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# Multi-scale assessment of distribution and density of procellariiform seabirds within the northern Antarctic Peninsula marine ecosystem

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## Abstract

The Antarctic Peninsula is one of the most rapidly warming regions on earth, and it is likely that the abundance and distribution of marine predators in this region will change as a result. Additionally, other anthropogenically induced factors, such as competition with commercial fisheries, may result in changes to the availability of prey for marine predators. Procellariiform seabirds are highly mobile marine predators, which target specific habitat characteristics associated with the underlying distribution of prey and areas of increased prey availability. We use surveys conducted from ship transects and hurdle models, to estimate the summer distribution and relative density of 11 seabird species within the northern Antarctic Peninsula marine ecosystem. Covariates included in hurdle models differed among species, however sea surface temperature and depth were frequently associated with seabird occurrence and had the greatest explanatory power across many species. Null models based on observation data were always better at predicting seabird density than models which included environmental covariates. This suggests that the main driver of distribution patterns is the broad-scale habitat features, and more fine scale aggregations within these broad-scale ranges are harder to predict. Our seabird distribution models reflect previously known habitat associations, species hotspots, and community organization relative to oceanic and coastal marine processes. Application of species distribution models will benefit the assessments of critical habitat and also potential responses to climate change and anthropogenic disturbance, which will provide insight into how these species may change in polar ecosystems.

Key words: Climate change, ecosystem oceanography, Fisheries management, Species Distribution modelling,

## Introduction

The Antarctic Peninsula region is one of the most-rapidly changing parts of our planet with a rise in atmospheric temperature of nearly 3°C since 1951, coinciding with a decrease in winter sea-ice duration (Meredith and King 2005, Stammerjohn et al. 2008, Cook et al. 2016). Such large-scale physical changes have the potential to cause alteration to the marine ecosystem across a range of trophic levels (Loeb et al. 2009, Ducklow et al. 2013). Indeed, changes in sea surface temperature as a result of climate change, are sometimes associated with demographic changes, and in some cases population collapse (Morrison et al. 2015, Rilov 2016). This may be a result of underlying changes in the distribution of prey, resulting in reduced prey availability to predators. Alternatively, some species may shift their distributions in response to a changing environment (Parmesan and Yohe 2003). For an area that is changing rapidly, documenting such processes will be important when evaluating how Antarctic and other geographic regions may change in the future. Long-term declines are hard to detect and the key to our understanding will be the establishment of baseline data describing the at-sea distribution of all species. For some groups, seabirds in particular, information about the location and size of breeding colonies in the region is sparse, leading to gaps in our baseline understanding of their distribution and abundance.

As the environment continues to change along the Antarctic Peninsula region, some species of seabird may decline. For example, Hinke et al. (2017b) predict continued declines in Adélie penguin (*Pygoscelis adeliae*) populations at the northern extent of their range. For many species, it is likely that regional warming will result in population range shifts (Grémillet and Boulinier 2009). In addition to warming, other anthropogenically induced factors, such as competition with commercial fisheries, can also result in changes to prey availability for marine predators, leading to changes in their foraging ecology (Bertrand et al. 2012), which can then lead to subsequent population change (Bost et al. 2015). In the Antarctic Peninsula region, there is concern regarding the commercial harvest of Antarctic krill *Euphausia superba*, the main prey item for many Southern Ocean seabirds (Trathan et al. 2018). Krill biomass in this region may already be in decline (Atkinson et al. 2004, Atkinson et al. 2008, , but see Cox et al. 2018a). Although many species in this study have much broader diets that include squid and mesopelagic fish, many of these mid-trophic dietary items also consume krill (Nemoto et al. 1988, Barrera-Oro 2002). Indeed, there is concern that krill depletion in key foraging areas for krill-dependent predators may cause population-level impacts (Mangel and Switzer 1998, Hinke et al. 2017a).

Seabirds are considered indicators of the status of the marine environment (Parsons et al. 2008). For example, demographic traits of seabird populations can be used to improve understanding about

ocean climate variability (Jenouvrier et al. 2003), the localised abundance of prey items (McLeay et al. 2009) and to detect illegal fisheries (Weimerskirch et al. 2020). Procellariiforms are pelagic seabirds comprising albatrosses, petrels, shearwaters and storm-petrels, and are the most abundant flying seabird group in Antarctica. They are colonial breeders and usually return to the same breeding site annually. One egg is laid annually (or every two years for some species of albatross), and parents alternate nest attendance during egg incubation and chick provisioning. Many procellariiforms employ a dual foraging strategy of interspersed long (6 – 29 days) and short (1 – 5 days) foraging trips providing for their young whilst maintaining adult body condition during the chick-rearing period (e.g. Weimerskirch et al. 1994, Granadeiro et al. 1998, Baduini and Hyrenbach 2003). Procellariiform seabirds are highly mobile, frequently associated with specific habitat characteristics, and forage across a range of spatio-temporal scales (Hunt Jr and Schneider 1987). Broad-scale habitat characteristics such as sea surface temperature may restrict the wider distribution of seabirds, whilst high density aggregations may occur at a finer scale within a species' broad range (Becker and Beissinger 2003, Waggitt et al. 2020). For example, sea surface temperature limits the broad-scale distribution of short-tailed albatrosses (*Phoebastria albatrus*), whilst their fine- and meso-scale foraging behaviour coincides with proxies of prey availability such as chlorophyll-a and topography (Suryan et al. 2006). These multi-scale distribution patterns are a result of prey patchiness, which is often associated with fine- (< 10 kms) or meso- scale (10s to ~100 kms) oceanographic features including eddies and fronts (van Franeker et al. 2002, Bost et al. 2015). The distribution of breeding seabirds is also limited by the distance from the breeding site, given that breeding individuals must return frequently to incubate eggs or provision chicks. Finally, both inter- and intra- specific competition for prey may help to shape seabird distribution at-sea (Wakefield et al. 2011, Trathan et al. 2018).

Areas of increased species abundance, diversity or trophic transfer can be referred to as biological hotspots (Santora et al. 2017a). By identifying seabird hotspots, we can provide information on the size, persistence, and distribution of areas that are key to trophic coupling in the marine ecosystem (Santora et al. 2017b). The use of animal-borne biologging devices such as GPS has improved our understanding of the at-sea distribution of some species of seabirds in the region, but does not provide information about relative density (animals km<sup>-2</sup>), community or population structure. The assessment of ecosystem and climate dynamics would benefit from an improved understanding about the at-sea density and distribution patterns of other seabird species, as well as species associations and interactions. Shipboard surveys provide a means to improve our understanding but are expensive and are on the decline compared to biologging studies. Thus, the gaps in our knowledge of seabird

abundance and distribution in the Antarctic Peninsula region leads to challenges when trying to quantify the impacts of disturbance or change within the area.

To better understand the implications of environmental change and anthropogenic disturbances, such as the commercial krill harvest in the Antarctic Peninsula region on seabirds, we can apply species distribution models to quantify habitat use. As such, observations of at-sea seabird distribution and abundance, derived from ship-based surveys, can be used to estimate species density and distribution (e.g. Oppel et al. 2012, Santora et al. 2017b, Waggitt et al. 2020). Ecological models that link seabird observations to environmental characteristics are frequently used to describe the distribution within, and predict the distribution beyond, the survey region (e.g. Clarke et al. 2003, Oppel et al. 2012), although extrapolations outside of the survey region will provide less robust estimations, and thus should be interpreted with caution (Yates et al. 2018). Here, we use seabird surveys conducted from extensively replicated ship transects, and multi-scale ecological models, to estimate the summer distribution and relative density of procellariiformes in the northern Antarctic Peninsula marine ecosystem. We use presence/absence models to estimate the broad-scale range of each species and to gain a baseline understanding about how their distributions are limited by broad-scale environmental characteristics. We then use density models to estimate the mesoscale areas within these broad species ranges where increased abundance of individuals are observed. These relative density models highlight the areas which may provide improved foraging habitats within the broader species range. The implications of these distribution patterns may be used for future assessments including understanding important areas for krill consumption or evaluating the potential impacts of the krill fishery.

## **Materials and methods**

### **Data collection**

The U.S. Antarctic Marine Living Resources (AMLR) Program conducted annual ecosystem monitoring surveys throughout the northern Antarctic Peninsula region during summer (January-March) from 2003 to 2011. A fixed grid of stations and transects were sampled (average spacing of 20 nm) to assess physical and biological oceanographic conditions and to map the distribution and abundance of krill and air-breathing predators. The survey area was partitioned by the South Shetland Islands, with oceanic waters to the north, and coastal waters to the south towards the Antarctic Peninsula. Ocean circulation in this region reflects inputs from the Antarctic Circumpolar Current (ACC), outflow from the Weddell Sea Gyre, and inflow from coastal upstream regions along the western Antarctic Peninsula that enter through western Bransfield Strait (Amos 2001, Moffat and Meredith 2018,

Trathan et al. 2018). Rugged bathymetry, including the peri-insular shelf around islands, deep basins of the Bransfield Strait, and the South Shetland Trench and the Shackleton Fracture Zone ridge in Drake Passage, are important drivers of hydrographic variability in the region (Figure. 1; Orsi et al. 1995; Thompson et al. 2009; Moffat and Meredith 2018). The southern ACC front (SACCF) is the southernmost deep-reaching front of the ACC, while the southern ACC Boundary (SACCB) defines the southernmost limit of ACC-derived waters (Orsi et al 1995). Additionally, the southern SACCF and SACCB are important hydrographic features that concentrate primary production, krill and top predators (Santora and Veit 2013, Force et al. 2015, Loeb and Santora 2015).

Standardized visual surveys of seabirds were conducted on 14 shipboard surveys during 2003-2011. The extensive shipboard tracklines covered open water, bays, inlets and passages between islands. Nine surveys were conducted in mid-summer (early January to early February) and 5 were in late-summer (mid-February to mid-March; Santora et al. (2009), Santora and Veit (2013), Santora et al. (2017b)). Strip-transect methods were used to estimate relative seabird abundance and counts were made within a 90° arc out to 300 m on the side of the trackline with the best visibility (i.e. no glare). A rangefinder was used to identify the 300m transect width (Heinemann 1981). All birds were counted whether flying or resting on the water. Strip-transect methods assume that all seabirds up to a pre-defined distance are detected because observers concentrate on a small area. All of the species included in these analyses, aside from storm-petrels which would not be sitting on the water, are relatively large and distinctively marked. As such, it is likely that almost all birds within the area were sighted and distance sampling and associated detection functions were not performed (Buckland et al. 2001). Therefore no distance or bearing data were collected for seabirds. Sea-surface conditions and visibility were continuously monitored and survey effort was ceased in poor conditions (e.g. Beaufort sea state >6; heavy fog impacting visibility of the 300 m strip transect, and storms). However, because we cannot guarantee that birds were not missed, or account for these, we present relative rather than absolute densities. Ship speed during transits between sampling stations was generally 10 knots (18.6 km hr<sup>-1</sup>) and observers used hand-held binoculars (10x50 and 20x60) to scan from a height of 13 m above sea level. Birds were detected with binoculars and by eye, and binoculars were used when necessary for identification. Each sighting was assigned a time and spatial position (of the vessel) and entered into a computer synchronized with the ship's navigational and acoustic system.

## Data analysis

The purpose of our analysis was to estimate a spatio-temporal average of seabird distributions, and to identify which habitats were consistently targeted. As such, we did not investigate inter-annual

variation in habitat use. To remove issues with unequal survey effort ship transects were divided into 3 km segments ( $n(3\text{km segments}) = 8,019$ ), and for each species the number of sightings and the number of individuals observed along each segment were summed. Segments which were not 3 km in length (i.e. the end of transects where the remaining length is  $< 3$  km) were removed from the dataset, as is commonly done in these analyses (removing  $< 3\%$  of all sightings). The position of the midpoint of each segment was recorded along with the corresponding time that the ship reached this location. For each species, the frequency of individuals sighted each year was plotted and examined visually to ensure that there was no major trend in abundance across the survey years. Species which were observed only in one or two years, or for which there were few sightings, were excluded from the analysis. Species which were observed but excluded from the analyses comprise: Antarctic petrel (*Thalassoica antarctica*), common diving petrel (*Pelecanoides urinatrix*), Kerguelen petrel (*Aphrodroma brevirostris*), light-mantled sooty albatross (*Phoebastria palpebrata*), northern giant petrel (*Macronectes halli*), royal albatross (*Diomedea epomophora*), snow petrel (*Pagodroma nivea*), sooty shearwater (*Ardenna grisea*) and soft-plumaged petrel (*Pterodroma mollis*). Almost all observed individuals were identified to species level. However, 49 % of prions were unidentified, and thus Antarctic (*Pachyptila desolata*,  $n=3,345$ ), thin-billed (*Pachyptila belcheri*,  $n=125$ ) and unidentified ( $n=3,401$ ) prions were grouped together.

#### Covariate data

Both static and dynamic environmental variables were used in our analysis (Table 1, Figure S1). The distance from the breeding colony may limit the foraging range of breeding seabirds during the breeding season when birds are constrained to return to the nest frequently to incubate eggs or provision young (e.g. Wakefield et al. 2011, Warwick-Evans et al. 2018). This variable was calculated using R package *gDistance* (van Etten 2012) for species documented to breed on the Antarctic Peninsula or South Shetland Islands, comprising: Wilson's storm-petrel *Oceanites oceanicus*, southern fulmar *Fulmarus glacialis*, Southern giant petrel *Macronectes giganteus* and cape petrel *Daption capense*. Breeding locations were obtained from the Agreement on the Conservation of Albatrosses and Petrels (ACAP), Harris et al. (2015) and (Creuwels et al. 2007). It was not possible to weight the distance from the breeding colony by population size because population counts for most of these species are not available for this region. Chlorophyll-a (Chl) provides a proxy for primary productivity, which is frequently used to indicate areas of increased prey biomass (Suryan et al. 2012). Sea Surface Temperature (SST) may limit the broad-scale distribution of seabirds as a result of the underlying distribution of prey (e.g. Hunt et al. 1992). Seabed depth (Depth) may shape the distribution of some species as a result of variation in prey availability (Paiva et al. 2010). For instance, interactions between

rough seabeds and currents in shallow water create flows which entrain and aggregate prey (Hunt 1999, Benjamins et al. 2015). Current speed (C) was included to identify areas where such flows are particularly prevalent (Waggitt et al. 2016). Eddy kinetic energy (EKE) and sea level anomalies (MSLA) provide measures of the relative intensity of mesoscale ocean circulation and are indicators of dynamic oceanic features such as eddies and large-scale fronts. Fronts and eddies can be associated with regions of strong upwelling, increasing the availability of nutrients in surface waters and consequently increasing phytoplankton production, and prey abundance (Owen 1981, Martin et al. 2002). Additionally, the convergent currents associated with frontal jets can concentrate prey, increasing its availability for foraging seabirds (e.g. Becker and Beissinger 2003, Sokolov et al. 2006, Santora et al. 2017b). Wind speed was included to account for variation in detectability due to weather conditions. High wind speeds create greater sea-surface featuring which may impact the detection rate for seabirds (Camphuysen et al. 2004). Wind speed was recorded throughout the survey transects, and thus represents real-time conditions. The remotely sensed dynamic covariates were based on daily or monthly mean values. Remote sensing data may be subject to data gaps from cloud cover. In cases where daily values were subject to missing data, monthly mean values were used. In addition, environmental climatologies were calculated for each of the dynamic covariates independently, by calculating the average value across January to March from 2003 to 2011. Contemporaneous and climatological values for each dynamic covariate, were extracted for each data point during the corresponding time period, and each of these were evaluated as described below. Covariates which were correlated by  $> 0.7$  were not included in the same models. Instead, the variable which improved model performance the most (as described below) was included.

### *Species distribution modelling*

Hurdle models were used to model the relationship between seabird sightings and environmental covariates, for each species independently. Hurdle models comprise two component models; firstly the probability of occurrence is modelled using a presence-absence General Linear Model (GLM) with a binomial error structure, and secondly the relative density of individuals, conditional on their presence, is modelled using a GLM with a zero truncated negative binomial error structure (Zuur et al. 2009). The hurdle approach helps to overcome the statistical challenges associated with zero-inflation and overdispersion of data, as well as providing multi-scale estimates of habitat use (e.g. Goetz et al. 2012, Sveegaard et al. 2012, Waggitt et al. 2020): This approach provides two descriptors of habitat use, discriminating between persistent presence of small groups and occasional presence of large groups (Waggitt et al. 2020). The biogeographical range of most seabird species may reflect the environmental conditions which influence the distribution of their prey, such as temperature and



depth, and this is represented by the presence-absence model. The clusters of animals within this range may be associated with the environmental conditions which aggregate prey, such as eddies and fronts, represented by MSLA, EKE and C, and modelled using the density model (Waggitt et al. 2020). To account for non-linear relationships between the environmental predictors and the response variables the second order polynomial of the values of environmental covariates were evaluated in addition to the linear value. Second order polynomials were chosen rather than polynomials of a higher order, or General Additive Models, to ensure ecologically plausible relationships and reduce the likelihood of overfitting models (Waggitt et al. 2020). Model residuals were checked for spatial and temporal autocorrelation using Morans-I tests and Auto Correlation Function (ACF) plots. Uncertainty in model predictions was calculated for the final models using a non-parametric bootstrapping approach (Davison and Hinkley 1997, Fieberg et al. 2020). For each species in turn, the data were resampled with replacement, selecting the same number of data points as in the original data frame. The resampled data were input into the final hurdle model, and the model was then used to predict the relative density of individuals across the study area, and the predicted density in each grid cell was stored. This was repeated 1000 times for each species and the difference between the 5% and 95% quantiles of predicted density in each grid cell was calculated. This provides a measurement of absolute uncertainty i.e. the absolute difference between upper and lower boundaries. For comparisons between species, this was standardised by dividing by the maximum value. In order to identify the covariates which provided the highest predictive power for the final hurdle model, we evaluated each component of the hurdle model independently (i.e. the presence-absence and the density models), before combining the two models.

#### Presence-absence models

The presence or absence of a species in each cell was modelled as a function of each of the specified broad-scale covariates independently (distance to colony, SST, Depth, Chl, Wind speed) using GLMs with a binomial error structure. As all data points represented 3 km of transect, there was no need to account for variation in effort between data points.

Model performance was evaluated using six-fold cross validation (Wood 2006). Data were split into six equal groups randomly and models were developed using five of the six groups. The models were then used to predict presence or absence of the excluded group, and predictions were evaluated to determine the predictive power of the model. This process was carried out six times to ensure that all groups were used for model evaluation, and the mean Area Under the Curve (AUC) value across groups was taken. AUC was generated by producing a ROC curve using R package *pROC* (Robin et al. 2011)

and used to evaluate the predictive power of the models. AUC values may range from 0.5 to 1.0, where a value of 0.5 is no different than random and 1.0 indicates a perfect model (Fielding and Bell 1997). AUC values of 0.5–0.7 are often taken to indicate low accuracy, values of 0.7–0.9 indicate useful applications and values of > 0.9 indicate high accuracy (Swets 1988). Model selection followed the forwards stepwise selection approach. Each of the covariates was ranked according to the AUC value, and the highest-ranking covariate was selected. Each of the remaining covariates was added in turn, to the best model, and the covariate was retained in the model if the AUC value increased. This process continued, adding more covariates until the AUC value no longer increased (Wakefield et al. 2017).

#### Density models

The relative density of individuals (individuals m<sup>-2</sup>) when encountered was modelled as a function of each of the specified mesoscale variables (MSLA, EKE, Currents, Wind speed) using GLMs with a truncated negative binomial error structure in R package *countreg* (Zeileis and Kleiber 2008). We also evaluated null models (i.e. no covariate) to ensure that no incorrect associations were identified for species where there was low variation in the relative density of individuals. All points where no individuals were observed were removed from this part of the analysis as this part of the model deals with the density of individuals where they do occur (both models are combined when calculating the final predictions). As with presence-absence models, six-fold cross validation was used for model selection and forward stepwise selection was used to find the best model. Covariate models were compared to null models to identify whether adding an environmental covariate provided a better fit. Density models were evaluated using normalised root mean square error (instead of AUC) where lower values indicate less residual variance.

Once the presence-absence model and the density model with the highest predictive power had been identified, they were combined into a hurdle model using the *hurdle* function in package *pscl* (Zeileis et al. 2008). These models were then applied to predict the distribution of each species across the survey area at a scale of 1 km<sup>2</sup>. Areas in the southern Drake Passage where SST exceeded 2.5 °C (the maximum value in the sampled area) were excluded from the prediction data frame to avoid predicting beyond the range of the sampled data, and potentially inflating predictions in this area. The maximum values in the prediction data for all other covariates did not exceed those in the sampled area, negating concerns associated with extrapolating outside of the environmental conditions surveyed. For species where wind speed was included, the mean value was used for predictions, to represent typical survey conditions. The predictions were cropped to exclude areas covered by the mean summer sea-ice extent (> 15% concentration, generally to the north east of the peninsula)

across the survey years, given that some of these species will be unable to forage in these conditions (Ainley et al. 1993). Finally, in order to identify areas with higher densities of seabirds in the region, we combined the predicted relative densities for each species: Firstly, the predicted relative densities were normalised by dividing the value for each grid cell by the sum of all grid cells for that species. Subsequently, the normalised predictions for each species were summed together to produce an overall plot of seabird density in the region.

## Results

Sightings of 11 species of procellariiform seabird occurred frequently across the survey years (Table 2, Figure 2). An average of 3,575 km (sd: 1,483 km) of surveys occurred each year with a maximum of 6,246 km of survey effort during 2011, and a total of 32,181 km across the nine years (this includes repeats of transects surveyed across multiple years). Survey effort occurred between seabed depths of ~20 m and ~5000 m, and sampling effort was highest in areas < 1 km deep (Figure S2). Surveys occurred in areas with SST between -0.05 °C and 2.5 °C, reflecting offshore ACC water and coastal Antarctic current and Weddell Sea surface water, and was highest in areas where temperature was ~2 °C (Figure S3). The frequencies of each species sighted across these temperature, depth and wind speed ranges are shown in Figures S2, S3 and S4. Although all species showed inter-annual variation in abundance, there appeared to be no major trend in abundance across survey years, except perhaps for wandering albatross and black-bellied storm petrel for which there appeared to be fewer sightings in recent years (Figure 2). For all species the majority of observations occurred during January, with considerably fewer individuals observed during February and March. This likely reflects the increased survey effort in January (ship transects across all years = 24,399 km) compared with February (5,577 km) and March (2,205 km).

The models which predict the presence-absence of seabirds performed moderately to well, with AUC values ranging from 0.60 to 0.80 (Table 3). There was little evidence of spatial or temporal autocorrelation in model residuals for most species, although negligible levels were detected for some species (southern fulmar, prions, Wilson's storm-petrel, black-bellied storm-petrel, Table S1, Figure S5). The models which predicted the relative density of individuals within this region were more variable, with NRMSE values ranging from 0.06 (good) to 0.26 (less good). In all cases the null models provided lower NRMSE values than the covariate models. MSLA and current speed were correlated (0.92) and thus only the highest scoring covariate was included in our final models. The best predictors for the presence-absence models varied among species (Table 3, Figure S6), however SST and depth were the covariates most frequently associated with seabird occurrence.

The distribution of species within the survey area was highly variable within and among seabird groups (Figure 3). However, the distribution of seabirds reflected two ecoregions: the southern Drake Passage (oceanic), and coastal Antarctic waters (coastal). Depth, SST and distance to breeding sites increase with distance from the Peninsula with highest values in the southern Drake Passage. When all species were standardised and combined the areas of highest density were in oceanic areas and in the mid Bransfield Strait (Figure 4). The uncertainty in model predictions were species specific, with blue petrels showing the highest uncertainty. For most species areas of highest uncertainty generally coincided with areas of highest density. For some species (e.g. southern giant petrel, southern fulmar and black-bellied storm-petrel) uncertainty was also high in areas where predicted density was lowest.

#### *Albatrosses*

Wandering and grey-headed albatross both showed an oceanic distribution (Figure 3) and were associated with increased SST, Chlorophyll (wandering and grey-headed) and depth (grey-headed, Figure S6), with the highest probability of occurrence over deeper, warmer water with higher chlorophyll concentration. Black-browed albatrosses showed a more widespread distribution, concentrated around the South Shetland Islands as well as further offshore, and associated with increased SST and shallower water (Figure S6).

#### *Petrels*

Southern giant petrels and southern fulmars showed a coastal distribution whereas prions, blue petrels, cape petrels and white-chinned petrels had an oceanic distribution (Figure 3). For southern giant petrels the probability of occurrence was highest in shallower waters with low chlorophyll-a concentrations near the colony (Figure S6). Fulmars had a quadratic association with depth and SST, peaking at depths of 2 – 3 km and SST  $\sim 0.5^{\circ} - 1^{\circ} \text{C}$  (Figure S6) with peak concentrations within the Bransfield Strait, over deep basins. Prions and cape petrels both had a quadratic relationship with depth, peaking  $\sim 3$  km deep, and were negatively associated with chlorophyll-a concentration. Prions were also associated with wind speed, peaking around 25 knots. Blue petrels preferred warm waters  $> 2^{\circ} \text{C}$  with wind speeds of 20-30 knots, and white-chinned petrels preferred deeper waters with SST  $\sim 2^{\circ} \text{C}$  (Figure S6).

#### *Storm-petrels*

Wilson's storm-petrel showed a highly coastal distribution (Figure 3) with probability of occurrence highest in areas closer to the colony, with SST of  $\sim 1^{\circ} \text{C}$  and low chlorophyll concentration and wind

speed (Figure S6). Conversely, black-bellied storm-petrels showed a more heterogenous distribution (Figure 3), with higher abundance in the Bransfield Strait and further offshore, yet very low abundance in areas close to the coast. Black-bellied storm-petrels showed an increased probability of occurrence at depths of ~3 km and in areas with low chlorophyll-a concentration and high wind speed (Figure S6).

## Discussion

Seabirds are an important component of the marine ecosystem, and without a better understanding of their baseline distributions in remote locations, we are limited in our ability to understand the impacts of anthropogenic disturbance and environmental change on their abundance and distribution. Specifically, relative seabird density can inform reference points in fishery management plans as well as scoping planning of Marine Protected Areas. In general, our models performed well in describing seabird distributions, although we noted some model limitations discussed below.

### Seabird habitat associations

The models highlighted two distinct patterns of species distribution; oceanic - concentrated in the southern Drake Passage and coastal – concentrated around the Peninsula and South Shetland Islands. Seabirds in the Southern Ocean feed predominantly on squid, fish and krill (Croxall et al. 1985), and the availability of species-specific key prey items is likely to drive the distribution of seabirds. As such, it might be expected that species with similar dietary preferences could utilise habitats with similar characteristics. This was observed to some extent in this study. For example, grey-headed and wandering albatrosses both feed on a variety of prey items, including a high proportion of squid (Croxall et al. 1985), also consuming a small proportion of krill in their diet. These species showed similar distributions, with higher concentrations predicted in the southern Drake Passage region than in the coastal Antarctic waters. Conversely, the predicted distribution of black-browed albatrosses, which generally consume a higher proportion of fish and krill and fewer squid, was more widely distributed occurring in both the oceanic and coastal Antarctic region (Santora et al. 2017b). Traditionally, black-browed albatrosses are considered a shelf-feeding species, often found in neritic or upper-shelf waters (Burg and Croxall 2001, Wakefield et al. 2011), and grey-headed and wandering albatrosses have a more oceanic distribution (Wood et al. 2000). It has been well documented that segregation of diet and foraging locations between these species provides a means of resource partitioning to reduce competition (Weimerskirch et al. 1988).

Another example is for the Wilson's and black-bellied storm-petrel, which are morphometrically similar, yet the Wilson's storm-petrel has a diet higher in krill, whereas the black-bellied storm petrel

consumes a larger amount of fish (Hahn 1998, Quillfeldt 2002). Our models indicate that Wilson's storm-petrel remain in coastal regions, whereas black-bellied storm petrels are also abundant further offshore and into the Drake Passage region. Whilst both species breed across the South Shetland Islands, previous studies have shown little overlap in their foraging distributions and abundance hotspots (Santora and Veit 2013). It is likely that these sympatrically breeding species are also resource partitioning in order to reduce competition. However, in other cases where one might expect to see similarities in the distribution of species with similar diets, this is not the case. For example, southern fulmars, blue petrels and cape petrels all consume 80 – 85 % krill in their diet (Croxall et al. 1985). Yet, observations and models suggest that fulmars remain in coastal waters, in areas with lower SST whereas blue petrels and cape petrels were concentrated in the Drake Passage region in areas of higher SST. Both cape petrels and fulmars breed in the region, and this may be constraining the distribution of fulmar to more coastal regions, although it is unclear why this constraint does not apply to cape petrels. It is plausible that species that consume high proportions of krill may prefer foraging further offshore, yet in some cases may be constrained to coastal areas during the breeding season. Since 87 % of krill has been estimated to be distributed in oceanic environments (Atkinson et al. 2008), this could explain the preferences we found for seabirds with a high proportion of krill in their diet. We recognise that species distributions change during the course of breeding, with brood or early chick-rearing being particularly constrained. For some management applications understanding how distributions differ during these periods will be important. Nevertheless, our analyses included data across different periods of breeding, enabling us to capture important generalities about species distributions.

#### *Broad scale presence-absence models*

SST and depth were the covariates in the most parsimonious models describing the broad-scale distribution of flying seabirds in this region, and at least one of these covariates was included in models for all species (Table 3, Figure S6). All models for species for which SST was important and with an oceanic distribution showed a positive or quadratic relationship between seabird occurrence and SST peaking at 2° C or above. For coastal species, this relationship peaked below 1° C. The relationship between oceanic species and depth was always positive or quadratic peaking at ~3 km or above. For coastal species this was more variable. Adding chlorophyll or distance to breeding colony into the models improved the predictive power of some models from both groups. Black-browed albatrosses did not clearly fit either group, instead showing more heterogeneous distributions across both habitat types. Some of the covariates selected in the final models showed some level of correlation (Table S2), so it is challenging to disentangle the exact effect each has on species distributions. However, the

addition of each variable in the final models improved their predictive power and as such we can be confident that they can be associated with the distribution of these species. Previous studies (e.g. Weimerskirch et al. 1997, Wakefield et al. 2011, Scales et al. 2016) provide strong evidence that all of these variables can influence the at-sea distribution of seabirds. We discuss how each of the variables in our final models may be associated with the distribution of seabirds.

SST is frequently associated with the preferred foraging habitats of many species of seabirds (e.g. Wakefield et al. 2011, Scales et al. 2016, Dehnhard et al. 2019), with birds generally showing a preference for a specific thermal range within their available habitat. This may be associated with the thermal requirements of the prey items sought, and the availability of lower trophic level species for consumption by these prey items. Alternatively, SST may indicate different water masses, for example oceanic versus regions of coastal influences. The broad scale variation of SST within the NAP reflects water masses among ACC, coastal currents in Bransfield Strait and Weddell Sea that are critical for zooplankton communities and krill species distribution (Loeb and Santora 2015).

Seabird foraging areas are often associated with bathymetric aspects, with many species showing clear preferences for neritic, shelf-break or upper shelf-slope waters (e.g. Weimerskirch et al. 1997, Warwick-Evans et al. 2016, Santora et al. 2017b), and others showing a preference for deeper oceanic waters (Hunt Jr et al. 1990). Upwelling associated with shelf edges and bathymetric features transport nutrients to the surface (Bakun 1996), increasing phytoplankton growth, and consequently concentration of higher trophic level species (Prézelin et al. 2000, Croxall and Wood 2002, Cox et al. 2018b). Additionally bathymetric features such as canyons, in nearshore areas may aggregate prey, or interact with mesoscale features leading to prey retention, increasing its availability to seabirds (Lavoie et al. 2000, Embling et al. 2012). In deeper waters mesoscale ocean dynamics may be enhancing biological processes, and aggregating prey. For example, mesoscale fronts and eddies such as the SACCF occur in the deeper oceanic waters, and it is likely that seabirds foraging in the oceanic environment are targeting such features. Indeed, models suggest that krill is advected in the ACC (Atkinson et al. 2008), with most of the population further offshore.

Chlorophyll concentration is frequently used to provide a measure of phytoplankton in the water column. Phytoplankton provides a key food supply for many prey species and, as such, chlorophyll concentration is frequently used as a proxy for prey availability. Relationships between chlorophyll concentration and the abundance of marine predators have been observed (e.g. Scott et al. 2010,

Santora et al. 2017a, Cox et al. 2018b), however, these relationships are complex for a number of reasons.

Firstly, it is likely that a time-lag exists between elevated chlorophyll concentration and elevated predator abundance, given that predators do not feed directly on phytoplankton, but on the small fish and krill which consume phytoplankton. As such it has been suggested that chlorophyll concentration may not adequately reflect prey abundance (Santora et al 2012). A time-lag between the abundance of predators and prey may also exist if prey move away from areas as predators arrive (Fauchald 2009). Additionally, areas of increased prey abundance may be less important to predators than areas of predictable prey availability (e.g. Boyd et al. 2015, Boyd et al. 2017, Waggitt et al. 2018), particularly for those species considered in this study, which generally forage in the upper water column and surface waters. Seabirds may also show hierarchical foraging behaviour between prey patches of varying spatial scales (Fauchald et al. 2000, Fauchald and Erikstad 2002), which may further explain why prey abundance does not always correspond with predator abundance at some spatial scales. Additionally, in some cases sub-surface maximum chlorophyll values (the maximum chlorophyll recorded at any depth across the water column) provides a better indication of increased prey abundance, than near-surface measurements (Scott et al. 2010, Embling et al. 2012). Finally, uncertainty in estimations of chlorophyll concentration using remote sensing approaches is often high in turbid waters, particularly in coastal areas (Cui et al. 2020), potentially biasing inferred relationships.

In this study the relationship between chlorophyll concentration and seabird density for those species for which chlorophyll was included in the final model, was negative. The majority of these species were those whose distribution was concentrated offshore in the ACC, and not in the coastal waters of the Bransfield Strait where chlorophyll concentrations were higher, due to spring ice break up. It is plausible that by including sub-surface chlorophyll concentration in these models, these relationships may be different, however we highlight the complexity in relating predator abundance with chlorophyll concentration.

Distance to the breeding colony was included in the final models for two of the five species breeding in this location and for which breeding locations are known. The probability of occurrence was negatively associated with distance to the colony for all models in which it was included. This covariate is frequently associated with distributions of central place foragers, including seabirds, which are constrained to return frequently to the breeding site to provision offspring (e.g. Wakefield et al. 2011, Warwick-Evans et al. 2018). As such, they are limited in the distance they may travel during foraging



trips, and it is not surprising that distance to the breeding colony is included in the models for some of these species. For three of the five species which breed in the region no association with distance to the colony were observed. This may be because not all Antarctic colony locations are known for many species, and species which are particularly wide ranging and oceanic are unlikely to remain near breeding colonies.

Wind speed showed variable relationships with the presence of four species of seabirds. Prions and blue petrels were associated with wind speed with peak relative density at wind speeds of 20 – 30 knots, whereas for Wilson’s storm-petrel this association was negative, and for black-bellied storm-petrel was positive. Peaks in relative density at mid-range wind speeds (non-linear) do not imply that wind speed is a significant driver as it is likely to be a result of interactions with habitat and location within the study area. Additionally, birds may be more flighty as wind speed increases, but this may be masked by waves at higher wind-speeds. The variation between two similar storm-petrel species, with varying distributions (oceanic and coastal), is an indication that it is indeed a result of interactions between habitat and location within the study area. It remains plausible that the association between Wilson’s storm-petrel and wind speed is related to detectability. This highlights the difficulty of accounting for detection when it naturally varies across environmental gradients of interest.

#### *Mesoscale density models*

In all of the density models the null model provided an equally good predictive power as the covariate model, and thus null models for density were used in the final hurdle models for all species. It is likely that different environmental drivers operate within coastal and oceanic domains given the complex bathymetry in coastal areas. Pelagic ecosystems are highly dynamic systems, and it is likely that the influence of ocean dynamics is variable across habitat types. Finer-scale and species-specific models may enable the detection of associations within one of these two habitats (coastal versus oceanic). This requires knowledge of the foraging behaviour and prey of the species, and the key physical processes in the region. The dynamic aggregation and dispersion of birds means that group size can be highly variable. As such, perhaps presence/absence is then driving distributions given that a species range is more or less consistent.

#### **Model performance and limitations**

Our AUC values indicate that the models generally performed moderately to well in their power to predict the presence or absence of seabirds within the study area. AUC values for covariate models for all species were higher than the null models, which supports the inclusion of these covariates in

the models. There was more variation in the NRMSE values, indicating that the density of individuals was more predictable in some species than others. The inclusion of null models instead of covariate models for density suggests that variations in aggregation size for many species are not linked to prominent environmental variations (that we have identified and included) in the region, and the main driver of distribution patterns is the variation in the presence-absence of a species rather than variations in their numbers. There may also be mismatch in scale between the fine-scale aggregations of birds, and the mesoscale biophysical variables that were included in the models, and as such, these may not accurately reflect prey distributions at smaller scales (e.g. Santora et al. 2009, Santora and Reiss 2011, Abrahms et al. 2018). Given the AUC values associated with the models we can be confident that the broad-scale prediction of the distributions of most species are indicative of their overall distributions. However, for species with a higher NRMSE value we should be cautious of over-interpreting the finer-scale densities within the broad species range.

The uncertainty in model predictions were highly species specific, although the highest uncertainty generally coincided with areas of highest density. This is likely to be a result of high variability in the abundance of individuals in each grid cell. For example, if the data selected in some model iterations included cells with high abundance, the predicted distributions in these areas would be large, however, if in other iterations cells with much lower abundance were sampled, these predictions would be lower. As a result the uncertainty would be higher in habitat types where these high abundances were observed. For some species (e.g. southern giant petrel, southern fulmar and black-bellied storm-petrel) uncertainty was also high in areas where predicted density was lowest. This may be because species which aggregate may have greater uncertainty due to variable group size, whereas solitary animals may have greater uncertainty in presence/absence.

Biases may be introduced into ecological models in a variety of ways, and we highlight two potential causes of bias in this study. Firstly, strip-transect methods assume that all seabirds up to a pre-defined distance are detected because observers concentrate on a small area. However, there is a chance that during some weather conditions birds may be missed. In this study, the majority of species are large and distinctly marked, and we believe that it is unlikely that many were missed. However, because we cannot guarantee this, we present relative rather than absolute densities. Additionally, in order to account for variation in detection rates between species as a result of weather conditions we have included wind speed as a sample covariate. As such we do not believe that our models are biased in this way. Secondly, there is potential inflation or deflation of apparent abundance due to movement

of flying birds relative to the survey platform. Seabirds may be attracted or repelled by the presence of the vessel (Bailey and Bourne 1972), with some species, including some albatross species being persistent ship-followers (Tasker et al. 1984). Additionally, the majority of flying birds travel faster than ship speed, and thus the chance of them entering the observers field is a function of their speed relative to the ship (Gaston et al. 1987). As such, Tasker et al. (1984), suggest a snapshot method is used to correct for bird movement, whereby a series of instantaneous counts of all flying birds occur regularly throughout the transect. In this study we used a continuous method to count flying seabirds and thus there is a chance that ship attraction or flux may introduce bias in our results. Accounting for all biases and errors remains challenging given that individuals cannot be uniquely identified. There was some evidence of autocorrelation in model residuals for four species, although this was deemed negligible (Diniz-Filho et al. 2003, Hawkins et al. 2007, Gaspard et al. 2019).

In order to conduct robust analyses with large sample sizes we included all seabird sightings in the analyses, including both flying and resting individuals. As such, we may have inadvertently included individuals that were commuting through a particular habitat type and not foraging. However, given that the majority of sightings recorded during transects were of individuals in flight, it was necessary to include all sightings. Many of these species are visual pursuit foragers, which forage after spotting prey items from the air, and thus an individual observed flying seabird is considered foraging. Additionally, previous studies show that time spent in an area can be used to infer foraging behaviour (Warwick-Evans et al. 2015), and it is likely that species were observed in areas where they were spending more time, and thus are likely to have been foraging.

For all species, using climatological values for SST and contemporaneous values for Chlorophyll-a concentration provided the best model fit. Mannocci et al. (2017) provide a detailed evaluation of the appropriate scales of environmental covariates for species distribution models. They conclude that associations with large scale and persistent oceanographic features are best modelled with climatological covariates. However, for mesoscale features where ephemeral processes are present or interannual variability occurs, contemporaneous covariates should be used, whereas if seasonal processes dominate and interannual variability is weak then climatologies are best (Mannocci et al. 2017). Given that all models that we evaluated which included SST provided the highest predictive power when climatological covariates were included, it is likely that SST is persistent across years, and represents different water masses. Conversely, contemporaneous values were the better predictors for all models which included chlorophyll-a concentration, which is likely to represent seasonality in

productivity, and be more variable. However, as previously discussed, relationships with chlorophyll are complex and not always intuitive.

One of the limitations of this type of survey is that it is not possible to determine the life history stage (sex or breeding status) of individuals observed, and for many species the distribution of non-breeders or failed breeders can vary greatly from that of breeding individuals (Phillips et al. 2005). Thus, these distributions may represent breeders from some species and non-breeders from others, and it is plausible that these broad scale habitat preferences may vary if models were conducted independently for specific life-history stages.

## **Conclusion**

Our models provide a baseline understanding of important habitat characteristics for Southern Ocean seabirds. This study describes how seabirds generally show either coastal or oceanic distributions, and that patterns are driven by presence/absence, with large scale climatic variables. Finer-scale processes, such as those at fronts or within submarine canyons, not associated with prominent environmental gradients could influence aggregations within these species' ranges. By gaining this baseline understanding, we can further enhance understanding about the impacts that environmental change and anthropogenic disturbance may have on ecologically important species in this remote and iconic environment. Understanding which species are likely to overlap with fisheries, and where these overlaps may occur, is an important consideration when establishing fisheries management frameworks. Baseline estimates of species distributions will be particularly important as the Antarctic marine ecosystem changes in the future, particularly in the context of ongoing climate change, but also with increased competition from recovering marine mammal populations, and as the potential impacts of krill harvesting are realised.

## **Data Availability**

All seabird sightings data are available upon request from Dr. Jarrod Santora at the National Oceanic and Atmospheric Administration, National Marine Fisheries Services, Southwest Fisheries Science Center; e-mail: jarrod.santora@noaa.gov.

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## Figure legends

Figure 1. Seabed depths and some of the key bathymetric features in the study area. Survey transects are indicated in white.

Figure 2. The number of individuals observed per km of effort during seabird observations throughout the survey period.

Figure 3. Observed and predicted distributions of flying seabirds around the western Antarctic Peninsula and South Shetland Islands; a) gridded observations of species density obtained from ship surveys, aggregated across all years of the survey (models were fitted to data for each year independently) b) number of individuals observed in each grid cell obtained from ship surveys and overlaid on predicted distributions of the relative density of species from model predictions. White dots indicate cells surveyed, black dots indicate observed individuals, increasing in size with the number of individuals observed c) predicted distributions of the relative density of species from models, d) Uncertainty in model predictions, calculated using a non-parametric bootstrapping approach.

Figure 4. The combined predicted distributions of eleven species of procellariiform around the northern Antarctic Peninsula and South Shetland Islands. Predicted relative densities for each species were normalised prior to summing all species.

# Figures

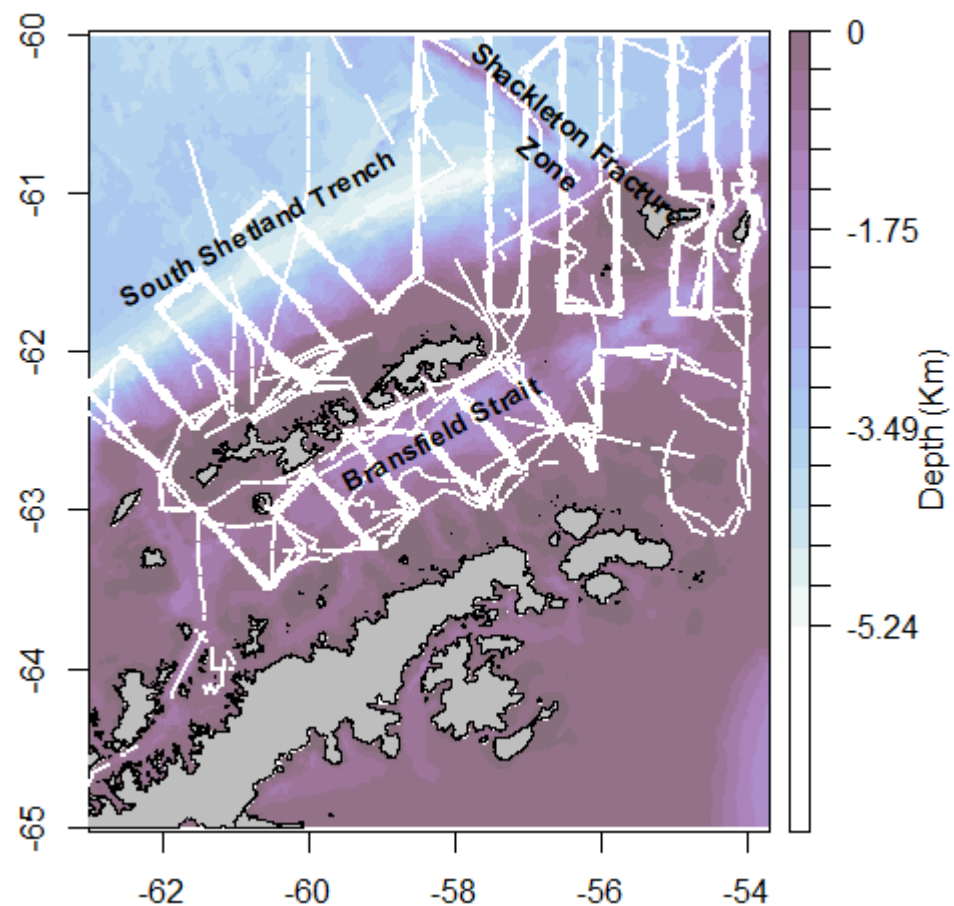
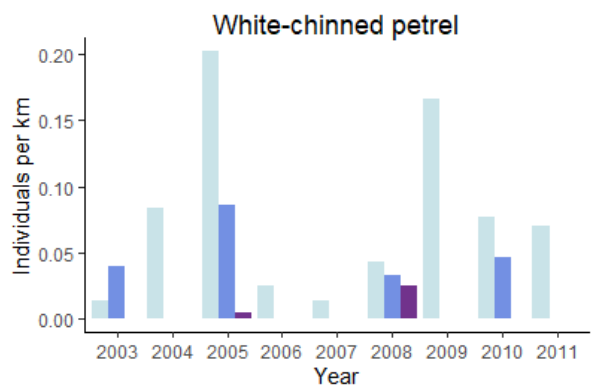
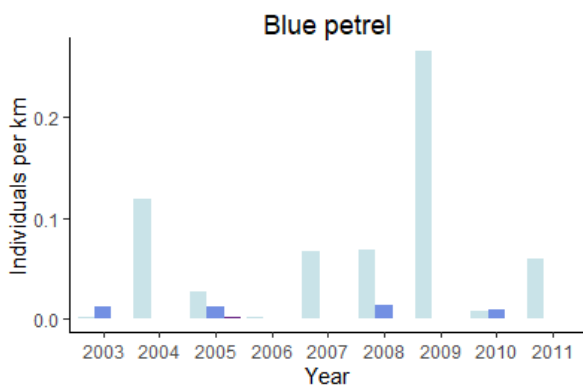
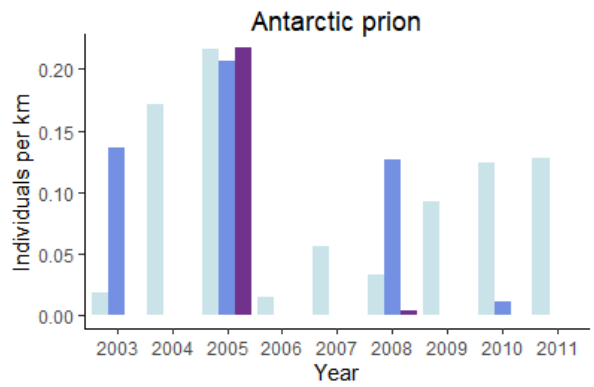
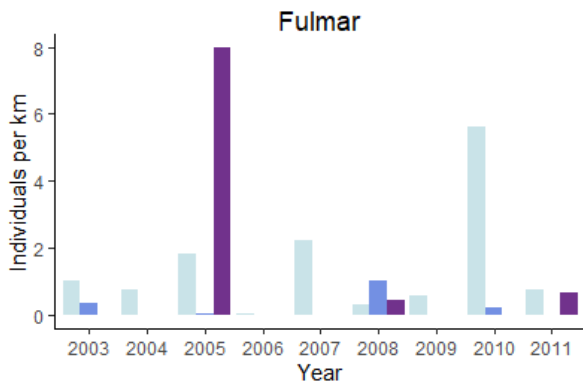
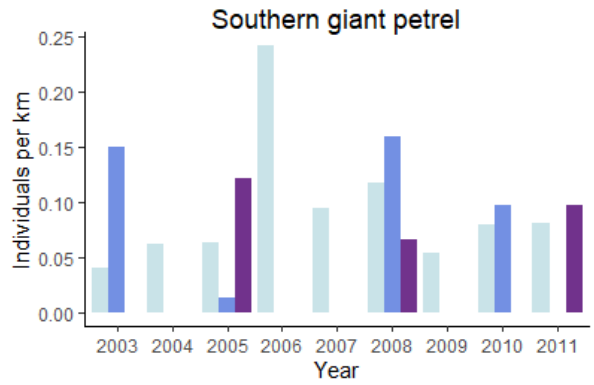
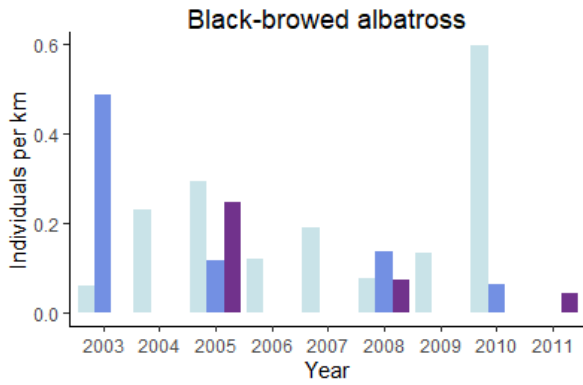
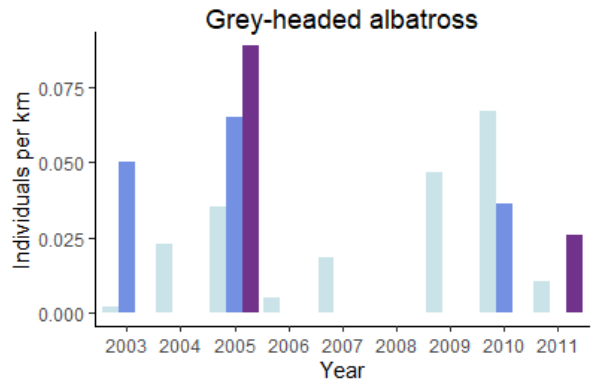
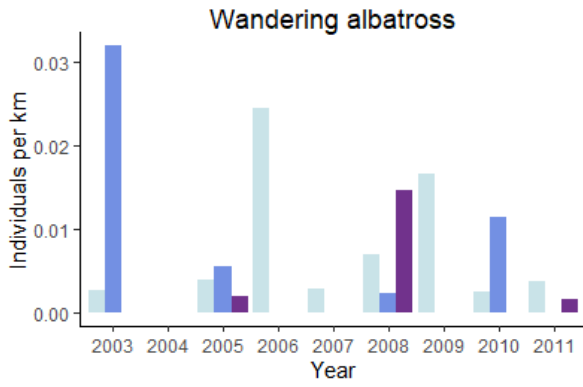
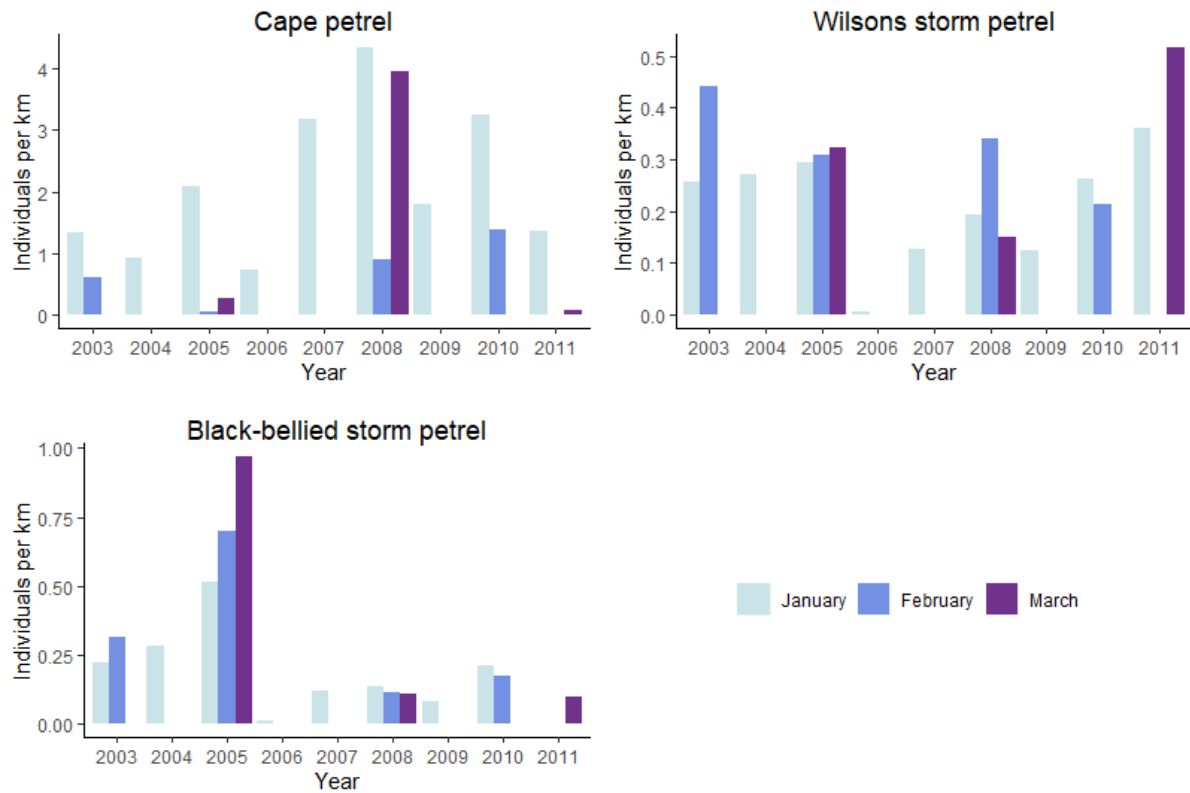


Figure 1. Seabed depths and some of the key bathymetric features in the study area. Survey transects are indicated in white.



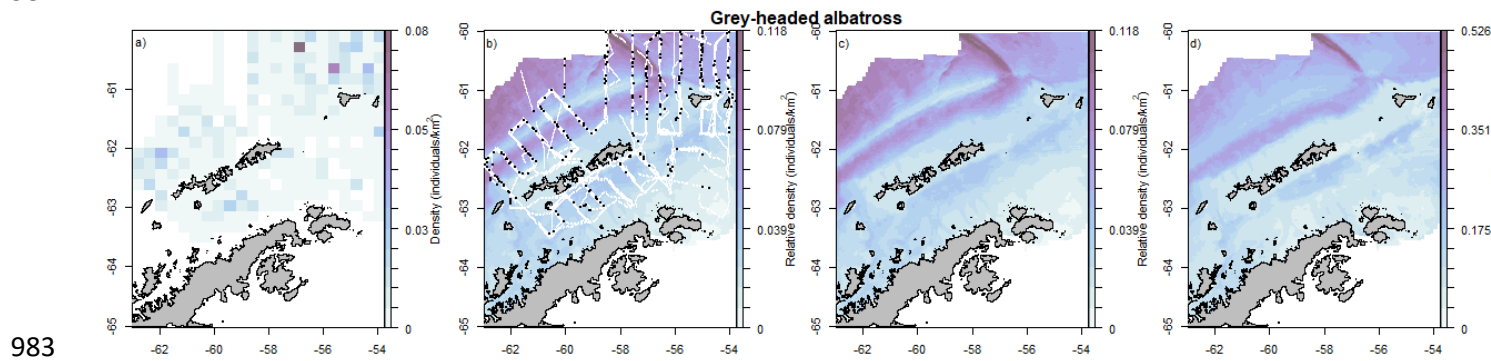
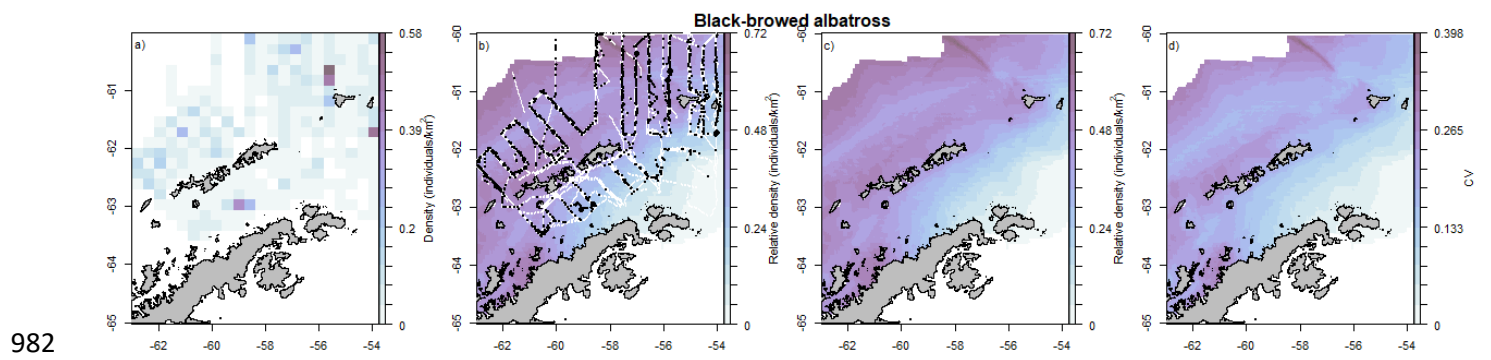
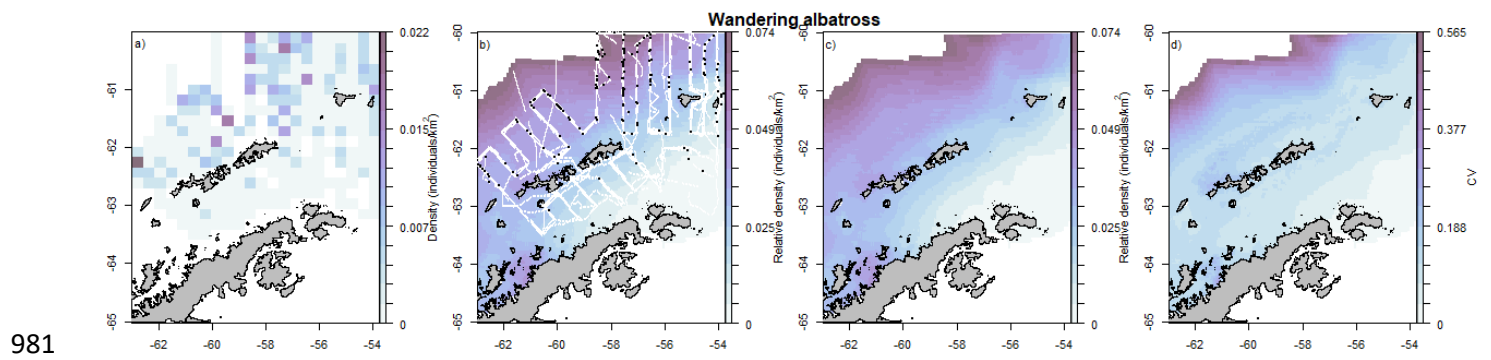


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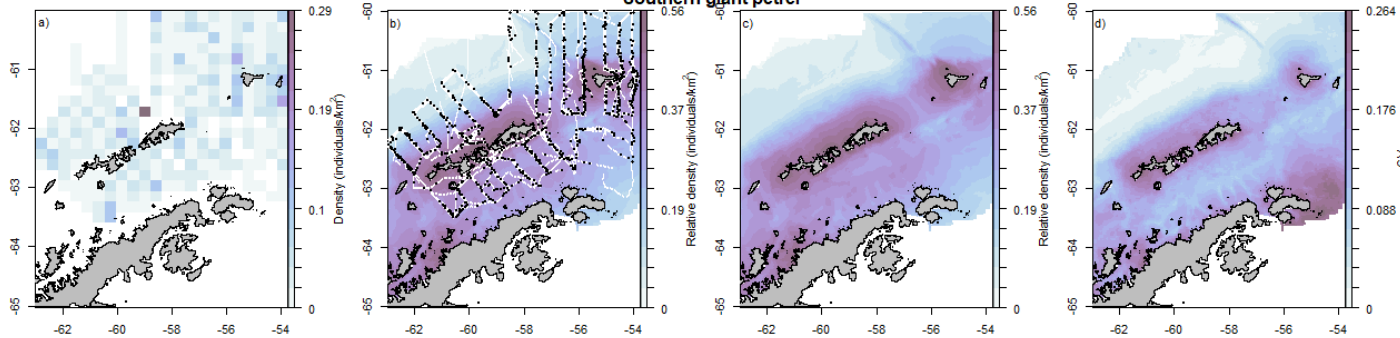
975 Figure 2. The number of individuals observed per km of effort during seabird observations  
 976 throughout the survey period.

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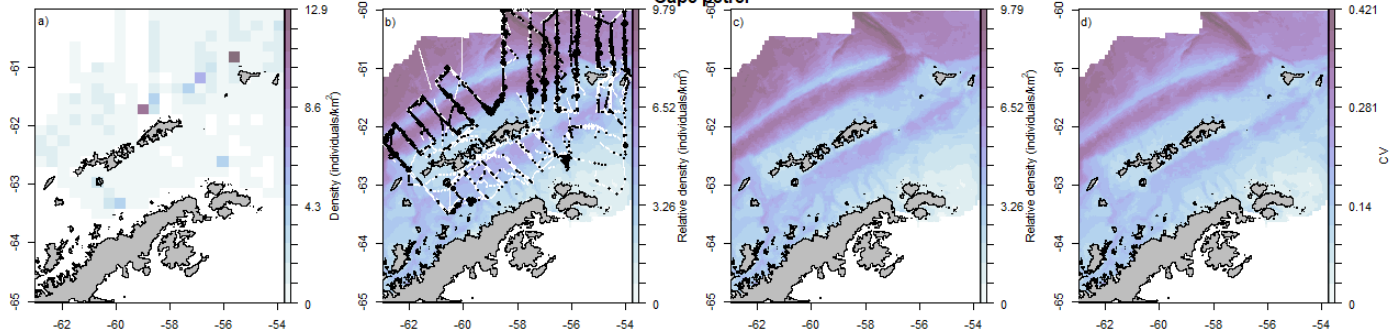
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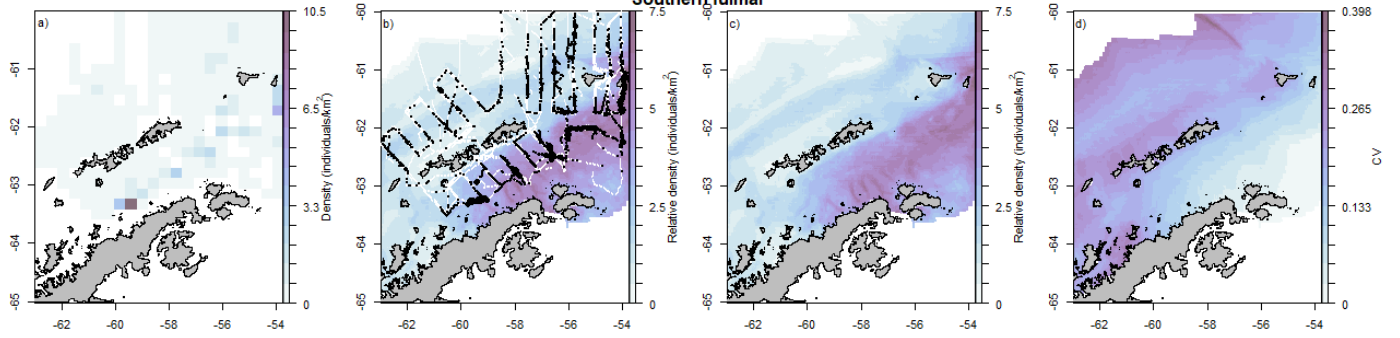
**Southern giant petrel**



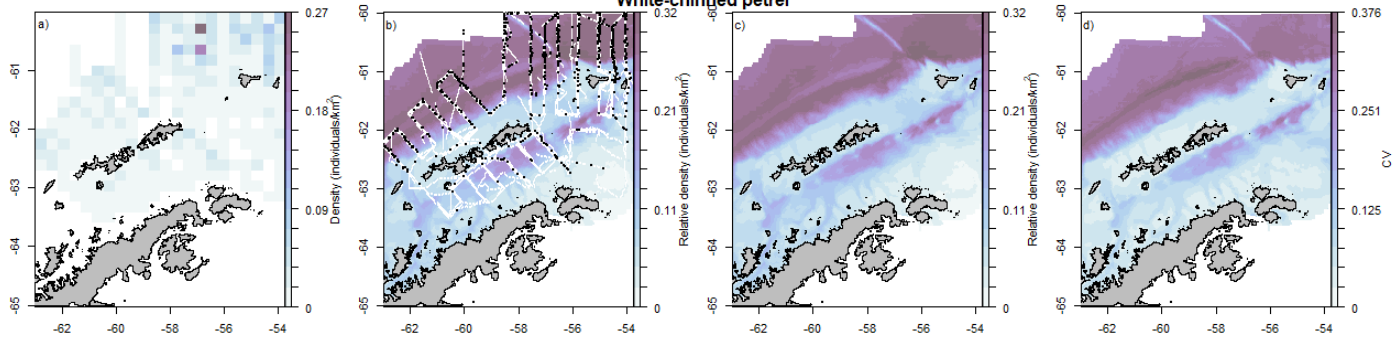
**Cape petrel**



**Southern fulmar**



**White-chinned petrel**





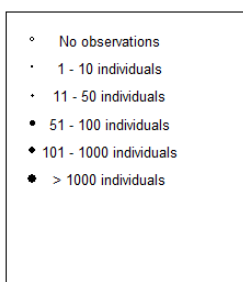
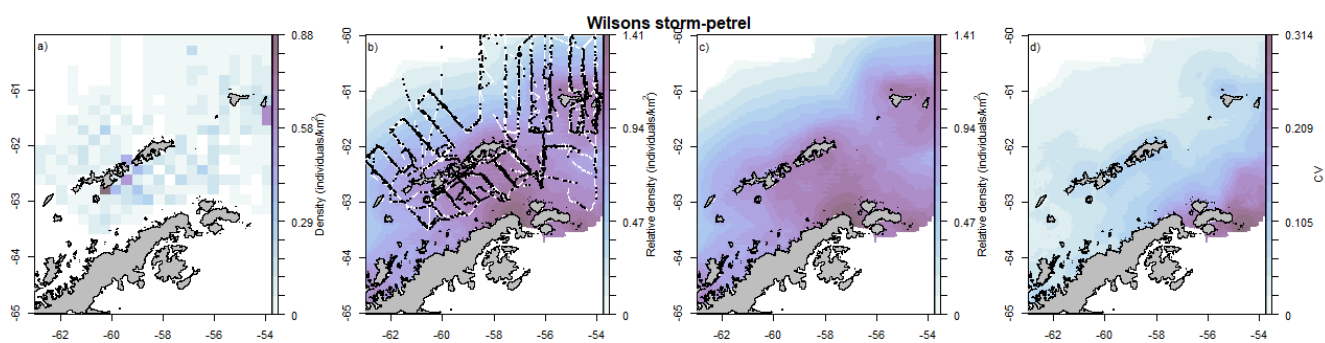
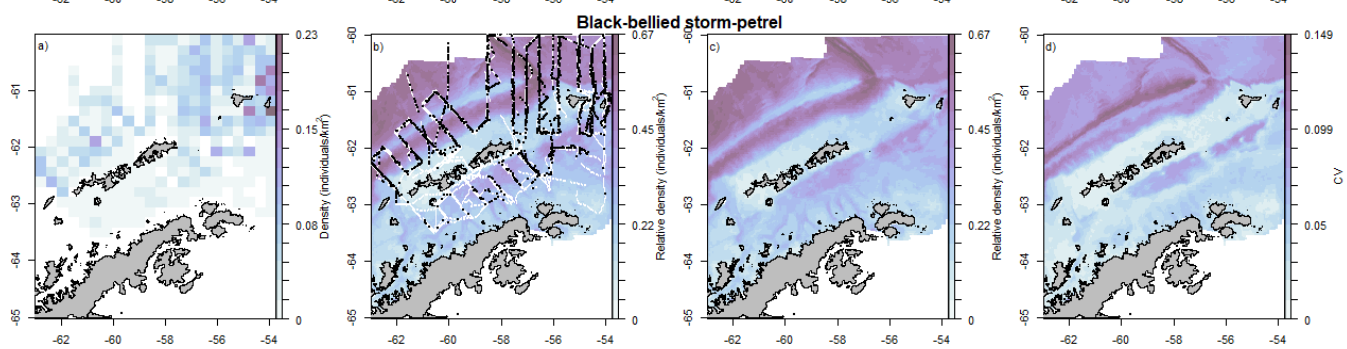
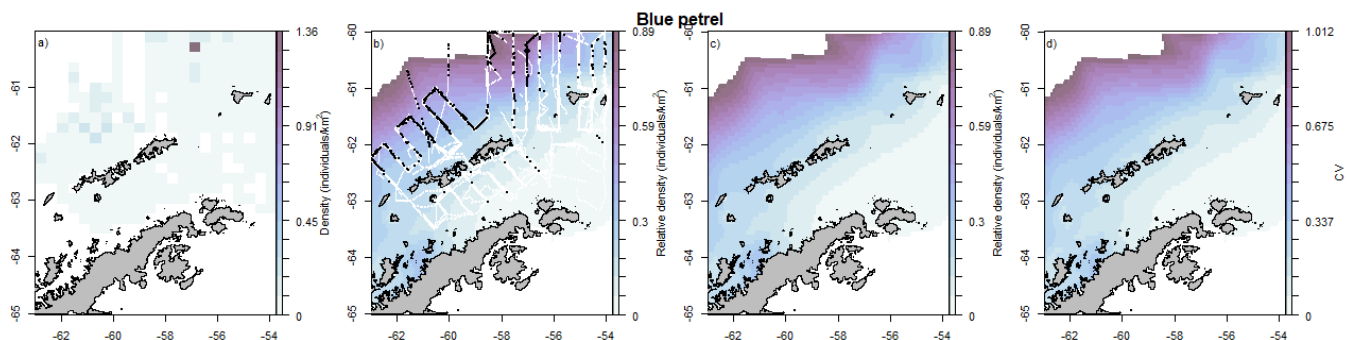
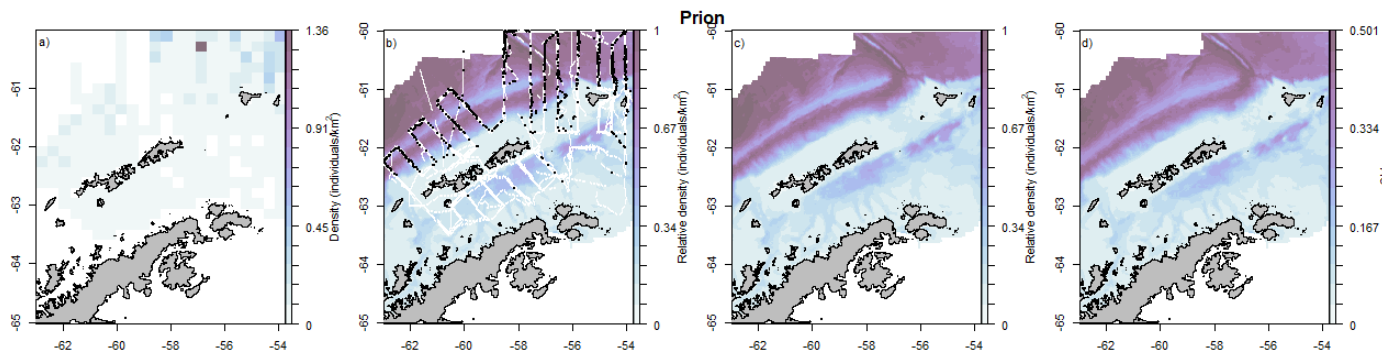
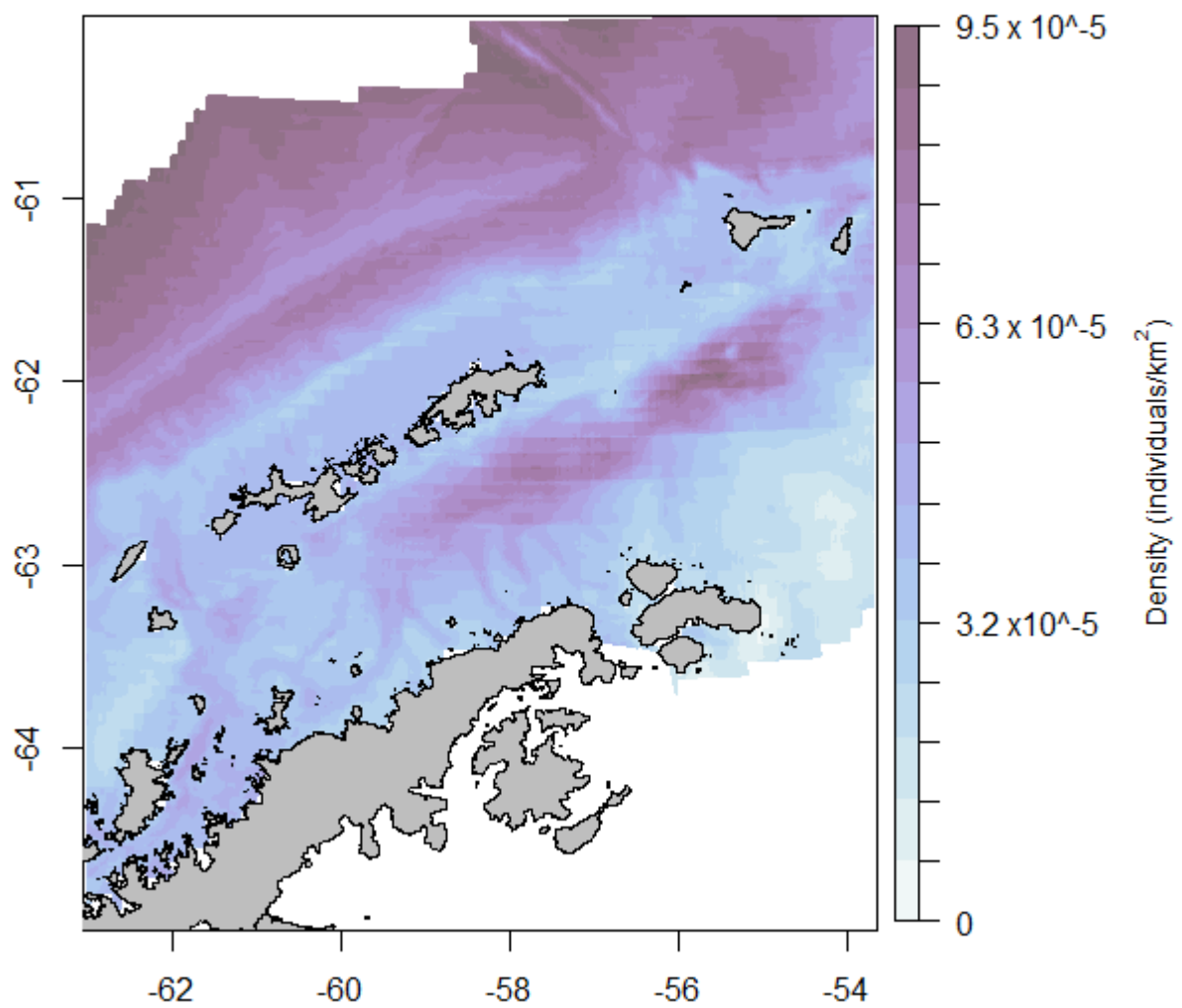


Figure 3. Observed and predicted distributions of flying seabirds around the western Antarctic Peninsula and South Shetland Islands; a) gridded observations of species density obtained from ship surveys, aggregated across all years of the survey (models were fitted to data for each year independently) b) number of individuals observed in each grid cell obtained from ship surveys and overlaid on predicted distributions of the relative density of species from model predictions. White dots indicate cells surveyed, black dots indicate observed individuals, increasing in size with the number of individuals observed c) predicted distributions of the relative density of species from models, d) Uncertainty in model predictions, calculated using a non-parametric bootstrapping approach.

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Figure 4. The combined predicted distributions of eleven species of procellariiform around the northern Antarctic Peninsula and South Shetland Islands. Predicted relative densities for each species were normalised prior to summing all species.

## Tables

**Table 1.** The explanatory variables used in statistical models to predict the at-sea distribution of seabirds

| Model            | Covariate                          | Spatial resolution | Temporal resolution | Source   |
|------------------|------------------------------------|--------------------|---------------------|--|
| Presence-absence | Distance from colony ( $D_{col}$ ) | 0.3 (km)           | NA                  | Calculated using R package <i>Raster</i>   |
|                  | Bathymetry (Depth)                 | 0.3 (km)           | NA                  | <a href="http://www.marine.copernicus.eu">www.marine.copernicus.eu</a>                                 |
|                  | Chlorophyll-a (Chl)                | 4.0 (km)           | Monthly             | <a href="http://www.marine.copernicus.eu">www.marine.copernicus.eu</a>                                 |
|                  | Sea Surface Temperature (SST)      | 0.083 degree       | Daily               | <a href="http://www.marine.copernicus.eu">www.marine.copernicus.eu</a>                                 |
|                  | Wind Speed (W)                     |                    | Continuous          | Collected throughout transects   |
| Density model    | Mean Sea Level Anomaly (MSLA)      | 0.25 degree        | Daily               | <a href="http://www.marine.copernicus.eu">www.marine.copernicus.eu</a>                                 |
|                  | Eddy Kinetic Energy (EKE)          | 0.083 degree       | Daily               | Calculated in R using data from <a href="http://www.marine.copernicus.eu">www.marine.copernicus.eu</a> |
|                  | Current Speed (C)                  | 0.083 degree       | Daily               | Calculated in R using data from <a href="http://www.marine.copernicus.eu">www.marine.copernicus.eu</a> |
|                  | Wind Speed (W)                     |                    | Continuous          | Collected throughout transects   |

**Table 2.** A summary of the number of sightings and individuals observed during ship surveys of seabirds in the west Antarctic Peninsula, and South Shetland Island region.

| Species                    | Number of individuals observed | Number of sightings observed | Mean (sd) flock size | Median (Interquartile range) flock size | Max flock size |
|----------------------------|--------------------------------|------------------------------|----------------------|---|----------------|
| Wandering albatross        | 260                            | 208                          | 1.1 (0.5)            | 1 (0)                                   | 6              |
| Black-browed albatross     | 4,728                          | 1,968                        | 1.7 (5.5)            | 1 (0)                                   | 120            |
| Grey-headed albatross      | 729                            | 556                          | 1.2 (1.4)            | 1 (0)                                   | 30             |
| Southern giant petrel      | 2,955                          | 1,448                        | 1.7 (2.8)            | 1 (0)                                   | 60             |
| Southern fulmar            | 34,633                         | 2,582                        | 5.7 (32)             | 1 (2)                                   | 1300           |
| Prions                     | 6,871                          | 3,073                        | 2.2 (9.8)            | 1 (1)                                   | 500            |
| Blue petrel                | 1,894                          | 629                          | 2.0 (16)             | 1 (0)                                   | 500            |
| White-chinned petrel       | 2,059                          | 1,183                        | 1.4 (3.1)            | 1 (0)                                   | 100            |
| Cape petrel                | 51,685                         | 2,992                        | 7.6 (63)             | 2 (2)                                   | 2500           |
| Wilson's storm-petrel      | 7,802                          | 3,215                        | 1.5 (2.5)            | 1 (0)                                   | 100            |
| Black-bellied storm-petrel | 6,114                          | 2,718                        | 1.4 (1.1)            | 1 (0)                                   | 30             |

**Table 3.** A summary of the explanatory variables used in the final models to describe the distribution of seabirds in the west Antarctic Peninsula, and South Shetland Island region. Quantitative evaluation of presence-absence and density predictions using area under the curve (AUC) and normalised root mean squared error (NRMSE). NRMSE values for intercept only (null) models only are provided.

| Species                    | Presence-absence model   | AUC  | NRMSE<br>null<br>model |
|----------------------------|--|------|------------------------|
| Wandering albatross        | T + Chl  | 0.7  | 0.26                   |
| Black browed albatross     | T + Depth  | 0.6  | 0.09                   |
| Grey-headed albatross      | Depth + Depth <sup>2</sup> + Chl + Chl <sup>2</sup> + T                  | 0.64 | 0.21                   |
| Southern giant petrel      | Depth + Chl + Chl <sup>2</sup> + Distance to colony                      | 0.63 | 0.12                   |
| Southern fulmar            | Depth + Depth <sup>2</sup> + T + T <sup>2</sup> + Chl + Chl <sup>2</sup> | 0.74 | 0.07                   |
| Prions                     | Depth + Depth <sup>2</sup> + W + W <sup>2</sup> + Chl                    | 0.75 | 0.15                   |
| Blue petrel                | T + W + W <sup>2</sup>   | 0.80 | 0.18                   |
| White-chinned petrel       | Depth + T + T <sup>2</sup>   | 0.67 | 0.1                    |
| Cape petrel                | Depth + Depth <sup>2</sup> + Chl + T                                     | 0.64 | 0.06                   |
| Wilson's storm petrel      | T + T <sup>2</sup> + Chl + Chl <sup>2</sup> + Distance to colony + W     | 0.7  | 0.06                   |
| Black-bellied storm petrel | Depth + Depth <sup>2</sup> + Chl + W                                     | 0.63 | 0.11                   |

## Supplementary Information

Table S1. Estimates for autocorrelation in model residuals using global *Morans I* tests

| Species                | Moran I<br>estimate | P-value |
|------------------------|---------------------|---------|
| White-chinned petrel   | 0.02                | 0.08    |
| Wandering albatross    | 0.017               | 0.12    |
| Prions                 | 0.17                | <0.001  |
| Grey-headed albatross  | 0.006               | 0.32    |
| Southern giant petrel  | 0.16                | 0.11    |
| Southern fulmar        | 0.08                | <0.001  |
| Black-browed albatross | 0.006               | 0.67    |

|                            |                      |        |
|----------------------------|----------------------|--------|
| Cape petrel                | 0.01                 | 0.15   |
| Blue petrel                | $4.1 \times 10^{-4}$ | 0.36   |
| Black-bellied storm-petrel | 0.11                 | <0.001 |
| Wilson's storm-petrel      | 0.09                 | <0.001 |

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**Table S2.** Pearson's correlation coefficients between the environmental explanatory variables used in model selection. Covariates for which correlation was > 0.7 were not included in the same models.

| var 1   | var 2      | correlation |
|---|------------|-------------|
| EKE   | Wind speed | 0.1         |
| MSLA  | Wind speed | 0.09        |
| Current                                       | Wind speed | 0.13        |
| SST   | Wind speed | 0.14        |
| Chl   | Wind speed | 0.15        |
| Distance to fulmar colony                     | Wind speed | 0.12        |
| Distance to Wilson's storm-petrel colony      | Wind speed | 0.21        |
| Distance to southern giant petrel colony      | Wind speed | 0.20        |
| Distance to cape petrel colony                | Wind speed | 0.19        |
| Distance to black-bellied storm-petrel colony | Wind speed | 0.08        |
| Depth   | Wind speed | 0.17        |
| EKE   | Depth      | 0.28        |
| MSLA  | Depth      | 0.33        |
| Current                                       | Depth      | 0.56        |
| SST   | Depth      | 0.62        |
| Chl   | Depth      | 0.68        |
| Distance to fulmar colony                     | Depth      | 0.60        |
| Distance to Wilson's storm-petrel colony      | Depth      | 0.57        |
| Distance to southern giant petrel colony      | Depth      | 0.58        |
| Distance to cape petrel colony                | Depth      | 0.58        |
| Distance to black-bellied storm-petrel colony | Depth      | 0.35        |
| MSLA  | EKE        | 0.18        |
| Current                                       | EKE        | 0.90        |
| SST   | EKE        | 0.41        |
| Chl   | EKE        | 0.31        |
| Distance to fulmar colony                     | EKE        | 0.31        |
| Distance to Wilson's storm-petrel colony      | EKE        | 0.10        |
| Distance to southern giant petrel colony      | EKE        | 0.09        |
| Distance to cape petrel colony                | EKE        | 0.11        |
| Distance to black-bellied storm-petrel colony | EKE        | 0.04        |
| Current                                       | MSLA       | 0.21        |
| SST   | MSLA       | 0.34        |
| Chl   | MSLA       | 0.09        |



|   |         |      |
|---|---------|------|
| Distance to fulmar colony                     | MSLA    | 0.10 |
| Distance to Wilson's storm-petrel colony      | MSLA    | 0.11 |
| Distance to southern giant petrel colony      | MSLA    | 0.01 |
| Distance to cape petrel colony                | MSLA    | 0.11 |
| Distance to black-bellied storm-petrel colony | MSLA    | 0.24 |
| SST   | Current | 0.52 |
| Distance to fulmar colony                     | Current | 0.35 |
| Distance to Wilson's storm-petrel colony      | Current | 0.11 |
| Distance to southern giant petrel colony      | Current | 0.1  |
| Distance to cape petrel colony                | Current | 0.12 |
| Distance to black-bellied storm-petrel colony | Current | 0.08 |
| Chl   | Current | 0.30 |
| Chl   | SST     | 0.29 |
| Distance to fulmar colony                     | SST     | 0.45 |
| Distance to Wilson's storm-petrel colony      | SST     | 0.16 |
| Distance to southern giant petrel colony      | SST     | 0.01 |
| Distance to cape petrel colony                | SST     | 0.14 |
| Distance to black-bellied storm-petrel colony | SST     | 0.03 |
| Distance to fulmar colony                     | Chl     | 0.42 |
| Distance to Wilson's storm-petrel colony      | Chl     | 0.64 |
| Distance to southern giant petrel colony      | Chl     | 0.63 |
| Distance to cape petrel colony                | Chl     | 0.65 |
| Distance to black-bellied storm-petrel colony | Chl     | 0.47 |

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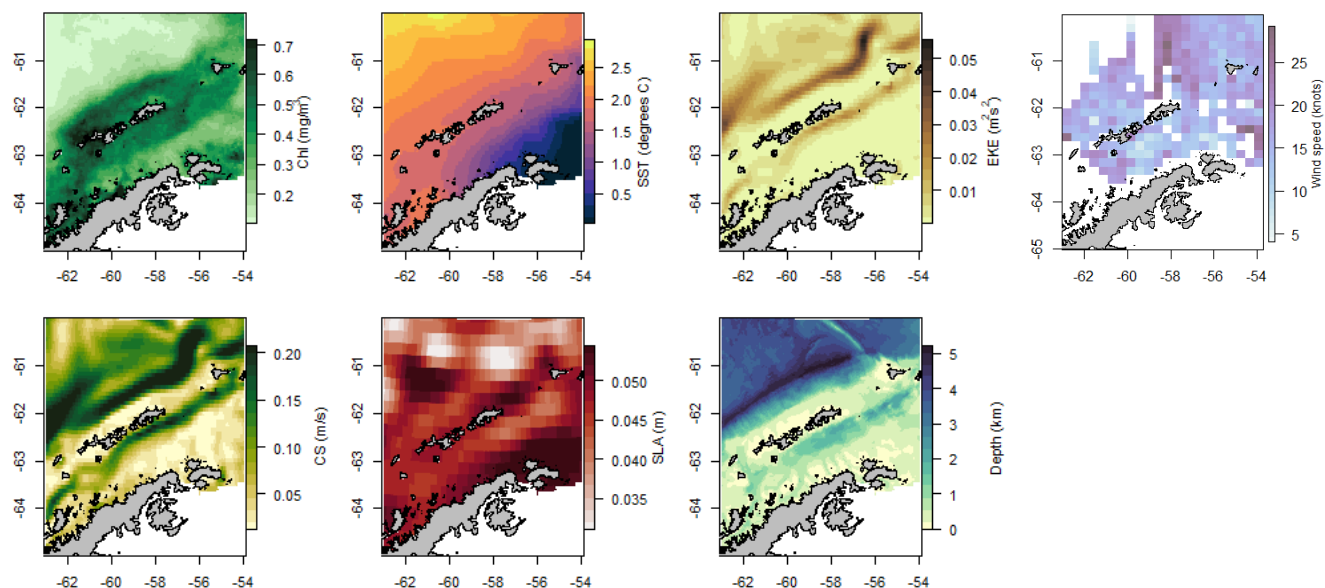
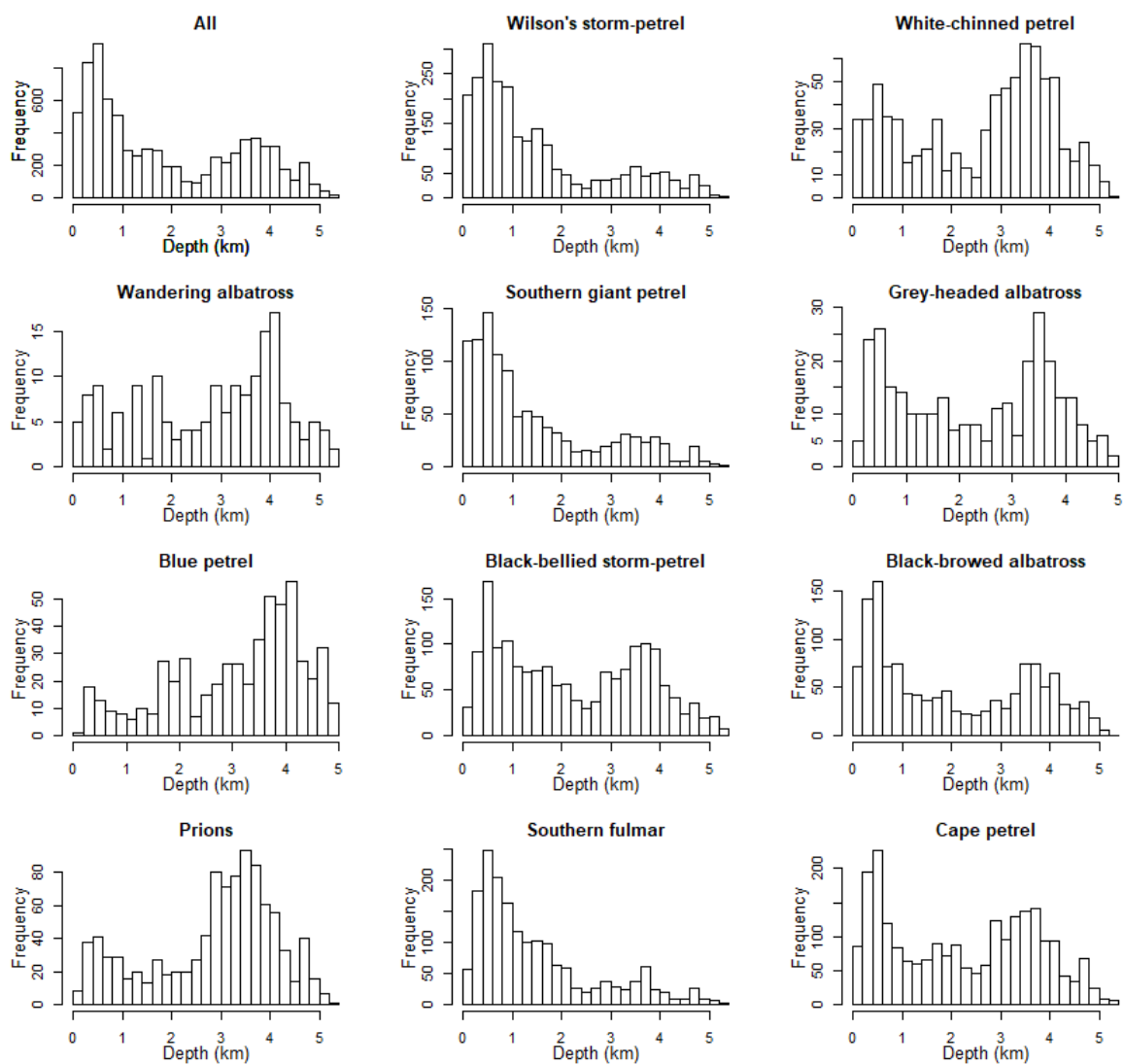


Figure S1. Climatological values for the environmental variables used in hurdle models to describe the at-sea distribution of seabirds: a) Chlorophyll-a concentration, b) Sea surface temperature, c) Eddy kinetic energy, d) Current speed, e) Sea level anomaly, f) Depth.

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1052 Figure S2. The frequency of sampling points a the frequency of individuals of each species sighted at  
1053 different depths.

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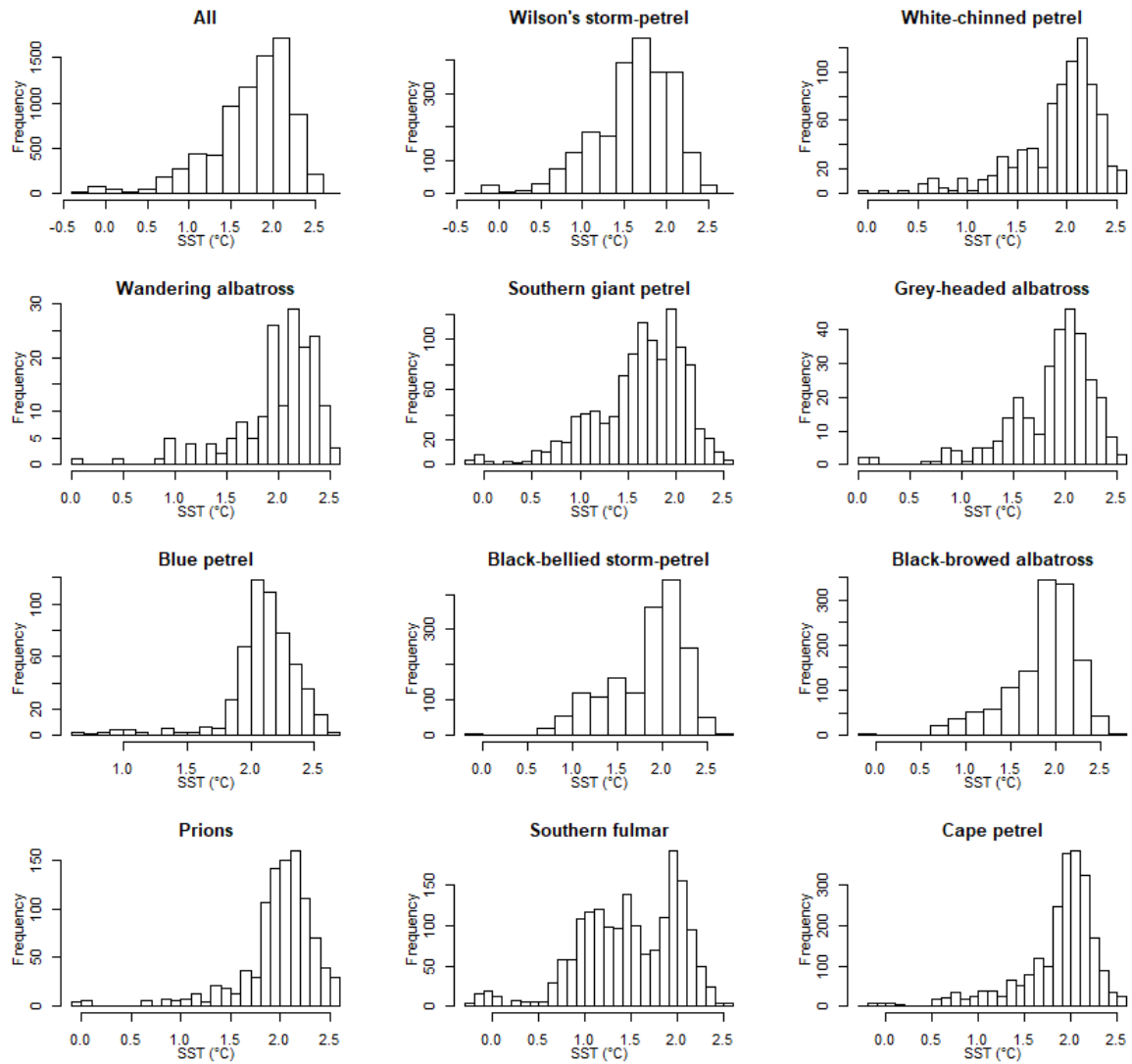
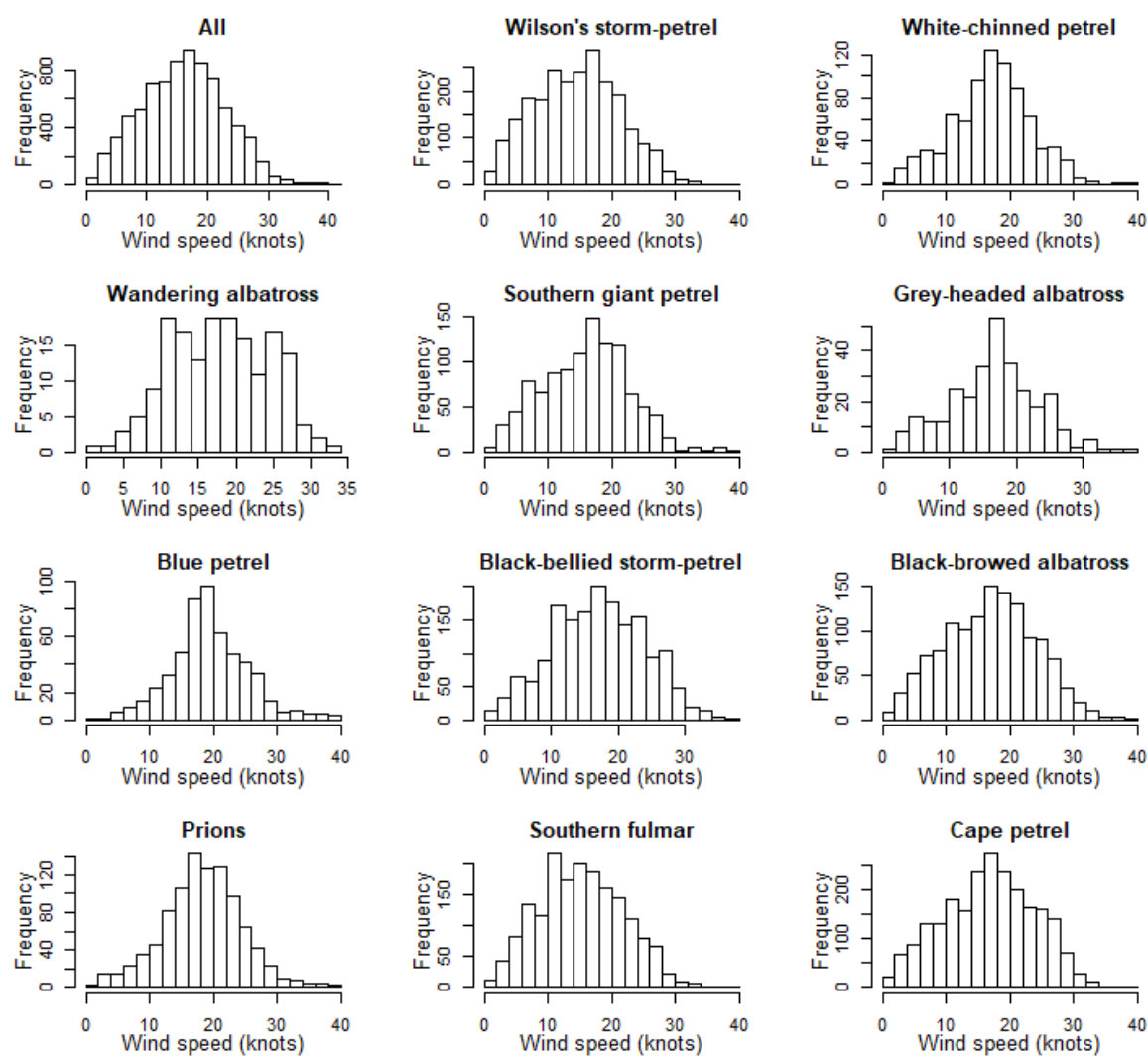


Figure S3. The frequency of sampling points and frequency of individuals of each species sighted across the temperature range.

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1063 Figure S4. The frequency of sampling points and frequency of individuals of each species sighted  
1064 across the wind speed range.

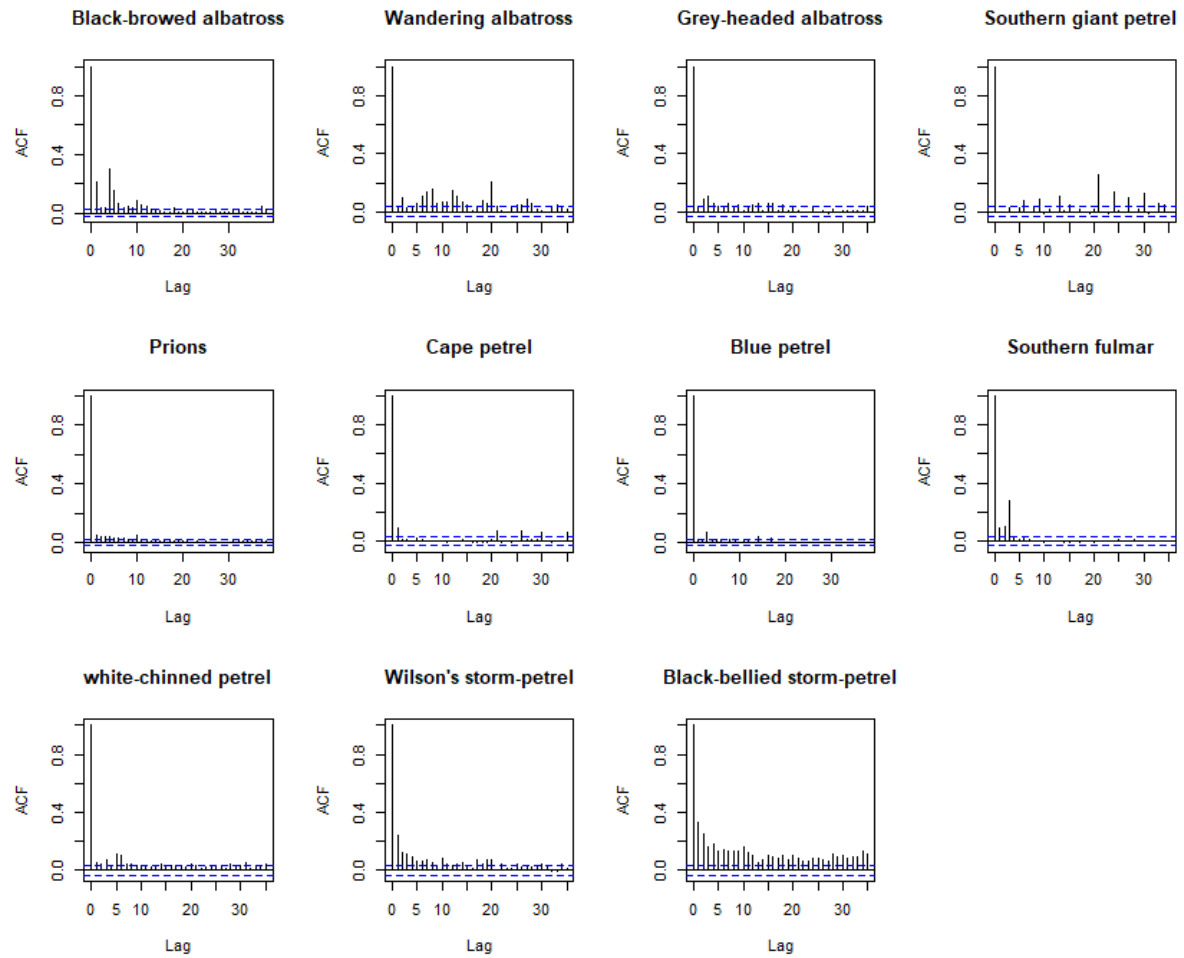
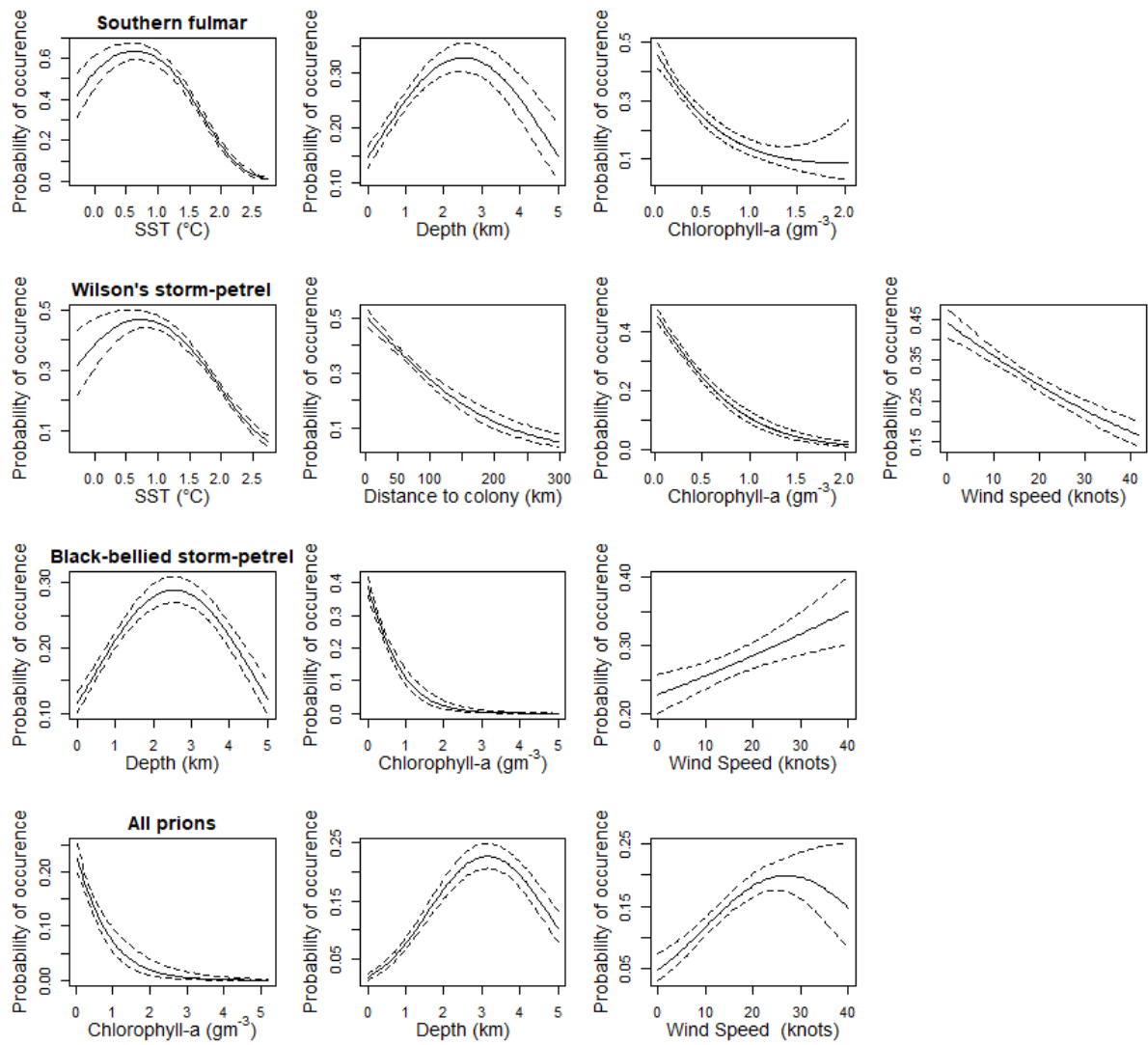
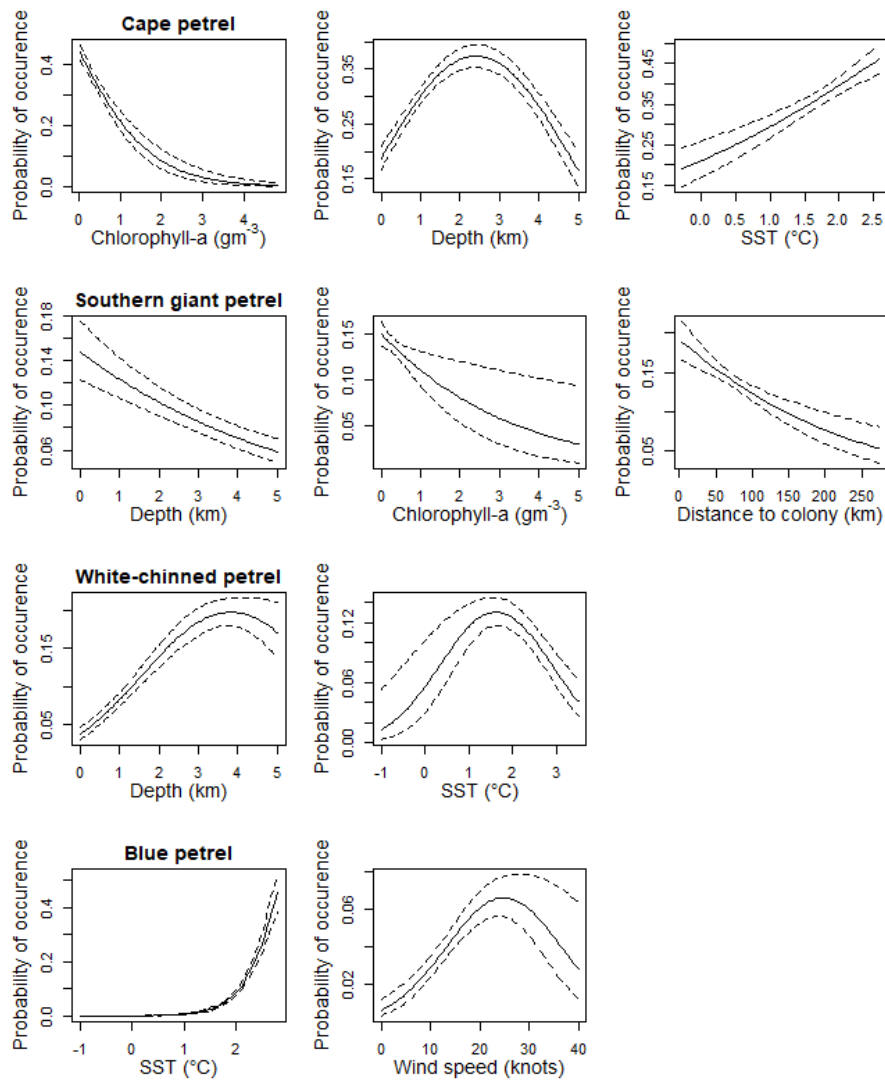


Figure S5. Autocorrelation plots for model residuals



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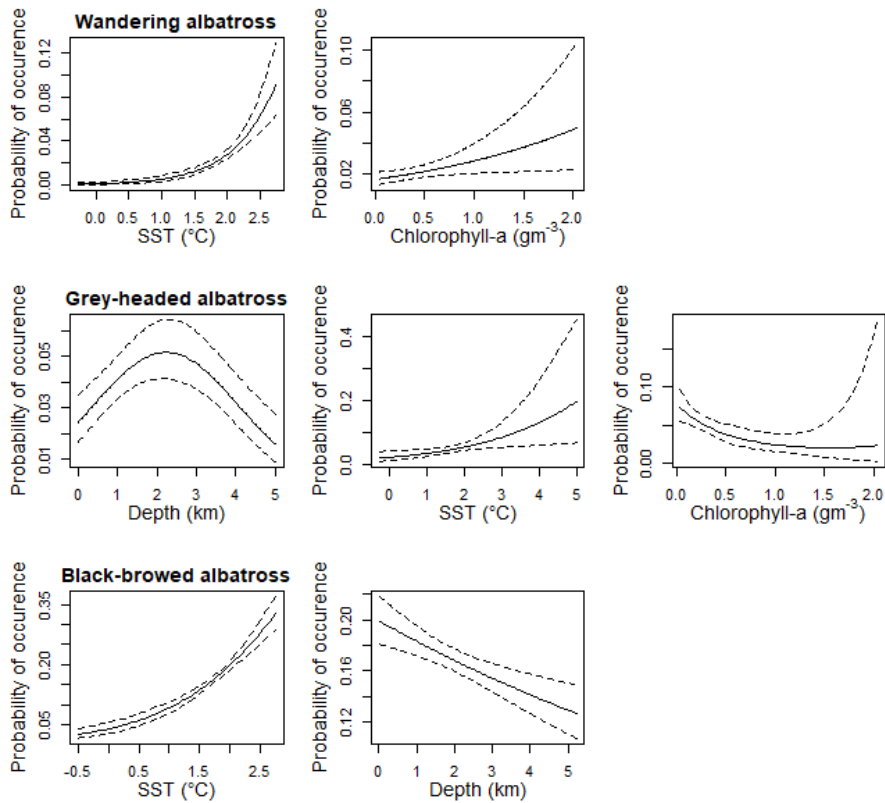


Figure S6. Response curves describing relationships between environmental variables and the at-sea distribution of seabirds around the Antarctic Peninsula, and South Shetland Islands, created using hurdle models. Confidence intervals were calculated using the standard errors provided with the model output.