Habitat-based density estimates for cetaceans in the California Current Ecosystem based on 1991–2018 survey data

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HABITAT-BASED DENSITY ESTIMATES FOR CETACEANS IN THE CALIFORNIA CURRENT ECOSYSTEM BASED ON 1991-2018 SURVEY DATA

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Introduction

The 2018 California Current Ecosystem Survey (CCES) was conducted between 26 June and 4 December 2018 as a joint project of the Marine Mammal and Turtle Division (MMTD) and the Fisheries Resources Division (FRD) of NOAA’s Southwest Fisheries Science Center (SWFSC). One of the primary objectives of this line-transect survey was to collect marine mammal sighting data to support the derivation of cetacean density estimates for the California Current Ecosystem (CCE) study area. Given the heterogeneity of the 2018 survey coverage in the CCE study area (Henry et al. 2020), density estimation required model-based (rather than design-based) analytical approaches for updating population size estimates for US West Coast marine mammal stocks. This report summarizes the results of the cetacean habitat modeling effort.

Habitat models, or species distribution models (SDMs), have been recognized as valuable tools for estimating the density and distribution of cetaceans and assessing potential impacts from a wide range of anthropogenic activities (e.g., Abrahms et al. 2019; Gilles et al. 2011; Goetz et al. 2012; Hammond et al. 2013; Redfern et al. 2013). SDMs for cetaceans have been developed for US West Coast waters from systematic ship survey data collected by SWFSC since 1991 (Barlow et al. 2009; Becker et al. 2010, 2014, 2016, 2018, 2020; Forney 2000; Forney et al. 2012). The most recent models provide spatially-explicit density predictions at a 0.1° (approximately 10km x 10km) grid resolution (Becker et al. 2020), and multi-year average density surfaces have been used by the US Navy to assess potential impacts on cetaceans as required by US regulations such as the Marine Mammal Protection Act and Endangered Species Act (U.S. Department of the Navy 2013, 2015, 2017).

The overall goal of this study was to include the 2018 survey data in the previous 1991–2014 modeling dataset in order to improve SMDs for the CCE study area. Specific objectives included:

- Generating multi-year average density surfaces for the Navy and others to use in their long-term (2–7 year) environmental planning efforts; and
- Providing updated abundance and “minimum population size (Nmin)” estimates as defined in the Guidelines for Assessing Marine Mammal Stocks (National Oceanic and Atmospheric Administration 2016).

To develop improved SDMs and to update US West Coast cetacean stock abundance estimates, sighting data from CCES 2018 were combined with previous line-transect survey data collected within the CCE to create a robust modeling database spanning more than 25 years (1991–2018). Habitat models were developed based on previously established methods that allow for the incorporation of segment-specific estimates of detection probability and included dynamic covariates from an ocean model calibrated to the CCE study area (Becker et al. 2016). In addition, recently-developed techniques for deriving more comprehensive estimates of uncertainty in SDM predictions (Miller et al. In Prep.) were used to provide variance estimates for the model-based abundance estimates. SDMs were developed for long-beaked common dolphin (*Delphinus delphis bairdii*), short-beaked common dolphin (*Delphinus delphis delphis*), Risso’s dolphin (*Grampus griseus*), Pacific white-sided dolphin (*Lagenorhynchus obliquidens*), northern right whale dolphin (*Lissodelphis borealis*), striped dolphin (*Stenella coerulealba*), common bottlenose dolphin (*Tursiops truncatus*), Dall’s porpoise (*Phocoenoides dalli*), sperm
whale (*Physeter macrocephalus*), blue whale (*Balaenoptera musculus*), fin whale (*B. physalus*), humpback whale (*Megaptera novaeangliae*), Baird’s beaked whale (*Berardius bairdii*), and a “small beaked whale guild” that included Mesoplodons (*Mesoplodon* spp.) and Cuvier’s beaked whale (*Ziphius cavirostris*). Sample sizes were also sufficient to develop the first model-based density estimates for minke whale (*B. acutorostrata*) in this study area.

The habitat-based models of cetacean density developed in this study represent an improvement over the previous models described by Becker et al. (2020) because they included additional sighting data over the continental shelf and slope that were surveyed more sparsely in previous years, providing better representation of these important habitat regions. In addition, the model-based abundance estimates more accurately account for uncertainty than prior iterations owing to methodological improvements.
Methods

Survey data

Cetacean sighting data used to build the SDMs were collected within waters of the CCE from 1991–2018 (Table 1) using line-transect methods (Buckland et al. 2001). The 1991–1993 surveys covered waters off the state of California while the 1996–2008 and 2014 surveys covered waters off the entire west coast of the United States, with all surveys extending approximately 300 nautical miles offshore (Barlow and Forney 2007). The 2009 survey was a finer-scale survey that focused on waters off central and southern California, as well as the west coast of Baja California (Carretta et al. 2011). The 2018 survey covered waters along the west coasts of southern Canada (Vancouver Island), the west coast of the United States, and Baja California out to a distance of approximately 200 nautical miles offshore (Henry et al. 2020). When combined across years, the surveys provided comprehensive coverage of waters throughout the CCE study area, although the spatial heterogeneity of the 2018 survey is clearly apparent (Figure 1). Only on-effort data collected in Beaufort sea state conditions ≤5 within the study area were used in model development.

The survey protocols were the same for all years (see Barlow 2006; Kinzey et al. 2000) and are briefly summarized here. Each survey used a NOAA research vessel and a team of six experienced visual observers. For each rotation, three observers stationed on the flying bridge of the ship visually searched for and recorded cetacean sightings between 0 and 90 degree to port and starboard using standard line-transect protocols. Port and starboard observers searched with pedestal-mounted 25 × 150 binoculars and a center-stationed third observer searched by eye or with handheld 7 × 50 binoculars. When cetaceans were detected within 3 nautical miles (5.6 km) of the trackline, the sighting was recorded (along with distance and direction from the vessel, from which perpendicular sighting distance was calculated), and the ship would then typically divert from the transect line and go “off-effort” to approach the animals and enable more accurate estimation of group size and species identification. All observers independently provided best, high, and low group size estimates. If the sighting included more than one species, the observers also estimated the percentage of each species in the group. The best estimate from each observer or the best estimate multiplied by the percentage of each species was averaged (i.e., arithmetic mean) to obtain a single group size estimate for each sighting.

Systematic survey effort was conducted along predetermined tracklines at a target survey speed of 18.5 km/hr. During transit between tracklines, transits to or from port, or deviations from predetermined tracklines for other purposes, the visual observers generally maintained standard data collection protocols. Although such non-systematic effort is generally not used to derive encounter rate for design-based density estimates, it is incorporated into the SDMs as the uneven distribution of effort can be accounted for within the statistical framework (Hedley and Buckland 2004).

Environmental predictor data

To create samples for modeling, continuous portions of on-effort survey tracklines were divided into approximate 5-km segments using methods described by Becker et al. (2010). The total number of species-specific sightings and associated average group size estimates were assigned to each segment and habitat covariates were derived based on the segment’s geographical
midpoint. To maintain consistency with the species-specific effective-strip-width estimates derived for this study based on methods described in Barlow et al. (2011a) and used to estimate cetacean densities, sighting data were truncated at a distance of 5.5 km perpendicular to the trackline for the delphinids and large whales, 4.0 km for small whales (Mesoplodons, minke whale, and Cuvier’s beaked whale), and at 3.0 km for Dall’s porpoise (Buckland et al. 2001).

Environmental variables from a data-assimilative CCE implementation of the Regional Ocean Modeling System (ROMS), produced by the University of California Santa Cruz Ocean Modeling and Data Assimilation group (Moore et al. 2011), were used as dynamic predictors as they have proven effective in similar SDMs for this study area (Becker et al. 2016, 2018, 2020). Daily averages for each variable at the 0.1 degree (~10 km) horizontal resolution of the ROMS output were used in the models. The suite of potential dynamic predictors included sea surface temperature (SST) and its standard deviation (sd(SST)), calculated for a 3 × 3-pixel box around the modeling segment midpoint, mixed layer depth (MLD, defined by a 0.5°C deviation from the SST), sea surface height (SSH), and sd(SSH). Water depth (m) was also included as a potential predictor, derived from the ETOPO1 1-arc-min global relief model (Amante and Eakins 2009) and obtained for the midpoint of each transect segment. In addition, distance to the 200-m isobath derived from the geomorphic feature map of the global ocean (Harris et al. 2014) was included in model selection as it represents the edge of the shelf break for much of the U.S. west coast and can be a distinguishing habitat feature for many cetacean species (Becker et al. 2010; Fiedler et al. 1998, 2018). In addition, for those species known to primarily inhabit offshore waters (beaked whales, sperm whale, striped dolphin), distance to the 2,000-m isobath was also included in the list of potential predictor variables, as this depth roughly represents the transition from the continental slope to the continental rise. To differentiate continental shelf, slope, and rise waters, negative values of the distance to isobath terms were used for waters shallower than the 200m or 2,000m isobath. Although the modeling framework applied in our analysis (mgcv; see ‘Habitat Models’ section below) is robust to correlated variables (Wood 2008), distance to the two isobath terms and depth (absolute correlation = 0.75–0.85) were considered separately in the models to avoid any confounding effects.

A spatial term (bivariate spline of longitude and latitude) was also included in the suite of potential predictors because SDMs that explicitly account for geographic effects have exhibited improved explanatory performance as they often account for unmeasured static variables that might be important for driving species distributions (Becker et al. 2018; Cañadas and Hammond 2008; Forney et al. 2015; Hedley and Buckland 2004; Tynan et al. 2005; Williams et al. 2006). The inclusion of a spatial term can result in more robust models but invalidates predictions outside the study area.

A continuous year term was also included as a potential predictor in the models to capture population trends both for species whose abundance has changed substantially during the time period considered in our analyses, and for species for which distribution shifts have resulted in abundance changes over time. For example, increases in population have been documented for both fin whale (Moore and Barlow 2011) and humpback whale (Barlow et al. 2011b), while notable shifts in distribution over the last few decades have resulted in a decline in the number of blue whales (Monnahan et al. 2015), and an increase in the number of short-beaked common dolphins (Barlow 2016; Becker et al. 2018) in the CCE study area. The degrees of freedom for the year term were constrained (i.e., < the maximum of 8 available) in order to capture linear or
thresholds in the response curves rather than simply tracking the variable encounter rates over the survey periods. In addition, since environmental covariates are often correlated with time, and year can serve as a proxy for unmeasured habitat variables, the functional forms of all the other dynamic variables were inspected during the modeling process to ensure they remained stable with the addition of the year term.

**Correction factors**

During CCES 2018, operational requirements necessitated that some of the effort be conducted in passing mode (i.e., when a cetacean/cetacean group is sighted the ship continues on course and is not diverted to the vicinity of the sighting for species identification or group size enumeration). This led to a high proportion of recorded “unidentified large whale” and “Delphinus spp.” sightings, when observers could not confirm which species of large whale or common dolphin subspecies was present, respectively. Omitting these sightings from the modeling dataset would have resulted in an underestimation of animal density for blue, fin, and humpback whales, as well as both long-and short-beaked common dolphins. To reduce this potential downward bias, species-specific correction factors were applied to account for unidentified animals, using the methods described in Becker et al. (2017) and summarized below.

For both the large whale and common dolphin groups, the correction factor $c$ was estimated from the 2018 sighting data according to the simplified formula:

$$
c = 1 + \frac{t_{unid}}{t_{tgt} + t_{oth}}
$$

(1)

where $t_{tgt}$ is the number of individuals identified as the target species, $t_{oth}$ is the number of individuals identified as other species within the broader species group, and $t_{unid}$ is the number of unidentified individuals in that species group. Due to the potential effect of Beaufort sea state on detectability (Barlow et al. 2001, 2011a; Barlow 2015), the correction factors were evaluated to determine if they varied by sea state. If so, separate correction factors were developed by sea state; otherwise a single correction factor was applied. The correction factors were applied to the numbers of animals estimated per segment in the SDMs for the common dolphin and large whale species (see equation 2 below).

The protocol for estimating sperm whale group size changed over the course of the 1991–2018 survey period, with less effort spent estimating group size during the three surveys conducted in the 1990’s. Group size estimates for larger sperm whale groups (> 2 animals) are now known to have been underestimated in the earlier surveys, and a correction factor has been estimated to account for this bias (Moore and Barlow 2014). Prior to modeling, this correction factor (2.3x) was applied to the average group size estimates for observed sperm whale group sizes > 2 for the 1991–1996 surveys. No group size corrections were applied to the other species.

**Habitat models**

Generalized Additive Models (GAM; Wood 2017) were developed in R (v. 3.4.1; R Core Team, 2017) using the package “mgcv” (v. 1.8-31; Wood 2011). Methods largely followed those described in Becker et al. (2016) and are summarized here. One of two modeling frameworks was used for each species, depending on its group size characteristics. For the two *Delphinus*
species that have very large and variable group sizes (e.g., 1 to 2,000 animals per sighting), separate encounter rate and group size models were developed. Encounter rate models were built using all transect segments, regardless of whether they included sightings, using the number of sightings per segment as the response variable and a Tweedie distribution to account for overdispersion (Miller et al. 2013). Group size models were built using only those segments that included sightings, using the natural log of group size as the response variable, and a Gaussian link function. For the rest of the species, GAMs were fit using the number of individuals of the given species per transect segment as the response variable using all transect segments, and a Tweedie distribution to account for overdispersion. The full suite of potential habitat predictors was offered to both the encounter rate and single response GAMs. A tensor product smooth of latitude and longitude (Wood 2003) was the only predictor variable included in the Delphinus group size models.

In all models, restricted maximum likelihood (REML) was used to obtain parameter estimates (Marra and Wood 2011). The shrinkage approach of Marra and Wood (2011) was used to potentially remove terms from each model by modifying the smoothing penalty, allowing the smooth effect to be shrunk to zero. Additionally, to avoid overfitting, an iterative forwards/backwards selection process was used to remove variables that had P-values > 0.05 (Redfern et al. 2017; Roberts et al. 2016). The natural log of the effective area searched (described below) was included as an offset in both the single response and encounter rate models.

Predictions from the final model were incorporated into the standard line-transect equation (Buckland et al. 2001) to estimate density ($D$; number of animals per km$^2$):

$$D_i = \frac{n_i \cdot s_i \cdot c_i}{A_i}$$  \hspace{1cm} (2)

where $i$ is the segment, $n$ is the number of sightings on segment $i$, $s$ is the average group size (i.e., number of a given species present in a group) on segment $i$, $c$ is the species-specific correction factor for unidentified common dolphins or large whales (derived in equation 1 and assumed to be 1 for all other species) based on sea state conditions on segment $i$, and $A$ is the effective area searched for segment $i$:

$$A_i = 2 \cdot L_i \cdot ESW_i \cdot g(0)_i$$  \hspace{1cm} (3)

where $L_i$ is the length of the effort segment $i$, $ESW_i$ is the effective strip half-width, and $g(0)_i$ is the probability of detection on the transect line. Following the methods of Becker et al. (2016), species-specific and segment-specific estimates of both $ESW$ and $g(0)$ were incorporated into the models based on the recorded detection conditions on that segment and using coefficients estimated specifically for the CCE dataset based on methods of Barlow et al. (2011a) for $ESW$ and Barlow (2015) for $g(0)$. For those segments where the average Beaufort sea state was 0 (<1% of the segments), $g(0)$ was assumed to be 1, i.e., that all animals directly on the transect line were detected, for all species except Cuvier’s beaked whale ($g(0) = 0.584$) and Mesoplodon spp. ($g(0) = 0.813$), which were <1 based on dive behavior (Barlow 2015).

In equation (3) above, the effective area searched is multiplied by two to account for observers searching on both sides of the transect line. During the 2018 survey, coastal fog and other
conditions occasionally prohibited visual observations on one side of the ship, so that cetacean sighting data were collected on only one side of the transect line. These portions of reduced effort were systematically recorded in the dataset and the effective area searched was reduced accordingly along these segments, i.e., the constant was changed to a “1” in equation (3) above.

Model performance was evaluated using established metrics, including the percentage of explained deviance, the area under the receiver operating characteristic curve (AUC; Fawcett 2006), the true skill statistic (TSS; Allouche et al. 2006), and visual inspection of predicted and observed distributions during the 1991–2018 cetacean surveys (Barlow et al. 2009; Becker et al. 2010, 2016; Forney et al. 2012). AUC measures the accuracy of predicting observed presences and absences; values range from 0 to 1, where a score > 0.5 indicates better than random skill. TSS accounts for both false negative and false positive errors and ranges from -1 to +1, where +1 indicates perfect agreement and values of zero or less indicate a performance no better than random. To calculate TSS, the sensitivity-specificity sum maximization approach (Liu et al. 2005) was used to obtain thresholds for species presence. In addition, the model-based abundance estimates for the CCE study area based on the sum of individual modeling segment predictions were compared to standard line-transect estimates derived from the same dataset used for modeling in order to assess potential bias in the habitat-based model predictions. The standard line-transect estimates were derived from the 1991–2018 survey data using equations (2) and (3) above, but without the inclusion of habitat predictors (i.e., observed rather than predicted densities).

Spatially-explicit density values for the CCE study area were derived from model predictions on the environmental conditions specific to the 1991–2018 CCE effort periods at a 0.1° (approximately 10km x 10km) grid resolution. Model predictions were made on separate environmental conditions for each day encompassing the survey periods, thus taking into account the varying oceanographic conditions during the 1991–2018 cetacean surveys. The separate daily predictions thus provide a dataset from which averages can be derived for any temporal period of interest. In past years, the Navy has used a “multi-year average” of predicted daily cetacean species densities to assess potential impacts on cetaceans as required by U.S. regulations such as the MMPA and ESA (U.S. Department of the Navy 2015, 2017). To ensure that the multi-year average reflects more recent conditions and is based on those survey years that more comprehensively covered the study area, predictions for 1991, 1993, and 2009 were not included in the multi-year average. Further, for the two species with documented population increases in the study area (i.e., fin and humpback whales), the year covariate was set to 2018 to decrease the potential for biased-low density estimates derived from the multi-year average surfaces. The daily predictions were also used to create individual yearly averages for 1996–2018. The prediction grid was clipped to the boundaries of the approximate 1,141,800-km² study area to ensure that predictions were not extrapolated outside the region used for model development.

The model-based abundance estimates were calculated as the sum of the individual grid cell abundance estimates, which were derived by multiplying the cell area (in km²) by the predicted grid cell density, exclusive of any portions of the cells located outside the CCE study area or on land. Area calculations were completed using the R packages geosphere and gpclib in R (version 2.15.0).
In highly dynamic ecosystems such as the California Current, variation in environmental conditions has been shown to be one of the greatest sources of uncertainty when predicting density as a function of habitat variables, and this source has been used to provide spatially-explicit variance measures for past CCE SDM model predictions (Barlow et al. 2009; Becker et al. 2016, 2018, 2020; Forney et al. 2012). Recently, Miller et al. (In Prep.) developed techniques for deriving more comprehensive measures of uncertainty in GAM predictions that, in addition to environmental variability, also account for the uncertainty from the GAM parameters, $ESW$, and $g(0)$. These techniques include generating multiple daily density surfaces taking into account model parameter uncertainty and providing a range of density estimates from which variance can be calculated.

Preliminary analyses in our study, however, revealed that the simulated model parameter draws can – for some species – result in a subset of unrealistic simulated surfaces (i.e., surfaces that infer high densities of a species in habitats where the species is not generally found), so this method was not yet deemed suitable for estimating spatially explicit uncertainty estimates for the pixel-based densities. The method did, however, confirm that environmental variability contributes the most substantial source of uncertainty in the CCE model predictions. Therefore, the methods of Becker et al. (2016, 2018) were applied to estimate spatially-explicit measures of uncertainty based on environmental variability, calculated as pixel-specific standard errors using the set of daily predictions that went into the multi-year average density estimates. The pixel-based variance estimates are thus under-estimated to some degree, but the dominant source of uncertainty (environmental variability) was accounted for.

The methods described in Miller et al. (In Prep.) were found to be suitable for estimating uncertainty in the overall model-based abundances for the entire CCE study area, and thus were used to derive variance estimates that included the combined uncertainty from environmental variability, the GAM parameters, and $ESW$. Study area variance was estimated based on the average values of each of the 200 simulations within each year, thereby providing an overall measure of uncertainty associated with the individual yearly average density surfaces for 1996–2018. One additional source of uncertainty in abundance estimates is introduced by $g(0)$, the probability of detecting animals directly on the trackline. The estimates of $g(0)$ developed by Barlow (2015) are based on segment-specific Beaufort sea state conditions, but they were not compatible with the Miller et al. (In Prep.) methods of incorporating $g(0)$; therefore, this source of uncertainty was handled separately. An overall estimate of uncertainty in $g(0)$ was derived using the variance estimates for this parameter weighted by the proportion of survey effort conducted within each of the Beaufort sea state categories and estimated based on 10,000 bootstrap values. Barlow (2015) did not provide $g(0)$ estimates for northern right whale dolphin, and the result for Pacific white-sided dolphin was considered an outlier (Barlow 2015), so for both species the $g(0)$ estimates for Delphinus spp. were used. Delphinus spp. was considered a suitable surrogate for Pacific white-sided dolphin since they have similar sighting characteristics. In addition, the Delphinus spp. $g(0)$ values were similar to the average of all the delphinids and were thus selected as a surrogate for northern right whale dolphin as well. The weighted $g(0)$ uncertainty was combined into the study area variance estimates using the delta method (Seber 1982).

For purposes of calculating Potential Biological Removal (PBR) of US West Coast cetacean stocks, the pooled average of the 2014 and 2018 model-predicted study area abundance estimates
and associated variance estimates, as well as minimum abundance estimates, were also calculated (National Oceanic and Atmospheric Administration 2016). Abundance estimates were based on the arithmetic mean of the model-predicted estimates for 2014 and 2018. Study area variance was estimated based on the methods described above for individual years but including data specific to 2014 and 2018.
Results

Habitat-based density models were developed for 14 species and one guild (Mesoplodonts and Cuvier’s beaked whale) using 92,214 km of on-effort survey data collected between 1991 and 2018 within the CCE study area. The number of sightings within the species-specific truncation distances and available for modeling ranged from 39 to 1,034 (Table 2).

Correction factors for unidentified large whales were applied separately by Beaufort sea state for the 2018 blue, fin, and humpback whale sightings, because the proportion of unidentified whales increased with increasing sea state. For blue and humpback whales, these correction factors were 1.03, 1.04, 1.05, 1.20, and 1.26 for Beaufort sea states 0-1, 2, 3, 4, and 5, respectively, and 1.04, 1.08, 1.10, 1.30, and 1.46 for fin whales. For the common dolphin group, higher multipliers were not associated with higher sea states, so a uniform correction factor of 1.71 was applied across all sea states for the 2018 sightings of both long- and short-beaked common dolphins.

Consistent with past modeling studies in the CCE study area (Becker et al. 2016, 2018, 2020), the most commonly selected predictor variables for the encounter rate models of groups (long- and short-beaked common dolphins) or individuals (all other species) included SST, MLD, and the smooth of latitude and longitude (Table 3). SSH and depth were also selected in many of the models. The group size model for both subspecies of common dolphin included a bivariate spline of longitude and latitude, consistent with other studies that have demonstrated significant spatial variation in group size, particularly for Delphinids (Barlow 2015; Ferguson et al. 2006). The functional forms of the key predictor variables were also consistent with those of SDMs built with subsets of the modeling dataset used for this study (Becker et al. 2016, 2018, 2020; Appendix A).

A year covariate was included in the final fin and humpback whale models, and both captured the documented increasing population trends for these species in the CCE study area (Moore and Barlow 2011; Barlow et al. 2011a; Calambokidis et al. 2017). A year term was also included in the models for short-beaked common dolphin and blue whale, consistent with observed northern shifts in the relative distribution of these two species that have resulted in increasing numbers of short-beaked common dolphins and decreasing numbers of blue whales in the CCE study area (Barlow 2016; Becker et al. 2018; Monnahan et al. 2015). A year term was also included in the SDMs for Risso’s, striped, and common bottlenose dolphins, as well as Dall’s porpoise (Table 3). The functional forms for the year term in all but the striped dolphin model suggest a decreasing trend in the numbers of these species in the CCE study area during the course of the survey period (Appendix A). For all three species, year represents a significant but very small effect as indicated by the range of values on the y-axis (i.e., relative to the other covariates the y-axis value for year is <1; Figures A3,A7, A8). The functional form of the year term in the striped dolphin model fluctuates throughout the 1991–2018 survey period (Figure A6), consistent with the highly variable abundance estimates for this species for each of the individual survey years (Barlow 2016; Becker et al. 2018).

Deviance explained by the models was variable, ranging from approximately 7% to 57% (Table 3). With the exception of sperm whale, AUC values for all models were greater than 0.7 and the majority were greater than 0.8, indicating that the models did a good job predicting true positives and negatives. The TSS values, which account for both omission and commission errors, were
more variable, ranging from 0.18 (sperm whale) to 0.90 (long-beaked common dolphin). All models had observed: predicted density ratios higher than 0.7, with the majority higher than 0.9, indicating that the sum of the segment-based density predictions captured overall abundance in the study area as derived from design-based line-transect methods.

The 1996–2018 multi-year average density surface maps generally captured observed distribution patterns as illustrated by actual sightings during the surveys (Figure 2). For the two species with documented population increases in the study area (i.e., fin and humpback whales), the density estimates were scaled to the 2018 abundance to decrease the potential for biased-low density estimates derived from the multi-year average surfaces (Figures 2l and 2m). The CVs, which were based on the environmental variability of the daily predictions, showed substantial variation among the species, with a few individual pixel values as high as 6.0 (e.g., common bottlenose dolphin and fin whale, Figures 2g and 2l).

The yearly average density surface maps show high annual variability for some species (e.g., short-beaked common dolphin, striped dolphin, Dall’s porpoise, blue whale, fin whale) and less so for other species (e.g., minke whale, Baird’s beaked whale) (Figure 3). There is almost no variability in the yearly density plots for sperm whale (Figure 3i), due to the overwhelming contribution of the distance to 2,000m isobath term. The pixel-based CVs were generally highest in 2005, suggesting that there was substantial variability in the habitat covariates within this year. For the majority of the species, the yearly sightings match well with the density predictions. However, given the heterogeneity of survey coverage in 2018, sighting data from this survey are not as useful for cross validation since survey coverage needs to be taken into account when assessing the accuracy of the density predictions. For example, the models for both short-beaked common and striped dolphins predict high density in the southwestern portion of the CCE study area in 2018 (Figures 3b and 3f), where there was no survey effort (Figure 1).

The model-based yearly abundance estimates were highly variable for the majority of the species considered here, particularly for those with documented trends due to either changes in abundance or shifts in distribution (i.e., fin, humpback, and blue whales, and short-beaked common dolphin; Table 4). Even for those species for which a year term did not enter the model, substantial variability in the annual model-predicted abundance values were apparent, particularly for the most recent survey years (e.g., long-beaked common dolphin, northern right whale dolphin, Baird’s beaked whale). Interestingly, the most stable mean abundance estimates over the 1991–2018 survey period were for sperm whale and the small beaked whale guild (Table 4), the two SMDs that generally had the worst performance metrics among all the species models (Table 3).

Four sources of uncertainty (i.e., environmental variability, GAM parameters, ESW, and g(0)) were combined to provide an overall measure of variance for the model-based study area abundance estimates (Table 4). Uncertainty estimates from the combination of environmental variability, GAM parameters, and ESW estimates (“CVm (Model)” in Table 4) were variable, ranging from 0.078 for sperm whale to 0.782 for northern right whale dolphin. The final model for sperm whale included only two predictors, of which one was dynamic (Table 3), so the low “Model” CVs are likely due to low parameter variability. Conversely, the final model for northern right whale dolphin included five predictors with large standard error bands around four (Table 3 and Figure A-5), resulting in high variability in the parameter simulations used to derive
the variance estimates. Uncertainty due to the Beaufort-weighted g(0) values was quite high for many of the species, particularly Dall’s porpoise (CV = 0.518) and minke whale (CV = 0.787). When combined, overall measures of CV for the study area abundance estimates were highly variable among the species, ranging from 0.127 (Risso’s dolphin) to 0.799 (minke whale). Similar to the yearly estimates, CVs for the pooled 2014 and 2018 abundance estimates were also variable among species (Table 5).


**Discussion and Conclusions**

During the last 20 years, subsets of the 1991–2018 SWFSC survey data have been used to model the relationship between habitat predictors and species density, both to improve abundance estimates and to gain valuable insight on spatial and temporal changes in species distributions (Barlow et al. 2009; Becker et al. 2010, 2014, 2016, 2018, 2020; Forney 2000; Forney et al. 2012). With each added year of survey data, the models for most species have become more robust, as increased numbers of sightings collected over a broader range of oceanic conditions have been able to better inform the models. The key functional forms for many of the species have become stable over time, suggesting that at this decadal temporal scale relationships with certain habitat predictors have not changed, despite changing oceanic conditions (e.g., Becker et al. 2018). For example, the functional form of SST in the Dall’s porpoise GAM consistently shows a threshold effect at approximately 16°C (Figure A8), apparent in previous GAMs built with only the 1991 and 1996 survey data (Forney 2000). The relationship between SST and fin whale density has also remained constant throughout the 1991 to 2018 period, with the highest densities of whales in waters between about 14°C and 18°C (Figure A12), consistent with GAMs developed with only four years of survey data (1991–2001; Becker et al. 2010). Although high seasonal and interannual variability in cetacean abundance and distribution patterns have been observed and predicted from habitat models developed for the CCE study area (Barlow and Forney 2007; Becker et al. 2014, 2017, 2018; Forney and Barlow 1998; Forney et al. 2012), the multi-year average density plots for the majority of species are broadly similar over the 1991–2018 time period, demonstrating consistency in “average” distribution patterns. These density estimates represent a composite view for the summer/fall survey months (typically July through November) and should not be extrapolated outside of these seasons, given the seasonality of the California Current Ecosystem.

Since a main objective of this study was to produce robust average multi-year density surfaces, a bivariate spline of longitude and latitude was included in the SDMs to increase their explanatory performance (Cañadas and Hammond 2008; Forney et al. 2015; Hedley and Buckland 2004; Tynan et al. 2005; Williams et al. 2006). As Becker et al. (2018) demonstrated, however, for many species the inclusion of a spatial term does not improve a model’s novel predictive power, suggesting that these models may not provide the best nowcasts or forecasts.

For Risso’s dolphin, sperm whale, and the small beaked whale guild, previous SDMs have not performed well, and there has generally been poor correlation between predicted density patterns and the sighting data used to build the models (Becker et al. 2010, 2020; Forney et al. 2012). Sightings of Risso’s dolphins within the CCE study area are concentrated either along the continental shelf (mainly south of 38°N) or in offshore deep waters, with a distinct longitudinal absence between these two areas (Barlow 2016; Barlow and Forney 2007). In the present study, this observed spatial pattern was captured quite well (Figure 2c), likely due to the addition of the CCES 2018 survey data, which contributed an additional 39 sightings to the modeling dataset and provided improved sampling of the continental shelf habitat.

Conversely, models for both sperm whale and the small beaked whale guild showed little to no improvement, with some of the worst model metrics among all species and predicted distribution patterns that match poorly to actual sightings during the surveys (Table 3, Figures 2h, 2n). The addition of the CCES 2018 survey data did not improve either of these models, likely due to the
very sparse sampling of offshore waters where both sperm and small beaked whales are typically found. These results also suggest that the current suite of environmental variables offered to the models are not effective proxies for their habitat and prey. Model improvements for these deep-diving species may only be realized by identifying an available proxy that better captures the ecological processes driving their distribution or by using alternative data (e.g., acoustics) for model input.

Unlike previous modeling efforts where a year term was considered only for those species with documented population increases or decreases in the CCE study area, a year term was included in the list of potential predictors for all the SDMs in this study. To ensure that year did not simply track the variable encounter rates over the 1991–2018 survey period, this term was constrained (i.e., the degrees of freedom were reduced) in the GAMs in order to identify a trend or threshold effect. Consistent with past modeling efforts, the year term entered the SDMs for those species with documented increases in population in the study area (fin and humpback whales; Moore and Barlow 2011; Barlow et al. 2011a; Calambokidis et al. 2017) and for those species with documented distribution shifts that have resulted in substantial changes in the number of animals present in the study area (blue whale and short-beaked common dolphin; Barlow 2016; Becker et al. 2018; Monnahan et al. 2015). A year term was also included in the striped dolphin GAM, indicating fluctuating numbers of this species in the study area over the survey period (Figure A6). This result is consistent with past studies that suggest that available striped dolphin habitat fluctuates substantially with changing ocean conditions (Barlow 2016; Becker et al. 2018, 2020), and since the range of this species extends continuously from the study area south to waters offshore Mexico (Perrin et al. 1985; Mangels and Gerrodette 1994), there can be a large increase or decrease of animals in the study area in any single year.

A year term was also included in the models for Risso’s dolphin, common bottlenose dolphin, and Dall’s porpoise, suggesting a decreasing trend in the numbers of these species in the CCE study area during the course of the survey period (Figures A3, A7, A8). A negative year trend indicates that the numbers of these species in the study area has decreased either due to a true change in population or to a distribution shift out of the study area. Boyd et al. (2018) demonstrated that the amount of suitable Dall’s porpoise habitat within the CCE study area changed substantially during the 1991–2008 survey period, so perhaps this could be driving the apparent decrease in numbers of this species over time. The yearly density predictions for Dall’s porpoise do not appear consistent with a shift in distribution to the north, however, but rather imply a contraction of suitable habitat centered off Oregon and northern California (Figure 3h). Bayesian hierarchical approaches have been used to improve population trend analyses for fin, sperm, and beaked whales in the CCE (Moore and Barlow 2011, 2014, 2017). Similar trend analyses that incorporate the additional 2018 survey data are needed to resolve what is driving the apparent decrease in abundance indicated by the GAMs for Risso’s dolphin, common bottlenose dolphin, and Dall’s porpoise.

The modeling framework used in the present analysis was largely the same as that used in Becker et al. (2016), but incorporated updated measures of uncertainty in the study area abundance estimates based on a modification of the methods described in Miller et al. (In Prep.). This is an improvement from past studies that only accounted for uncertainty due to environmental variability. Uncertainty estimates for the overall study-area abundance estimates based on the combined sources of environmental variability, GAM parameters, and ESW were
generally lower for species with high sighting numbers and lower variability in encounter rates such as short-beaked common dolphin, Dall’s porpoise, and blue, fin, and humpback whales (Table 4). Uncertainty due to the Beaufort-weighted $g(0)$ values was quite high for many of the species, and served to increase uncertainty in the overall study area abundance estimates. This is not surprising given the nontrivial uncertainty estimates associated with the Beaufort-specific $g(0)$ values calculated by Barlow (2015) and used in this study. Similar to past studies, the pixel-based variance estimates presented here account for uncertainty due to environmental variability and are thus under-estimated to some degree. Methods to derive spatially-explicit variance measures that also account for uncertainty in the GAM parameters, $ESW$, and $g(0)$ are currently in development (Miller et al. *In Prep.*).

For all species, abundance estimates derived from the habitat-based models are more stable than previous design-based estimates for each of the 1996–2014 survey years (Barlow 2016). Design-based estimates are based on the realized encounter rates within each year, and are thus subject to high variation due to sampling error and patchiness in both the environment and animal distribution. This generally results in highly variable single year abundance estimates that often appear inconsistent with long-term trends in animal abundance (Moore and Barlow 2014). Conversely, habitat models establish relationships between environmental predictors and species density based on the full, multi-year dataset, and yearly abundance estimates derived from the models are based on the temporally-specific environmental conditions throughout the study area, thus serving to smooth across the annual variation in observed encounter rates along transect lines. This results in less variability in model-based abundance estimates between years, as much of the remaining variance is largely attributed to environmental variability rather than to low single year sample size (Barlow et al. 2009; Forney et al. 2012). The most variable yearly design-based estimates are thus typically for those species with the highest variation in encounter rates (Barlow 2016), and these tend to differ most from the more stable model-based estimates (e.g., common bottlenose dolphin, Table 4).

GAMs are able to effectively deal with spatial heterogeneity of survey coverage within the statistical framework (Hedley and Buckland 2004), and thus the CCES 2018 survey contributed valuable data to the CCE modeling dataset and allowed for population size updates for many of the US West Coast cetacean stocks. Offshore waters were undersampled, however, and SDMs for species that primarily inhabit these regions did not improve with the addition of the CCES 2018 data (i.e., sperm whale, beaked whales). One of the greatest strengths of the SWFSC dataset is the broad, consistent survey coverage of the CCE study area over multiple years, which has supported novel analyses and methodological improvements in SDM development. For example, the SWFSC CCE dataset has supported the evaluation of different modeling approaches, different sampling scales, different interpolation methods, and different sources of habitat data (Barlow et al. 2009; Becker et al. 2010, 2016, 2020; Forney et al. 2012; Redfern et al. 2008). This extensive dataset has also supported studies evaluating the predictive ability of SDMs to provide nowcasts, forecasts, and across-season predictions (Becker et al. 2012, 2014, 2018), as well as allow for robust trend analyses (Moore and Barlow 2011, 2014, 2017). While systematic regional surveys or those that cover only portions of the CCE study area provide valuable data, routine survey coverage of the full study area is required to maintain and increase the utility of this unique dataset.
As additional data are collected on future surveys, model improvements are expected to continue, both from increased sample sizes and ideally from surveys conducted in more anomalous conditions that will allow for an even broader range of habitat conditions to be represented. Model improvements are also expected from the availability of additional habitat variables that are more relevant to the cetaceans than the proxy variables used here. Improvements to ocean model products may in turn produce more robust cetacean SDMs, particularly if the ocean model outputs can be produced at finer spatial resolutions. Continued methodological improvements are also expected, with active research aimed at developing robust methods for combining data from different sources, e.g., visual line-transect, passive acoustics, tagging data, etc. For those species that exhibit substantial distribution shifts in and out of the CCE study area, e.g., striped dolphin, long-beaked common dolphin, and Dall’s porpoise (Becker et al. 2018; Boyd et al. 2018; Carretta et al. 2011), SDMs that incorporate survey data that better sample the broader distribution range of these species should provide greater insight into observed abundance changes within the study area. SDMs that incorporate data from portions of the CCES 2018 survey that covered waters along the west coasts of southern Canada and Baja California will help in this regard, and SDMs for waters off Baja California are currently in development.
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Tables

Table 1. Cetacean and ecosystem assessment surveys and effort conducted within the California Current Ecosystem study area during 1991–2018. CA/OR/WA = California/Oregon/Washington, CenCA = central California, SoCA = southern California, Baja = Baja California. DSJ = David Starr Jordan.

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<td>1617/1619</td>
<td>Jul-Dec 2001</td>
<td>McArthur/DSJ</td>
<td>CA/OR/WA</td>
</tr>
<tr>
<td>1627/1628</td>
<td>June-Dec 2005</td>
<td>McArthur II/DSJ</td>
<td>CA/OR/WA</td>
</tr>
<tr>
<td>1642</td>
<td>Jul-Nov 2008</td>
<td>McArthur II</td>
<td>CA/OR/WA</td>
</tr>
<tr>
<td>1635</td>
<td>Sept-Dec 2009</td>
<td>McArthur II</td>
<td>CenCA/SoCAL/Baja</td>
</tr>
<tr>
<td>1647</td>
<td>Aug-Dec 2014</td>
<td>Ocean Starr*</td>
<td>CA/OR/WA</td>
</tr>
<tr>
<td>2017</td>
<td>June-Dec 2018</td>
<td>Reuben Lasker</td>
<td>Canada/CA/OR/WA/Baja</td>
</tr>
</tbody>
</table>

*Previously the David Starr Jordan

Table 2. Number of sightings and average group size (Avg. GS) of cetacean species observed in the California Current Ecosystem study area during the 1991–2018 shipboard surveys for which habitat-based density models were developed. All sightings were made while on systematic and non-systematic effort in Beaufort sea states ≤5 within the species-specific truncation distances (see text for details).

<table>
<thead>
<tr>
<th>Common name</th>
<th>Taxonomic name</th>
<th>No. of sightings</th>
<th>Avg. GS</th>
</tr>
</thead>
<tbody>
<tr>
<td>Long-beaked common dolphin</td>
<td>Delphinus delphis bairdii</td>
<td>160</td>
<td>291.82</td>
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<tr>
<td>Short-beaked common dolphin</td>
<td>Delphinus delphis delphis</td>
<td>1,034</td>
<td>155.73</td>
</tr>
<tr>
<td>Risso’s dolphin</td>
<td>Grampus griseus</td>
<td>249</td>
<td>18.57</td>
</tr>
<tr>
<td>Pacific white-sided dolphin</td>
<td>Lagenorhynchus obliquidens</td>
<td>296</td>
<td>54.70</td>
</tr>
<tr>
<td>Northern right whale dolphin</td>
<td>Lissodelphis borealis</td>
<td>147</td>
<td>45.31</td>
</tr>
<tr>
<td>Striped dolphin</td>
<td>Stenella covertaolba</td>
<td>153</td>
<td>39.38</td>
</tr>
<tr>
<td>Common bottlenose dolphin</td>
<td>Tursiops truncatus</td>
<td>66</td>
<td>14.48</td>
</tr>
<tr>
<td>Dall’s porpoise</td>
<td>Phocoenoides dalli</td>
<td>678</td>
<td>3.72</td>
</tr>
<tr>
<td>Sperm whale</td>
<td>Physeter macrocephalus</td>
<td>105</td>
<td>6.67</td>
</tr>
<tr>
<td>Minke whale</td>
<td>Balaenoptera acutostrata</td>
<td>49</td>
<td>1.13</td>
</tr>
<tr>
<td>Blue whale</td>
<td>Balaenoptera musculus</td>
<td>316</td>
<td>1.66</td>
</tr>
<tr>
<td>Fin whale</td>
<td>Balaenoptera physalus</td>
<td>558</td>
<td>2.06</td>
</tr>
<tr>
<td>Humpback whale</td>
<td>Megaptera novaeangliae</td>
<td>967</td>
<td>1.70</td>
</tr>
<tr>
<td>Baird’s beaked whale</td>
<td>Berardius bairdii</td>
<td>39</td>
<td>7.46</td>
</tr>
<tr>
<td>Small beaked whale guild</td>
<td>Mesoplodon spp. &amp; Ziphius cavirostris</td>
<td>92</td>
<td>2.12</td>
</tr>
</tbody>
</table>
Table 3. Summary of the final models built with the 1991–2018 survey data. Variables are listed in the order of their significance and are as follows: SST = sea surface temperature, SSTsd = standard deviation of SST, MLD = mixed layer depth, SSH = sea surface height, SSHsd = standard deviation of SSH, depth = bathymetric depth, shelf = distance to shelf, d2000 = distance to the 2,000m isobath, LON = longitude, and LAT = latitude. Separate encounter rate (ER) and group size (GS) models were built for long- and short-beaked common dolphins due to large and variable group sizes. All single response and encounter rate models were corrected for effort with an offset for the effective area searched (see text for details). Performance metrics included the percentage of explained deviance (Exp.Dev.), the area under the receiver operating characteristic curve (AUC), the true skill statistic (TSS), and the ratio of observed to predicted density for the study area (Obs:Pred).

<table>
<thead>
<tr>
<th>Species</th>
<th>Predictor variables</th>
<th>Expl.Dev.</th>
<th>AUC</th>
<th>TSS</th>
<th>Obs:Pred</th>
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</thead>
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<tr>
<td>Long-beaked common dolphin</td>
<td>ER: LON:LAT + SST + SSHsd + SSH</td>
<td>52.50</td>
<td>0.98</td>
<td>0.90</td>
<td>0.95</td>
</tr>
<tr>
<td></td>
<td>GS: LON:LAT</td>
<td>6.55</td>
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<td></td>
<td></td>
</tr>
<tr>
<td>Short-beaked common dolphin</td>
<td>ER: LON:LAT + year + SST + SSH + MLD</td>
<td>17.00</td>
<td>0.77</td>
<td>0.40</td>
<td>0.95</td>
</tr>
<tr>
<td></td>
<td>GS: LON:LAT</td>
<td>11.10</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Risso's dolphin</td>
<td>LON:LAT + SST + MLD + year + SSTsd</td>
<td>22.40</td>
<td>0.76</td>
<td>0.41</td>
<td>0.87</td>
</tr>
<tr>
<td>Pacific white-sided dolphin</td>
<td>LON:LAT + shelf + SST + SSH + MLD</td>
<td>51.70</td>
<td>0.87</td>
<td>0.62</td>
<td>0.86</td>
</tr>
<tr>
<td>Northern right whale dolphin</td>
<td>LON:LAT + SST + depth + MLD + SSTsd</td>
<td>44.40</td>
<td>0.83</td>
<td>0.51</td>
<td>0.92</td>
</tr>
<tr>
<td>Striped dolphin</td>
<td>depth + LON:LAT + SST + year + MLD</td>
<td>33.20</td>
<td>0.76</td>
<td>0.41</td>
<td>0.72</td>
</tr>
<tr>
<td>Common bottlenose dolphin</td>
<td>LON:LAT + MLD + SSTsd + SST + year</td>
<td>51.20</td>
<td>0.92</td>
<td>0.74</td>
<td>0.94</td>
</tr>
<tr>
<td>Dall's porpoise</td>
<td>LON:LAT + SSH + year + SST + SSHsd + SSTsd</td>
<td>32.20</td>
<td>0.89</td>
<td>0.63</td>
<td>0.95</td>
</tr>
<tr>
<td>Sperm whale</td>
<td>d2000 + MLD</td>
<td>13.30</td>
<td>0.61</td>
<td>0.17</td>
<td>0.91</td>
</tr>
<tr>
<td>Minke whale</td>
<td>shelf + SST + LON:LAT</td>
<td>7.73</td>
<td>0.85</td>
<td>0.59</td>
<td>1.00</td>
</tr>
<tr>
<td>Blue whale</td>
<td>LON:LAT + year + SSH + depth + SST + MLD</td>
<td>23.90</td>
<td>0.78</td>
<td>0.42</td>
<td>0.94</td>
</tr>
<tr>
<td>Fin whale</td>
<td>LON:LAT + SST + SSH + year + MLD + depth</td>
<td>22.40</td>
<td>0.75</td>
<td>0.39</td>
<td>0.88</td>
</tr>
<tr>
<td>Humpback whale</td>
<td>LON:LAT + year + depth + SST + MLD</td>
<td>57.40</td>
<td>0.94</td>
<td>0.75</td>
<td>0.98</td>
</tr>
<tr>
<td>Baird's beaked whale</td>
<td>LON:LAT + depth + MLD + SSH</td>
<td>46.00</td>
<td>0.90</td>
<td>0.65</td>
<td>0.96</td>
</tr>
<tr>
<td>Small beaked whale guild</td>
<td>shelf + MLD + SST + LON:LAT</td>
<td>8.19</td>
<td>0.73</td>
<td>0.39</td>
<td>0.97</td>
</tr>
</tbody>
</table>
Table 4. Annual model-predicted mean estimates of abundance, density (animals km\(^{-2}\)), and corresponding coefficient of variation (CV) within the CCE study area. Annual estimates are predicted from the full model using the habitat characteristics in that year. CV\(_m\) (Model) represents the combined uncertainty from three sources: GAM parameters, ESW, and environmental variability. CV\(_{\text{Tot}}\) is the total CV from CV\(_m\) (Model) and CV\(_{g0}\) derived using the Delta method (see text for details). Log-normal 95% confidence intervals (Low and High 95% CIs) apply to abundance estimates. Also shown is the 20\(^{\text{th}}\) percentile for the abundance estimate, corresponding to the “minimum population size (Nmin)” as defined in the Guidelines for Assessing Marine Mammal Stocks, and calculated as the log-normal 20\(^{\text{th}}\) percentile of the mean abundance estimate using standard formulae.

<table>
<thead>
<tr>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
<th></th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Long-beaked common dolphin</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Abundance</td>
<td>57,623</td>
<td>53,044</td>
<td>52,356</td>
<td>58,624</td>
<td>58,794</td>
<td>83,379</td>
</tr>
<tr>
<td>Density</td>
<td>0.0506</td>
<td>0.0465</td>
<td>0.0459</td>
<td>0.0514</td>
<td>0.0516</td>
<td>0.0732</td>
</tr>
<tr>
<td>CV(_m) (Model)</td>
<td>0.151</td>
<td>0.128</td>
<td>0.146</td>
<td>0.087</td>
<td>0.101</td>
<td>0.140</td>
</tr>
<tr>
<td>CV(_{g0})</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
</tr>
<tr>
<td>CV(_{\text{Tot}})</td>
<td>0.224</td>
<td>0.209</td>
<td>0.220</td>
<td>0.187</td>
<td>0.193</td>
<td>0.216</td>
</tr>
<tr>
<td>Low 95% CI</td>
<td>37,370</td>
<td>35,381</td>
<td>34,170</td>
<td>40,799</td>
<td>40,380</td>
<td>54,823</td>
</tr>
<tr>
<td>High 95% CI</td>
<td>88,851</td>
<td>79,524</td>
<td>80,221</td>
<td>84,236</td>
<td>85,605</td>
<td>126,809</td>
</tr>
<tr>
<td>Nmin</td>
<td>47,841</td>
<td>44,574</td>
<td>43,587</td>
<td>50,170</td>
<td>50,031</td>
<td>69,636</td>
</tr>
<tr>
<td><strong>Short-beaked common dolphin</strong></td>
<td></td>
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<td></td>
<td></td>
</tr>
<tr>
<td>Abundance</td>
<td>328,134</td>
<td>391,356</td>
<td>394,610</td>
<td>433,628</td>
<td>880,425</td>
<td>1,056,308</td>
</tr>
<tr>
<td>Density</td>
<td>0.2879</td>
<td>0.3434</td>
<td>0.3462</td>
<td>0.3804</td>
<td>0.7724</td>
<td>0.9267</td>
</tr>
<tr>
<td>CV(_m) (Model)</td>
<td>0.145</td>
<td>0.196</td>
<td>0.139</td>
<td>0.163</td>
<td>0.090</td>
<td>0.125</td>
</tr>
<tr>
<td>CV(_{g0})</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
</tr>
<tr>
<td>CV(_{\text{Tot}})</td>
<td>0.220</td>
<td>0.256</td>
<td>0.216</td>
<td>0.232</td>
<td>0.188</td>
<td>0.207</td>
</tr>
<tr>
<td>Low 95% CI</td>
<td>214,423</td>
<td>238,750</td>
<td>259,781</td>
<td>276,866</td>
<td>611,073</td>
<td>707,020</td>
</tr>
<tr>
<td>High 95% CI</td>
<td>502,146</td>
<td>641,507</td>
<td>599,417</td>
<td>679,148</td>
<td>1,268,504</td>
<td>1,578,155</td>
</tr>
<tr>
<td>Nmin</td>
<td>273,320</td>
<td>316,497</td>
<td>329,739</td>
<td>357,612</td>
<td>752,592</td>
<td>888,971</td>
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<tr>
<td><strong>Risso’s dolphin</strong></td>
<td></td>
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<tr>
<td>Abundance</td>
<td>15,761</td>
<td>15,462</td>
<td>12,044</td>
<td>11,657</td>
<td>8,153</td>
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<td>Density</td>
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<td>0.0106</td>
<td>0.0102</td>
<td>0.0072</td>
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</tr>
<tr>
<td>CV(_m) (Model)</td>
<td>0.116</td>
<td>0.087</td>
<td>0.123</td>
<td>0.128</td>
<td>0.189</td>
<td>0.190</td>
</tr>
<tr>
<td>CV(_{g0})</td>
<td>0.093</td>
<td>0.093</td>
<td>0.093</td>
<td>0.093</td>
<td>0.093</td>
<td>0.093</td>
</tr>
<tr>
<td>CV(_{\text{Tot}})</td>
<td>0.149</td>
<td>0.127</td>
<td>0.154</td>
<td>0.158</td>
<td>0.211</td>
<td>0.212</td>
</tr>
<tr>
<td>Low 95% CI</td>
<td>11,796</td>
<td>12,059</td>
<td>8,918</td>
<td>8,565</td>
<td>5,419</td>
<td>5,957</td>
</tr>
<tr>
<td>High 95% CI</td>
<td>21,060</td>
<td>19,826</td>
<td>16,265</td>
<td>15,865</td>
<td>12,265</td>
<td>13,528</td>
</tr>
<tr>
<td>Nmin</td>
<td>13,916</td>
<td>13,896</td>
<td>10,586</td>
<td>10,211</td>
<td>6,841</td>
<td>7,527</td>
</tr>
<tr>
<td><strong>Pacific white-sided dolphin</strong></td>
<td></td>
<td></td>
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</tr>
<tr>
<td>Abundance</td>
<td>37,147</td>
<td>38,533</td>
<td>39,008</td>
<td>37,369</td>
<td>28,901</td>
<td>34,999</td>
</tr>
<tr>
<td>Density</td>
<td>0.0326</td>
<td>0.0338</td>
<td>0.0342</td>
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<td>0.0254</td>
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<tr>
<td>CV(_m) (Model)</td>
<td>0.230</td>
<td>0.235</td>
<td>0.506</td>
<td>0.323</td>
<td>0.292</td>
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<td>CV(_{g0})</td>
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<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
<td>0.165</td>
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<tr>
<td></td>
<td>Abundance</td>
<td>Density</td>
<td>CV&lt;sub&gt;m&lt;/sub&gt; (Model)</td>
<td>CV&lt;sub&gt;g0&lt;/sub&gt;</td>
<td>CV&lt;sub&gt;Tot&lt;/sub&gt;</td>
<td>Low 95% CI</td>
</tr>
<tr>
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<td>-----------------</td>
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<tr>
<td><strong>Northern right whale dolphin</strong></td>
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<tr>
<td>Abundance</td>
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<td>21,558</td>
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<td>0.0240</td>
<td>0.0158</td>
<td>0.0158</td>
<td>22,194</td>
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<td><strong>Striped dolphin</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
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<tr>
<td>Abundance</td>
<td>17,758</td>
<td>9,481</td>
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<td>0.309</td>
<td>17,758</td>
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<td>0.0263</td>
<td>0.0421</td>
<td>0.0615</td>
<td>0.0332</td>
<td>26,215</td>
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<tr>
<td><strong>Common bottlenose dolphin</strong></td>
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<tr>
<td>Abundance</td>
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<td>9,826</td>
<td>0.504</td>
<td>0.098</td>
<td>0.309</td>
<td>6,198</td>
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<td>Density</td>
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<td>0.0230</td>
<td>0.0537</td>
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<td>0.159</td>
<td>5,408</td>
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<td><strong>Dall's porpoise</strong></td>
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<tr>
<td>Abundance</td>
<td>49,811</td>
<td>17,398</td>
<td>0.244</td>
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<td>0.573</td>
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<td>Density</td>
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<td>0.0319</td>
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<td>0.0186</td>
<td>44,418</td>
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<tr>
<td><strong>Sperm whale</strong></td>
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<tr>
<td>Abundance</td>
<td>2,783</td>
<td>2,208</td>
<td>0.518</td>
<td>0.518</td>
<td>0.518</td>
<td>2,783</td>
</tr>
<tr>
<td>Density</td>
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<td>0.0076</td>
<td>0.0319</td>
<td>0.0319</td>
<td>0.0319</td>
<td>2,896</td>
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</table>

26
<table>
<thead>
<tr>
<th></th>
<th>Density</th>
<th>Abund</th>
<th>Nmin</th>
<th>High 95% CI</th>
<th>Low 95% CI</th>
<th>CV</th>
</tr>
</thead>
<tbody>
<tr>
<td><strong>Baird's beaked whale</strong></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>CV\textsubscript{g0}</td>
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<td>0.285</td>
<td>0.285</td>
<td>0.285</td>
<td>0.285</td>
<td></td>
</tr>
<tr>
<td>CV\textsubscript{Tot}</td>
<td>0.295</td>
<td>0.305</td>
<td>0.306</td>
<td>0.299</td>
<td>0.340</td>
<td>0.315</td>
</tr>
<tr>
<td>Low 95% CI</td>
<td>1,578</td>
<td>1,614</td>
<td>1,497</td>
<td>1,617</td>
<td>1,388</td>
<td>1,425</td>
</tr>
<tr>
<td>High 95% CI</td>
<td>4,907</td>
<td>5,197</td>
<td>4,836</td>
<td>5,090</td>
<td>5,082</td>
<td>4,765</td>
</tr>
<tr>
<td>Nmin</td>
<td>2,181</td>
<td>2,253</td>
<td>2,092</td>
<td>2,243</td>
<td>2,010</td>
<td>2,011</td>
</tr>
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</table>

| **Minke whale**    |         |       |      |             |            |     |
| Abundance          | 847     | 812   | 819  | 804         | 1,062      | 915 |
| Density            | 0.0007  | 0.0007| 0.0007| 0.0007       | 0.0009     | 0.0008|
| CV\textsubscript{m} (Model) | 0.139 | 0.110 | 0.110| 0.113       | 0.109      | 0.085|
| CV\textsubscript{g0}  | 0.787   | 0.787 | 0.787| 0.787       | 0.787      | 0.787|
| CV\textsubscript{Tot} | 0.799   | 0.795 | 0.795| 0.795       | 0.795      | 0.792|
| Low 95% CI         | 214     | 206   | 208  | 204         | 270        | 233 |
| High 95% CI        | 3,358   | 3,200 | 3,227| 3,170       | 4,184      | 3,590|
| Nmin               | 469     | 451   | 454  | 446         | 589        | 509 |

| **Blue whale**     |         |       |      |             |            |     |
| Abundance          | 1,946   | 1,657 | 1,042| 919         | 1,077      | 670 |
| Density            | 0.0017  | 0.0015| 0.0009| 0.0008       | 0.0009     | 0.0006|
| CV\textsubscript{m} (Model) | 0.224 | 0.139 | 0.149| 0.227       | 0.273      | 0.299|
| CV\textsubscript{g0}  | 0.309   | 0.309 | 0.309| 0.309       | 0.309      | 0.309|
| CV\textsubscript{Tot} | 0.382   | 0.339 | 0.343| 0.383       | 0.412      | 0.430|
| Low 95% CI         | 945     | 868   | 542  | 445         | 495        | 299 |
| High 95% CI        | 4,009   | 3,162 | 3,004| 1,899       | 3,542      | 1,502|
| Nmin               | 1,427   | 1,255 | 787  | 673         | 771        | 474 |

| **Fin whale**      |         |       |      |             |            |     |
| Abundance          | 3,804   | 5,733 | 7,319| 7,606       | 10,139     | 4,065|
| Density            | 0.0033  | 0.0050| 0.0064| 0.0067       | 0.0089     | 0.0097|
| CV\textsubscript{m} (Model) | 0.200 | 0.212 | 0.250| 0.303       | 0.175      | 0.333|
| CV\textsubscript{g0}  | 0.230   | 0.230 | 0.230| 0.230       | 0.230      | 0.230|
| CV\textsubscript{Tot} | 0.305   | 0.313 | 0.340| 0.381       | 0.289      | 0.405|
| Low 95% CI         | 2,120   | 3,149 | 3,828| 3,699       | 5,817      | 5,156|
| High 95% CI        | 6,826   | 10,439| 13,994| 15,640      | 17,672     | 23,747|
| Nmin               | 2,959   | 4,432 | 5,540| 5,580       | 7,986      | 7,970|

| **Humpback whale** |         |       |      |             |            |     |
| Abundance          | 1,181   | 1,364 | 1,575| 1,727       | 2,178      | 4,784|
| Density            | 0.0010  | 0.0012| 0.0014| 0.0015       | 0.0019     | 0.0042|
| CV\textsubscript{m} (Model) | 0.147 | 0.081 | 0.113| 0.175       | 0.271      | 0.118|
| CV\textsubscript{g0}  | 0.283   | 0.283 | 0.283| 0.283       | 0.283      | 0.283|
| CV\textsubscript{Tot} | 0.319   | 0.294 | 0.305| 0.333       | 0.392      | 0.307|
| Low 95% CI         | 642     | 775   | 878  | 915         | 1,038      | 2,658|
| High 95% CI        | 2,173   | 2,400 | 2,824| 3,259       | 4,568      | 8,609|
| Nmin               | 909     | 1,070 | 1,226| 1,315       | 1,584      | 3,717|

<p>| <strong>Baird's beaked whale</strong> |         |       |      |             |            |     |
| Abundance          | 739     | 730   | 590  | 681         | 977        | 1,363|
| Density            | 0.0006  | 0.0006| 0.0005| 0.0006       | 0.0009     | 0.0012|</p>
<table>
<thead>
<tr>
<th></th>
<th>CV&lt;sub&gt;m&lt;/sub&gt; (Model)</th>
<th>CV&lt;sub&gt;g0&lt;/sub&gt;</th>
<th>CV&lt;sub&gt;Tot&lt;/sub&gt;</th>
<th>Low 95% CI</th>
<th>High 95% CI</th>
<th>Nmin</th>
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<td>0.628</td>
<td>0.521</td>
<td>0.423</td>
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<tr>
<td></td>
<td>0.562</td>
<td>0.543</td>
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<td>265</td>
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<td>169</td>
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<td>366</td>
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<tr>
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<td>2,064</td>
<td>1,976</td>
<td>2,057</td>
<td>2,065</td>
<td>2,608</td>
<td>3,634</td>
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<tr>
<td></td>
<td>475</td>
<td>476</td>
<td>345</td>
<td>423</td>
<td>641</td>
<td>894</td>
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**Small beaked whale guild**

<table>
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<tr>
<th></th>
<th>Abundance</th>
<th>Density</th>
<th>CV&lt;sub&gt;m&lt;/sub&gt; (Model)</th>
<th>CV&lt;sub&gt;g0&lt;/sub&gt;</th>
<th>CV&lt;sub&gt;Tot&lt;/sub&gt;</th>
<th>Low 95% CI</th>
<th>High 95% CI</th>
<th>Nmin</th>
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<td>0.153</td>
<td>0.438</td>
<td>0.464</td>
<td>2,096</td>
<td>11,830</td>
<td>3,433</td>
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<td>0.113</td>
<td>0.438</td>
<td>0.452</td>
<td>2,447</td>
<td>13,281</td>
<td>3,964</td>
</tr>
<tr>
<td></td>
<td>4,399</td>
<td>0.0039</td>
<td>0.213</td>
<td>0.438</td>
<td>0.487</td>
<td>1,781</td>
<td>10,866</td>
<td>2,983</td>
</tr>
<tr>
<td></td>
<td>5,088</td>
<td>0.0045</td>
<td>0.201</td>
<td>0.438</td>
<td>0.482</td>
<td>2,078</td>
<td>12,461</td>
<td>3,463</td>
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<tr>
<td></td>
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<td>0.438</td>
<td>0.477</td>
<td>1,924</td>
<td>11,336</td>
<td>3,191</td>
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<tr>
<td></td>
<td>4,989</td>
<td>0.0044</td>
<td>0.211</td>
<td>0.438</td>
<td>0.486</td>
<td>2,023</td>
<td>12,306</td>
<td>3,385</td>
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</table>
Table 5. Arithmetic mean of the model-predicted 2014 and 2018 estimates of abundance and density (animals km\(^{-2}\)) within the CCE study area. The corresponding coefficient of variation (CV\(_{\text{Tot}}\)) is the total CV from four sources: environmental variability, GAM parameters, ESW, and \(g(0)\) (see text for details). Log-normal 95% confidence intervals (Low and High 95% CIs) apply to abundance estimates. Also shown is the 20\(^{\text{th}}\) percentile for the abundance estimate, corresponding to the “minimum population size (Nmin)” as defined in the Guidelines for Assessing Marine Mammal Stocks, and calculated as the log-normal 20\(^{\text{th}}\) percentile of the mean abundance estimate using standard formulae.

<table>
<thead>
<tr>
<th>Species</th>
<th>Abundance</th>
<th>Density</th>
<th>CV(_{\text{Tot}})</th>
<th>Low 95% CI</th>
<th>High 95% CI</th>
<th>Nmin</th>
</tr>
</thead>
<tbody>
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<td>Long-beaked common dolphin</td>
<td>71,087</td>
<td>0.0624</td>
<td>0.190</td>
<td>49,156</td>
<td>102,803</td>
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<td>Short-beaked common dolphin</td>
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<td>0.8496</td>
<td>0.192</td>
<td>667,050</td>
<td>1,405,792</td>
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<tr>
<td>Risso's dolphin</td>
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<td>0.0075</td>
<td>0.209</td>
<td>5,713</td>
<td>12,841</td>
<td>7,197</td>
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<tr>
<td>Pacific white-sided dolphin</td>
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<td>0.0280</td>
<td>0.249</td>
<td>19,769</td>
<td>51,636</td>
<td>25,996</td>
</tr>
<tr>
<td>Northern right whale dolphin</td>
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<td>0.612</td>
<td>7,836</td>
<td>71,428</td>
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<tr>
<td>Striped dolphin</td>
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<td>0.314</td>
<td>27,454</td>
<td>91,237</td>
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</tr>
<tr>
<td>Common bottlenose dolphin</td>
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<td>10,117</td>
<td>3,374</td>
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<tr>
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<td>Sperm whale</td>
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<td>0.324</td>
<td>1,415</td>
<td>4,891</td>
<td>2,016</td>
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<tr>
<td>Minke whale</td>
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<td>0.0009</td>
<td>0.793</td>
<td>251</td>
<td>3,874</td>
<td>548</td>
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<tr>
<td>Blue whale</td>
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<td>0.0008</td>
<td>0.396</td>
<td>414</td>
<td>1,845</td>
<td>634</td>
</tr>
<tr>
<td>Fin whale</td>
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<td>0.328</td>
<td>5,670</td>
<td>19,824</td>
<td>8,103</td>
</tr>
<tr>
<td>Humpback whale</td>
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<td>0.320</td>
<td>1,888</td>
<td>6,417</td>
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</tr>
<tr>
<td>Baird’s beaked whale</td>
<td>1,170</td>
<td>0.0010</td>
<td>0.501</td>
<td>463</td>
<td>2,956</td>
<td>786</td>
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<tr>
<td>Small beaked whale guild</td>
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<td>0.0042</td>
<td>0.481</td>
<td>1,976</td>
<td>11,804</td>
<td>3,290</td>
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</table>
Figure 1. Completed transects for the Southwest Fisheries Science Center systematic ship surveys conducted between 1991 and 2018 in the California Current Ecosystem study area. The lines (green = 1991–2014 surveys, red=2018 survey) show on-effort transect coverage in Beaufort sea states of 0-5.
Figure 2a-b. Predicted mean density (animals km\(^2\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for (a) long-beaked common dolphin, and (b) short-beaked common dolphin. Panels show the multi-year average density based on predicted daily cetacean species densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the SWFSC 1996-2018 summer/fall ship surveys for the respective species.
Figure 2c-d. Predicted mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for (c) Risso’s dolphin, and (d) Pacific white-sided dolphin. Panels show the multi-year average density based on predicted daily cetacean species densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the SWFSC 1996-2018 summer/fall ship surveys for the respective species.
Figure 2e-f. Predicted mean density (animals km\(^{-2}\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for (e) northern right whale dolphin, and (f) striped dolphin. Panels show the multi-year average density based on predicted daily cetacean species densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the SWFSC 1996-2018 summer/fall ship surveys for the respective species.
Figure 2g-h. Predicted mean density (animals km\(^{-2}\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for (g) common bottlenose dolphin, and (h) Dall’s porpoise. Panels show the multi-year average density based on predicted daily cetacean species densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the SWFSC 1996-2018 summer/fall ship surveys for the respective species.
Figure 2i-j. Predicted mean density (animals km\(^{-2}\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for (i) sperm whale, and (j) minke whale. Panels show the multi-year average density based on predicted daily cetacean species densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the SWFSC 1996-2018 summer/fall ship surveys for the respective species.
Figure 2k-l. Predicted mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for (k) blue whale, and (l) fin whale. Panels show the multi-year average density based on predicted daily cetacean species densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the SWFSC 1996-2018 summer/fall ship surveys for the respective species.
Figure 2m-n. Predicted mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for (m) humpback whale, and (n) Baird’s beaked whale. Panels show the multi-year average density based on predicted daily cetacean species densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the SWFSC 1996-2018 summer/fall ship surveys for the respective species.
Figure 2o. Predicted mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for (o) small beaked whale guild. Panels show the multi-year average density based on predicted daily cetacean species densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the SWFSC 1996-2018 summer/fall ship surveys for the respective species.
Figure 3a. Predicted annual (1996-2018) mean density (animals km\(^{-2}\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for long-beaked common dolphin. Panels show the yearly average density based on predicted daily long-beaked common dolphin densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3b. Predicted annual (1996-2018) mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for short-beaked common dolphin. Panels show the yearly average density based on predicted daily short-beaked common dolphin densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3c. Predicted annual (1996-2018) mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for Risso’s dolphin. Panels show the yearly average density based on predicted daily Risso’s dolphin densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3d. Predicted annual (1996-2018) mean density (animals km\(^{-2}\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for Pacific white-sided dolphin. Panels show the yearly average density based on predicted daily Pacific white-sided dolphin densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3e. Predicted annual (1996-2018) mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for northern right whale dolphin. Panels show the yearly average density based on predicted daily northern right whale dolphin densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3f. Predicted annual (1996-2018) mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for striped dolphin. Panels show the yearly average density based on predicted daily striped dolphin densities covering the 1996–2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3g. Predicted annual (1996-2018) mean density (animals km\(^{-2}\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for common bottlenose dolphin. Panels show the yearly average density based on predicted daily common bottlenose dolphin densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3h. Predicted annual (1996-2018) mean density (animals km\(^{-2}\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for Dall’s porpoise. Panels show the yearly average density based on predicted daily Dall’s porpoise densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3i. Predicted annual (1996-2018) mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for sperm whale. Panels show the yearly average density based on predicted daily sperm whale densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3j. Predicted annual (1996-2018) mean density (animals km\(^{-2}\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for minke whale. Panels show the yearly average density based on predicted daily minke whale densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
(k) Blue whale

Figure 3k. Predicted annual (1996-2018) mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for blue whale. Panels show the yearly average density based on predicted blue whale densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3l. Predicted annual (1996-2018) mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for fin whale. Panels show the yearly average density based on predicted fin whale densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^{2}$). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3m. Predicted annual (1996-2018) mean density (animals km$^{-2}$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for humpback whale. Panels show the yearly average density based on predicted humpback whale densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3n. Predicted annual (1996-2018) mean density (animals km\(^{-2}\)) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for Baird’s beaked whale. Panels show the yearly average density based on predicted Baird’s beaked whale densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km\(^2\)). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Figure 3o. Predicted annual (1996-2018) mean density (animals km$^2$) and associated coefficients of variation (CV) from the 1991–2018 habitat-based density models for the small beaked whale guild (Mesoplodonts and Cuvier’s beaked whale). Panels show the yearly average density based on predicted small beaked whale guild densities covering the 1996-2018 survey periods (summer/fall). Predictions are shown for the study area (1,141,800 km$^2$). White dots in the average plots show actual sighting locations from the respective SWFSC summer/fall ship surveys.
Appendix A: SDM functional plots

Final SDM response curves for (1) long-beaked common dolphin, (2) short-beaked common dolphin, (3) Risso’s dolphin, (4) Pacific white-sided dolphin, (5) northern right whale dolphin, (6) striped dolphin, (7) common bottlenose dolphin, (8) sperm whale, (9) minke whale, (10) blue whale, (11) fin whale, (12) humpback whale, (13) Baird’s beaked whale, and (14) the small beaked whale guild (*Mesoplodon* spp. and Cuvier’s beaked whale). The suite of environmental and geographic covariates included: SST = sea surface temperature, sdSST = standard deviation of SST, MLD = mixed layer depth, SSH = sea surface height, sdSSH = standard deviation of SSH, depth = bathymetric depth, dShelf = distance to the 200m isobath, d2000 = distance to the 2,000m isobath, mlat = latitude, mlon = longitude, and yearCoVar = year. Models were constructed with both linear terms and smoothing splines. Degrees of freedom for single variables are shown in the parentheses on the y-axis. Variables for the interaction terms are shown on the x- and y-axes. For single variables the y-axes represent the term’s (linear or spline) function. Zero on the y-axes corresponds to no effect of the predictor variable on the estimated response variable. Scaling of y-axis varies among predictor variables to emphasize model fit. The shading reflects 2x standard error bands (i.e., 95% confidence interval); tick marks (‘rug plot’) above the X axis show data values. For the interaction terms, yellow indicates higher prediction densities and red lower predicted densities.
Long-beaked common dolphin

Figure A 1. Functional plot for long-beaked common dolphin (*Delphinus delphis bairdii*) encounter rate model.
Figure A 2. Functional plot for short-beaked common dolphin (*Delphinus delphis delphis*) encounter rate model.
Figure A 3. Functional plot for Risso’s dolphin (*Grampus griseus*) model.
Figure A 4. Functional plot for Pacific white-sided dolphin (*Lagenorhynchus obliquidens*) model.
Figure A 5. Functional plot for northern right whale dolphin (*Lissodelphis borealis*) model.
Figure A 6. Functional plot for striped dolphin (*Stenella coeruleoalba*) model.
Figure A 7. Functional plot for common bottlenose dolphin (*Tursiops truncatus*) model.
Dall’s porpoise

Figure A 8. Functional plot for Dall’s porpoise (*Phocoenoides dalli*) model.
Sperm whale

Figure A 9. Functional plot for sperm whale (*Physeter macrocephalus*) model.
Minke whale

Figure A 10. Functional plot for minke whale (*Balaenoptera acutorostrata*) model.
Blue whale

Figure A 11. Functional plot for blue whale (*Balaenoptera musculus*) model.
Figure A 12. Functional plot for fin whale (*Balaenoptera physalus*) model.
Figure A 13. Functional plot for fin whale (*Megaptera novaeangliae*) model.
Figure A 14. Functional plot for Baird’s beaked whale (*Berardius bairdii*) model.
Small beaked whale guild

Figure A 15. Functional plot for the small beaked whale guild (*Mesoplodon* spp. & *Ziphius cavirostris*) model.