorbill from MRSea across the survey area for months with sufficient observations between August 2022 and February 2023





Figure 146. Spatial coefficient of variation of predicted densities of all razorbill from MRSea across the survey area for months with sufficient observations between March and April 2023





Figure 147. Autocorrelation test for razorbill density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)







Figure 148. Fitted (MRSea predictions) versus observed counts of all razorbill

| Variable            | Degrees of Freedom | Chi-square | P-value |
|---------------------|--------------------|------------|---------|
| Aspect              |                    | 2.26       | 0.52    |
| Slope               |                    | 3.66       | 0.3     |
| Bathymetry          |                    | 5.12       | 0.163   |
| Sandeel Density     |                    | 3.52       | 0.318   |
| Distance to Colony  |                    | 16.79      | <<0.001 |
| Distance to Turbine |                    | 14.68      | 0.002   |
| X/Y (location)      |                    | 97.17      | <<0.001 |

Table 18. ANOVA results from the best MRSea model for all razorbill as selected by cross-validation




Figure 149. Partial dependence plots for significant variables for all razorbill from MRSea

#### 3.6 Puffin

Table 19. Candidate and final covariates for puffin MRSea model



Distribution maps generated using MRSea (Figure 149-151) suggest that puffin are distributed more generally towards the eastern half of the study area during the breeding season (01 April - 15 August). In the non-breeding season, with the exception of September - October 2021 and September 2022, puffins are mostly absent from the study area (less than 20 observations). The highest densities were observed in the center and southern regions of the study area in August 2021.

Model fit was better for Puffin than for most species, with a marginal R squared value of 0.17 and root mean squared error of 0.77. Cumulative residuals in the model showed that there was a moderate relationship between predicted and observed values across most of the range of predicted value, but the model tended to underpredict for densities above 1.75 birds/km2 (Figure 160).





Figure 150. Median density of all puffin in the survey area for months with sufficient observations between May and September 2021





Figure 151. Median density of all puffin in the survey area for months with sufficient observations between October 2021 and July 2022





Figure 152. Median density of all puffin in the survey area for months with sufficient observations between August and September 2022





Figure 153. Lower confidence limit of density of all puffin in the survey area for months with sufficient observations between May and September 2021





Figure 154. Lower confidence limit of density of all puffin in the survey area for months with sufficient observations between October 2021 and July 2022

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Figure 155. Lower confidence limit of density of all puffin in the survey area for months with sufficient observations between August and September 2022





Figure 156. Upper confidence limit of density of all puffin in the survey area for months with sufficient observations between May and September 2021





Figure 157. Upper confidence limit of density of all puffin in the survey area for months with sufficient observations between October 2021 and July 2022

188





Figure 158. Upper confidence limit of density of all puffin in the survey area for months with sufficient observations between August and September 2022





Figure 159. Spatial coefficient of variation of predicted densities of all puffin from MRSea across the survey area for months with sufficient observations between May and September 2021





Figure 160. Spatial coefficient of variation of predicted densities of all puffin from MRSea across the survey area for months with sufficient observations between October 2021 and July 2022





Figure 161. Spatial coefficient of variation of predicted densities of all puffin from MRSea across the survey area for months with sufficient observations between August and September 2022



Figure 162. Autocorrelation test for puffin density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)





Figure 163. Fitted (MRSea predictions) versus observed counts of all puffin Table 20. ANOVA results from the best MRSea model for all puffin as selected by crossvalidation





Figure 164. Partial dependence plots for significant variables for all puffin from MRSea. (Left: Distance to turbine; right: daily standard deviation of sea surface temperature.)

 $\frac{8}{5}$ 

 $0.2$ 

 $0.4$ 

 $0.6$ 

 $0.8$ 

SST\_sd

# 3.7 Great Black-backed Gull

 $10$ 

 $20$ 

turbine\_distance

30

40

### 3.7.1 All birds model

 $\circ$ 

Table 21. Candidate and final covariates for all great black-backed gull MRSea model



Daily mean of Sea Surface Temperature

Due to a limited number of observations, it was more appropriate to model great black-backed gulls by survey year rather than month. Distribution maps generated using MRSea (Figure 164) suggest that all great black-backed gulls are minimally distributed throughout the study area for both years of surveys. Higher densities are estimated along the central eastern boundary of the study area.

Model fit was moderate with a marginal R squared value of 0.10 and root mean squared error of 0.01. Cumulative residuals in the model showed that there was poor relationship between predicted and observed values where the model tended to under predict above values of ~0.25 birds/km<sup>2</sup> (Figure 169).

<u>n mangan mana sa sa pasan</u>

 $1.2$ 

 $1.0$ 



Figure 165. Median density of all great black-backed gull in the survey area for each year of surveys.



Figure 166. Lower confidence limit of density of all great black-backed gull in the survey area for each year of surveys





Figure 167. Upper confidence limit of density of all great black-backed gulls in the survey area for each year of surveys





Figure 168. Spatial coefficient of variation of predicted densities of all great black-backed gull from MRSea across the survey area for each year of surveys





Figure 169. Autocorrelation test for great black-backed gull density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)





**Fitted Values** 

<sup>10</sup><br>Observed Values

Figure 170. Fitted (MRSea predictions) versus observed counts of all great black-backed gull

| Table 22. ANOVA results from the best MRSea model for all great black-backed gull as selected by |  |  |  |
|--|--|--|--|
| cross-validation   |  |  |  |





Figure 171. Partial dependence plots for significant variables for all great black-backed gull from MRSea (Clockwise from top left: distance to colony, distance to turbine, daily mean of sea surface temperature)

## 3.7.2 Flying birds model

Table 23. Candidate and final covariates for flying great black-backed gull MRSea model



Due to a limited number of observations, it was more appropriate to model great black-backed gulls by survey year rather than month. Distribution maps generated using MRSea (Figure 168) suggest that flying great black-backed gulls are very minimally distributed throughout the study area for both years. Year one has slightly higher densities along the central western boundary of the study area. Year two has lower densities in that region, but higher densities in the northern corner of the study area.

Model fit was poor with a marginal R squared value of 0.004 and root mean squared error of 0.01. Cumulative residuals in the model showed that there was moderate relationship between predicted and observed values across most of the range of predicted values, but residuals were bounded around 0 across the whole (Figure 173).





Figure 172. Median density of flying great black-backed gull in the survey area for each year of surveys.





Figure 173. Lower confidence limit of density of flying great black-backed gull in the survey area for each year of surveys





Figure 174. Upper confidence limit of density of flying great black-backed gulls in the survey area for each year of surveys





Figure 175. Spatial coefficient of variation of predicted densities of flying great black-backed gull from MRSea across the survey area for each year of surveys





Figure 176. Autocorrelation test for great black-backed gull density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0. 0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)





Figure 177. Fitted (MRSea predictions) versus observed counts of flying great black-backed gull









Figure 178. Partial dependence plots for significant variables for flying Great Black-backed Gull from MRSea (left: distance to colony, right: distance to tubine)

## 3.8 Herring Gull

Table 25. Candidate and final covariates for herring gull MRSea model



Due to a limited number of observations, we decided to model herring gulls by survey year rather than month. Distribution maps generated using MRSea (Figure 178) suggest that herring gull occur in minimal numbers throughout the study area. The highest densities were modeled to be along the southern edge during the first year.

Model fit was relatively poor with a marginal R squared value of 0.03 and root mean squared error of 0.03. Cumulative residuals in the model showed that there was a poor relationship between predicted and observed values across most of the range of predicted values, especially above densities of 0.1 (Figure 183). Due to there being so few observations, we had to aggregate across the all the months of surveys and therefore the SD was very high, and our confidence in this model is low.





Figure 179. Median density of all herring gull in the survey area for each year of surveys





Figure 180. Lower confidence limit of density of all herring gull in the survey area for each year of surveys





Figure 181. Upper confidence limit of density of all herring gulls in the survey area for each year of surveys





Figure 182. Spatial coefficient of variation of predicted densities of all herring gull from MRSea across the survey area for each year of surveys




Figure 183. Autocorrelation test for herring gull density surface models when using transect as a blocking feature in MRSea showing no significant correlation. A Runs test on the data prior to using transect as a blocking feature gave a p-value of << 0.0001 (i.e., that the data were significantly autocorrelated when not using a blocking feature)





 $\frac{40}{7}$  $\overline{0.0}$  $\frac{1}{0.3}$ <br>Predicted  $0.5$  $\overline{5000}$ 10000  $15000$  $0.1$  $0.2$  $0.4$ ć Index

Figure 184. Fitted (MRSea predictions) versus observed counts of all herring gull

| Table 26. ANOVA results from the best MRSea model for all herring gull as selected by cross- |  |  |  |  |  |  |
|--|--|--|--|--|--|--|
| validation   |  |  |  |  |  |  |





Figure 185. Partial dependence plots for significant variables for all herring gull from MRSea (Clock wise from top left: Bathymetry, distance to turbine, daily standard deviation of sea surface temperature)



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# **Exploratory Data Analysis**

### I.1 Introduction

In order to properly model the data we first performed an exploratory data analysis to understand how many of each species were in each survey, and how they were spatially distributed throughout the study area. We did this by examining the observation data by species, then mapping species distribution in the study area for the smallest temporal scale that the number of observations would allow. The target temporal scale was by survey, but if the number of observations was too low, we try breeding season, then survey year, and finally for all surveys. MRSea models can fail with a minimal number of observations, therefore we used a threshold of 20 observations per species, Where possible, we mapped the observations plotting behaviours as sitting (red) or flying (blue). We found 624 observations of animals that were not identified to species level, which could supplement some data-poor surveys for some species through apportioning. We were confident we have sufficient data to model seven out of thirteen of the key species (See table I-1). For quillemot, puffin, and razorbill, we did not suggest modelling unique behaviours (sitting/flying) as these species do not fly high enough to be considered at risk for collision with a turbine. We summarise our findings in table I-1, then provide more detail and plots below.

# I.2 Summary Table



Table I-1. Summary of all key species, including whether they have sufficient observations for modelling, at which temporal scale (survey, breeding season, survey year, or all surveys) we suggest that they should be modelled (# surveys that can be modelled / # total surveys are in brackets), and which behaviours we can model. Number of available surveys are in brackets.



# I.3 Results by Species

### I.3.1 Guillemot

Guillemot are the most common species throughout all surveys consisting of almost half of all observations (N=20135). Sufficient observations were recorded for 22/24 surveys. Surveys 8 and 19 (Dec 2021, Nov 2022) do not have enough observations (more than 20), and will not be included in modelling. We could also model guillemot by breeding season (April - 15 August). The August surveys from both years fall within the breeding season (01 Aug 2021, 10 Aug 2022). We suggest modelling all behaviours (flying & sitting) together as guillemot do not fly high enough to be at risk for collision with turbines.







































































### I.3.2 Razorbill

We suggest modelling razorbill (N=2542) by survey for the surveys that had sufficient observations (15/24). We could also model razorbill by breeding season (April - 15 August). The August surveys from both years fall within the breeding season (01 Aug 2021, 10 Aug 2022). We suggest modelling all behaviours (flying & sitting) together as razorbill do not fly high enough to be at risk for collision with turbines.















































### **Razorbill**





#### I.3.3 Kittiwake

We suggest modelling kittiwake (N=5366) by survey for the surveys that had sufficient observations (19/24). We have sufficient observations of kittiwake to include behaviour in the model. We could also model by breeding season (15 April - August). The April 2023 survey does fall outside of the breeding season (4 Apr 2023) and is therefore accounted for as "Non-Breeding." The April 2022 survey (26 Apr 2022), however does fall within the breeding season.


























































#### **Kittiwake**





# I.3.4 Fulmar

We suggest modelling fulmar (N=4611) by survey because the majority of surveys had sufficient observations (22/24). We have sufficient observations of fulmar to include behaviour in the model. We could also model by breeding season (April - 15 September). The September surveys from both years fall within the breeding season (14 Sep 2021, 11 Sep 2022).



















































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**Fulmar** 





#### I.3.5 Puffin

We suggest modelling puffin (N=2054) by survey for the surveys that had sufficient observations (13/24). All but three surveys with enough observations occur during the breeding season, so we could also model puffin by breeding season (April - 15 August). The August surveys from both years fall within the breeding season (01 Aug 2021, 10 Aug 2022). We suggest modelling all behaviours (flying & sitting) together as puffin do not fly high enough to be at risk for collision with turbines.





































**Puffin** 





#### I.3.6 Gannet

We suggest modelling gannet (N=1042) by survey for the surveys that had sufficient observations (10/24). All but two surveys with enough observations occur during the breeding season, so we could also model gannet by breeding season (15 March - September). Neither of the March surveys (02 Mar 2022, 10 Mar 2023) occured within the the breeding season, and are therefore considered "Non-Breeding." For months with enough observations of flying birds we will also suggest including behaviour in the model.















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**Gannet** 





#### I.3.7 Great Black-backed Gull

We suggest modelling Great Black-backed Gull (N=540) by survey-year because only 8/24 surveys had enough observations to model, and all occurred outside of the breeding season (April - August). Some surveys have minimal observations of flying birds, therefore behaviour would be better captured on the larger temporal unit of 'survey year'. We also noticed that there are about 64 observations of this species where the behaviour is listed as "perched". Depending on what the birds are perched upon we may be able to use these observations. For example, if the gulls are perched on turbines, we would not be able to model them with a distance to turbine covariate.



























## I.3.8 Herring Gull

With minimal observations, we will attempt to model herring gull (N=99) for all behaviours and surveys combined. This would constitute a "hot spot" analysis, which would identify potential areas of importance for this species within the study area, however this may be at a temporal scale too large to be very useful.







## I.3.9 Great Skua

Great skua observations were too few to be modeled (N=22).





## I.3.10 Manx Shearwater

Common tern observations were too few to be modelled (N=28).





# I.3.11 Common Gull

Common gull observations were too few to be modelled (N=13).





# I.3.12 Arctic Tern

Arctic tern observations were too few to be modelled (N=15).




# I.3.13 Common Tern

Common tern observations were too few to be modelled (N=3).





# II Random Forests

### II.1 Kittiwake

#### II.1.1 Model assessment

The top five models had RMSE values between 15.662 and 15.656 with R squared values between 0.0485 and 0.0499. These values were higher than the MRSea model, which had an Rsquared value of 0.0426. The best model had an mtry value of 7, with a minimum node size of 80 (Figure II-1, Table II[-1\)](#page-289-0).



Figure II-1: Root mean squared error when varying mtry and min.node.size parameters in random forests

Table II-1: The top 5 models selected by the random forests analysis showing root mean squared error (RMSE) and R-squared values as calculated by 5-fold cross validation

<span id="page-289-0"></span>

#### II.1.2 Variable importance

The top predictor variables were daily sea surface temperature, northing (y.pos), and monthly mean sea surface temperature, followed by survey ID. This shows that most of the signal in the data come from variables that represent temporal variability (Table II[-2\)](#page-289-1).

<span id="page-289-1"></span>Table II-2: The top 5 predictor variables from the random forests model and overall importance





## II.1.3 Distributional response

The partial relationship of distance to turbines on densities of kittiwake demonstrates an inverse relationship in that predicted densities decreased from 0 to 20km distance from turbines. The relationship reaches an inversion point at approximately 23km, where between 20 and 40km, there is a positive relationship between distance to turbine and density (Figure II-2). This suggests that turbines could be attracting kittiwake as the highest densities seems to be in areas nearer to turbines. However, this could simply be that kittiwake were more abundant around turbines for other reasons that we are unable to measure.



Figure II-2: Partial dependence plot of distance to turbine from the random forests model. The red line depicts the loess curve

#### II.1.4 Population estimates

Population estimates derived from the random forests baseline scenario (i.e., Existing turbines) fell well within the range of estimates derived from MRSea, sometimes only differing by 2 - 3%. We are



therefore confident that the random forests kittiwake model is reliable, particularly in the context of the MRSea analysis.

Populations of kittiwake in the survey area were predicted to increase in all three scenarios due to the relationship between distance to turbine and density. The highest population increase was predicted for scenario 3, with percent change from baseline varying from -7 to 47% across all surveys (Table II[-3\)](#page-291-0).

<span id="page-291-0"></span>Table II-3: Population estimates from the random forests models for the 4km buffer zone plus windfarm footprint survey area for the baseline scenario (i.e., based on currently installed turbines) and three potential turbine scenarios. Percent change from baseline is calculated for each scenario



#### II.1.5 Distributions

The broad distribution of kittiwake in the baseline random forests models were nearly identical to those in the MRSea models. For Kittiwake, the models predict the highest densities in the southwest part of the survey area. In almost all cases, the densities of kittiwake increase through the site in relation to the locations of proposed turbines. This distributional response suggests a potential increase in the numbers of birds in the windfarm site (Figures II[-3](#page-292-0) – II[-18\)](#page-307-0).





<span id="page-292-0"></span>Figure II-3: Random forests baseline predictions of Kittiwake from May 2021 to September 2021



Figure II-4: Random forests baseline predictions of Kittiwake from October 2021 to March 2022





Figure II-5: Random forests baseline predictions of Kittiwake from April 2022 to August 2022



Figure II-6: Random forests baseline predictions of Kittiwake from September 2022 to April 2023





Figure II-7: Random forests predictions from turbine scenario 1 of Kittiwake from May 2021 to September 2021 with turbines presented as black dots





Figure II-8: Random forests predictions from turbine scenario 1 of Kittiwake from October 2021 to March 2022 with turbines presented as black dots





Figure II-9: Random forests predictions from turbine scenario 1 of Kittiwake from April 2022 to August 2022 with turbines presented as black dots



Figure II-10: Random forests predictions from turbine scenario 1 of Kittiwake from September 2022 to April 2023 with turbines presented as black dots





Figure II-11: Random forests predictions from turbine scenario 2 of Kittiwake from May 2021 to September 2021 with turbines presented as black dots





Figure II-12: Random forests predictions from turbine scenario 2 of Kittiwake from October 2021 to March 2022 with turbines presented as black dots





Figure II-13: Random forests predictions from turbine scenario 2 of Kittiwake from April 2022 to August 2022 with turbines presented as black dots



Figure II-14: Random forests predictions from turbine scenario 2 of Kittiwake from September 2022 to April 2023 with turbines presented as black dots







Figure II-15: Random forests predictions from turbine scenario 3 of Kittiwake from May 2021 to September 2021 with turbines presented as black dots





Figure II-16: Random forests predictions from turbine scenario 3 of Kittiwake from October 2021 to March 2022 with turbines presented as black dots





Figure II-17: Random forests predictions from turbine scenario 3 of Kittiwake from April 2022 to August 2022 with turbines presented as black dots



Figure II-18: Random forests predictions from turbine scenario 3 of Kittiwake from September 2022 to April 2023 with turbines presented as black dots

# <span id="page-307-0"></span>II.2 Fulmar

#### II.2.1 Model assessment

The top five models had RMSE values between 9.888 and 9.867 with R squared values between 0.144 and 0.141. These values were similar to those from the MRSea model, which had an Rsquared value of 0.1437. The best model had an mtry value of 8, with a minimum node size of 80 (Figure II-19, Table II[-4\)](#page-308-0).



Figure II-19: Root mean squared error when varying mtry and min.node.size parameters in random forests

Table II-4: The top 5 models selected by the random forests analysis showing root mean squared error (RMSE) and R-squared values as calculated by 5-fold cross validation

<span id="page-308-0"></span>

#### II.2.2 Variable importance

The top predictor variables were monthly mean sea surface temperature, northing (y.pos), and survey ID. This shows that most of the signal in the data come from variables that represent temporal variability (Table II[-5\)](#page-308-1).

Table II-5: The top 5 predictor variables from the random forests model and overall importance

<span id="page-308-1"></span>



## II.2.3 Distributional response

The partial relationship of distance to turbines on densities of fulmar demonstrates a displacement effect with fewer birds expected in areas close to turbines. The relationship reaches a slight inflection point at approximately 15km, where the curve begins to flatten (Figure II-20). This does not mean that there is a displacement effect out to 15km or more, simply that birds have been predicted further away from turbines, which could be driven by other factors.



Figure II-20: Partial dependence plot of distance to turbine from the random forests model. The red line depicts the loess curve

# II.2.4 Population estimates

Population estimates derived from the random forests baseline scenario (i.e., Existing turbines) fell well within the range of estimates derived from MRSea, sometimes only differing by 2 - 3%. We are therefore confident that the random forests fulmar model is reliable, particularly in the context of the MRSea analysis.

Populations of fulmar in the survey area were predicted to decrease in all three scenarios due to the relationship between distance to turbine and density. The biggest population decrease was predicted for scenario 3, with percent change from baseline varying from -1 to -22% across all surveys (Table II[-6\)](#page-309-0).

<span id="page-309-0"></span>Table II-6: Population estimates from the random forests models for the 4km buffer zone plus windfarm footprint survey area for the baseline scenario (i.e., based on currently installed turbines) and three potential turbine scenarios. Percent change from baseline is calculated for each scenario



#### II.2.5 Distributions

The broad distribution of fulmar in the baseline random forests models were nearly identical to those in the MRSea models. For fulmar, the models predict the highest densities in the southwest part of the survey area for most months. In all cases, the densities of fulmar decrease through the site in relation to the locations of proposed turbines. This distributional response suggests a potential displacement effect which could push beyond the boundary of the wind farm site (Figures  $II-21 - II-40$  $II-21 - II-40$  $II-21 - II-40$ .





<span id="page-311-0"></span>Figure II-21: Random forests baseline predictions of Fulmar from May 2021 to September 2021



Figure II-22: Random forests baseline predictions of Fulmar from November 2021 to March 2022





Figure II-23: Random forests baseline predictions of Fulmar from April 2022 to August 2022



Figure II-24: Random forests baseline predictions of Fulmar from September 2022 to February 2023





Figure II-25: Random forests baseline predictions of Fulmar from March 2023 to April 2023





Figure II-26: Random forests predictions from turbine scenario 1 of Fulmar from May 2021 to September 2021 with turbines presented as black dots





Figure II-27: Random forests predictions from turbine scenario 1 of Fulmar from November 2021 to March 2022 with turbines presented as black dots





Figure II-28: Random forests predictions from turbine scenario 1 of Fulmar from April 2022 to August 2022 with turbines presented as black dots



Figure II-29: Random forests predictions from turbine scenario 1 of Fulmar from September 2022 to February 2023 with turbines presented as black dots





Figure II-30: Random forests predictions from turbine scenario 1 of Fulmar from March 2023 to April 2023 with turbines presented as black dots





Figure II-31: Random forests predictions from turbine scenario 2 of Fulmar from May 2021 to September 2021 with turbines presented as black dots





Figure II-32: Random forests predictions from turbine scenario 2 of Fulmar from November 2021 to March 2022 with turbines presented as black dots





Figure II-33: Random forests predictions from turbine scenario 2 of Fulmar from April 2022 to August 2022 with turbines presented as black dots


Figure II-34: Random forests predictions from turbine scenario 2 of Fulmar from September 2022 to February 2023 with turbines presented as black dots





Figure II-35: Random forests predictions from turbine scenario 2 of Fulmar from March 2023 to April 2023 with turbines presented as black dots





Figure II-36: Random forests predictions from turbine scenario 3 of Fulmar from May 2021 to September 2021 with turbines presented as black dots





Figure II-37: Random forests predictions from turbine scenario 3 of Fulmar from November 2021 to March 2022 with turbines presented as black dots





Figure II-38: Random forests predictions from turbine scenario 3 of Fulmar from April 2022 to August 2022 with turbines presented as black dots



Figure II-39: Random forests predictions from turbine scenario 3 of Fulmar from September 2022 to February 2023 with turbines presented as black dots





Figure II-40: Random forests predictions from turbine scenario 3 of Fulmar from March 2023 to April 2023 with turbines presented as black dots

# II.3 Gannet

## II.3.1 Model assessment

The top five models had RMSE values that were all around 3.835 with R squared values between 0.0442 and 0.0448. These R squared values were better than those from the MRSea model, which had an R-squared value of 0.0395. The best model had an mtry value of 6, with a minimum node size of 80 (Figure II-41, Table II[-7\)](#page-331-0).



Figure II-41: Root mean squared error when varying mtry and min.node.size parameters in random forests

Table II-7: The top 5 models selected by the random forests analysis showing root mean squared error (RMSE) and R-squared values as calculated by 5-fold cross validation

<span id="page-331-0"></span>

## II.3.2 Variable importance

The top predictor variables were distance to colony, density of sandeel, northing (y.pos), bathymetry and presence of sandeel. This shows that most of the signal in the data come from variables that represent ecological parameters related to prey (Table II[-8\)](#page-331-1).

Table II-8: The top 5 predictor variables from the random forests model and overall importance

<span id="page-331-1"></span>

| <b>Variable</b>     | Overall importance |
|---------------------|--------------------|
| dist2col            | 100.00             |
| sandeel_pr_density  | 93.43              |
| y.pos               | 85.49              |
| Bathymetry          | 71.91              |
| sandeel_pr_presence | 69.76              |



## II.3.3 Distributional response

The partial relationship of distance to turbines on densities of gannet demonstrates that there was an inverse relationship between density and distance to turbine from 0 to approximately 18km away from the turbines. After this, the curve has an inflection point where the relationship reverses, and more birds are predicted further away. (Figure II-42). The relationship between 0 - 18km only contributes to a partial dependence of approximately 0.03 and when comparing to the overall scale of the figure of 0.64 to 0.72, it suggests little evidence of a displacement effect. However, this relationship could be driven by other environmental covariates that we were unable to measure.



Figure II-42: Partial dependence plot of distance to turbine from the random forests model. The red line depicts the loess curve

## II.3.4 Population estimates

Population estimates derived from the random forests baseline scenario (i.e., Existing turbines) fell well within the range of estimates derived from MRSea, sometimes only differing by 0.6 to 2%. We are therefore confident that the random forests gannet model is reliable, particularly in the context of the MRSea analysis.

There was no consistency in population increases or decreases between turbine scenarios vesus baseline. However, the largest variations were in scenario 3 (ranging from -9.75% to 19.94%) (Table II[-9\)](#page-332-0).

<span id="page-332-0"></span>Table II-9: Population estimates from the random forests models for the 4km buffer zone plus windfarm footprint survey area for the baseline scenario (i.e., based on currently installed turbines) and three potential turbine scenarios. Percent change from baseline is calculated for each scenario



## II.3.5 Distributions

The broad distribution of gannet in the baseline random forests models were nearly identical to those in the MRSea models. For gannet, the models predict the highest densities in the southern part of the survey area for most months (with the exception of October 2021). The densities of gannet change variably between surveys in relation to the locations of proposed turbines. This distributional response suggests a variable displacement effect (Figures II[-43](#page-334-0) – II[-50\)](#page-341-0).





<span id="page-334-0"></span>Figure II-43: Random forests baseline predictions of Gannet from June 2021 to October 2021





Figure II-44: Random forests baseline predictions of Gannet from May 2022 to October 2022







Figure II-45: Random forests predictions from turbine scenario 1 of Gannet from June 2021 to October 2021 with turbines presented as black dots





Figure II-46: Random forests predictions from turbine scenario 1 of Gannet from May 2022 to October 2022 with turbines presented as black dots

II.3.5.3 Turbine scenario 2



Figure II-47: Random forests predictions from turbine scenario 2 of Gannet from June 2021 to October 2021 with turbines presented as black dots





Figure II-48: Random forests predictions from turbine scenario 2 of Gannet from May 2022 to October 2022 with turbines presented as black dots







Figure II-49: Random forests predictions from turbine scenario 3 of Gannet from June 2021 to October 2021 with turbines presented as black dots





Figure II-50: Random forests predictions from turbine scenario 3 of Gannet from May 2022 to October 2022 with turbines presented as black dots

# <span id="page-341-0"></span>II.4 Puffin

## II.4.1 Model assessment

The top five models had RMSE values between 3.999 and 4.011 with R squared values between 0.163 and 0.159. The R squared values were slightly lower than those from the MRSea model, which had an R-squared value of 0.1669. The best model had an mtry value of 8, with a minimum node size of 80 (Figure II-51, Table II[-10\)](#page-342-0).



Figure II-51: Root mean squared error when varying mtry and min.node.size parameters in random forests

Table II-10: The top 5 models selected by the random forests analysis showing root mean squared error (RMSE) and R-squared values as calculated by 5-fold cross validation

<span id="page-342-0"></span>

## II.4.2 Variable importance

The top predictor variables were survey ID, standard deviation of monthly sea surface temperature, and monthly mean sea surface temperature. This shows that most of the signal in the data come from variables that represent temporal variability (Table II[-11\)](#page-342-1).

Table II-11: The top 5 predictor variables from the random forests model and overall importance

<span id="page-342-1"></span>

| Variable    | Overall importance |
|-------------|--------------------|
| SurveyID.Q  | 100.00             |
| SST sd      | 98.34              |
| SurveyID^13 | 7972               |
| SST mean    | 74.95              |
| SurveyID^23 | 70.23              |



## II.4.3 Distributional response

The partial relationship of distance to turbines on densities of puffin demonstrates a displacement effect with fewer birds expected in areas close to turbines. The relationship reaches a slight inflection point at approximately 20km, where the curve begins to flatten (Figure II-52). This does not indicate that there is a displacement effect out to 20km, simply that the model predicts more birds further away from the turbines, which could be due to relationships with other covariates.



Figure II-52: Partial dependence plot of distance to turbine from the random forests model. The red line depicts the loess curve

# II.4.4 Population estimates

Population estimates derived from the random forests baseline scenario (i.e., Existing turbines) fell well within the range of estimates derived from MRSea, sometimes only differing by 0.6 - 2%. We are therefore confident that the random forests puffin model is reliable, particularly in the context of the MRSea analysis. There were notably higher populations in August 2021 and September 2022, likely associated with post-breeding dispersal.

Populations of puffin in the survey area were mostly predicted to decrease in all three scenarios due to the relationship between distance to turbine and density, with the exception of the September 2021, where the population was predicted to increase by 11.69% in scenarios 2 and 3. The biggest population change was predicted for scenario 3, with percent change from baseline varying from -14% to 11.69% across all surveys (Table II[-12\)](#page-344-0).





<span id="page-344-0"></span>Table II-12: Population estimates from the random forests models for the 4km buffer zone plus windfarm footprint survey area for the baseline scenario (i.e., based on currently installed turbines) and three potential turbine scenarios. Percent change from baseline is calculated for each scenario

## II.4.5 Distributions

The broad distribution of puffin in the baseline random forests models were nearly identical to those in the MRSea models. For most surveys, the models predicted the highest densities in the eastern and southern parts of the survey area, with the exception of August 2021 and September 2022 where high densities of birds were identified through the whole region. In all cases, the densities of puffin decrease through the site in relation to the locations of proposed turbines. This distributional response suggests a potential displacement effect which could push individuals beyond the boundary of the wind farm site (Figures II[-53](#page-345-0) – II[-64\)](#page-356-0).





<span id="page-345-0"></span>Figure II-53: Random forests baseline predictions of Puffin from May 2021 to September 2021





Figure II-54: Random forests baseline predictions of Puffin from October 2021 to July 2022





Figure II-55: Random forests baseline predictions of Puffin from August 2022 to September 2022





Figure II-56: Random forests predictions from turbine scenario 1 of Puffin from May 2021 to September 2021 with turbines presented as black dots





Figure II-57: Random forests predictions from turbine scenario 1 of Puffin from October 2021 to July 2022 with turbines presented as black dots





Figure II-58: Random forests predictions from turbine scenario 1 of Puffin from August 2022 to September 2022 with turbines presented as black dots

II.4.5.3 Turbine scenario 2



Figure II-59: Random forests predictions from turbine scenario 2 of Puffin from May 2021 to September 2021 with turbines presented as black dots





Figure II-60: Random forests predictions from turbine scenario 2 of Puffin from October 2021 to July 2022 with turbines presented as black dots

![](_page_353_Figure_0.jpeg)

![](_page_353_Figure_1.jpeg)

Figure II-61: Random forests predictions from turbine scenario 2 of Puffin from August 2022 to September 2022 with turbines presented as black dots

![](_page_354_Figure_1.jpeg)

![](_page_354_Figure_2.jpeg)

Figure II-62: Random forests predictions from turbine scenario 3 of Puffin from May 2021 to September 2021 with turbines presented as black dots

![](_page_355_Figure_0.jpeg)

![](_page_355_Figure_1.jpeg)

Figure II-63: Random forests predictions from turbine scenario 3 of Puffin from October 2021 to July 2022 with turbines presented as black dots

![](_page_356_Figure_0.jpeg)

![](_page_356_Figure_1.jpeg)

Figure II-64: Random forests predictions from turbine scenario 3 of Puffin from August 2022 to September 2022 with turbines presented as black dots

# <span id="page-356-0"></span>II.5 Razorbill

## II.5.1 Model assessment

The top five models had RMSE values between 4.125 and 4.127 with R squared values between 0.0892 and 0.0903. The R squared values were higher than those from the MRSea model, which had an R-squared value of 0.0776. The best model had an mtry value of 6, with a minimum node size of 80 (Figure II-65, Table II[-13\)](#page-357-0).

![](_page_357_Figure_0.jpeg)

Figure II-65: Root mean squared error when varying mtry and min.node.size parameters in random forests

Table II-13: The top 5 models selected by the random forests analysis showing root mean squared error (RMSE) and R-squared values as calculated by 5-fold cross validation

<span id="page-357-0"></span>![](_page_357_Picture_161.jpeg)

## II.5.2 Variable importance

The top predictor variables were monthly mean sea surface temperature, northing| (y.pos) and survey ID. This shows that most of the signal in the data come from variables that represent temporal variability (Table II[-14\)](#page-357-1).

Table II-14: The top 5 predictor variables from the random forests model and overall importance

<span id="page-357-1"></span>![](_page_357_Picture_162.jpeg)

![](_page_358_Picture_115.jpeg)

## II.5.3 Distributional response

The partial relationship of distance to turbines on densities of razorbill demonstrates that higher densities of birds are predicted in areas close to turbines. The relationship reaches an inflection point at approximately 18km (Figure II-66). However, this effect seems quite small overall with the relative contribution towards predictions ranging between 1.05 and 1.13 (0.08). This does not necessarily mean that razorbill are attracted to turbines, but that they are predicted to be occurring at higher densities closer to turbines, which could be due to relationships with other environmental features.

![](_page_358_Figure_3.jpeg)

Figure II-66: Partial dependence plot of distance to turbine from the random forests model. The red line depicts the loess curve

## II.5.4 Population estimates

Population estimates derived from the random forests baseline scenario (i.e., Existing turbines) fell well within the range of estimates derived from MRSea, sometimes only differing by 1 - 3%. We are therefore confident that the random forests razorbill model is reliable, particularly in the context of the MRSea analysis.

Populations of razorbill in the survey area increased and decreased variably in all three scenarios due to the relationship between distance to turbine and density. The biggest population change was predicted for scenario 3, with percent change from baseline varying from -5% to 46% across all surveys (Table II[-15\)](#page-359-0).

![](_page_359_Picture_277.jpeg)

![](_page_359_Picture_278.jpeg)

<span id="page-359-0"></span>Table II-15: Population estimates from the random forests models for the 4km buffer zone plus windfarm footprint survey area for the baseline scenario (i.e., based on currently installed turbines) and three potential turbine scenarios. Percent change from baseline is calculated for each scenario

## II.5.5 Distributions

The broad distribution of razorbill in the baseline random forests models were nearly identical to those in the MRSea models. For most surveys, the models predicted the highest densities in the southern parts of the survey area, with the exception of May and July 2022 where high densities of birds were identified through the whole region. The distributional response was varied between the surveys, but is most extreme in scenario 3 (Figures II[-67](#page-360-0) – II[-82\)](#page-375-0).




Figure II-67: Random forests baseline predictions of Razorbill from May 2021 to September 2021





Figure II-68: Random forests baseline predictions of Razorbill from February 2022 to July 2022



Figure II-69: Random forests baseline predictions of Razorbill from August 2022 to February 2023





Figure II-70: Random forests baseline predictions of Razorbill from March 2023 to April 2023





Figure II-71: Random forests predictions from turbine scenario 1 of Razorbill from May 2021 to September 2021 with turbines presented as black dots





Figure II-72: Random forests predictions from turbine scenario 1 of Razorbill from February 2022 to July 2022 with turbines presented as black dots



Figure II-73: Random forests predictions from turbine scenario 1 of Razorbill from August 2022 to February 2023 with turbines presented as black dots





Figure II-74: Random forests predictions from turbine scenario 1 of Razorbill from March 2023 to April 2023 with turbines presented as black dots

II.5.5.3 Turbine scenario 2



Figure II-75: Random forests predictions from turbine scenario 2 of Razorbill from May 2021 to September 2021 with turbines presented as black dots





Figure II-76: Random forests predictions from turbine scenario 2 of Razorbill from February 2022 to July 2022 with turbines presented as black dots



Figure II-77: Random forests predictions from turbine scenario 2 of Razorbill from August 2022 to February 2023 with turbines presented as black dots





Figure II-78: Random forests predictions from turbine scenario 2 of Razorbill from March 2023 to April 2023 with turbines presented as black dots





Figure II-79: Random forests predictions from turbine scenario 3 of Razorbill from May 2021 to September 2021 with turbines presented as black dots





Figure II-80: Random forests predictions from turbine scenario 3 of Razorbill from February 2022 to July 2022 with turbines presented as black dots



Figure II-81: Random forests predictions from turbine scenario 3 of Razorbill from August 2022 to February 2023 with turbines presented as black dots





Figure II-82: Random forests predictions from turbine scenario 3 of Razorbill from March 2023 to April 2023 with turbines presented as black dots

# II.6 Guillemot

#### II.6.1 Model assessment

The top five models had RMSE values between 33.386 and 33.095 with R squared values between 0.215 and 0.199. The R squared values were higher than those from the MRSea model, which had an R-squared value of 0.1843. The best model had an mtry value of 8, with a minimum node size of 80 (Figure II-83, Table II[-16\)](#page-376-0).



Figure II-83: Root mean squared error when varying mtry and min.node.size parameters in random forests

Table II-16: The top 5 models selected by the random forests analysis showing root mean squared error (RMSE) and R-squared values as calculated by 5-fold cross validation

<span id="page-376-0"></span>

#### II.6.2 Variable importance

The top predictor variables were northing (y.pos), survey ID, monthly mean sea surface temperature, and standard deviation of monthly sea surface temperature. This shows that most of the signal in the data come from variables that represent temporal variability (Table II[-17\)](#page-376-1).

Table II-17: The top 5 predictor variables from the random forests model and overall importance

<span id="page-376-1"></span>

| Variable                | Overall importance |
|-------------------------|--------------------|
| y.pos                   | 100.00             |
| SurveyID^10             | 71.47              |
| SST_mean                | 62.77              |
| SurveyID <sup>^12</sup> | 62.52              |
| SST_sd                  | 61.24              |



### II.6.3 Distributional response

The partial relationship of distance to turbines on densities of guillemot demonstrates a slight inverse relationship between 0 - 10 km (Figure II-84). The relationship with the partial dependence from 0 to 20 km varies between 6.3 and 6.5, whereas the overall scale is from 6.3 to 7.6 which indicates that it is not as strong an effect within that x-axis range. If interpreting this as the figure depicts, there is relatively no distributional response associated with turbines for guillemot.



Figure II-84: Partial dependence plot of distance to turbine from the random forests model. The red line depicts the loess curve

## II.6.4 Population estimates

Population estimates derived from the random forests baseline scenario (i.e., Existing turbines) mostly fell well within the range of estimates derived from MRSea, sometimes only differing by 2 - 3%. The largested discrepancy was the July 2021 survey which predicted a median population estimate 16704 birds, while the random forests analysis predicted 22587 birds. However, this estimate of 22587 fell within the confidence limits of the MRSea analysis. We are therefore confident that the random forests guillemot model is reliable, particularly in the context of the MRSea analysis.

Populations of guillemot in the survey area were mostly predicted to decrease in all three scenarios due to the relationship between distance to turbine and density, with the exception of May 2022, where the population was predicted to increase by 26.99%, and December 2022 where the population was predicted to increase by 11.94% in scenario 3. The biggest population change was predicted for scenario 3, with percent change from baseline varying from -19% to 27% across all surveys (Table II[-18\)](#page-378-0).





<span id="page-378-0"></span>Table II-18: Population estimates from the random forests models for the 4km buffer zone plus windfarm footprint survey area for the baseline scenario (i.e., based on currently installed turbines) and three potential turbine scenarios. Percent change from baseline is calculated for each scenario

#### II.6.5 Distributions

The broad distribution of guillemot in the baseline random forests models were nearly identical to those in the MRSea models. For most surveys, the models predicted the highest densities in the south western parts of the survey area, although the species was mostly ubiquitous throughout the site in all surveys. In nearly all cases, the densities of guillemot decrease through the site in relation to the locations of proposed turbines. This distributional response suggests a potential displacement effect which could push individuals beyond the boundary of the wind farm site (Figures II[-85](#page-379-0) – II[-104\)](#page-398-0).





<span id="page-379-0"></span>Figure II-85: Random forests baseline predictions of Guillemot from May 2021 to September 2021



Figure II-86: Random forests baseline predictions of Guillemot from October 2021 to February 2022





Figure II-87: Random forests baseline predictions of Guillemot from March 2022 to July 2022



Figure II-88: Random forests baseline predictions of Guillemot from August 2022 to December 2022



Figure II-89: Random forests baseline predictions of Guillemot from January 2023 to April 2023





Figure II-90: Random forests predictions from turbine scenario 1 of Guillemot from May 2021 to September 2021 with turbines presented as black dots





Figure II-91: Random forests predictions from turbine scenario 1 of Guillemot from October 2021 to February 2022 with turbines presented as black dots





Figure II-92: Random forests predictions from turbine scenario 1 of Guillemot from March 2022 to July 2022 with turbines presented as black dots



Figure II-93: Random forests predictions from turbine scenario 1 of Guillemot from August 2022 to December 2022 with turbines presented as black dots



Figure II-94: Random forests predictions from turbine scenario 1 of Guillemot from January 2023 to April 2023 with turbines presented as black dots

II.6.5.3 Turbine scenario 2



Figure II-95: Random forests predictions from turbine scenario 2 of Guillemot from May 2021 to September 2021 with turbines presented as black dots





Figure II-96: Random forests predictions from turbine scenario 2 of Guillemot from October 2021 to February 2022 with turbines presented as black dots





Figure II-97: Random forests predictions from turbine scenario 2 of Guillemot from March 2022 to July 2022 with turbines presented as black dots



Figure II-98: Random forests predictions from turbine scenario 2 of Guillemot from August 2022 to December 2022 with turbines presented as black dots



Figure II-99: Random forests predictions from turbine scenario 2 of Guillemot from January 2023 to April 2023 with turbines presented as black dots





Figure II-100: Random forests predictions from turbine scenario 3 of Guillemot from May 2021 to September 2021 with turbines presented as black dots





Figure II-101: Random forests predictions from turbine scenario 3 of Guillemot from October 2021 to February 2022 with turbines presented as black dots




Figure II-102: Random forests predictions from turbine scenario 3 of Guillemot from March 2022 to July 2022 with turbines presented as black dots

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Figure II-103: Random forests predictions from turbine scenario 3 of Guillemot from August 2022 to December 2022 with turbines presented as black dots

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Figure II-104: Random forests predictions from turbine scenario 3 of Guillemot from January 2023 to April 2023 with turbines presented as black dots



## III Hot spot analysis

## III.1 Results

The hot-spot analysis revealed no persistent hot spots across the site for the entire set of density surface models. Most of the site was found to have moderate relative densities (see table 2 for a definition of the classes). The south-west portion of the site was the only region that had hot spots, though they were variable in nature. This is in line with the broad distributional patterns noted in the density surface models (Figure III-1).



Figure III-1: Classified grid cells from the hot-spot analysis in the Caledonia Offshore wind farm area.



Data has the power to change how we interact with the natural world; we can explore its complexities and nuances then make changes for the better

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