Whale distribution in a breeding area: spatial models of habitat use and abundance of western South Atlantic humpback whales

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ABSTRACT: The western South Atlantic humpback whale population was severely depleted by commercial whaling in the late 19th and 20th centuries, and today inhabits a human-impacted environment in its wintering grounds off the Brazilian coast. Here, we identify distribution patterns related to environmental features and provide new estimates of population size, which can inform future management actions. We fitted spatial models to line transect data from two research cruises conducted in 2008 and 2012 to investigate (1) habitat use and (2) abundance of humpback whales wintering in the Brazilian continental shelf. Potential explanatory variables were year, depth, seabed slope, sea-surface temperature (SST), northing and easting, current speed, wind speed, distance to coastline and to the continental shelf break, and shelter (a combination of wind speed and SST categories). Whale density was higher in slower currents, at shorter distances to both the coastline and shelf break, and at SSTs between 24 and 25°C. The distribution of whales was also strongly related to shelter. For abundance estimation, easting and northing were included in the model instead of SST; estimates were 14,264 whales (CV = 0.084) for 2008 and 20,389 (CV = 0.071) for 2012. Environmental variables explained well the variation in whale density; higher density was found to the south of the Abrolhos Archipelago, and shelter seems to be important for these animals in their breeding area. Estimated distribution patterns presented here can be used to mitigate potential human-related impacts, such as supporting protection in the population’s core habitat near the Abrolhos Archipelago.

Keywords: shelter, conservation, density surface model, cetacean, line transect, reproduction

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INTRODUCTION

The Brazilian coast is inhabited every winter and spring by the western South Atlantic (WSA) humpback whale (*Megaptera novaeangliae*) population (also referred to as breeding stock A by the International Whaling Commission). Whales aggregate in coastal waters along the central and northeastern coasts of Brazil to mate and give birth before migrating to feeding areas (Martins et al. 2001, Zerbini et al. 2006). This population was severely exploited by whaling between the late nineteenth and mid-twentieth centuries (Zerbini et al. 2011; Morais et al. 2017), to the point of near extinction in the 1950s, but has since been recovering (Andriolo et al. 2010, Zerbini et al. 2011, Bortolotto et al. 2016a). The Red List of the International Union for Conservation of Nature and Natural Resources (IUCN) lists the conservation status of this species as “Least Concern” (Reilly et al. 2008). Recent abundance estimates from ship-based line transect surveys suggest that the WSA population size was near 20,000 animals in 2012 (Bortolotto et al. 2016a). However, that estimate was not computed for the entire area currently recognized as the typical distribution range of these animals during the breeding season. This increasing population faces today an environment modified by human activities such as marine traffic (Bezamat et al. 2015), fishing (Rocha-Campos et al. 2011, Moura et al. 2013, Ott et al. 2016), coastal water pollution (Moura et al. 2013, Ott et al. 2016), noise pollution (Rossi-Santos 2015), and activities related to the oil industry (Iversen et al. 2009, Martins et al. 2013, Ronconi et al. 2015, Rossi-Santos 2015, Brasil 2017a). Specifically, there is an increasing interest for oil and gas production activities in the area; according to the Brazilian National Agency of Petroleum, Natural Gas and Biofuels (Agência Nacional do Petróleo, Gás Natural e Biocombustíveis, ANP) the majority of the Brazilian petroleum reserves is found in the marine environment (Brasil 2017a).

Human-related activities in the area are expected to increase and negative interactions with humpback whales are likely to become more frequent (Andriolo et al. 2010, Martins et al. 2013). Existing marine protected areas (MPAs) alone provide very limited effective protection in the breeding grounds for this population, because they only cover a small fraction of the range of these whales (Castro et al. 2014). Therefore, a broad understanding of their distribution patterns and habitat use is fundamental to inform management actions. Area-based management, with the objective of protecting this charismatic flagship species, may also enhance biodiversity protection, because populations occupy relatively large and biodiversity-rich marine habitats.

For seasonal migratory animals such as many baleen whale species, the environmental factors expected to be important in habitat selection differ between feeding areas, where prey distribution is the primary driver (e.g., MacLeod et al. 2004, Friedlaender et al. 2006), and breeding areas (Corkeron & Connor 1999). During the breeding season, large whales select habitat according to their breeding status (Rayment et al. 2015), presence of calves in groups (Cartwright et al. 2012) and other
reproduction-related characteristics (Ersts & Rosenbaum 2003, Craig et al. 2014, Lindsay et al. 2016).

In this context, sheltered waters, bathymetric features, distance to the shore and sea-surface temperature (SST) are important factors for habitat usage of humpback whales in breeding areas (e.g., Taber & Thomas 1982, Smultea 1994, Rasmussen et al. 2007, Felix & Botero-Acosta 2011, Cartwright et al. 2012, Trudelle et al. 2016). Understanding and explaining key features of the ecology of migratory whale populations, such as habitat use, distribution and abundance, may provide important information to evaluate the impacts of human use of the environment inhabited by them.

WSA humpback whales are found in their breeding area, the Brazilian continental shelf between Natal (5°S) and Cabo Frio (23°S) (Fig. 1), during winter and spring every year, and animals concentrate on the Abrolhos Bank (~18°S) (Zerbini et al. 2006, Andriolo et al. 2010). The few previous studies that formally investigated their distribution relative to environmental variables (Wedekin 2011, Pavanato et al. 2017), or how they use the available habitat (Martins et al. 2001), indicate that bathymetric features (i.e., depth) may play an important role in how WSA whale groups are distributed.

Here we provide new insights into the distribution and density of WSA humpback whales in relation to environmental features in their breeding grounds, and present new abundance estimates for this population. We applied density surface models (DSMs) to line transect data (Miller et al. 2013) from ship-based surveys conducted in 2008 and 2012 (Bortolotto et al. 2016a) and fitted spatial models focusing on two main objectives: (1) to investigate habitat use and (2) to calculate model-based abundance estimates.

The new information should inform management actions to conserve humpback whales on their Brazilian breeding grounds. More specifically, new abundance estimates may be used to update this population’s conservation status, and the distribution results to evaluate areas where this population may be at higher risk of being affected by human-related activities, such as oil and gas exploration and production activities.

**Methods**

Shipboard visual line transect surveys were conducted in 2008 and 2012 during research cruises aboard the R/V Atlântico Sul (Universidade Federal do Rio Grande, FURG). Cruises were part of the Monitoring Whales by Satellite Project (Projeto Monitoramento de Baleias por Satélite, PMBS). PMBS main objectives were to deploy satellite-link tags on humpback whales to track their movements, to understand their space-use patterns in breeding and feeding grounds and characterize their migratory routes (Zerbini et al. 2006).

The survey area corresponded to the Brazilian continental shelf, between the shore and the shelf break (defined here as up to the 500 m isobath) from Cabo de São Roque (5°S), in Rio Grande do...
Norte State, to Cabo Frio (23°S), in Rio de Janeiro State (Fig. 1). Surveys were conducted from 25 August to 23 September in 2008 and from 7 August to 3 September in 2012, during the expected annual peak of occurrence of humpback whales in the area (August–September; Martins et al. 2001, Morete et al. 2003). Lines were designed to survey the full extent of this population’s breeding area and data collection followed the distance sampling methodology (Buckland et al. 2001). Trackline design, observation effort and data collection details are described in previous work (Bortolotto et al. 2016a, Bortolotto et al. 2016b).

Correcting for imperfect detection: detection function modelling
We used a detection function to correct for whales that were not detected when lines were surveyed (Buckland et al. 2001). Because other large whale species where rarely seen during the survey, sightings that were attributed to “unidentified large whales” were pooled with those of confirmed humpback whales. It is very unlikely that unidentified whale sightings were not of humpback whales, as discussed in Bortolotto et al. (2016a).

Detection functions were fitted to perpendicular distance data using R (version 3.2.1; R Core Team 2015) and “Distance” package (version 0.9.6; Miller 2016). Factor covariates sea conditions (“calm” for Beaufort 0–3 and “moderate” for Beaufort 4–6), detection cue (splash, body, blow or “other”), detection method (binoculars or naked eye) and year (2008 or 2012), and the continuous covariate group size (from 1 to 7) were considered. Variance in the detection function parameters was estimated using Fisher’s information matrix (Buckland et al. 2001, p. 61–68).

Data for spatial modelling
Survey tracklines were divided into 8 km segments using QGIS software (version 2.8.3; QGIS Development Team 2015). Standard segment length was chosen to be twice the truncation distance (= 4 km), resulting in 8 by 8 km squares for most segments. During line segmentation, some segments at the end of lines were shorter than 8 km. In those cases, segments less than 4 km long were merged with the previous one and those longer than 4 km were considered as an independent new segment. A few segments (5 out of 516) that were less than 4 km long, and that could not be merged with another line, were excluded from the analysis. The response variable used to model whale distribution was the whale counts in each segment, which were corrected using the detection function described above.

Based on previous studies on the distribution of cetaceans in breeding areas and environmental data availability, covariates considered as potential explanatory variables were: current speed close to the surface, depth, distance to coast, distance to the shelf break, SST, seabed slope, wind speed at the surface, geographic position (northing and easting) and year (Table 1). Additionally, to represent a combination of environmental conditions that may be related to energy saving for the calf, six
categories for shelter (Table 1) were created by combining three categories of wind speeds at the surface (“light” for values between 0.94 and 5.15 m s\(^{-1}\); “moderate” for values between 5.15 and 6.67 m s\(^{-1}\); “strong” for values between 6.67 and 9.16 m s\(^{-1}\)) and two categories of SST (“cold” for values between the minimum of 20.2\(^\circ\) and 24.7\(^\circ\)C; “warm” for values between 24.7\(^\circ\) and the maximum 26.9\(^\circ\)C). The wind and SST categories were delimited by quantiles of wind speed (33\(^{rd}\) percentile = 5.15 m s\(^{-1}\) and 66\(^{th}\) percentile = 6.67 m s\(^{-1}\)) and SST (median = 24.7\(^\circ\)C).

Values for depth were extracted from the global model of land topography and ocean bathymetry ETOPO1 (Amante & Eakins 2009). Circular buffers (radius = 4 km) were created around segment midpoints in QGIS and the average of depth values within the buffer zone was computed for each segment. This procedure was adopted because the resolution of ETOPO1 was much finer than the size of segments and buffers (between 13 and 16 ETOPO1 cells were included in the 50 km\(^2\) buffers and used to compute mean depth values). After mean depth values extraction, 25 out of 511 segments gave values greater than 500 m and were excluded from the analysis because the study area was previously defined as the continental shelf, from the shore up to the 500 m isobath. Slope values were derived from ETOPO1 data and were obtained in the same way, i.e., extracting mean values using the same circular buffers.

Distances to physical features (distance to coast and distance to shelf break) were calculated in QGIS or R as the shortest distance between the segment midpoint and the feature. For the distance to coast variable, the Brazilian coastline was obtained from a shapefile provided by SisCom (IBAMA 2011). To represent the continental shelf break, the 500 m isobath was generated from ETOPO1 in ArcGIS software using the “contour tool” function (Esri 2011).

SST was extracted from “MUR Global Foundation Sea Surface Temperature Analysis” dataset (JPL MUR MEaSUREs Project 2010) and ocean currents from “OSCAR” dataset (ESR 2009), both available from PO.DAAC/NASA website. Wind speed data were extracted from “ERA-Interim” dataset (ECMWF; Dee et al. 2011). With the exception of SST, the resolution of these datasets was too coarse when compared to the size of the circular buffers, so segment midpoints were used to extract covariate values in R software (“raster” package; Hijmans 2016). For SST, the circular buffers previously described were used to obtain mean values (around 40 SST values per buffer).

Spatial models and model selection

An initial investigation was performed to assess correlation among explanatory variables, and those that were highly correlated (i.e., a pair of variables that presented Pearson’s correlation coefficient greater than 0.7, or clear correlation identified via pair plots) were not included in the same model at the same time. Interaction terms, combining year and other covariates, were not tested because part of the study area was not surveyed in 2012, which would make the comparison severely unbalanced.
The quasi-Poisson distribution with logarithmic link function was assumed for the response variable (negative binomial and Tweedie distributions were also tested). An offset of ln(segment length) was included in all models. Generalized Additive Models (GAMs) were fitted using the “dsm” R package (version 2.2.14; Miller et al. 2017). Smooth functions were fitted to covariates, with a bivariate smooth for geographic position, since this included easting and northing. The basis dimension parameter \( k \) for the geographic position smooth term was set to 20, and for the univariate smooth terms it was set to 8 (see Wood 2006, p. 161, for an explanation on setting the dimension parameter). Model selection was conducted using a forward approach (i.e., adding one variable at a time), starting with a set of models, each with only one candidate explanatory variable. The model selected at each step was chosen by looking for an improvement in the Restricted Maximum Likelihood (REML) (Harville 1977) score. This score was used to minimize problems with parameter estimation that other potential scores (e.g., UBRE and GCV) may present when applying DSMs, following the recommendation in Miller et al. (2013). The auto-correlation in the residuals (ordered by the time of data collection) of spatial models was checked using the “acf” function (“stats” R package; R Core Team 2015). Model performance was assessed with model diagnostic plots (function “gam.check”, “dsm” R package) and 10-fold cross validation (Refaelzadeh, Tang & Liu 2009).

Two modelling exercises were undertaken, each considering a different set of covariates and having different objectives:

1. Habitat Use Model (HUM): to explain habitat use in a way that could be interpreted biologically. All variables, except geographic position (northing/easting), were considered;
2. Abundance Estimation Model (AEM): to compute abundance estimates from the spatial model and all available variables were considered.

The HUM was designed to investigate which environmental variables were more related to distribution, while the AEM was designed to obtain the best density surface prediction, possibly including northing/easting, which could explain variability that was not explained by the environmental covariates.

**Predictions**

A prediction grid formed by 8 by 8 km cells was created over the entire study area using QGIS. The size of the prediction grid cells was chosen to match that of the segments used in the models. Covariate values for each grid cell were obtained in a similar way of that described for segments, using cell midpoints or 4 km buffers around midpoints. For covariates that varied in time within each survey (e.g., SST), the mean of values for the survey period was used for predictions.

The model-based abundance estimates for 2008 and 2012 were obtained from the sums across all grid cells of predicted values from the AEM, for each year. Maps showing patterns of distribution
(density surface) were created using the AEM predictions in QGIS. Variances were obtained with the delta method, combining the variance from the detection function and the spatial models, using the “dsm.var” function of the “dsm” R package. Maps of uncertainty in model predictions (standard deviation surface) were also created with the variance calculated for each grid cell (Fig. S6). Predictions in 2012 were extrapolated to the area to the north of Salvador (~13°S), which was not surveyed in 2012 (Fig. 1) because of poor weather conditions (Bortolotto et al. 2016a).

RESULTS

Survey effort used in the analysis totaled 2,350 km in 2008 and 1,700 km in 2012. The number of whale groups (including mother-calf pairs and solitary animals) in the data was 493 (416 humpbacks and 77 unidentified large whales) and 737 (557 humpbacks and 180 unidentified large whales) in 2008 and 2012, respectively.

Detection function

Perpendicular distances were truncated at 4 km, resulting in 81 (out of 1230) detections being excluded from the detection function analysis. The best-fitting detection function was a hazard rate model with covariates cue, year and sea conditions (Fig. 2; Table S1). The average probability of detection $p$ was estimated as 0.482 (CV = 0.044) and the goodness of fit tests showed a good fit (Kolmogorov-Smirnov test statistic = 0.016, p-value = 0.930; Cramer-von Mises test (unweighted) statistic = 0.036, p-value = 0.952).

Spatial models

Model diagnostics (Fig. S1 and S2) indicated the quasi-Poisson distribution to be adequate and to provide a better fit than the other distributions that were considered. Cross-validation yielded root-mean-square errors of 6.932 (SD = 1.116) for 2008 and 7.981 (SD = 0.967) for 2012 (Table S7). SST was found to be highly correlated with geographic position. Depth, slope and distance to the shelf break were also correlated to each other. Therefore, if one of the above variables was selected at a model selection step, those correlated were not considered in subsequent steps of model selection.

The selected HUM included variables distance to the coast, distance to the shelf break, SST, current speed and shelter, and presented 54.1% of deviance explained. The variable with the most pronounced effect was SST, with a peak around 24–25°C (Fig. 3). Whale density was positively related to distance to the coast and distance to the shelf break, but negatively related to current speed, apparent from around 0.2 m s⁻¹ and greater. Shelter coefficients indicated differences in whale densities between shelter categories, with significantly (at $a = 0.05$) higher densities in relatively cold waters with light winds (Table 2; Tables S2 and S3).
The selected AEM included variables distance to the coast, distance to the shelf break, current speed, shelter and geographic position, and had an explained deviance of 66.8%. This model was used for plotting purposes here, because this model presented a larger portion of explained deviance and the distribution patterns are likely better represented. Very weak signs of auto-correlation were found in the residuals of HUM and no signs of auto-correlation were present in the residuals of AEM (ACF plots; Fig. S1 and S2).

**Abundance estimates**

Estimated abundances for prediction grid cells ranged from 0.139 to 53.0 animals (mean = 7.47, SD = 8.90) in 2008 and from 0.144 to 60.9 animals (mean = 10.7, SD = 12.7) in 2012. Model-based abundance estimates were 14,264 whales (CV = 0.084) for 2008 and 20,389 (CV = 0.071) for 2012 (Table S6). Surface maps for predicted density showed higher numbers in the Abrolhos Bank region, with a concentration area to the south of the Abrolhos Archipelago, which was more pronounced for 2012 (Fig. 4). Other areas also showed relatively high densities, such as the coast of Alagoas and Sergipe States (Fig. S4), and near the city of Salvador, Bahia State (Fig. S5).

**DISCUSSION**

Systematically collected sightings data were used to model the distribution and abundance of humpback whales in their wintering areas off the coast of Brazil. The suite of environmental covariates tested included powerful predictors of whale density across the study area, with SST and geographic position being the most powerful explanatory terms. The effect of year was not selected in the spatial models, suggesting that differences in the distribution patterns from 2008 to 2012 were better explained by the variation in the spatial covariates than by temporal changes between survey years.

These sighting data were previously used to estimate abundance of humpback whales off the coast of Brazil in 2008 and 2012 using design-based methods (Bortolotto et al. 2016a). However, the realized effort in that study did not conform exactly to the designed lines. For example, because of unfavorable weather conditions in 2012, there were no data available for areas to the north of Salvador, Bahia State (Fig. 1). Consequently, the abundance estimate previously presented for that year was computed for only part of what is currently known to be the typical breeding area for WSA humpback whales. Because of logistical restrictions, our results likely represent WSA humpback distribution during the annual peak of their occurrence in the area (August–September) and it is not possible to infer intra-season variations.

Migratory whales show marked differences in habitat preferences according to different age classes, sexes, reproductive-related individual characteristics and/or group composition (Best 1990, Craig & Herman 2000, Martins et al. 2001, Ersts & Rosenbaum 2003, Elwen & Best 2004a, Oviedo
& Solis 2008, Cartwright et al. 2012, Craig et al. 2014, Rayment et al. 2015, Lindsay et al. 2016), and for specific group types (Elwen & Best 2004b, Felix & Botero-Acosta 2011) when in breeding areas. However, the passing mode data collection procedure adopted here prevented obtaining more specific data on individual whales, such as sex, age class or accurate group composition. Because of this, results presented here are representative of the population as a whole, not of any particular sex, age or group. Although some of the results may be consistent with what could be expected for habitat preferences of breeding or/and calving animals in the area, such as the importance of shelter as a predictor of density, it is not possible to make robust inferences for specific reproductive stages. A study to investigate the distribution and habitat use of WSA humpback whales based on data from satellite tagging of individual whales (Zerbini et al. 2006) is underway, which is expected to provide information on predictors of distribution and habitat use in relation to sex and group composition. Because the procedure of attaching tags requires close proximity to the animals, collection of individual and group information is possible at the moment of tagging.

**Spatial modelling**

The covariates retained in the models explained a high portion of the variation in whale density across the surveyed area (deviance explained = 54.1% for HUM; 66.8% for AEM). In addition to this increase in explained deviance, in the AEM the residual auto-correlation in the HUM (“ACF” plots; Fig. S1 and S2) was no longer apparent (although the auto-correlation in the residuals of the HUM was not high and required no further action; see Wood 2006 for concerns about residual auto-correlation of GAMs). It is likely, therefore, that the bivariate smooth for easting/northing included in the AEM is acting as a proxy for unmodelled environmental or social characteristics. For example, because it was highly correlated with SST, which was not included in the AEM, easting/northing may be representing not only SST but also some other environmental feature(s). This may explain the increase in percentage of explained deviance when SST is substituted by easting/northing in the AEM.

Shelter (a combination of SST and wind speed) was created as an environmental feature that could be important to whales that are calving, for example to represent conditions that may be related to energy saving for the calf (Corkeron & Connor 1999). Because the effects of wind speed on detectability have been accounted for in the estimation of detection probability, no confounding with the effects of wind in the shelter variable is expected. The response variables in the detection function model and the habitat use/abundance estimation spatial models are completely different; in the detection process it is the perpendicular distance (in relation to the trackline) and in the spatial models the response variable is abundance (corrected count per segment). Furthermore, wind speed may influence both the detectability of animals and how animals use their habitat, which is supported by
the present results. Indeed, a major advantage of density surface modelling using data from distance
sampling surveys is that the effects of variables on detectability and on abundance can be teased apart.

The density surface modelling approach permitted inference and extrapolation from the AEM
to the area not surveyed in 2012 by Bortolotto et al. (2016a), resulting in a 2012 abundance estimate
for a larger part of the breeding ground distribution than would otherwise be available. The lack of
data to the north of Salvador in 2012 implies that the effect of the bivariate smooth for
easting/northing on the predictions for that area is largely influenced by data from 2008. However,
the other variables retained in the model were responsible for the large majority of the explained
deviance, as illustrated by the percentage of explained deviance of the HUM (54.1%), so this is not
considered to be an important limitation for our inferences about abundance.

Model-based abundances for humpback whales breeding off the coast of Brazil (14,264,
CV = 0.084 for 2008; 20,389, CV = 0.071 for 2012) were estimated to be close to those computed by
design-based methods (16,410, CV = 0.228 for 2008; 19,429, CV = 0.101 for 2012; Bortolotto et al.
2016a). This similarity could be expected because both estimates are derived from the same data. The
higher precision in the model-based abundance estimates (CV = 0.084 vs 0.228 for 2008; CV = 0.071
vs 0.101 for 2012) is mainly because the covariates explained some of the variability in the data,
demonstrating the value of the analysis.

Habitat use

The main reasons for SST to be considered an important factor in explaining the distribution of
migratory whales in their breeding grounds are likely related to presence of calves, which are not as
efficient in conserving their body temperature as older animals (Corkeron & Connor 1999). SST was
the most important variable selected in the HUM and it was highly correlated with geographic
position (northing/easting). The overall relation between whale density and SST was positive,
peaking at 24 to 25°C. This result for SST may reflect habitat selection of calving females for the
reason stated above. The habitat use of North Atlantic right whales in their calving grounds off south-
eastern United States was also observed to be strongly related to SST (Keller et al. 2006), however
differences in species characteristics (e.g., latitudinal range) should be taken into account in any
comparison. Trudelle et al. (2016) did not find a relationship between SST and humpback whale
movements in their Madagascar coastal breeding area, possibly because of the relatively low variation
in SST in the area. Although a temporal change in distribution was not supported by our models, long
term monitoring should provide important insights on this, as the effects of climate change (Walther
et al. 2002), for example, may impact the distribution of marine animals.

Shelter, which incorporated SST, was consistently retained in our spatial models and therefore
can be considered an important factor to explain this population’s distribution in the breeding area.
The fitted relationship for this covariate suggests that relatively slow and moderate surface winds had a significant positive effect on density, when the water was relatively colder. Because wind speed was not selected in the spatial models, our results suggest that wind may be an important habitat feature for WSA humpback whales only when the water temperature is relatively cool. A possibility is that, because temperature is one of the most important features for these animals in the area, they tolerate a range of wind speeds that is not their preferred, when the SST is relatively warmer. As mentioned above, because calves may benefit from an environment where they can save body energy reserves, calm conditions at the water surface are likely preferable for calves to swim and to surface to breathe (Taber & Thomas 1982, Cartwright et al. 2012). At a daily scale study of habitat use, Felix & Botero-Acosta (2011) found that mother-calf humpback whale pairs in Ecuador preferred shallower waters during the afternoon hours, when wind speeds in the area tended to increase and the sea to become rougher. The combination of water temperature and wind at the surface seems to be an important factor for WSA humpback whale habitat selection in breeding grounds. Rayment et al. (2015), to the best of our knowledge, was the only study that incorporated a variable to explicitly represent shelter in habitat use models for breeding migratory whales. These authors investigated the influence of shelter in breeding right whales distribution and found that wave exposure and distance to shelter (defined as areas with lower wind exposure) influenced habitat selection of right whale groups with calves.

It is still unclear which environmental features really represent shelter for breeding whales and how this may vary among different species. Martins et al. (2001) showed that the occurrence of WSA humpback whales groups containing calves increased with the proximity to the Abrolhos Archipelago, which may represent shelter for these animals, with the archipelago presence perhaps creating a calmer environment. Also, Zerbini et al. (2004) observed that WSA mothers-calf groups were more frequently found closer to the shore than other group types off the north-eastern coast of Brazil. Our results add to this discussion of which environmental variables may combine to create a sheltered environment that benefits migratory whale species in their breeding grounds. While several other covariates could have been included or combined to create a spatial covariate to represent shelter (e.g., speed and direction of ocean currents), the simple combination that we present here for shelter permits easy interpretation of model results. A complicated combination of several covariates would likely produce results that would be difficult to interpret biologically.

The relationships between whale density and environmental covariates revealed by our models are consistent with what could be expected for mothers, which may prefer a secure environment for the development of their calves in sheltered waters. However, Trudelle et al. (2016) noted that while the movements of female humpback whales in a breeding area off the Madagascar coast are influenced by environmental features such as depth and distance to the shore, male movements are
probably more influenced by social factors, such as female occurrence. Despite the fact that their distribution may also be influenced by the presence of other males (Herman 2017), adult males are indeed likely to seek receptive females, not those that are about to or have just given birth. Calving females may prefer shallow waters where the chances of being harassed by males are lower; their habitat selection may be driven primarily by avoidance of males (Craig et al. 2014). Humpback whale groups containing calves have been found significantly more frequently in shallower waters than groups without calves in Brazilian breeding grounds (Martins et al. 2001, Zerbini et al. 2004). Thus, bathymetric features may also be related to what may represent shelter for the whales.

Overall, this discussion highlights the importance of having data on the sex and reproductive status of individuals and not only on environmental features to understand the distribution of large whales in breeding areas. For example, we did not consider bathymetry as part of shelter to facilitate interpretation of results, but if such individual data were available it could be informative to investigate a wider range of covariate combinations representing shelter in models of habitat use. Future studies could also investigate in detail the conditions of the marine environment in areas surrounding the Abrolhos Archipelago. For example, the presence of coral reefs may be related to (or contribute to) shelter from rough water (Lindsay et al. 2016).

The positive relationship between whale density and distance to both the coast and the continental shelf break could mean that humpback whales off the coast of Brazil prefer to be in the middle part of the shelf, or that they prefer to avoid the shelf boundaries. Trudelle et al. (2016) suggested that the distance to the coast was one of the most important factors affecting the movement patterns of female humpback whales off the Madagascar breeding grounds and other studies have shown that calving humpback whales are associated with areas close to the shore (Martins et al. 2001, Zerbini et al. 2004, Felix & Botero-Acosta 2011). Avoidance of the shelf edge could be in response to the risk of predation by large predators in offshore waters, such as large shark species (Smultea 1994). Areas too close to the shore could be avoided because they are too shallow for swimming (Oviedo & Solis 2008) or because of disturbances that were not considered here, such as noise from human activities.

The estimated negative effect on predicted whale numbers of current speeds greater than 0.2 m s\(^{-1}\) is not very well supported by the data (95% confidence interval widens with increasing current speed). In a study that supports the importance of the current for large whales in breeding areas, Trudelle et al. (2016) found that differences in current speed between shelf and oceanic waters influenced the movement patterns of humpback whales in their Madagascar breeding area. Whales of both sexes swim faster in slower currents and the authors suggest that when animals are engaged in mate-searching-related movements close to the coast, the current speed probably did not have an important effect. Therefore, it is likely that data on the behavioral status and/or movements of
individual animals are needed to better understand the effects of current speed on habitat use of humpback whales off Brazil. In addition, the resolution of this covariate (5-day bins and 0.33x0.33° of latitude/longitude; Table 1) was likely unable to capture fine scale variability, particularly around complex coastlines.

**Implications for conservation and management**

The predicted distributions support previous work showing that WSA humpback whales have a strong preference for the Abrolhos Bank region during their breeding season in coastal waters of Brazil (Siciliano 1997, Andriolo et al. 2010, Wedekin 2011, Martins et al. 2013, Pavanato et al. 2017). However, other areas also had relatively high predicted densities, such as near Salvador and off the coasts of Sergipe and Alagoas States (Figs. S4 and S5). Little is known about their distribution or habitat use in these areas (Zerbini et al. 2004, Baracho-Neto et al. 2012), but relatively recent observations indicate that the distribution of WSA humpback whales in Brazil may be broader than currently recognized (e.g., Wedekin et al. 2014, Bortolotto et al. 2016c, Pavanato et al. 2017).

The Abrolhos Archipelago is included in the Abrolhos Marine National Park, which is a national “Conservation Unit” (abbreviated as UC in Portuguese) area of 880 km² (ICMBio 2017). According to the Brazilian Ministry of Environment (Brasil 2017b) this is a federal UC of “integral protection” where only scientific research and educational, recreational and small-scale ecotourism activities are permitted. All of these activities are regulated by the Chico Mendes Institute for Biodiversity Conservation (ICMBio), the federal body responsible for protected areas in Brazil. Commercial activities are therefore mostly limited to those related to small-scale ecotourism. The nearby Environmental Protection Area of Ponta da Baleia is regulated by the Bahia State and is in the category of “sustainable use area” (INEMA 2017). These protected areas cover a very small portion of the area predicted to have the highest concentration of animals (Fig. 5). Our results support the conclusions of Castro et al. (2014) who used satellite tracked movement data to show that MPAs only cover a very small portion of the areas most used by WSA humpback whales in their breeding grounds.

The Abrolhos Bank is a region of high biodiversity (Werner et al. 2000) and expanding the area under protection could benefit not only cetaceans but also other marine organisms, such as the unique coral reefs in the area (Francini-Filho & Moura 2008). Because most humpback whale births are expected to occur on or near Abrolhos Bank (Martins et al. 2001), expanding the protected area during the period when whales are present consistently (winter–spring), could reduce the risk of anthropogenic impact especially for calves that are known to be more vulnerable to disturbance (Schaffar et al. 2013). To conserve marine species in the area, past management actions have included the cancellation of seismic activity on the Bank during humpback breeding season and other oil and
gas exploitation activities (Engel et al. 2004, Marchioro et al. 2005). However, there is increasing interest from the oil and gas industry to explore for oil on the Bank (Brasil 2017a). Because young animals are more vulnerable to stressors (Schaffar et al. 2013, Ott et al. 2016, Dunlop et al. 2017) and we did not include group composition in this study, future studies aiming to provide information for conservation should investigate the distribution of different group types at a finer scale and include potentially stressors and displacement factors associated with human presence in the marine environment, with special attention to the Abrolhos Bank region.

Abundance estimates presented here (14,264, CV = 0.084 for 2008 and 20,389, CV = 0.071 for 2012) provide additional confirmation that the WSA humpback whale population is growing (Zerbini et al. 2011). A new population status assessment in the framework of Zerbini et al. (2011) is currently underway, which will take the present results and new catch history data (Morais et al. 2017) into account to provide an updated understanding of this population’s recovery, more than four decades after whaling ceased in 1973 in this area.

Going forward, it is important that efforts to monitor potential threats are intensified, because our current knowledge on this is very limited (Bezamat et al. 2015, Bortolotto et al. 2016c, Ott et al. 2016). To evaluate adequately the need for improvement or adjustment of current conservation strategies and management actions, such as enhancing protection in the area (Castro et al. 2014), it is essential to assess the conservation status of WSA humpback whales and to take into account the current and future potential impacts on the population. The distribution results presented here may also be used in evaluating areas of higher risk for this population by investigating sources of impact by human-related activities in the areas predicted to be most used by the animals.

ACKNOWLEDGEMENTS

The Monitoring Whales by Satellite Project (Projeto Monitoramento de Baleias por Satélite, PMBS) research cruises were sponsored by Shell Brasil. The Universidade Federal de Rio Grande (FURG) and the N/Pq Atlântico Sul crew provided essential support during fieldwork. The Brazilian Inter-Ministerial Commission for the Resources of the Sea (CIRM) supplied the diesel for the ship. GA Bortolotto PhD is funded by the Brazilian National Council for Scientific and Technological Development (Conselho Nacional de Desenvolvimento Científico e Tecnológico, CNPq; Science Without Borders, scholarship #208203/2014-1). Cetacean International Society granted GA Bortolotto with small grants which contributed to the development of this study. Franciele Castro, Marco Aurélio Crespo, Juliana Di Tullio, Daniela Godoy, Ygor Geyer, Natália Mamede, Igor Morais, Paulo Ott, Jonatas Prado, Eduardo Secchi, Suzana Stutz, Federico Sucunza, and Janaina Wickert assisted with data collection. We are especially thankful for the contribution of Artur Andriolo to this
work. David Miller provided valuable advice on statistical analysis. This manuscript was improved thanks to suggestions of three anonymous reviewers.

**LITERATURE CITED**


Fig. 1. Survey lines in 2008 and 2012. Planned (dashed grey lines) and completed effort (black thick lines) are shown. A black triangle indicates the location of the Abrolhos Archipelago.
Fig. 2. Detection function curve (red line) from a hazard rate model fitted to the perpendicular distances (in meters) of humpback whale groups detected. Different dotted curves represent different combinations of covariates sea conditions, cue and year. Each point represents the predicted value for observation.
Fig. 3. Model terms for the Habitat Use Model (HUM) of humpback whales off the coast of Brazil. Smooth terms’ effective degrees of freedom are shown inside brackets in the vertical axis. The shelter coefficients are presented relative to the intercept. (wa = warm SST, co = cold SST, li = light wind, mo = moderate wind, st = strong wind).
Fig. 4. Density surface maps for 2008 and 2012. Predictions were made with the Abundance Estimation Model (AEM).
Fig. 5. Density surface maps for 2008 and 2012 for the Abrolhos Bank region. Predictions were made with the Abundance Estimation Model (AEM). A black triangle shows the location of the Abrolhos archipelago. Red polygons represent the Abrolhos Marine National Park and the brown polygon represents the Ponta da Baleia MPA.
Table 1. Explanatory variables tested in Generalized Additive Models to model the density of humpback whales off the coast of Brazil.

<table>
<thead>
<tr>
<th>Variables</th>
<th>Description</th>
<th>Resolution*</th>
<th>Unit</th>
<th>Reference/Data source</th>
</tr>
</thead>
<tbody>
<tr>
<td>curr.sp</td>
<td>Speed of the water current close to the surface</td>
<td>5-day; 0.33 x 0.33° (latitude x longitude)</td>
<td>m s⁻¹</td>
<td>OSCAR dataset (ESR 2009)</td>
</tr>
<tr>
<td>depth</td>
<td>Depth</td>
<td>0.1 x 0.1° (latitude x longitude)</td>
<td>m</td>
<td>ETOPO1 (Amante &amp; Eakins 2009)</td>
</tr>
<tr>
<td>dist.coast</td>
<td>Distance to the coastline</td>
<td>–</td>
<td>m</td>
<td>SisCom (IBAMA 2011)</td>
</tr>
<tr>
<td>dist.shelf</td>
<td>Distance to the 500 meter isobath</td>
<td>–</td>
<td>m</td>
<td>500 meter isobath created from ETOPO1 in GIS software</td>
</tr>
<tr>
<td>shelter</td>
<td>Category according to values of wind.sp and sst</td>
<td>–</td>
<td>–</td>
<td>–</td>
</tr>
<tr>
<td>slope</td>
<td>Seabed slope: percentage of elevation over distance</td>
<td>0.1 x 0.1° (latitude x longitude)</td>
<td></td>
<td>Derived from ETOPO1</td>
</tr>
<tr>
<td>sst</td>
<td>Temperature at the surface of the sea</td>
<td>1-day; 0.011 x 0.011° (latitude x longitude)</td>
<td>°C</td>
<td>JPL-L4UHfind-GLOB-MUR dataset (JPL MUR MEaSUREs Project 2010)</td>
</tr>
<tr>
<td>wind.sp</td>
<td>Speed of wind at the surface</td>
<td>6-hour (the daily mean was used); 80 x 80 km</td>
<td>m s⁻¹</td>
<td>ERA-Interim dataset (Dee et al. 2011)</td>
</tr>
<tr>
<td>x</td>
<td>Easting</td>
<td>–</td>
<td>m</td>
<td>Survey GPS</td>
</tr>
<tr>
<td>y</td>
<td>Northing</td>
<td>–</td>
<td>m</td>
<td>Survey GPS</td>
</tr>
<tr>
<td>year</td>
<td>Year of survey</td>
<td>–</td>
<td>year</td>
<td>Survey data</td>
</tr>
</tbody>
</table>

*Spatial and/or temporal resolution, depending on covariate nature.*
Table 2. Generalized Additive Model results for the HUM (Habitat Use Model) and AEM (Abundance Estimation Model). Variables are described in Table 1. Effective degrees of freedom for smooth terms are presented inside brackets. Blank spaces represent variables not selected and a dash represents a covariate not considered in the model selection. (*S* = smooth term, *F* = factor)

<table>
<thead>
<tr>
<th>Variable</th>
<th>HUM</th>
<th>AEM</th>
</tr>
</thead>
<tbody>
<tr>
<td>curr.sp</td>
<td><em>S</em>(3.315)</td>
<td><em>S</em>(3.294)</td>
</tr>
<tr>
<td>Depth</td>
<td></td>
<td></td>
</tr>
<tr>
<td>dist.coast</td>
<td><em>S</em>(2.401)</td>
<td><em>S</em>(5.528)</td>
</tr>
<tr>
<td>dist.shelf</td>
<td><em>S</em>(0.975)</td>
<td><em>S</em>(0.940)</td>
</tr>
<tr>
<td>Shelter</td>
<td><em>F</em></td>
<td><em>F</em></td>
</tr>
<tr>
<td>Slope</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Sst</td>
<td><em>S</em>(3.766)</td>
<td></td>
</tr>
<tr>
<td>wind.sp</td>
<td></td>
<td></td>
</tr>
<tr>
<td>x, y</td>
<td>—</td>
<td><em>S</em>(15.865)</td>
</tr>
<tr>
<td>year</td>
<td></td>
<td></td>
</tr>
<tr>
<td>% Deviance explained</td>
<td>54.1</td>
<td>66.8</td>
</tr>
<tr>
<td>-REML score</td>
<td>718.5</td>
<td>678.0</td>
</tr>
</tbody>
</table>
**The following supplement accompanies the article**

**Whale distribution in a breeding area: spatial models of habitat use and abundance of western South Atlantic humpback whales**

Guilherme A. Bortolotto*, Daniel Danilewicz, Philip S. Hammond, Len Thomas, Alexandre N. Zerbini

*Corresponding author: bortolotto.ga@gmail.com

**Detection function model results**

Table S1. Detection function parameters from a hazard-rate key-model fitted to 1149 perpendicular distance values for humpback whale sightings (data were truncated at 4000 m). Coefficient values are on the scale of the log link function. The intercept includes terms “cue blow”, “year 2008” and “sea state calm”.

<table>
<thead>
<tr>
<th>Scale Coefficients</th>
<th>Estimate</th>
<th>Standard error</th>
</tr>
</thead>
<tbody>
<tr>
<td>Intercept</td>
<td>7.097</td>
<td>0.125</td>
</tr>
<tr>
<td>Cue splash</td>
<td>0.535</td>
<td>0.162</td>
</tr>
<tr>
<td>Cue body</td>
<td>-0.470</td>
<td>0.164</td>
</tr>
<tr>
<td>Cue “other”</td>
<td>0.363</td>
<td>0.310</td>
</tr>
<tr>
<td>Year 2012</td>
<td>0.291</td>
<td>0.107</td>
</tr>
<tr>
<td>Sea state moderate</td>
<td>-0.220</td>
<td>0.107</td>
</tr>
</tbody>
</table>
Habitat Use Model (HUM) results

Table S2. Parametric coefficients in the Habitat Use Model (HUM). ($t = t$ distribution value)

<table>
<thead>
<tr>
<th>Coefficients</th>
<th>Estimate</th>
<th>Standard error</th>
<th>$t$</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Intercept</td>
<td>-15.704</td>
<td>0.116</td>
<td>-134.819</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>shelter.cold.moderate</td>
<td>-0.473</td>
<td>0.111</td>
<td>-4.272</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>shelter.cold.strong</td>
<td>-1.122</td>
<td>0.271</td>
<td>-4.138</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>shelter.warm.light</td>
<td>-0.760</td>
<td>0.193</td>
<td>-3.942</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>shelter.warm.moderate</td>
<td>-1.140</td>
<td>0.261</td>
<td>-4.364</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>shelter.warm.strong</td>
<td>-0.524</td>
<td>0.242</td>
<td>-2.164</td>
<td>0.031</td>
</tr>
</tbody>
</table>

*Significant at $\alpha = 0.05$

Table S3. Smooth terms in the Habitat Use Model (HUM). (edf = effective degrees of freedom, df = degrees of freedom, F = F distribution value)

<table>
<thead>
<tr>
<th>Smooth terms</th>
<th>edf</th>
<th>Reference df</th>
<th>F</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>s(sst)</td>
<td>3.766</td>
<td>7</td>
<td>6.347</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>s(dist.shelf)</td>
<td>0.975</td>
<td>7</td>
<td>5.041</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>s(coast)</td>
<td>2.401</td>
<td>7</td>
<td>4.918</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>s(curr.sp)</td>
<td>3.315</td>
<td>7</td>
<td>2.535</td>
<td>&lt; 0.001*</td>
</tr>
</tbody>
</table>

*Significant at $\alpha = 0.05$
Abundance Estimation Model (AEM) results

Table S4. Parametric coefficients in the Abundance Estimation Model (AEM). (t = t distribution value)

<table>
<thead>
<tr>
<th>Coefficients</th>
<th>Estimate</th>
<th>Standard error</th>
<th>t</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Intercept</td>
<td>-16.007</td>
<td>0.105</td>
<td>-153.078</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>shelter.cold.moderate</td>
<td>-0.279</td>
<td>0.109</td>
<td>-2.559</td>
<td>0.011*</td>
</tr>
<tr>
<td>shelter.cold.strong</td>
<td>-0.830</td>
<td>0.247</td>
<td>-3.364</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>shelter.warm.light</td>
<td>-0.484</td>
<td>0.148</td>
<td>-3.268</td>
<td>0.001*</td>
</tr>
<tr>
<td>shelter.warm.moderate</td>
<td>-0.532</td>
<td>0.221</td>
<td>-2.402</td>
<td>0.012*</td>
</tr>
<tr>
<td>shelter.warm.strong</td>
<td>-0.470</td>
<td>0.207</td>
<td>-2.272</td>
<td>0.024*</td>
</tr>
</tbody>
</table>

*Significant at α = 0.05.

Table S5. Smooth terms in the Abundance Estimation Model (AEM). (edf = effective degrees of freedom, df = degrees of freedom, F = F distribution value)

<table>
<thead>
<tr>
<th>Smooth terms</th>
<th>Edf</th>
<th>Reference df</th>
<th>F</th>
<th>p-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>s(x,y)</td>
<td>15.865</td>
<td>19</td>
<td>9.911</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>s(curr.sp)</td>
<td>3.294</td>
<td>7</td>
<td>4.009</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>s(coast)</td>
<td>5.528</td>
<td>7</td>
<td>4.283</td>
<td>&lt; 0.001*</td>
</tr>
<tr>
<td>s(dist.shelf)</td>
<td>0.940</td>
<td>7</td>
<td>2.155</td>
<td>&lt; 0.001*</td>
</tr>
</tbody>
</table>

*Significant at α = 0.05
Abundance estimates results

Table S6. Summaries of uncertainty in a density surface model (Abundance Estimation Model, AEM) calculated analytically for GAM, with delta method, for 2008 and 2012.

<table>
<thead>
<tr>
<th></th>
<th>2008</th>
<th>2012</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Approximate asymptotic confidence interval</td>
<td></td>
</tr>
<tr>
<td></td>
<td>2.5%</td>
<td>Mean</td>
</tr>
<tr>
<td></td>
<td>2.5%</td>
<td>Mean</td>
</tr>
<tr>
<td></td>
<td>12,108</td>
<td>14,264</td>
</tr>
</tbody>
</table>

Abundance

<table>
<thead>
<tr>
<th></th>
<th>2008</th>
<th>2012</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Point estimate</td>
<td>14,264</td>
</tr>
<tr>
<td></td>
<td>CV of detection function</td>
<td>0.044</td>
</tr>
<tr>
<td></td>
<td>CV from GAM</td>
<td>0.071</td>
</tr>
<tr>
<td></td>
<td>Total standard error</td>
<td>1,195</td>
</tr>
<tr>
<td></td>
<td>Total coefficient of variation</td>
<td>0.084</td>
</tr>
</tbody>
</table>
Cross-validation results

Table S7. Root-Mean-Squared-Errors (RMSEs) for 10-fold cross-validation of Abundance Estimates Model (AEM) and Habitat Use Model (HUM).

<table>
<thead>
<tr>
<th>Cross-validation fold</th>
<th>AEM</th>
<th>HUM</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>7.429</td>
<td>8.413</td>
</tr>
<tr>
<td>2</td>
<td>7.295</td>
<td>7.185</td>
</tr>
<tr>
<td>3</td>
<td>8.855</td>
<td>8.875</td>
</tr>
<tr>
<td>4</td>
<td>5.517</td>
<td>6.998</td>
</tr>
<tr>
<td>5</td>
<td>6.696</td>
<td>8.046</td>
</tr>
<tr>
<td>6</td>
<td>6.813</td>
<td>7.422</td>
</tr>
<tr>
<td>7</td>
<td>8.271</td>
<td>9.705</td>
</tr>
<tr>
<td>8</td>
<td>6.852</td>
<td>8.563</td>
</tr>
<tr>
<td>9</td>
<td>5.213</td>
<td>6.500</td>
</tr>
<tr>
<td>10</td>
<td>6.382</td>
<td>8.097</td>
</tr>
<tr>
<td>Mean</td>
<td>6.932</td>
<td>7.981</td>
</tr>
<tr>
<td>Standard deviation</td>
<td>1.116</td>
<td>0.967</td>
</tr>
</tbody>
</table>
Fig. S1. Habitat Use Model (HUM) diagnostic plots from “gam.check” R function and autocorrelation regression plot from “acf” function.
Fig. S2. Abundance Estimation Model (AEM) diagnostic plots from “gam.check” R function and auto-correlation regression plot from “acf” function.
Fig. S3. Model terms for the Abundance Estimation Model (AEM) of humpback whales off the coast of Brazil. Smooth terms’ effective degrees of freedom are shown inside brackets in the vertical axis. The plot for $s(x,y)$ is not included here. The shelter coefficients are presented relative to the intercept. (wa = warm SST, co = cold SST, li = light wind, mo = moderate wind, st = strong wind).
Fig. S4. Density surface maps for 2008 and 2012 for the region of Sergipe and Alagoas coasts.

Predictions were made with the Abundance Estimation Model (AEM).
Fig. S5. Density surface maps for 2008 and 2012 for part of the coast of Bahia State. Predictions were made with the Abundance Estimation Model (AEM).
Fig. S6. Standard deviation surface maps for 2008 and 2012. Standard deviations from the Abundance Estimation Model (AEM).