# **Combining multiple visual surveys to model the habitat of deep-diving cetaceans at the basin scale**

### **Large scale modelling of deep-diving cetacean habitats**

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#### **Biosketch**

This work is part of Auriane Virgili's PhD project which aims to model distributions of rare marine species with a focus on deep-diving cetaceans. These species are rare and difficult to observe at the surface thus it was necessary to assemble datasets from different surveys to model their distribution in the North Atlantic Ocean and the Mediterranean Sea. This required

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# **Large scale modelling of deep-diving cetacean habitats**

## **Abstract**

 **Aim:** Deep-diving cetaceans are oceanic species exposed to multiple anthropogenic pressures including high intensity underwater noise, and knowledge of their distribution is crucial to manage their conservation. Due to intrinsic low densities, wide distribution ranges and limited presence at the sea surface, these species are rarely sighted. Pooling data from multiple visual surveys sharing a common line-transect methodology can increase sightings but requires accounting for heterogeneity in protocols and platforms. 3 Article type **Combining multiple visual surveys to model the habitat of**<br>3 **Groups and the highest relative densities were predicted along the densities were predicted and the higher scale in Alone Cape alone alone alone** 

- **Location:** North Atlantic Ocean and Mediterranean Sea
- **Time period:** 1998 to 2015
- **Major taxa:** *Ziphiidae*; *Physeteriidae*; *Kogiidae*

 **Methods:** About 1,240,000 km of pooled effort provided 630 sightings of ziphiids, 836 of physeteriids and 106 of kogiids. For each taxon, we built a hierarchical model to estimate the effective strip width depending on observation conditions and survey types. We then modelled relative densities in a Generalised Additive Modelling framework. Geographical predictions were limited to interpolations identified with a gap analysis of environmental space coverage.

 **Results:** Deeper areas of the North Atlantic gyre were mostly environmental extrapolation, thereby highlighting gaps in sampling across the different surveys. For the three species  the western North Atlantic Ocean where the Gulf Stream creates dynamic frontal zones and eddies.

 **Main conclusions:** Pooling a large number of surveys provided the first basin-wide models of distribution for deep-diving cetaceans, including several data-deficient taxa, across the North Atlantic Ocean and the Mediterranean Sea. These models can help the conservation of elusive and poorly known marine megafauna.

 **Keywords:** Beaked whales, Data-assembling, Deep-diving cetaceans, Habitat modelling, Kogiids, Sperm whales

### **1. Introduction**

 Deep-diving cetaceans, defined here as beaked whales (family *Ziphiidae*; *e.g. Ziphius cavirostris*, *Hyperoodon* spp. and *Mesoplodon* spp.) and sperm whales (families *Physeteridae* and *Kogiidae*), are distributed worldwide. They are oceanic species that feed in deep waters during long dives (close to or even longer than an hour; Perrin et al., 2009). Due to their offshore habitat and the short time they remain available at the sea surface, little is known about their synoptic distribution (especially for kogiids and ziphiids). Moreover, these species are threatened by anthropogenic activities, including bycatch, debris ingestion, ship collisions (Carrillo & Ritter, 2010; Madsen et al., 2014; Unger et al., 2016) and any activity producing high intensity noise (*e.g.* military sonars, seismic guns or techniques used on large maritime construction projects; Stone & Tasker, 2006). Recent studies have demonstrated the sensitivity of deep-diving cetaceans, and particularly beaked whales, to underwater noise pollution, with a number of unusual stranding events associated with the use of military sonars (Fernández et al., 2005; D'Amico et al., 2009). To mitigate the impact of these activities, accurate knowledge of the distribution and abundance of deep-diving cetaceans is crucial to Marine Spatial Planning to inform management measures at a national scale (Douvere, 2008). International initiatives, such as Important Marine Mammal Areas (IMMAs, Corrigan et al., 2014), are needed for these highly mobile species. However, any single survey often yields only a handful of sightings that are then restricted to areas too small compared to the large geographical scale needed for effective conservation planning. volution many coverage or wave being the many conservation in the matter of equality and the synthesising different and the quality and the quality of the process

 Data-assembling is increasingly used to model habitat preferences of cetaceans at the basin scale (Roberts et al., 2016; Rogan et al., 2017; Cañadas et al., 2018). Due to the various protocols, platform types and observation heights, species detectability and data quality vary with surveys. In addition, each survey may not collect the same information, particularly with regard to observation conditions. Some surveys only record Beaufort sea-state while others record additional parameters that also influence species detection, such as

 only variables common across all datasets can generally be retained in a broad scale analysis, which nevertheless needs to account for heterogeneity. Finally, to make basin-wide predictions from the assemblage of a number of local surveys, identifying areas of environmental extrapolations is crucial to bolster confidence in predicted maps (Mannocci et al., 2018).

 Our study aims to understand how deep-diving cetaceans are distributed at a large scale and to highlight areas of high relative densities for conservation purposes. To model the habitats of deep-diving cetaceans at a large scale, we assembled data from different surveys in the North Atlantic Ocean and the Mediterranean Sea from 15 organisations. To take into account heterogeneity in sighting protocols, we built a hierarchical model to estimate the effective strip width across platforms and observation conditions. We then modelled relative densities of three deep-diving cetacean taxa with Generalized Additive Models (GAM). Finally, we performed a gap analysis (Jennings, 2000, Mannocci et al., 2018) to assess the reliability of the predictions outside the surveyed area. 12 and to helphalma bottlendose of the method of the m

### **2. Methods**

#### **2.1. Data origin**

 The study area encompassed the North Atlantic Ocean and the Mediterranean Sea from the Guiana Plateau to Iceland, excluding the Baltic and Black Seas, the Gulf of Mexico and the Hudson Bay, both because of an absence of effort data and of ecological and environmental differences (Fig. 1A; Appendix S1 in Supporting Information). Four sub- regions were defined in the study area (Table 1; Fig. 1A): the northeast Atlantic Ocean (NE- ATL), the northwest Atlantic Ocean (NW-ATL), the tropics and the Mediterranean Sea (MED).

 We assembled visual shipboard and aerial surveys performed by 15 independent organisations in the North Atlantic Ocean and the Mediterranean Sea between 1998 and 2015 (Fig. 1; survey-specific information are detailed in Appendix S2 in Supporting Information). Except for the JNCC-ESAS surveys that use a 300m-strip-transect methodology, all surveys used line-transect methodologies that correct for non-detection bias with the estimation of an Effective Strip Width (ESW) from the measurement of the perpendicular distances to the sightings (Buckland et al., 2015; see below).

 To account for the difficulty in identifying deep-diving cetaceans to the species level (*e.g.* genera *Mesoplodon*, *Kogia*), we pooled species into three groups: (1) beaked whales, consisting of Cuvier's beaked whales (*Ziphius cavirostris),* mesoplodonts (*Mesoplodon* spp.)  *macrocephalus*), and (3) kogiids, including pygmy (*Kogia breviceps)* and dwarf sperm whales *(K. sima*).

#### **2.3. Data processing**

#### *2.3.1. Data-assembling*

 All survey datasets were standardised for units and formats (*e.g.* date, time and coordinates) and aggregated into a single common dataset. A specific coordinate projection encompassing the entire survey area was used for accurate distance computations (Albers equal-area conic defined from http://projectionwizard.org). Effort data were linearized and divided into 5 km segments using ArcGIS 10.3 (ESRI, 2016) and the Marine Geospatial Ecology Tools software (Roberts et al., 2010). The segment length represented a trade-off value across varying survey transect lengths, for example aerial surveys had transect lengths 115 of up to 100 km while shipboard surveys were often much shorter. Finally, for each species 116 group, sightings were linked to their respective 5 km segments. 23.3. *Columinates* corrects strip with a method of the decreasing detective strip with a strip correct in a correct strip of the decreasing the orrect strip equilibratic projection<br>131 equal-area cone defined for the trav

117 Encounter rates were calculated in each sub-region as: (number of encounters/ 118 total distance travelled)  $*100$ .

#### *2.3.2. Environmental variables*

 In habitat models, we tested the static and dynamic variables that were expected to influence the distributions of deep divers (Table 2). All variables were resampled at a 0.25° resolution because of the very large size of the study area and the spatial resolution of the variables (Table 2; Appendix S3 in Supporting Information). Spatial gradients of sea surface temperature (SST) were calculated as the difference between the minimum and maximum SST values in an eight-pixel buffer around a given pixel. Net primary production (NPP) was used as a proxy for prey availability.

 Dynamic variables, which relate to the movements of water masses or prey availability, were computed at a monthly resolution *i.e.* averaged over the 29 days prior to each sampled day to avoid gaps in remote sensing oceanographic variables. They were used in addition to static variables because they reveal the presence of time-stable structures such as temperature gradients or eddies when variables are averaged.

#### *2.3.3. Effective Strip Width estimation*

 Line-transect surveys are commonly used to estimate cetacean abundance (Hammond et al., 2013; Buckland et al., 2015). A key parameter to estimate this abundance is the  from the trackline. ESW is expected to depend on survey platform height, platform type, sea-state, species, etc… (Buckland et al., 2015).

 ESW estimation was a key step in the data-assembling process to take into account heterogeneity in effort per segment in the models and to directly compare the different surveys (Hedley & Buckland, 2004). ESWs are generally estimated for each survey (*i.e.* no pooling of information) by using the 'Distance' software (Thomas et al., 2010; Buckland et al., 2015). However, for deep-diving cetaceans, the majority of surveys contained insufficient 145 sightings to allow survey-specific detection functions to be fitted. Consequently, for each species group, we pooled sightings from the various surveys, taking into account survey heterogeneity. We built a hierarchical model in which survey identity was included as a random effect.

 In conventional distance sampling (Marques & Buckland, 2003; Buckland et al., 2015), factors such as the characteristics of the species being surveyed, search methods, search platform, environmental conditions can all affect ESW estimation. However, the different datasets did not always contain this information, especially regarding observation conditions. All surveys recorded environmental data such as Beaufort sea-state, cloud coverage and sun glare, although Beaufort sea-state was the only parameter recorded by all of them. Platform type, observation height and Beaufort sea-state were used as covariates in the hierarchical model.

157 Truncation distance w was first determined as the  $95<sup>th</sup>$  percentile of the set of perpendicular distances for each species group, *i.e.* the 5% most distant sightings were discarded from the analysis (Buckland et al., 2001, page 16). Then, we created classes to pool the different surveys; namely platform type (plane or boat), observation height (*e.g.* 0-5 m; 5-10 m…) and Beaufort sea-state (0-1; 1-2; 2-3 and 3-4; data collected beyond a Beaufort sea-state 4 being removed from the analysis). Hierarchical modelling was then performed in R-3.3.1 (R Core Team, 2016) in a Bayesian framework using JAGS version 4-6 and package 'rjags' (jags model in Appendix S4 in Supporting Information; Royle & Dorazio, 2008; Plummer, 2016). 214 soling the survey are survey in the mand is an universal to user survey random the survey random effects. Bivariate in the survey scenarios of the survey scenarios of the survey scenarios in the survey scenarios in th

 For each taxa, perpendicular distances of sightings were used to estimate a detection 167 function with a hazard key. For a sighting i made during survey s at height j under Beaufort 168 sea-state k, let  $d_{iks}^i$  denotes the perpendicular distance. The detection probability of sighting is:

$$
\begin{cases}\np_{ijk}^s = g_s(d_{ijk}) = 1 - \exp\left(-\left(\frac{d_{ijk}}{\sigma_{jks}}\right)^{-\nu_s}\right) \\
\log(\sigma_{jks}) = \beta_{j0} + \beta_{j1} \times k + \alpha_s\n\end{cases}
$$

170 where  $\beta_{j0}$  and  $\beta_{j1}$  are respectively random intercept and slope parameters for the effect of

 specified with a Cholesky decomposition and using priors for the Cholesky factors from Kinney & Dunson (2008). We used half Student-t distributions with 3 degrees of freedom and scale set to 1.5 as priors for dispersion parameters, and standard normal priors for all other parameters. Four chains were run with a warmup of 10,000 iterations, followed by another 10,000 iterations (with a thinning factor of 10). Parameter convergence was assessed with 177 Gelman-Rubin  $\hat{R}$  statistics. Posterior inferences were based on the pooled sample of 4,000 values (1,000 per chain).

 The advantage of setting a hierarchical model to estimate detection functions is to borrow strength across the different datasets to increase the precision of estimates. For each combination of survey – platform type – observation height – Beaufort sea-state, estimated detection functions are shrunk towards a common detection function (itself estimated from the data) according to the available data corresponding to this particular combination of survey – platform type – observation height – Beaufort sea-state. If, for a given combination of parameters, there were few sightings, the estimated detection function was very close to the common detection function, whereas if there were enough data, the estimated detection function could deviate from this common function. Upon model fitting and successful parameter estimation, the ESW for each combination of survey – platform type – observation height – Beaufort sea-state was computed: 202<br>
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ESW_{jks} = \int_0^w g_s(x) dx = \int_0^w \left[1 - \exp\left(-\left(\frac{x}{e^{\beta_{j0} + \beta_{j1} \times k + \alpha_s}}\right)^{-\nu_s}\right)\right] dx
$$

190 The posterior mean value of estimated ESW was then allocated to each segment with 191 respect to species group, survey, platform type, sea-state and observation height class.

192

#### 193 **2.4. Habitat modelling**

 To model habitat preferences of deep-divers, we fitted Generalised Additive Models (GAMs; Hastie & Tibshirani, 1986; Wood, 2006) with a Tweedie distribution to account for over-dispersion (Foster & Bravington, 2013) with the 'mgcv' R-package (R-3.3.1. version; Wood, 2013). GAMs extend Generalised Linear Models to allow for smooth nonlinear functions of predictor variables (Hastie & Tibshirani, 1986; Wood, 2006). The mean number 199 of individuals per segment  $\mu$  was modelled with a logarithmic link function:

$$
\log(\mu) = \alpha + \sum_{p} f(X_p)
$$

200 where  $f(X_n)$  are non-parametric smooth functions (thin plate regression splines) of the 201 covariates and  $\alpha$  is the intercept (Hastie & Tibshirani, 1986). To attenuate the scope for 202 over-fitting, the maximum number of knots was limited to 4 (macy parameter  $k = 4$ ; Wood, 203 2006). An offset equal to segment length multiplied by twice the ESW was included (except  removed combinations of variables with Spearman partial correlation coefficients higher than |0.7| (Dormann et al., 2013; Mannocci et al., 2014) and tested all models with combinations of one to four variables. A maximum of four covariates per model was used to avoid excessive complexity of models and difficulty in their interpretation (Mannocci et al., 2014). Model selection was done with the Akaike Information Criterion (AIC, the lower the better; Anderson & Burnham, 2002) and Akaike model weight (akaike.weights function from 'qpcR' package; Spiess, 2014).

 A key assumption of line-transect surveys is that animals on the trackline are always detected (Buckland et al., 2001). However, this assumption is not met with diving species 214 and trackline detection probability  $q(0)$  needs to be accounted for (Barlow, 2015). Observers on a plane spend less time in a given area and the following inequality is expected:  $g^{boat}(0) > g^{plane}(0)$ . Thus a segment of effort with zero sighting of deep-divers is more likely to be a false absence (non-detection of a diving animal present on the trackline) if that 218 segment comes from a plane survey rather than a boat survey. As detection probability  $q(0)$  was not available for every survey and is expected to differ between platforms, we calculated 220 the ratio of  $g(0)$  between the plane and boat platforms from Roberts et al. (2016) and obtained a ratio of approx. 1/5 for beaked whales, approx. 2/5 for sperm whales and approx. 1/3 for kogiids. These crude ratios were then used to weight plane segments with zero sightings when fitting GAMs. While this method does not fully correct for availability bias, it down-weights zeroes from plane surveys. 209 Moder selection was<br>
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 We fitted "year-round" models as the studied taxa have been reported to show little or no seasonal variation in their habitats (*e.g.* Wimmer & Whitehead, 2004; McSweeney et al., 2007). We did not model yearly variations because of little temporal overlap between surveys. Consequently, the year effect is confounded with survey heterogeneity.

 Predictions of relative densities (in number of animals per km²) were provided at 0.25° resolution. There was not enough data to fit a model by month or by season (the number of sightings in winter was too low) and we therefore produced averaged maps over the entire time period. These predictive maps provided the expected distribution of beaked whales, sperm whales and kogiids according to static and monthly environmental conditions to highlight relationships with static (canyons and seamounts) and time-stable structures (temperature gradients or eddies).

 Finally, coefficients of variation (CVs) were estimated for each 0.25° pixel. Coefficients of variations are a measure of the prediction uncertainty per cell, it is a standard error associated with the calculation of the prediction. Therefore, high CVs indicate high model uncertainties due to the lack of detection.

 Even though more than 1,240,000 km of effort was pooled, extensive geographical gaps remained. Predictions in the middle of the Atlantic Ocean are from geographical extrapolation (Fig. 1A) but not necessarily environmental extrapolations. The latter depends on the selected habitat models and covariates therein. We conducted a gap analysis on environmental space coverage to identify areas where habitat models could produce reliable predictions outside survey blocks, *i.e.* geographical extrapolation, whilst remaining within the ranges of surveyed conditions for the combinations of covariates selected by the models, *i.e*. areas of environmental interpolation (Jennings, 2000; Mannocci et al., 2018).

 From the selected models for each taxa, we estimated the convex hull defined by the environmental data used to fit habitat models (hereafter the calibration data). The convex 252 hull of a set of points is the smallest [convex envelop](https://en.wikipedia.org/wiki/Convex_set) that contains all these points. We then assessed whether a prediction from a set of environmental covariates with a given model fall inside or outside this convex hull (King & Zeng, 2007; Authier et al., 2016). We used climatological predictors instead of monthly predictors to lessen the computational burden.

 Due to the large number of data (more than 280,000 points in the calibration dataset), convex hulls were estimated by random sub-sampling with the 'WhatIf' R-package (Stoll et al., 2014). We randomly extracted a fraction of the calibration dataset (10,000 points) to estimate a convex hull and assess environmental extrapolation in the prediction dataset. A combination of climatological predictor values that fall inside the convex hull corresponds to an interpolation. Combinations of climatological predictor values that were classified as interpolations were set aside but other combinations were retained and further tested against another random sample of 10,000 points from the calibration data. This procedure was 264 carried out until the complete calibration dataset was examined. reinal space<br>Ins outside st<br>Ins outside starting the selecte<br>nental data u<br>set of points<br>d whether a r<br>outside thi<br>poical predict<br>to the large<br>nulls were es<br>4). We randc<br>a convex hu<br>a convex hu<br>tion of climat<br>polation. Cor

 The full procedure was conducted twice. In a simple approach, the full range of sampled variables was considered to identify all points of the whole study area where the actual combinations of environmental variables had been sampled in survey blocks. In a more 'precautionary approach', we excluded 5% of the extreme values of the sampled environmental variables to include in the interpolation areas only the points whose associated combinations of covariates fell within 95% of the core ranges sampled. This allowed the definition of two levels of confidence (hereafter 'simple' and 'precautionary') in 272 the predictions.

 Finally, we produced maps delineating the extent of the simple and precautionary interpolation areas and overlaid them with the relative density prediction maps to show areas with greater reliability.

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#### **3. Results**

#### **3.1. Encounter rates**

 The survey pool represented a total of 1,240,000 km of on-effort transects (*i.e.* following a transect at a specified speed and altitude with a specified level of visual effort) of which 58% were carried out by plane and 42% by boat (Fig. 1A, Table 1). Effort data with a Beaufort sea-state higher than 4, which represented 9% of the effort data, were removed from further analysis to only keep sightings collected during good to excellent detection conditions. Most sampling effort was performed in the northeast (37 %) and northwest (45 %) Atlantic Ocean. Surveys in the Mediterranean Sea and in the tropics represented 287 respectively only 16 % and 2 % of total sampling effort.

 A total of 630 sightings of beaked whales, 836 sightings of sperm whales and 106 sightings of kogiids, mainly distributed in the northeast and northwest Atlantic Ocean (north of the 35°N latitude) and in the northwest Mediterranean Sea, were assembled for the present study (Fig. 1B-D).

Overall encounter rates were very low with 0.05 sightings∙100 km<sup>-1</sup> for beaked whales, 0.07 sightings∙100 km<sup>-1</sup> for sperm whales and <0.01 sightings∙100 km<sup>-1</sup> for kogiids (Table 3). The highest encounter rates were recorded in the tropics for all three species groups, particularly for kogiids. There were no sightings of kogiids in the Mediterranean Sea.

#### **3.2. Effective strip width**

 Estimated ESWs varied across surveys and platform type and were on average narrower in aerial than shipboard surveys (Fig. 2). This is probably because aerial observers are more restricted to recording animals below the plane while shipboard observers can look further afield. ESWs were generally larger and more consistent between surveys using the same platform type, for sperm whales than for beaked whales. There were not enough kogiid sightings to estimate an ESW for each survey and particularly for shipboard surveys; consequently, we pooled all aerial surveys and estimated an ESW of 1.1 km that was then applied to all surveys (shipboard and aerial). 282 58% were carried of<br>
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 The outcomes from the hierarchical model were consistent with expectations (Fig. S4.1 in Supporting Information S4): a decrease in Beaufort sea-state (less wind-sea) resulted in a larger ESW (milder non-detection bias).

#### **3.3. Habitat modelling**

 For each species group, selected variables, explained deviances and Akaike weights are shown in Table 4.

 Highest relative densities were found in depth *ca.* 1,500 m, high values of slopes and SST and intermediate NPP. This resulted in high predicted relative densities of beaked whales along steep slope areas associated with deep depths and high gradients of temperature, particularly on the western side of the Atlantic Ocean. The lowest relative densities were predicted in the Mediterranean Sea (Fig. 3B).

 The gap analysis identified areas where the combination of the four variables selected by the best model had not been sampled. Reliable predictions were available for 94% of the study area under the simple approach and only 53% under the precautionary approach (Fig. 3B and 3C). This discrepancy was mostly due to low sampling effort in the oceanic zone. Coefficients of (temporal) variation were higher on the continental shelf associated with high gradients of SST, where beaked whales were not sighted in any of the surveys (Fig. S5.2A in Supporting Information S5). 330 effered areas (Fig. 43 effered areas (Fig. 45).<br>
330 anological contributed areas where the combination of the four variables of 94% of the<br>
331 bt and 30%. This discorpancy was mostly due to low sampling effort in the

#### **Sperm whales**

 Predicted relative densities of sperm whales increased in deep waters (> 2000 m) associated with high gradients of SST and high NPP. The highest relative densities were 331 also predicted on the western side of the Atlantic Ocean, along the Gulf Stream, although were lowest in the Mediterranean Sea (Fig. 4B).

 Reliable predictions for sperm whales were available for 84% of the study area under the simple approach and only 30% under the precautionary approach because of low survey effort in deeper areas. The highest predicted relative densities were predicted outside the precautionary interpolation zone (Fig. 4B and 4C). Coefficients of (temporal) variation were highest in non-sampled areas where uncertainty was therefore greatest (Fig. S5.2B in Supporting Information S5).

#### **Kogiids**

 As the Akaike weight was small for kogiids (0.17), we used model-averaging and generated predictions from the five first models (cumulative Akaike weight of 0.63) and because all predictions were very similar (see Appendix S6 in Supporting Information), we only kept the first model for practical reasons. The highest relative densities were found in deep waters associated with fronts, canyons and seamounts (Fig. 5B). The highest relative densities were predicted on the western side of the Atlantic Ocean, along the Gulf Stream (Fig. 5C).

 Reliable predictions for kogiids were available for 94% of the study area under the simple approach against only 55% under the precautionary approach because of low survey  waters and in the Mediterranean Sea where kogiids were not sighted in any of the surveys (Fig. S5.2C in Supporting Information S5).

#### **4. Discussion**

 Deep-diving cetaceans are species characterised by low sighting rates and modelling their habitats is particularly challenging. Our study pooled different surveys allowing us to capitalise on more than 1,240,000 km of survey effort deployed over the North Atlantic Ocean and the Mediterranean Sea in the past two decades. For each taxon, we built a hierarchical model to estimate the effective strip width depending on observation conditions and surveys. We investigated habitats of deep-divers using GAMs with a focus on quantifying how reliable the predictions were. The selected habitat models of deep-diving cetaceans included static environmental variables such as depth and slope as well as spatial gradients of temperatures, revealing the highest densities in the western North Atlantic Ocean. Deeper areas of the North Atlantic gyre were mostly areas of environmental extrapolation, thereby highlighting gaps in sampling across the different surveys.

#### **4.1. Methodological considerations**

 Over the past few years, data-assembling has been increasingly used for the study of top marine predators (Roberts et al., 2016; Rogan et al., 2017; Cañadas et al., 2018). Due to the very low sighting rates of deep-diving cetaceans, each survey taken separately cannot provide enough data to investigate the habitats of these rare species. In contrast to Rogan et al. (2017), we did not assemble data collected with similar protocols but data collected with different variants of the line transect distance sampling protocol which meant standardising the data according to their core communalities before developing a single spatial model. Ideally, at a time when shared databases are becoming increasingly important (*e.g.* OBIS SEAMAP -- http://seamap.env.duke.edu/, EMODnet -- http://www.emodnet.eu/), implementing standardised survey methods would greatly improve data compatibility, by enhancing the level of communalities in shared datasets, and helping to describe large-scale habitats and distributions of marine species. However, we realise this can lead to financial and logistical constraints and the work we present here could be a way to embrace and incorporate the diversity of data collection methods. 385 Doep awing redaceans are species characterised by low sighting rates and modelling<br>387 capitalise common the maintain of car and the next survey profile different survey and origins to study position and the Modelling

 Hierarchical modelling accommodates heterogeneity between surveys; it borrows strength across surveys ('partial pooling') when estimating survey-specific ESWs. The resulting estimates are biased (in proportion to the available data contributed by each survey) toward a common mean, although are more precise than those that would be  when the number of sightings per survey is large (Buckland et al., 2015; Laran et al., 2017; Redfern et al., 2017). Results from the hierarchical model were consistent with expectations and showed that a decrease in Beaufort sea-state values resulted in increased ESW estimates.

 The majority of environmental variables we used in habitat modelling describe the euphotic zone (upper layer) because variables that describe the deep-water column are difficult to obtain or simply do not exist at a basin-wide scale. As deep-diving cetaceans spend most of their time at depth and generally feed on mesopelagic to bathypelagic prey (*e.g.* Perrin et al., 2009; Spitz et al., 2011), the use of surface variables limits the ability to correctly infer their habitat. The identified relationships between deep-diving cetacean abundance and environmental variables may be indirect rather than causal (Austin, 2006). Although causation may be out of reach, prediction remains a worthy goal, especially for spatial planning and conservation (McShea, 2014).

 We took care in using appropriate statistical tools for modelling the habitat of species with few sightings (Virgili et al., 2018). Indeed, Virgili et al. (2018) showed that GAMs with a Tweedie distribution generated reliable habitat modelling predictions for rarely sighted marine predators. Here, the habitat models we selected had moderate to high levels of explained deviances (from 20.6% to 55.7%), suggestive of a good fit to the data. Nevertheless, the rather high explained deviance of the kogiid model (55.7%) might indicate some level of model over-fitting due to the small dataset, even if predictions were in general consistent with the known ecology of the species group (McAlpine, 2009).

#### **4.2. Large-scale deep-diver habitats**

 Depth and spatial gradients of sea surface temperature were consistently selected across deep-diving cetaceans, suggesting a major influence of topographic features and thermal fronts in structuring their habitats. As a result, higher relative densities of deep-divers were predicted in areas of strong gradients associated with thermal fronts in which deep- diver prey aggregates (Bost et al., 2009; Woodson & Litvin, 2015). Indeed, deep-divers typically feed on mesopelagic to bathypelagic species, such as pelagic cephalopods and benthic fishes (Spitz et al., 2011) that aggregate along continental slopes where temperature gradients are the strongest. Hence, the Gulf Stream, which is the most active frontal zone in 418 the study area compared to the eastern boundary currents that are broader and much slower, may explain the high predicted relative densities of deep-divers on the western side of the North Atlantic Ocean (Waring et al., 2001; Roberts et al., 2016). 232 curve for the transfer of the transfer of the cost of the basis and the state of the products and the anti-<br>232 cupholic 200s (upper layer) do not exist at a basin-wide scale. As deep-dwing cetaceans<br>334 diffecult to o

Despite commonalities, each studied taxon also showed specificities. Slope appeared to

 whales are more specific than those of sperm whales, which have broader prey size spectrum (Spitz et al., 2011), and their distribution is more driven by dynamic variables than by static features. Accordingly, the selected model for sperm whales included more dynamic variables such as NPP and SSH than for beaked whales. Canyons and seamounts were included in the selected model for kogiids, suggesting a more restricted habitat than for the other two groups of deep-divers conforming Staudinger et al.'s (2014) evidence of how kogiids' feeding areas concentrated on the deeper shelf and slope, particularly in the epipelagic and mesopelagic zones.

 Overall, our model predictions corroborated species distribution predictions of previous smaller-scale studies. In the Mediterranean Sea, our predictions were consistent with the documented presence of beaked whales and sperm whales in the Alborán, Tyrrhenian and Ligurian Seas (Praca & Gannier, 2008; Arcangeli et al., 2015; Lanfredi et al. 2016; Cañadas et al., 2018) and along the eastern coasts of the Mediterranean Sea (Podestà et al., 2006). In 436 the North Atlantic Ocean, the highest relative densities of beaked whales and sperm whales 437 were predicted along the slope, a result consistent with those of Rogan et al.'s (2017) and Roberts et al.'s (2016). In the Northwest Atlantic Ocean, higher relative densities of kogiids 439 were predicted in warmer and deeper waters, which is consistent with their known ecology (McAlpine, 2009) and the predictions of Mannocci et al. (2017) except for predictions off the coast of Florida. Our predictions could probably be improved by incorporating the NOAA SEFSC surveys of southeast US waters off Florida and Virginia. In contrast to beaked and sperm whales, we were not able to fit a hierarchical model on kogiid sightings and resorted to complete pooling of the plane data to estimate an ESW. This shortcoming probably resulted in a larger bias (with respect to the true density) in predicted relative density of kogiids compared to other deep-diving species. Given the paucity of information on kogiids, we think 447 that our results are tentative but important nonetheless. 427 Included in the selected induer of the selected induer of the selected induer of the selected induer in the selected induer in the Sogiids' feeding areas concentrated experience of beaked v Ligurian Seas (Praca & Ganni

 The gap analysis revealed large gaps in environmental space coverage across the study area, especially in the deeper and less productive waters of the central north Atlantic gyre and in tropical waters. High relative densities of deep-divers were predicted at the margin of the precautionary interpolation zone (Figs. 3-5) in particular because deeper waters and steeper slopes were within the upper 2.5% quantiles of aggregated survey coverage for these two physiographic covariates. This suggests that sampling effort was not sufficient in deeper and steeper areas and more intensive sampling effort performed in these areas could help better describe the habitat used by deep-divers.

 The management and conservation of species and ecosystems increasingly relies on habitat models (McShea, 2014; Hazen et al., 2016). The ability of these to predict species occurrence in non-sampled or poorly documented areas is useful (Fleishman et al., 2001; Lumaret & Jay-Robert, 2002) because the implementation of dedicated surveys is sometimes impracticable due to budgetary and logistical challenges. It is logistically challenging to carry out dedicated cetacean surveys in the middle of the North Atlantic 464 Ocean. However, by collecting data on both sides of the Atlantic Ocean, relative density maps were produced and our analyses indicated these predictions may be reliable (Figs. 3D, 4D, 5D).

 Here, we showed that deep-diving cetaceans are closely associated with stable topographic features, thus it could be possible to delineate marine protected areas that cover the principal habitats used by the species (*e.g.* Cañadas et al., 2005). However, these species are also responsive to temporally dynamic structures, such as thermal fronts, 471 implying that protected areas will need to be large enough to capture seasonal variation of 472 such features. In this context, Important Marine Mammal Areas, which are currently being discussed by the Marine Mammal Protected Areas Task Force and incorporate governmental and intergovernmental considerations (Corrigan et al., 2014), could help the delineation of sufficiently large protected areas. In addition, in a Marine Spatial Planning approach (Douvere, 2008), it would be worthwhile to overlay predicted density maps with anthropogenic pressure maps (Halpern et al., 2008) to define areas where pressures could be mitigated.

#### **5. Conclusion**

 Habitat modelling of rare species is particularly challenging because habitat models require large datasets, yet rare species typically yield low numbers of sightings. As a result, combining datasets is a useful strategy to model the large-scale habitats of deep-divers; beaked whales, sperm whales and kogiids, across the North Atlantic Ocean and the Mediterranean Sea. At a local scale, predicted relative densities of deep-diving cetaceans were consistent with previous studies. At a larger scale, a gradient in predicted relative 487 densities emerged, with the highest relative densities predicted on the western side of the study area. This pattern was evidenced thanks to assembling a large dataset and had not been detected previously. It highlighted the pronounced influence of active frontal zones, such as the Gulf Stream, on deep-diving cetaceans. Even though extensive gaps remain at a large scale, we were able to predict the habitats of these taxa throughout the North Atlantic Ocean and adjacent Mediterranean Sea, thus identifying potential habitats, including in non-493 sampled areas. The visit of the most wise of the study and visites of the Nativity and Ocean. However, the study of the Stamilto Cosan, relative density<br>444 Ocean. However, by collecting data on both sides of the Altan

 area represented geographical extrapolations and about half (mostly deeper waters) represented environmental extrapolations. Indeed, through an environmental space coverage gap analysis, we identified areas in tropical and deep oceanic waters where sampling effort was insufficient to predict habitats and needs to be increased to improve prediction reliability.

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662 All sighting and effort data used in this study are available in the OBIS SEAMAP 663 database: http://seamap.env.duke.edu/. All data providers can be contacted via the OBIS 664 SEAMAP website.

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#### 666 **Tables**

 **Table 1. Effort performed by platform type or Beaufort sea-state for all surveys in the North Atlantic Ocean and the Mediterranean Sea.** This table presents the total effort conducted in each sector broken down by platform type and Beaufort sea-state. Beaufort sea-state values reported with decimals in the surveys were rounded up. For the analyses, all segments with Beaufort sea-state > 4 were excluded. 'NE-ATL' means northeast Atlantic Ocean; 'NW-ATL' means northwest Atlantic Ocean and 'MED' means Mediterranean Sea.



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 **Table 2. Candidate environmental predictors used for the habitat modelling**. All variables were resampled at a 0.25° resolution. A: Depth and slope were derived from GEBCO-08 30 arc-second database (http://www.gebco.net/); 30 arc-second is approximately equal to 0.008°. B: Surface area per cell was calculated in ArcGIS 10.3 from the shapefile of canyons and seamounts provided by Harris et al. (2014). C: The mean, standard error and gradient of Sea Surface Temperature (SST) were calculated from the GHRSST Level 4 CMC  D: The Aviso ¼° DT-MADT geostrophic currents dataset was used to compute mean and standard deviation of Sea Surface Height (SSH) and Eddy Kinetic Energy (EKE; https://www.aviso.altimetry.fr/en/data/products/sea- surface-height-products/global/madt-h-uv.html). E: Net primary production (NPP) was derived from SeaWIFS and Aqua using the Vertically Generalised Production Model (VGPM; http://orca.science.oregonstate.edu/1080.by.2160.8day.hdf.vgpm.m.chl.m.sst.php).

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**Table 3. Encounter rates in sightings∙100 km-1** 698 **calculated for the entire study area and each sub-region of**  699 **the North Atlantic Ocean and the Mediterranean Sea.** 'NE-ATL' means northeast Atlantic Ocean; 'NW-ATL' 700 means northwest Atlantic Ocean and 'MED' means Mediterranean Sea.

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#### 706 **Table 4. Summary of the selected models by species group.**



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#### 708 **Figures**

 **Fig. 1. Study area divided into sub-regions showing assembled survey effort (A), along with the beaked whale (B), sperm whale (C) and kogiid (D) sightings recorded during all surveys.** The blue polygon delineates overall study area and other polygons delineate sub-regions. Surveys were carried out along transects following a line-transect methodology (survey details in Appendix S1 in Supporting Information). Sightings were classified by group sizes with each point representing one group of individuals and point size representing the number of animals in a group.

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#### 716 **Fig. 2. Beaked whale and sperm whale averaged ESWs estimated for each survey group and each**

717 **platform type.** For each survey group, the boxplot represents the extent of estimated ESWs depending on 718 Beaufort sea-states and observation heights recorded within the group.

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720 **Fig. 3. Functional relationships for the selected variable (A) and the predicted relative densities of beaked whales in individuals·km-2** 721 **(B and C).** A: Solid lines are the estimated smooth functions, and the shaded regions 722 represent the approximate 95% confidence intervals. The y-axes indicate the number of individuals on a log 723 scale, where zero indicates no effect of the covariate. The vertical lines indicate the  $2.5<sup>th</sup>$  and 97.5<sup>th</sup> quantiles of 724 the data. Black areas on prediction maps (B: without precautionary approach and C: with a 5% precautionary 725 approach) represent zones where we did not extrapolate the predictions. Percentages represent the proportion of 726 the study area defined as interpolation with the gap analysis.

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 **Fig. 4. Functional relationships for the selected variable (A) and the predicted relative densities of sperm whales in individuals·km-2 (B and C).** A: Solid lines are the estimated smooth functions, and the shaded regions represent the approximate 95% confidence intervals. The y-axes indicate the number of individuals on a log 731 scale, where zero indicates no effect of the covariate. The vertical lines indicate the 2.5<sup>th</sup> and 97.5<sup>th</sup> quantiles of the data. Black areas on prediction maps (B: without precautionary approach and C: with a 5% precautionary approach) represent zones where we did not extrapolate the predictions. Percentages represent the proportion of 734 the study area defined as interpolation with the gap analysis.

 **Fig. 5. Functional relationships for the selected variable (A) and the predicted relative densities of kogiids in individuals·km-2 (B and C).** A: Solid lines are the estimated smooth functions, and the shaded regions represent the approximate 95% confidence intervals. The y-axes indicate the number of individuals on a log 739 scale, where zero indicates no effect of the covariate. The vertical lines indicate the 2.5<sup>th</sup> and 97.5<sup>th</sup> quantiles of the data. Black areas on prediction maps (B: without precautionary approach and C: with a 5% precautionary approach) represent zones where we did not extrapolate the predictions. Percentages represent the proportion of 742 the study area defined as interpolation with the gap analysis. Figure 1.1 Figureshing the study are defined as interpolation with the gap analysis.<br>
The study area defined as interpolation with the gap analysis.<br>
T35<br>
Fig. 5. Functional relationships for the selected variable (weights

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### **Supporting Information**

**Appendix S1:** Characteristics of the study area.

- **Appendix S2:** Details of surveys used in the analyses. Total effort represents the total length of transects of each survey (without removing the transects with a Beaufort sea-state > 4). MED: Mediterranean Sea; NE-ATL: Northeast Atlantic Ocean; NW-ATL: Northwest Atlantic Ocean.
- **Appendix S3**: Monthly environmental conditions averaged over the study period (from 1998 to 2015).
- **Appendix S4:** Effective Strip Width estimation methodology.

**Appendix S5:** Supporting information for the models.

 **Appendix S6:** Comparison between predictions of the 5 best kogiid models, the average prediction of the 5 best models (Mean) and the average prediction of the 5 best models weighted by the Akaike weight (Weighted). The 5 models are described in the table at the bottom (mod: model; AIC: Akaike Information Criterion). "Mean" is the simple average of the predictions of the 5 best models. To calculate the "Weighted" prediction, we averaged the predictions of the 5 best models by weighting each prediction by the Akaike





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