



# Habitat Partitioning in Sympatric Delphinids Around the Falkland Islands: Predicting Distributions Based on a Limited Data Set

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Spatial modelling based on line transect data is a standard method for characterising marine mammal distributions and habitat preference. However, collecting the data required is costly and may be difficult in remote areas. Models based on habitat variables offer the potential to predict where the species will occur in areas outside the area of a localised survey. This has important implications for spatial management where decisions have to be made that affect wide areas over which comprehensive survey efforts may not be feasible. This study demonstrates that it is possible, using a spatially limited data set, to characterise habitat use and predict the distribution of two poorly known sympatric delphinids around the Falkland (Malvinas) Islands (FI), Commerson's dolphins (*Cephalorhynchus commersonii*) and Peale's dolphins (*Lagenorhynchus australis*). We used a Hurdle model approach to investigate the relationship between dolphin sightings (from a spatially restricted boat-based line transect survey) and environmental covariates. We then used the modelled relationships to predict the distribution and relative abundance of Commerson's and Peale's dolphins over the entire FI inshore waters. We compared the predicted distribution maps to independent sightings from a subsequent island-wide aerial line transect survey, and found a close match between predicted and observed distributions. Commerson's dolphins preferred nearshore waters with strong tidal mixing and were most numerous close to river mouths and in upper inlets or channels. In contrast, Peale's dolphins preferred deeper, well-stratified areas further from shore as well as nearshore waters with extensive kelp beds. While the two dolphin species are often considered sympatric, our results indicate fine-scale habitat partitioning based on specific habitat preferences, which is important to consider in further studies and marine spatial planning. We provide several methodological refinements to prepare transect sighting data for spatial analysis and implement Hurdle models more easily using the new "dshm" R-package. We also

show the usefulness of such refinements applied to a carefully chosen spatially limited dataset as a cost-effective approach to elucidating species distribution patterns. Our methodology and software implementations can be easily applied to transect survey data of other marine and terrestrial taxa.

**Keywords:** Falkland Islands, habitat partitioning, Hurdle model, line transect survey, predictive species distribution modelling, Commerson's dolphin, Peale's dolphin

## INTRODUCTION

Understanding how environmental factors shape species distributions is a key concept in ecology. Characterising species-environment relationships can facilitate accurate predictions of species' spatial distributions (Guisan and Zimmermann, 2000), and help identify ecological niches (Austin et al., 1990; Barragán-Barrera et al., 2019). Knowing where animals occur within a specific region is also crucial for effective management and for the implementation of conservation measures (Guisan et al., 2013). Ideally, the entire area of interest is surveyed in order to estimate the distribution of animals. However, survey data may be very costly to obtain and are restricted to times and areas where direct observations are possible. To overcome this spatial limitation, such data can be analysed together with associated habitat covariates, and the resulting models used to predict where animals could be, as long as the surveyed area has covered similar environmental conditions to that of the region of interest (Guisan and Zimmermann, 2000; Mannocci et al., 2014).

One such area where surveys in the complex marine environment are challenging, is around the Falkland (also known as Malvinas) Islands (FI) in the South Atlantic. The FI are situated on the Patagonian Shelf and include the two main islands, East Falkland and West Falkland, and 776 smaller islands. The coast is convoluted and lined by extensive kelp beds (*Macrocystis* and *Lessonia* spp.), which can extend up to 10 km offshore, and cover an area of around 1,600 km<sup>2</sup>. These inshore waters are important to a community of marine animals including meso predators such as Commerson's dolphins (*Cephalorhynchus commersonii*) and Peale's dolphins (*Lagenorhynchus australis*). The current lack of information on the distribution and habitat use of Commerson's and Peale's dolphins in FI waters limits the proper assessment of their conservation status at national, regional and international levels as well as their inclusion in local and regional management plans (Augé et al., 2018). Both are key species in the FI Species Action Plans (Otley, 2008), requiring population assessments and baseline understanding of their ecology so that they can be included in wider marine spatial planning initiatives. The shelf area around the FI has experienced a substantial increase in anthropogenic activities including oil exploration, shipping traffic, commercial fishing, aquaculture, and tourism (Otley, 2008; Augé et al., 2018). Such activities often negatively affect marine ecosystem stability and functioning, and thus should be carefully assessed and managed, requiring information on spatial overlap between species distributions and anthropogenic activities. Commercial fishing and aquaculture are still in their infancy in the FI inshore waters, providing a relatively rare

opportunity to investigate species distributions prior to the onset of these potentially damaging human activities.

Commerson's and Peale's dolphins inhabit continental shelf waters ranging from shallow nearshore to offshore waters up to 300 m deep (Dellabianca et al., 2016; Cipriano, 2018), and are often considered to be sympatric species in the South Atlantic, including the FI. Peale's dolphins make extensive use of kelp beds for foraging (Viddi and Lescauwaet, 2005), and their diet is known to consist of bottom and demersal fishes as well as octopus and squid species living on the continental shelf or associated with kelp (Schivavini et al., 1997). A similar diet composition was also found for Commerson's dolphins inhabiting the same region (Ricciardelli et al., 2010). Commerson's dolphins have been observed in estuarine zones feeding in shallow waters near river mouths where their distribution is influenced by tidal patterns (Garaffo et al., 2011; de Castro et al., 2013). Tidal fronts are a dominant oceanographic feature around the FI (Acha et al., 2004). Frontal areas are associated with high primary productivity, and usually attract diverse low and high trophic level consumers, including meso or top predators such as dolphins (Davis et al., 2002; Dellabianca et al., 2012). Modelled distribution for both Commerson's and Peale's dolphins matched previously described frontal zones on the Patagonian shelf (Acha et al., 2004; Dellabianca et al., 2016), but such relationships have not yet been investigated in other parts of the species' ranges.

The aim of this study was to make use of a spatially limited dataset to investigate habitat use of Commerson's and Peale's dolphins in the entire FI inshore waters. Firstly, we describe the relationships between habitat covariates and sightings of dolphins based on data from a systematic small-scale survey. Secondly, we predict the spatial distributions of both species throughout the FI inshore waters, using a Hurdle model approach. We use the model predictions to identify key areas for both species, and examine spatial overlap between preferred habitats. We demonstrate the usefulness of our methodology for predictive habitat modelling, which should be relevant to other complex areas and data-poor species.

## MATERIALS AND METHODS

### Sighting Data

Sightings of Commerson's and Peale's dolphins were collected during a 10-day vessel-based line transect survey conducted from the 26th of February to the 7th of March 2014 and implemented as part of the Darwin Initiative UK Overseas Territories Challenge Fund Project "Inshore Cetaceans of the

Falkland Islands” (Project Reference: EIDCF019; Thomsen, 2013). Surveys were conducted with a 15 m converted fishing vessel (MV Condor) which housed the observation team. During surveys, two dedicated observers scanned continuously ahead of the vessel and out to 90° of the track line using the naked eye aided by 7 × 50 binoculars when needed. For each sighting, species identification, estimated group size, behaviour, distance and angle to the sighting (using measuring sticks and angle boards, respectively) as well as sighting conditions were reported to a third observer inside the wheelhouse who recorded these data in the bespoke software Logger (Gillespie et al., 2010) along with the date, time and the vessel’s GPS location.

Due to time and budget constraints the survey could only cover a small but representative part of the FI inshore waters (Thomsen, 2013). The survey area comprised the different coastal habitat types around the FI, and encompassed approximately 2,000 km<sup>2</sup> split into nine smaller regions or strata containing 72 transect lines (extending from 0.3 to 10 km offshore), located in the northern part of West Falkland, the northern part of Falkland Sound, and the southern part of East Falkland (**Figure 1**). The survey was designed using the software Distance v.6 (Thomas et al., 2010). Average vessel steaming speed was 8 knots (Thomsen, 2014).

Aerial line-transect surveys with a fixed wing aircraft were conducted during nine days from March to May 2017 using two trained observers and one data recorder. Software Distance v.6.2. was used to design the survey to cover the entire FI inshore waters resulting in 217 parallel transect lines (extending from the coast to 10 km offshore; **Supplementary Figure 1**). The aircraft flew at a height of 150 m at a speed of 90 knots (~167 km/h).

All field work was carried out in accordance with the guidelines for the Treatment of Marine Mammals in Field Research (Society for Marine Mammalogy) and approved by the University of St Andrews, School of Biology Animal Ethics Committee.

## Habitat Covariates

To model dolphin habitat preference, we considered six environmental covariates: distance to coast (DC), distance to kelp (DK), distance to 100 m isobath (D100), distance to main river mouths (DRM), water depth, and water column stratification index (SI). These abiotic variables were considered proxies characterising biologically relevant yet unknown links between dolphins and their environment (e.g., via prey distribution) and had been shown or were suspected to be of importance to either species (Garaffo et al., 2011; de Castro et al., 2013; Dellabianca et al., 2016; Heinrich et al., 2019).

All distances were calculated as least cost distances using the “gdistance” R-package (van Etten, 2017). Water depth was obtained from the 2014 GEBCO 30 arc-second gridded bathymetry (equivalent to a spatial resolution of 0.9 km; Weatherall et al., 2015). We applied bilinear interpolation in QGIS (v 2.16.2) to resample covariates at a resolution of 50 m (Qgis Development Team, 2017).

We obtained kelp and coastline data from the South Atlantic Environmental Research Institute (SAERI) and validated them with satellite images in Google Earth Pro. The 100 m isobath

was extracted from the GEBCO gridded bathymetry while main river mouth locations were obtained from a nautical chart and validated in Google Earth Pro.

The water column stratification index (SI) used in this study was calculated using depth and the mean depth-averaged tidal speed, and indicated if the water column was more likely to be stratified (two layers) or vertically homogeneous (mixed) (Simpson and Hunter, 1974). The SI values were calculated using depths from the GEBCO 30 arc-second bathymetry and tidal currents predicted by the 30 arc-second Patagonian Shelf model using the Tidal Model Driver (Egbert and Erofeeva, 2002). The resulting stratification indices represented a static field unlikely to reflect exactly the highly dynamic *in situ* conditions at the time of the survey, and should be interpreted as representing the general stratification characteristics of a particular area.

Characteristically on the continental shelves SI values range from 1 to 5 with the boundary between stratified and mixed layers called tidal front (Simpson and Hunter, 1974; Hill et al., 2008). Stratification index values higher than 2.7 indicate a high possibility for a stratified water column, while lower values suggest a homogeneous (mixed) water column. Changes in water temperature and salinity across tidal fronts and between the top and bottom layers in the stratified water column are large enough to be detectable by marine mammals, and therefore make them a biologically relevant oceanographic feature for our study (Bost et al., 2009).

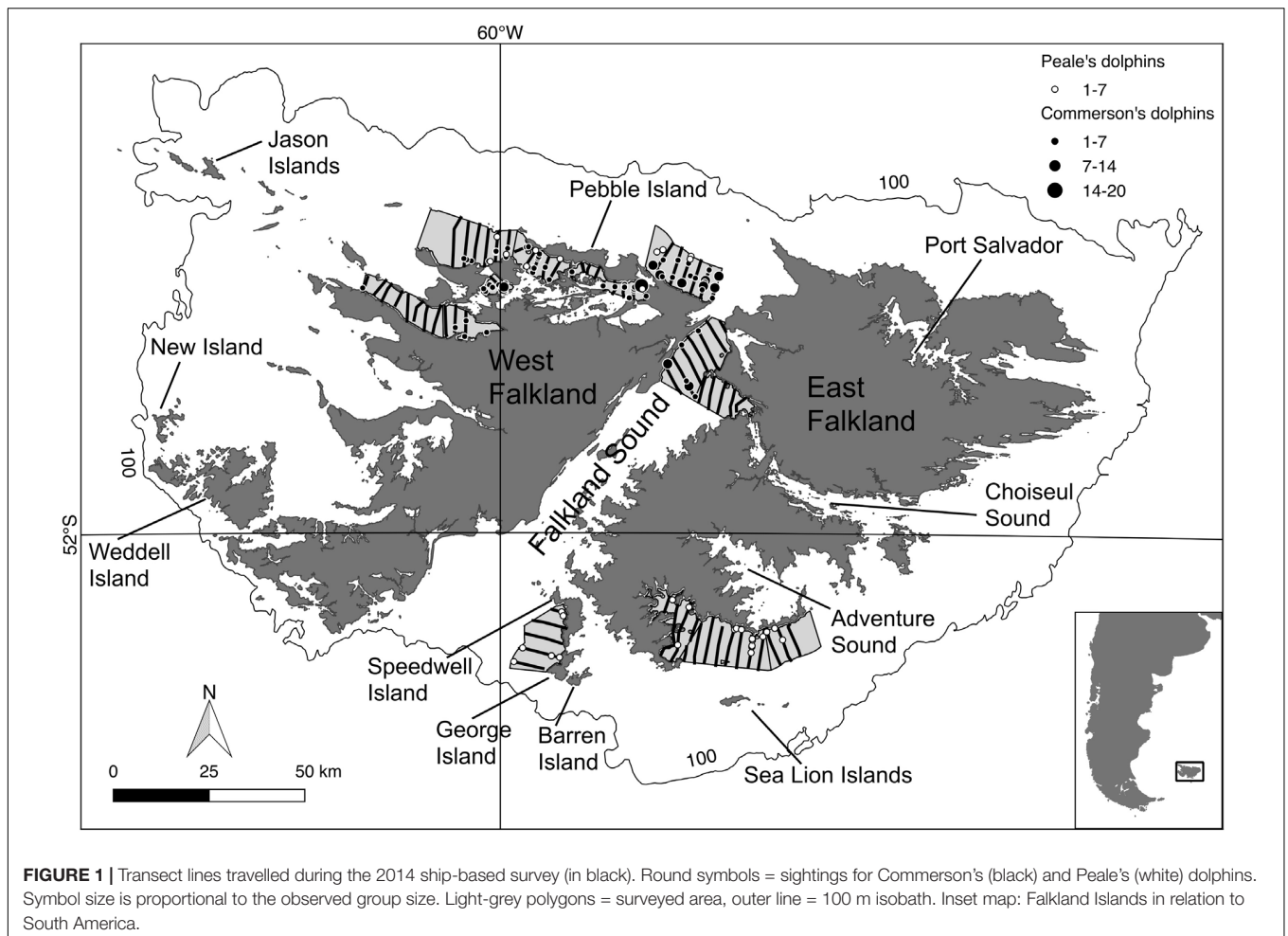
## Modelling

### General Approach

Dolphin distribution patterns were assessed by investigating the relationship between the six habitat covariates and dolphin sighting data collected in 2014. This was achieved with a modelling approach divided into five main steps: (i) Average detection probabilities were estimated for each species from all sightings along all transect lines. (ii) Transect lines were split into segments, with each segment having values for segment size (area), habitat covariates, and the number of dolphins observed. (iii) Models were fitted using the segment data, and then used to predict the expected numbers of dolphins over the entire FI inshore waters using gridded values of habitat covariates. (iv) The models were validated by qualitatively comparing predictions with dolphin sightings collected during an independent, island-wide aerial survey for cetaceans in 2017. (v) Finally, the uncertainty in the predictions was assessed through non-parametric bootstrapping.

### Setting Up Segments and the Prediction Grid

We split transect lines into segments with a minimum length of 5 km and added a buffer (i.e., transforming segments from lines to polygons). Using segments of sufficient length had the following advantages: avoiding excessive autocorrelation between data from adjacent segments; restricting the size of the data set to be fitted; and reducing the number of segments containing no dolphin sightings (counts = 0). This ensured satisfactory goodness-of-fit (see Results). Since dolphins were sighted at a range of distances from the vessel and were not always close to the transect lines, we selected a 1.5 km buffer in order to provide



a better representation of the habitat covariates associated with the actual sighting locations. We then calculated the segment area (used as an offset term in the modelling) and a median value of each habitat covariate for each segment. Finally, we built a 1 km grid to predict the distributions of both dolphin species across the entire FI inshore waters (see below). Median covariate values were calculated within each grid cell. We restricted the grid extent to the surveyed environmental space (i.e., we excluded grid cells where covariates were outside the original range of the data). This avoided predictions outside the calibration ranges, as these might be unreliable.

### Model Calibration, Predictions and Validation

We estimated the detection probability  $\hat{p}$  for each species by using perpendicular distances to all sightings (from all transect lines in the entire survey area) and a hazard-rate detection function in the “Distance” R-package (Thomas et al., 2010). Species-specific average detection probabilities entered the models as an offset term. We modelled dolphin sighting data using a Hurdle model approach consisting of a binomial sub-model for presence-absence (PA) and a zero-truncated Poisson sub-model for the number of dolphins detected conditional on presence (AB). Final

predictions from the Hurdle model were obtained by multiplying together PA and AB sub-model predictions.

Both sub-models assumed smooth GAM (Generalized Additive Model) relationships between response variable and habitat covariates. We used a shrinkage version of cubic splines to reduce overfitting, and a maximum knot number of 10 and 3 for PA and AB sub-models, respectively. We selected a lower knot number for the AB sub-model due to fitting instabilities (the algorithm did not converge with the smaller presence-only dataset). We fitted full models (i.e., containing all 6 habitat covariates) and checked them for concurvity, which occurs when one or more smooth terms in a GAM model can be approximated with one or more smooth terms within the same model (i.e., a non-parametric analogue for multi-collinearity). In case of concurvity indices close to 1 (i.e., full concurvity vs. 0 = no concurvity), all covariates with Pearson correlation coefficient ( $R$ )  $\geq 0.5$  or  $\leq -0.5$  were only considered in different sub-model variants (i.e., sub-models with different covariate structure; Ramsay et al., 2003). We then specified and fitted variants for each sub-model and we selected all variants yielding an AICc (i.e., Akaike Information Criterion corrected for smaller sample size) weight ( $\delta_w$ )  $\geq 0.1$ . If multiple variants were selected, we scaled their AICc weights by dividing the  $\delta_w$  of each  $i^{th}$  selected

variant ( $\delta_{w,i}$ ) by the sum of the weights of all  $n$  selected variants (i.e.,  $\sum_{i=1}^n \delta_{w,i}$ ). After checking selected variants for concurrency, we calculated the weighted average of their predictions using scaled  $\delta_w$ . Hurdle model goodness-of-fit was assessed through examination of the QQ-plot of the cumulative distribution function (CDF) vs. the empirical distribution function (EDF) and Kolmogorov–Smirnov (K–S) tests. Spatial autocorrelation in residuals was evaluated with correlograms. We used the selected Hurdle sub-models to estimate the expected number of dolphins per km<sup>2</sup> on the prediction grid, based on the covariate values within each 1 km<sup>2</sup> grid cell. Grid cells with higher values were interpreted to represent areas of better dolphin habitat and contribute to what we term key areas of distribution. Finally, model predictions were validated visually by overlaying the dolphin sightings from the 2017 aerial surveys on the predicted density surface. If the habitat preference model included an appropriate set of covariates and provided that the association between dolphin distribution and habitat variables remained consistent over time, our expectation was that the independent sighting data should provide a reasonably good match to the overall predicted distribution patterns, i.e., that areas where more dolphins were predicted to occur should have yielded more sighted dolphins than those areas where the species was predicted to be absent or occur in low numbers. Because of the nature of the different datasets we did not attempt to quantify their close correspondence, but were looking to corroborate the overall predicted distribution patterns for each species.

### Model Uncertainty

We assessed uncertainty in predictions for each grid cell with lower and upper limits of the 95% confidence interval (CI) calculated after stratified non-parametric bootstrapping. Briefly, we ran 1,000 simulations each made of two steps: (i) we randomly sampled (with replacement) rows of the observation dataset (i.e., sightings) to estimate detection probabilities; and (ii) we fitted the calibrated model and predicted number of dolphins on the prediction grid after randomly sampling (with replacement) rows of the segment dataset (i.e., segments) containing information on segment habitat covariates, area and average detection probability estimated in (i). Random sampling with replacement was stratified by stratum, i.e., we resampled segments within each stratum. Note that we fitted the same calibrated sub-model variants with the same AICc weights (in case of selection of multiple variants) and smooth term knot locations. All 1,000 prediction grids were subsequently converted into two grids for lower and upper limits of the 95% CI.

### Modelling Implementation

The R-package “dshM” (Density Surface Hurdle Modelling) was developed for this analysis<sup>1</sup> to undertake some of the data preparation and implement the Hurdle models. One of the advantages of the “dshM” R-package is that it contains a set of functions to quickly split transect lines into segments, add a buffer to them and calculate zonal statistics for each segment.

<sup>1</sup><https://github.com/FilippoFranchini/dshM>

This allows the user to explore how model goodness-of-fit and spatial correlation change with segment size, and thus to select a reasonable size to ensure model robustness.

## RESULTS

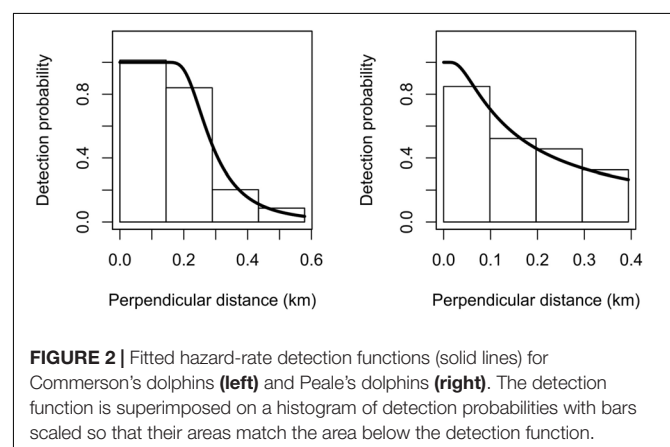
The total length of the 2014 boat survey transects was 567 km. The number of observed dolphin groups was 73 (total number of individuals: 266) and 33 (total number of individuals: 103) of Commerson’s and Peale’s dolphins, respectively (Figure 1).

### Detection Probabilities

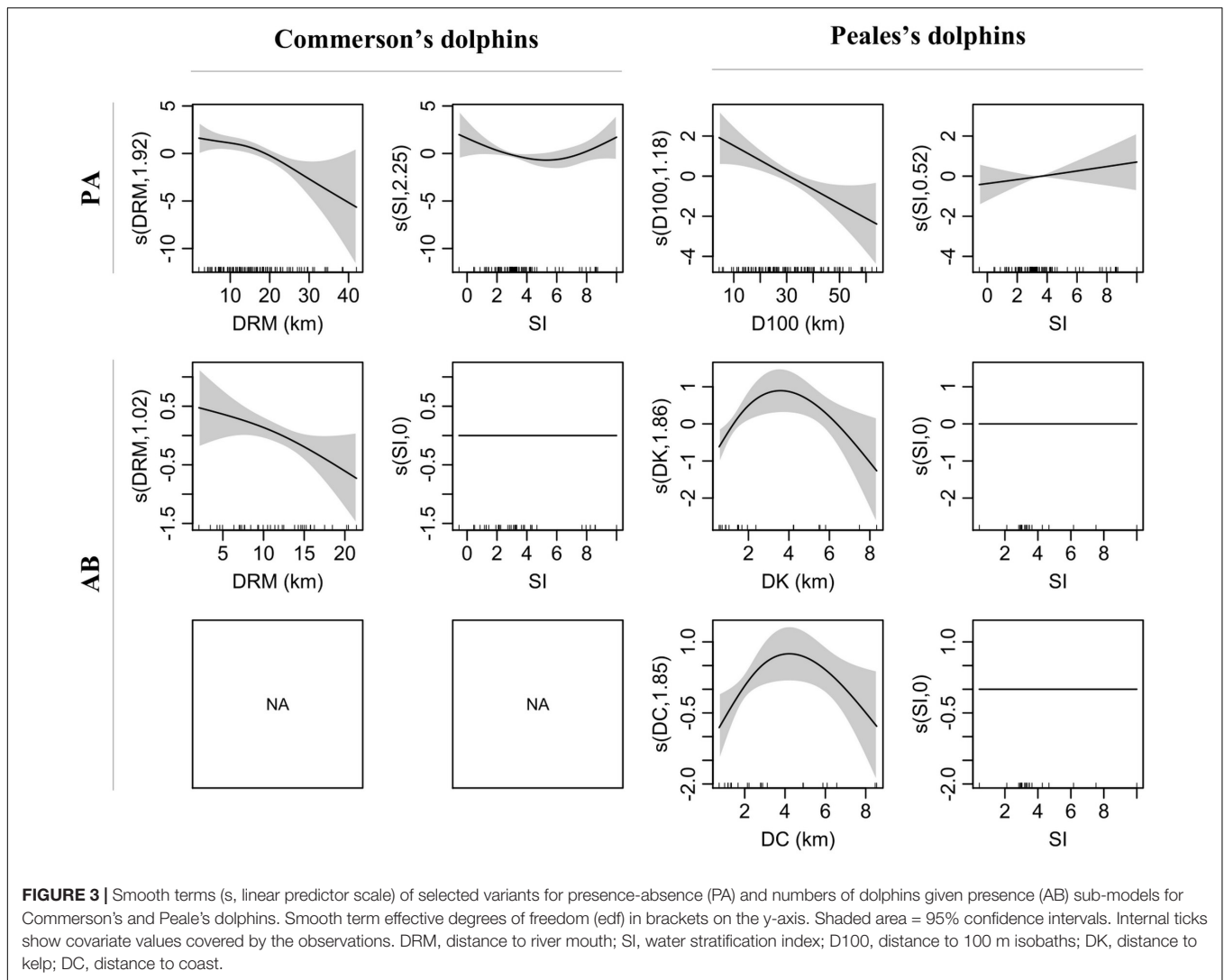
Due to the relatively limited number of sightings for both species, we decided to omit the 5% truncation distance that is usually applied to the right tail of the detection function (Buckland et al., 2001). The estimated detection probability  $\hat{p}$  was 0.54 (CV = 0.093) for Commerson’s and 0.54 (CV = 0.352) for Peale’s dolphins (Figure 2). The goodness-of-fit tests showed satisfactory fit for both species (Commerson’s dolphins: K–S test statistics = 0.08 and  $p = 0.76$ ; Peale’s dolphins: K–S test statistics = 0.09 and  $p = 0.94$ ).

### Spatial Hurdle Models

The full Hurdle model yielded smooth term concurrency indices ranging between 0.32 and 0.99 (Supplementary Table 1). As a consequence, pairs of covariates with Pearson correlation coefficient ( $r$ ) outside the interval including 0.5 and  $-0.5$  were not considered together in the same model (Supplementary Table 2). Water stratification index was not correlated with any of the other covariates ( $-0.2 \leq r \leq 0.3$ ), which instead were all correlated ( $-0.8 \leq r \leq -0.5$  and  $0.5 \leq r \leq 1$ ), and thus not included together in the same sub-model variant (i.e., reduced Hurdle sub-models with different covariate structure). Although the Pearson correlation coefficient for the distance to the 100 m isobaths and water depth was equal to 0.4, a visual check through scatterplots suggested a substantial degree of correlation (not shown). This is reasonable as water depth increases so D100 decreases on the coastal side. DC, DRM, and DK were also positively correlated since kelp beds line the coast, and river



**FIGURE 2** | Fitted hazard-rate detection functions (solid lines) for Commerson’s dolphins (left) and Peale’s dolphins (right). The detection function is superimposed on a histogram of detection probabilities with bars scaled so that their areas match the area below the detection function.

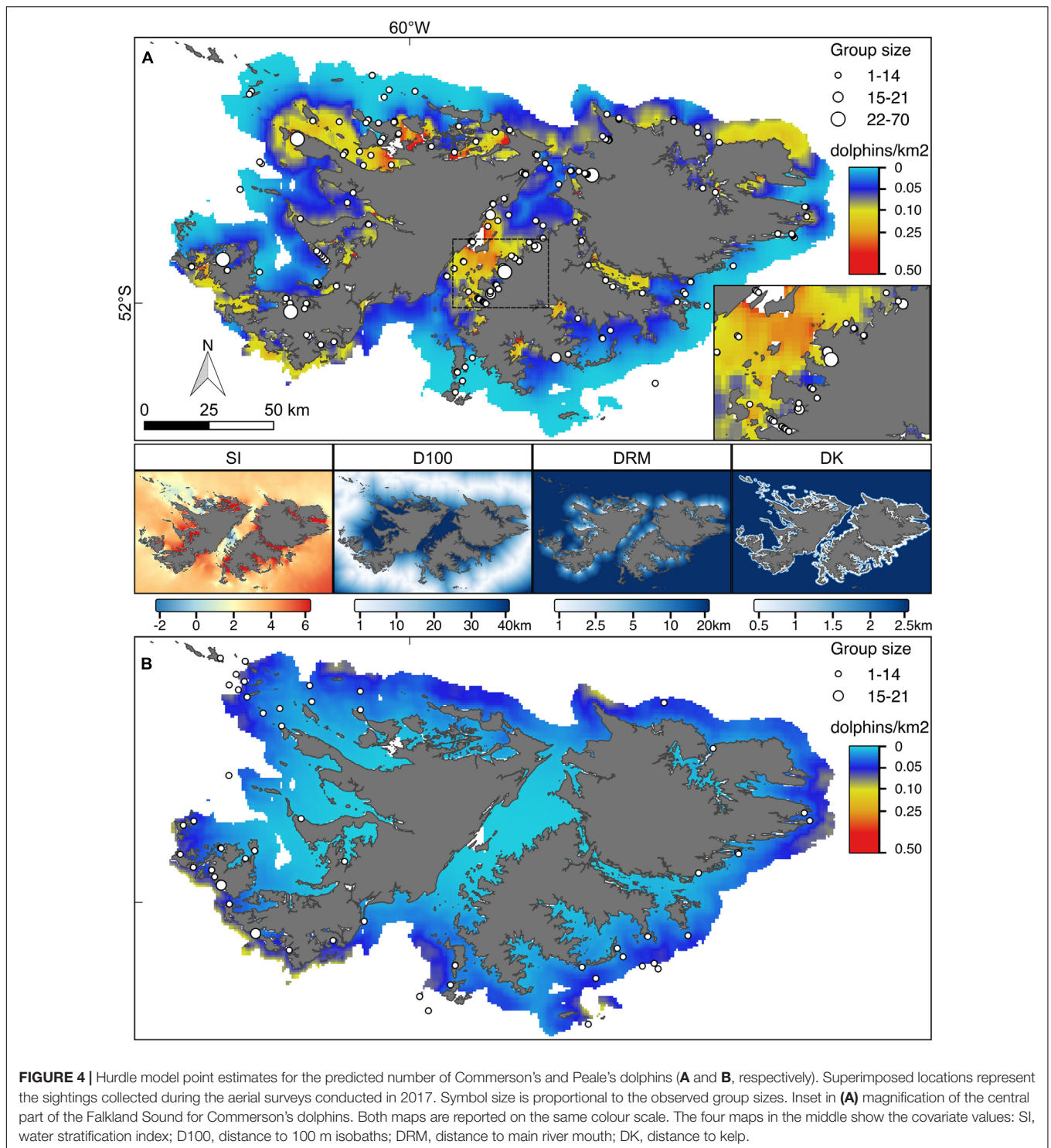


mouths are also located along the coastline. We therefore applied model selection to five reduced sub-model variants that included the water stratification index in combination with each of the remaining covariates (**Supplementary Table 3**).

The variant including DRM and SI yielded the best fit for both PA and AB Commerson's dolphin sub-models which explained 27.2 and 13.5% of the deviance, respectively (**Supplementary Table 3**). The covariate with the most pronounced effect on probability of presence was SI, which showed a U-shaped relationship with minimum values around 5 (**Figures 3, 4**). This relationship clearly showed that Commerson's dolphins were associated with deep, well-stratified waters ( $\text{SI} > 5$ ) as well as regions that experience very strong tidal mixing ( $\text{SI} < 2$ ). Commerson's dolphins were found close to tidal fronts as well as on both stratified and mixed sides. Both probability of presence and  $E(n|n > 0)$  (i.e., the estimated number of dolphins given presence) were inversely related to DRM, i.e., Commerson's dolphins appeared to prefer to stay close to river mouths and

tended to aggregate in such areas. Stratification index did not have an effect on estimated numbers of Commerson's dolphins.

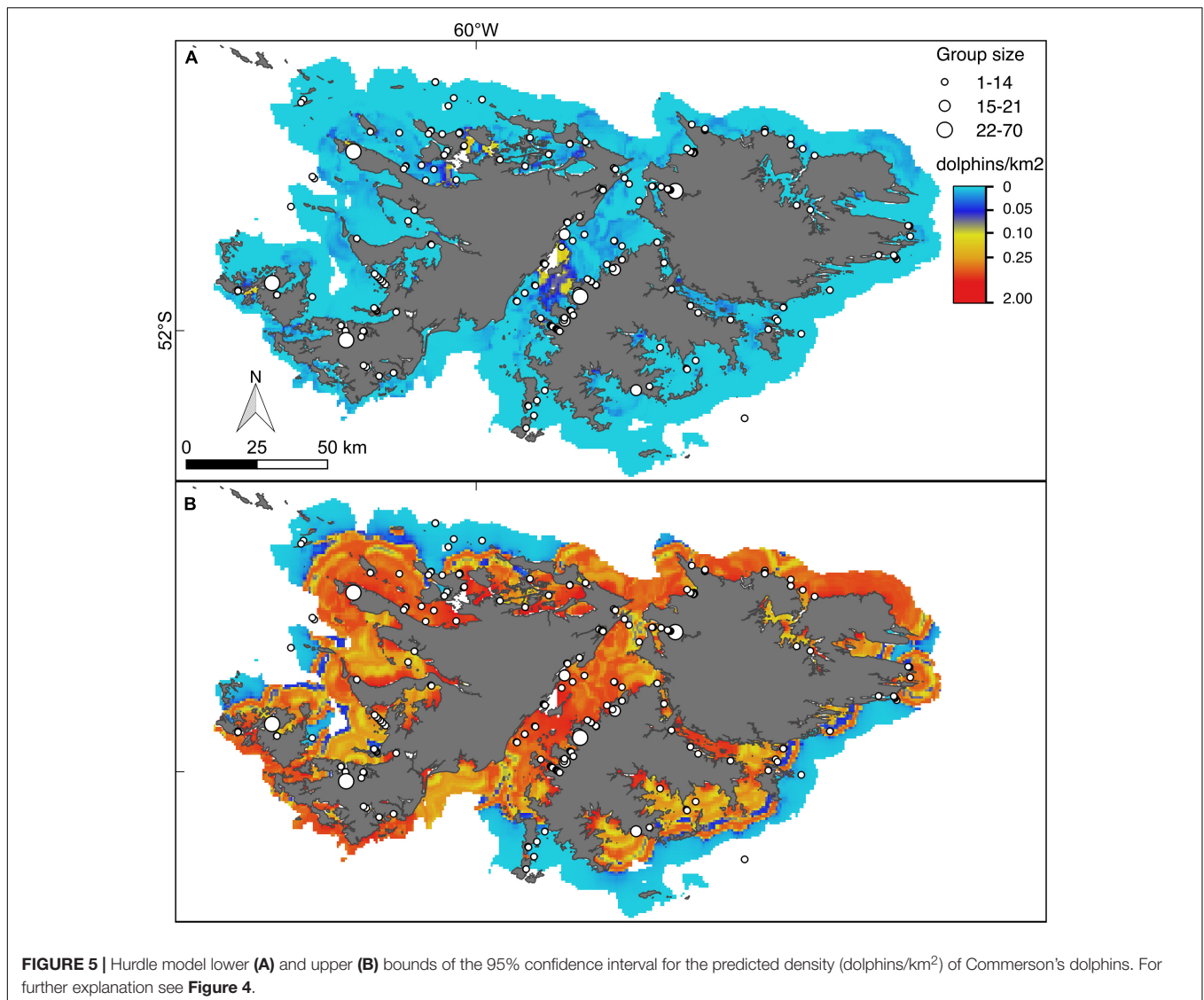
In addition to SI, the selected PA sub-model variant for Peale's dolphins included D100 and explained 17.1% of the deviance (**Supplementary Table 3**). In contrast to Commerson's dolphins, the AB sub-model included two variants with DK and DC explaining 38.1 and 33.3% of the deviance, respectively. Water stratification index had the lowest effect in the PA sub-model and no effect in the AB sub-model (**Figures 3, 4**). Probability of presence was inversely related to D100 and positively related to SI, while  $E(n|n > 0)$  for Peale's dolphins showed a similar inverted U-shaped relationship with both DK and DC. Thus, Peale's dolphins seemed to prefer offshore and well-stratified waters, but they tended to aggregate at approximately 2 km from kelp beds and coast. Since DK and DC belonged to two different variants, the degree of association of  $E(n|n > 0)$  for Peale's dolphins with these two covariates differed, with covariates with highest  $\delta_w$  having the highest degree of association. Thus,



the  $E(n|n > 0)$  for Peale's dolphins was more associated with DK ( $\delta_w = 0.81$ ) than with DC ( $\delta_w = 0.17$ ).

Concurvity indices for smooth terms within the selected sub-model variants for both dolphin species ranged between 0.08 and 0.31 (**Supplementary Table 1**). QQ-plots and goodness-of-fit tests showed a satisfactory fit of the Hurdle model

for Commerson's dolphins (Kolmogorov–Smirnov test statistics = 0.14,  $p = 0.06$ ) and a less satisfactory fit of that for Peale's dolphins (Kolmogorov–Smirnov test statistics = 0.15,  $p = 0.03$ ) (**Supplementary Figure 2**). Spatial correlation of Hurdle model residuals for Commerson's dolphins decreased with distance while that for Peale's showed a fluctuating pattern



across the whole range of distances (Supplementary Figure 3). The proportion of the correlation values within  $-0.1$  and  $0.1$  was 58% for Commerson's and 70% for Peale's dolphins.

### Hurdle Model Predictions, Uncertainty, and Validation

According to the fitted Hurdle model, Commerson's dolphins were found in all FI inshore waters (Figures 4–6). Larger numbers of Commerson's dolphins were predicted for the central part of the Falkland Sound. In contrast, Peale's dolphins were predicted to occur in more offshore waters and in lower densities (Figures 4–6). For both species predicted key areas matched the sighting locations of dolphins from the 2017 aerial surveys (SAERI unpublished data) which had covered 4,255 km in the entire inshore waters, with 195 and 55 groups of Commerson's and Peale's dolphins observed, respectively (Figures 4–6, Supplementary Figure 1). This finding suggests that the habitat model included an appropriate and sufficient set

of relevant predictor variables to make effective predictions, and supports the assumption that the association between dolphins and habitat covariates remained consistent over time.

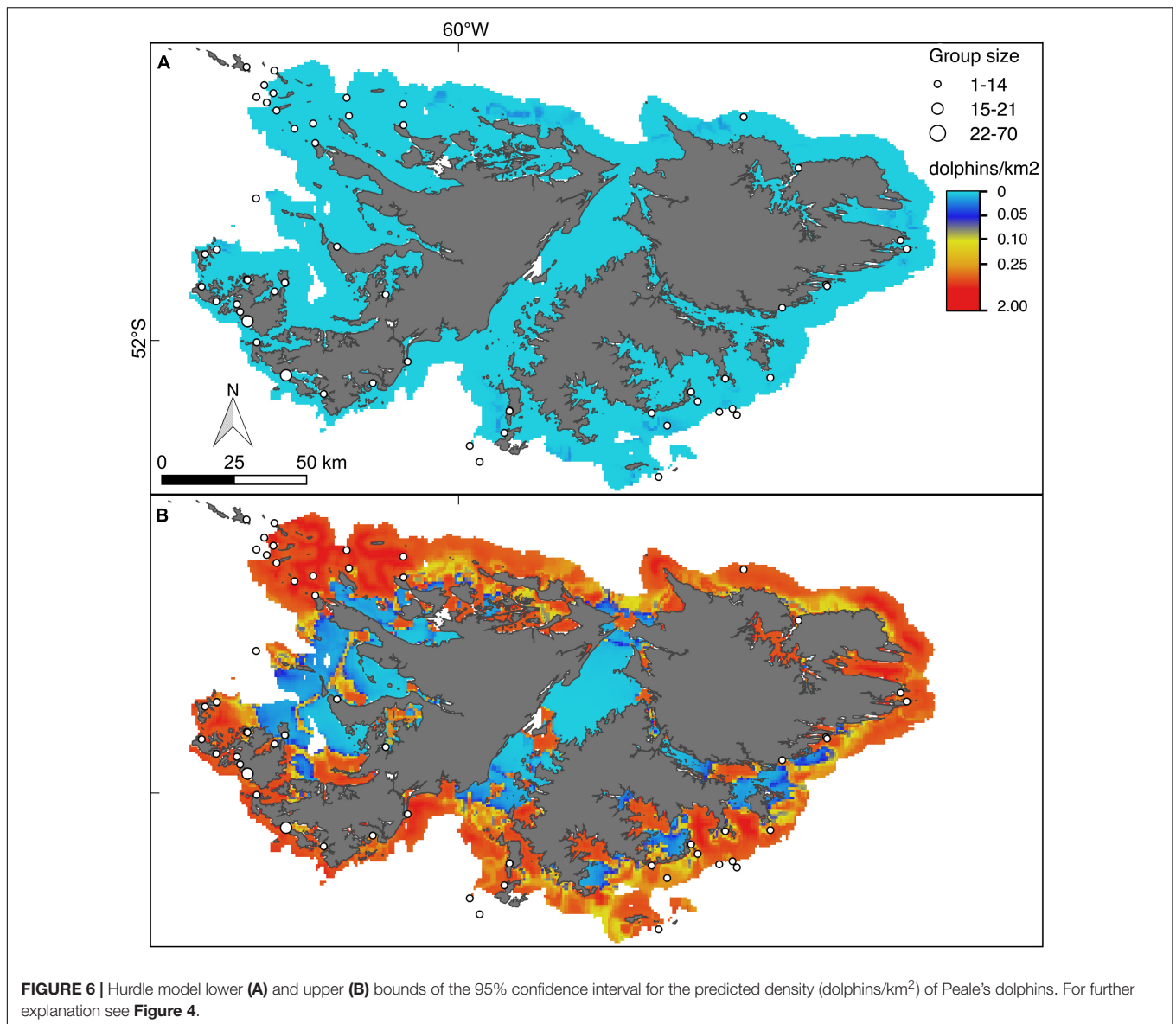
### DISCUSSION

This study succeeded in generating robust spatial predictions from a spatially limited survey conducted over a short time period and extrapolating the distribution and habitat use of two dolphin species to cover the entire Falkland archipelago. Our results suggest that while some preferred habitat is shared, there is spatial niche separation between the two species.

### Methodological Considerations

Representative sampling is an important consideration in modelling studies and requires some *a priori* knowledge of the species' ecology. We were able to draw on ecological knowledge





of each species from studies conducted in other parts of the species' ranges (Garaffo et al., 2011; de Castro et al., 2013; Dellabianca et al., 2016; Heinrich et al., 2019) to guide the selection of potentially relevant environmental variables. The covariate ranges surveyed during the boat-based survey were representative of the overall FI inshore waters, thus allowing predictions within environmental space but extrapolation in geographic space.

Habitat modelling requires decisions about a trade-off between high-resolution data and model robustness. Where data are limited, model robustness can only be assured at the expense of data resolution. The resolution or grain size (e.g., segment area in our study) of the model should not be confused with the resolution of the prediction grid (e.g., grid cell area). The use of an offset term for grain size in the model allows for predictions of animal occurrence or relative abundance on higher resolution

grid cells, while the underlying relationships are those originally fitted to the model grain size. Decreasing model resolution by taking large grain sizes also helps to reduce spatial autocorrelation in the response variable, which is a frequent problem in predictive habitat modelling (Guisan et al., 2007; Mannocci et al., 2017). Ideally one would like to know the best grain or segment size to ensure robustness and simultaneously maximise resolution (Mertes and Jetz, 2018). Splitting transects into segments can be a very tedious, time-consuming process that often requires the user to switch between different software packages. The newly developed "dshm" R-package greatly facilitates the splitting of transects into segments, the exploration of different grain sizes and the selection of a reasonable resolution to ensure model robustness.

This study presents five updates to the methodology currently used for predictive habitat modelling (Embling et al., 2010;

Hammond et al., 2013; Dellabianca et al., 2016): (1) Use of zonal statistics: Predictor variables are usually derived by using a point-sampling approach, in which the covariate values associated with the chosen spatial unit (e.g., segment) are those sampled just below the unit centroids. This procedure might be inappropriate since habitat features at a small specific location might not be representative of the area where the animals were observed. We used a zonal-statistics approach in which the values of each predictor variable for a segment were derived as the median of all that covariate's values available for that particular segment. Using the median rather than the mean decreased possible bias in covariate statistics for segments near the coastline. This is particularly important for heterogeneous environments, where covariate values might vary greatly over small spatial scales. (2) Use of buffered segments: Responsive movements are a common problem in boat-based surveys for cetaceans (Palka and Hammond, 2001) and can lead to biases in the detection probabilities (e.g., in our study both dolphin species showed strong attraction to the boat, leading to positive biases in the detection probabilities). Buffering the transect line by 1.5 km and using a zonal-statistics approach alleviated this attraction bias in our habitat modelling approach. The buffer size was chosen to reflect the range over which responsive movements were thought to occur. Thus, although the recorded spatial position of the animals might not represent their precise undisturbed position, the spatial integration of covariate values ensured representation of the natural habitat that the animals occurred in. (3) The binomial part of the Hurdle model allows zero inflation to be addressed. This is a common problem in animal surveys (Martin et al., 2005), and particularly so for rarely observed species (Welsh et al., 1996). Hurdle models are also ecologically sensible for scenarios where the processes driving the presence of a species differ from those affecting its abundance (Ridout et al., 1998). (4) Addressing GAM flexibility: The flexibility of the GAM framework is a “double-edged sword”: it enables one to model complex biological relationships but can also lead to overfitting (Morgan and Ridout, 2008). In this study the risk of overfitting was minimised by implementing a shrinkage version of cubic spline. (5) Stratum-specific bootstrapping: Model uncertainty was evaluated by using non-parametric bootstrapping with replacement, which is free from assumptions on parameter distributions (Fletcher et al., 2005). This approach, however, can be affected by fitting instabilities related to spatial gaps arising from resampling with replacement over the entire study area. We reduced such gaps by adopting a “stratified” approach in which resampling with replacement was limited to segments within the same stratum.

## Predicting Distributions of Sympatric Dolphins

Our model predictions showed that Commerson's and Peale's dolphins used different key habitats around the FI with only some spatial overlap in distributions (Figures 4–6). Commerson's dolphins were predicted to be most abundant in the central part of Falkland Sound, off the northern East Falkland coast as well as

in the interior part of sounds and inlets around West and East Falkland. In contrast, Peale's dolphins were predicted to occur mostly off the southern and north-western coasts of the Falkland archipelago. This difference in predicted spatial distributions resulted from the different modelled relationships that both species had with inshore habitat covariates. The environmental variables likely served as proxies for linked biological variables that could not be measured directly (such as the distribution or density of prey or predators) and might have reflected the differences in species-specific physiological and ecological responses (Kiszka et al., 2015; de Thoisy et al., 2016).

Probability of presence of Commerson's dolphins appeared to be linked to the water SI with areas of very low SI (e.g., Falkland Sound) or very high SI (e.g., northern West Falkland, most FI inlets) being those where Commerson's dolphins were most likely to occur (Figure 4). This suggests that Commerson's dolphins preferred to be close to tidal fronts as well as stay on the mixed or stratified sides. The importance of tidal fronts to the distribution of Commerson's dolphins has also been suggested for the Argentinian coast (Dellabianca et al., 2016). In contrast to presence, Commerson's dolphin abundance seemed to be mostly linked to distance to rivers with the highest numbers occurring in upper inlets, and close to river mouths. Such preference for nearshore neritic environments with strong tidal influence and estuarine zones is a defining feature for all species in the genus *Cephalorhynchus* (Heinrich et al., 2010; de Castro et al., 2013; Heinrich et al., 2019).

Presence of Peale's dolphins was mostly associated with D100, with the highest probability of observing the species close to the 100 m isobaths, i.e., shelf-waters further offshore. Although occurrence was predicted to be higher in deeper and well-stratified shelf waters, Peale's dolphins also tended to be more numerous in shallower waters close to the coast and near to kelp as shown by the importance of both DK and DC in the abundance model and the resulting predicted distribution (Figures 3, 4). Such overall plasticity in occurrence has now emerged as a feature consistent for Peale's dolphins across their entire distributional range (Dellabianca et al., 2016; Heinrich et al., 2019; this study). Peale's dolphins' reliance on nearshore habitat, at least during summer, is well documented from southern Argentina (De Haro and Iñiguez, 1997), with kelp forests thought to represent important feeding grounds (Schiavini et al., 1997; Viddi and Lescauwaet, 2005; Riccialdelli et al., 2010). The low number of Peale's dolphin sightings affected the model goodness-of-fit ( $p$ -value near the 5% threshold, Supplementary Figure 2). Despite these limitations the resulting predicted spatial distribution was well matched by sightings from the independent aerial survey suggesting that the model predictions succeeded in capturing the dolphins' overall habitat preference.

Although Commerson's and Peale's dolphins are fully sympatric across their South Atlantic range, our predicted distribution maps and other large-scale modelling studies (Dellabianca et al., 2016) indicate fine-scale habitat partitioning between the two species. Strategies that enable sympatric species to co-exist include spatial or temporal differences in habitat use, dietary divergence and specialisation, as well as differences in activity patterns and socially mediated behaviours

(Begon et al., 2006). Future research into niche partitioning of these two species should therefore focus on investigating ranging patterns, site fidelity, social interactions (e.g., using individual identification studies; e.g., Parra, 2006; Coscarella et al., 2011) and establishing diet and trophic niche overlap (using stable isotope or fatty acid analyses; e.g., Querouil et al., 2013; Giménez et al., 2018). The predicted distribution maps from this study should prove helpful to identify suitable FI study areas for such intensive, focal research endeavours.

## Conservation and Management Relevance

The most recent Falkland Islands Species Action Plan for Cetaceans (2008–2018) identifies a range of suspected and known threats to the marine ecosystem on the shelf around the FI, including pollution, bycatch, vessel traffic and climate change, and highlights novel and expanding activities with the potential for harmful interactions such as aquaculture, tourism and commercial fishing (Otley, 2008). Identifying potential areas of importance for these dolphin species as well as their overlap with human activities should serve as a guide for future research efforts and the current FI marine spatial planning activities.

The underlying data for such studies is, however, often very hard to obtain, especially as information from a range of species is needed to enable focussed management and conservation efforts. One can argue that meso and top predators often forage in aggregations with other species (hotspots) and that therefore the use of data from sentinel species are enough, but on smaller scales, as in coastal waters, niche separation might indicate reliance on different prey and/or areas. Relevant high-resolution data are therefore necessary to resolve such differences. Remote tracking technologies have enabled us to collect high-resolution information about movements and foraging behaviour, but while such data can provide an unprecedented amount of detail, studies are usually limited to one species, low numbers or are hampered by a lack of concurrent data in case of multi-species approaches (Baylis et al., 2019).

We have shown a practical and cost-effective approach using a spatially limited dataset to estimate multi-species distribution patterns. We have demonstrated the robustness of this method by comparing the patterns with sightings from another survey. The methodology and software implementations provided here can be easily applied to transect survey data of other mobile marine and terrestrial taxa where the area of interest is representative of but larger than the area surveyed. This will help us to address the challenges ahead in conservation and management of the marine environment.

## DATA AVAILABILITY STATEMENT

The datasets generated for this study are available on request to the corresponding author.

## ETHICS STATEMENT

The animal study was reviewed and approved by University of St Andrews, School of Biology Animal Ethics Committee.

## AUTHOR CONTRIBUTIONS

FF undertook the statistical modelling and GIS analysis. CB assisted with the GIS analysis. SS assisted with the statistical modelling. LB assisted with some of the covariate generation. GM and SH designed the study and collected the 2014 data. MC led the 2017 aerial surveys that provided the independent dataset for corroboration of predicted distributions. FF, SH, SS, and LB wrote the manuscript. All authors commented on the analyses and the final manuscript and approved the final manuscript to be published.

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## SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmars.2020.00277/full#supplementary-material>

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**Conflict of Interest:** Grant Munro was employed by company Austral Biodiversity Ltd.

The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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