



Weekly predictions of North Atlantic right whale *Eubalaena glacialis* habitat reveal influence of prey abundance and seasonality of habitat preferences

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ABSTRACT: Primary sources of mortality and serious injury to endangered North Atlantic right whales *Eubalaena glacialis* are vessel strikes and entanglement in fishing gear. All management plans depend on knowing when and where right whales are likely to be present. We tested the feasibility of a system designed to predict potential right whale habitat on a weekly time scale. The system paired right whale occurrence records with a collection of data layers including: results from a coupled biological–physical model of *Calanus finmarchicus* (the primary prey of right whales), satellite-derived sea surface temperature and chlorophyll, and bathymetry. Using these data, we trained seasonal habitat models and projected them onto environmental data for each 8 d period from January to June, 2002 to 2006. Two hypotheses were tested: (1) that right whale environmental preferences change from season to season and (2) that modeled prey concentration is an important predictor of the distribution of right whales. To test H_1 , we trained, tested, and compared models for 3 time periods: winter, spring, and winter and spring combined. To test H_2 , we trained and tested models with and without *C. finmarchicus*. Predictions of habitat suitability were highly dynamic within and across years. Our results support the hypothesis that right whale environmental preferences change between winter and spring. The inclusion of modeled *C. finmarchicus* abundance improved the accuracy of habitat suitability predictions.

KEY WORDS: Right whale · *Eubalaena glacialis* · *Calanus finmarchicus* · Species distribution model · Gulf of Maine · Transferability

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INTRODUCTION

North Atlantic right whales *Eubalaena glacialis* are endangered. Despite protective measures, vessel collisions with whales and entanglement in fishing gear continue to pose significant risk to these animals

(Knowlton & Kraus 2001, Kraus et al. 2005). This high level of threat is due, in part, to overlap of right whale habitat with commercially important shipping and fishing areas. Thus, there is an economic incentive to resist measures, especially blanket restrictions, designed to protect right whales from human en-

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counters. There is an ongoing effort to find solutions that will help to reduce risk to right whales without placing undue restrictions on shipping and fishing activities (Myers et al. 2007). Temporally and spatially dynamic protective measures could ease some of the tension between commercial and conservation interests.

We know where to find right whales within large spatial and temporal windows (Winn et al. 1986). During winter, females can be found giving birth in the coastal waters of the southeast USA (Kraus et al. 2007). From winter to mid-spring, a portion of the population can be found in Cape Cod Bay (Fig. 1) off the northeast US coast (Mayo et al. 2004). Beginning in mid-spring and extending into the summer, right whales can be found in the Great South Channel (CETAP 1982, Kenney & Wishner 1995). In the late summer and fall, right whales can be found in the Bay of Fundy and Roseway Basin (Brown et al. 2009). With few exceptions (e.g. Kenney 2001, Patrician & Kenney 2010), these patterns are predictable on seasonal timescales and at regional spatial scales. Protective measures such as Seasonal Area Management zones (NOAA 2002), the mandatory ship reporting system (Ward-Geiger et al. 2005), vessel speed restrictions (NOAA 2008), and the boundaries of the USA (NOAA 1994) and Canadian (Brown et al. 2009) Critical Habitats have been based upon these known distributional patterns.

A leading hypothesis to explain the distribution of right whales in the Gulf of Maine is that whales move to areas with high concentrations of prey, relative to

nearby regions. The aggregation of right whales in regions rich in late-stage *Calanus finmarchicus* (hereafter *Calanus*) has been well documented (Wishner et al. 1988, 1995, Murison & Gaskin 1989, Baumgartner et al. 2003a). Sighting records indicate that individual right whales utilize both Cape Cod Bay and the Great South Channel in the same year (NARWC 2011a, P. Hamilton pers. comm.). Pendleton et al. (2009) examined abundance of right whales with respect to 2 groups of prey: (1) *Calanus* and (2) *Pseudocalanus* spp. and *Centropages typicus*. The authors found that regional-scale mean concentration of prey is a statistically significant predictor of the relative abundance of right whales. Right whales in Cape Cod Bay appeared to respond more strongly to the concentration of *Pseudocalanus* spp. and *C. typicus* than to *Calanus*. However, in the Great South Channel, right whale abundance was more strongly correlated with *Calanus* than with other prey taxa. This suggests that (1) the environmental preferences of right whales are dynamic and change on seasonal time scales, and (2) the distribution and availability of prey are important factors in determining the distribution of right whales.

Pershing et al. (2009a) went on to build and ground-truth a model of *Calanus* abundance throughout the Gulf of Maine, and Pershing et al. (2009b) found a statistically significant relationship between modeled concentration of *Calanus* and the arrival date of right whales in the Great South Channel region. Thus, modeled concentration of *Calanus* was found to be a good predictor of right whale abundance at regional spatial scales. Finer-resolution estimates of where and when right whales are likely to occur, within and between years, are needed so that government managers can issue alerts to mariners regarding the likelihood of right whale occurrence.

For this study, we modeled the distribution of right whale habitat on small space (1 km) and short time (weekly) scales. Our objective was to build a model that could identify potential right whale habitat on a weekly time

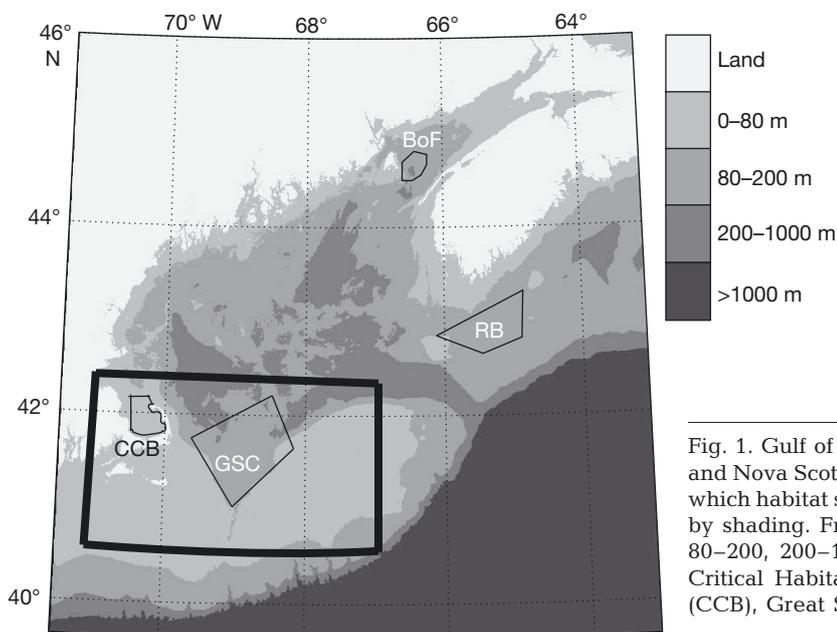


Fig. 1. Gulf of Maine region, with Cape Cod Bay at lower left and Nova Scotia at upper right. Bold rectangle is the region for which habitat suitability was estimated. Bathymetry is depicted by shading. From light to dark, shades represent land, 0–80, 80–200, 200–1000 and >1000 m. Gulf of Maine right whale Critical Habitats are identified by polygons: Cape Cod Bay (CCB), Great South Channel (GSC), Bay of Fundy (BoF), and Roseway Basin (RB)

scale, and that was sensitive to intra- and interannual variability in environmental conditions. Within that purview, we tested 2 hypotheses: H_1 : that right whale environmental preferences differ between seasons, and H_2 : that prey is an important predictor of the spatial and temporal distribution of right whales.

MATERIALS AND METHODS

Study region and species occurrence records

Habitat suitability was modeled in 2 important right whale Critical Habitats: Cape Cod Bay and the Great South Channel (Fig. 1). Right whale occurrence records were collected during right whale aerial surveys conducted by the Provincetown Center for Coastal Studies (Mayo et al. 2004) and the US National Marine Fisheries Service (Cole et al. 2007), and were obtained from the North Atlantic Right Whale Consortium database. Our study used occurrence records for the years 2002 to 2006. Records from opportunistic and directed surveys to known aggregations of right whales were excluded from our analysis. Only records from randomized or complete trackline surveys were used (Brown et al. 2007, Cole et al. 2007). All occurrence records and environmental data came from the region spanning 40.5° to 42.5° N latitude and 71.0° to 67.0° W longitude. In response to the vertical distribution of the prey, right whales often go on extended (8 to 10 min) feeding dives and spend a considerable amount of time underwater (Goodyear 1996). Visual surveys only detect whales at or near the surface; therefore, true absence cannot be inferred from visual absence. For this reason, right whale occurrences were treated as a presence-only dataset.

Environmental data layers

Three dynamic predictor variables and 1 static predictor variable were used to model right whale habitat suitability. Dynamic predictor variables change every 8 d, while static variables do not change. Dynamic predictor variables included 8 d mean sea surface temperature (SST), chlorophyll, and modeled *Calanus* abundance. The static predictor variable was a TOPEX-derived bathymetric grid, linearly interpolated to 1 km resolution, indicating water depth (Smith & Sandwell 1997).

SST measurements for the years 2002 and 2003 were obtained from the Advanced Very High Resolu-

tion Radiometer and were available as Level 3 coverages (4 km resolution) from the NOAA-NODC Pathfinder V5 Project. These data were downsampled to 1 km resolution to match the resolution of the majority of our satellite data. SST measurements for the years 2004 to 2006 came from the Moderate Resolution Imaging Spectrometer (MODIS) instrument on the Aqua satellite. Chlorophyll measurements were obtained from the Sea-viewing Wide Field-of-view Sensor (SeaWiFS) for the year 2002. In order to avoid periodic gaps coverage caused by SeaWiFS outages, chlorophyll data for 2003 to 2006 were obtained from the MODIS instrument on the Aqua satellite, which came online in mid-2002. SeaWiFS and MODIS-Aqua chlorophyll data sources are comparable (Zhang et al. 2006). SST and chlorophyll data from MODIS-Aqua and SeaWiFS were downloaded as Level 2 coverages (1 km resolution) from the Ocean Color Web (<http://oceancolor.gsfc.nasa.gov>). Daily satellite imagery was processed into 8 d means using a simple arithmetic mean, and was then interpolated to eliminate pixels with missing data, which were a consequence of cloud cover. We used the interpolation procedure described by Pershing et al. (2009a). All chlorophyll data were $\log_{10}(x + 1)$ transformed. Values of predictor variables at each pixel were georeferenced, and the collection of values for each environmental variable over the model domain is referred to as an environmental data layer.

The modeled *Calanus* abundance estimates used in our study were described in detail by Pershing et al. (2009a). Briefly, the model couples a *Calanus* life-history model to output from an ocean circulation model. For this study, we used climatological circulation fields from Naimie (1996). These fields contain realistic seasonal changes in circulation, but do not vary from year to year. The model was initialized with a climatological spatial *Calanus* distribution from January. The abundance of *Calanus* in January was adjusted in each year using data from the Gulf of Maine continuous plankton recorder survey (Jossi & Goulet 1993). The dynamical evolution of the *Calanus* abundance field was then determined by satellite estimates of SST, which determines development rate (Campbell et al. 2001), and chlorophyll, which determines egg production (Runge & Plourde 1996, Durbin et al. 2003). The model was run from 1 January through 31 May. After 1 June, many *Calanus* exhibit diel vertical migration, although this is highly variable (Durbin et al. 1995). Furthermore, many subadults (C5s) enter a state of reduced activity known as diapause, typically below 150 m. Both the daily and seasonal vertical migrations make it diffi-

cult for our model, which is forced mainly by surface data, to reproduce the dynamics of this population during the summer. Overall, the model does a good job capturing the seasonal development of the *Calanus* population in the Gulf of Maine and reproducing the interannual variability in right whale prey (Pershing et al. 2009a).

Although SST and chlorophyll data were used to force the *Calanus* model, all 3 products were used as predictors in our model and were treated as independent variables for the following reasons. Due to the long generation time of *Calanus* (30 to 100 d, depending on temperature), there is a highly variable and nonlinear time lag between the environmental variables and the modeled *Calanus* distribution. Temperature and chlorophyll act differently on each life stage of *Calanus*. Therefore, modeled *Calanus* abundance is not linearly related to SST or chlorophyll. The relationship between SST and chlorophyll inputs and *Calanus* output is further complicated because the *Calanus* life-history model is embedded in a circulation model. Thus, the modeled *Calanus* abundance at a particular location is a non-linear integration of past SST and chlorophyll conditions from a range of upstream locations.

Modeling algorithm

We used the maximum entropy method, or Maxent, to model the relationship between right whale occurrence and environmental covariates. Entropy is a measure of uniformity. The goal of the Maxent algorithm, applied in the context of species distribution modeling, is to produce an estimate of the unknown distribution describing the relationship between the species occurrences and a set of measured environmental predictor variables (i.e. covariates). The estimated distribution should maximize entropy subject to a set of constraints representing what is known (Phillips et al. 2006), or minimize relative entropy (Elith et al. 2011). Here we provide a conceptual overview of the algorithm, following Elith et al. (2011), but refer readers to additional publications for detailed explanations (Phillips et al. 2004, 2006, Phillips & Dudik 2008).

The core of the Maxent algorithm is an estimate of the ratio of the conditional density of the covariates at the occurrence locations to the unconditional density of covariates across the study area as measured by a random sample of the background. Maxent chooses the conditional density such that the distance between it and the unconditional density is mini-

mized, subject to a set of constraints, e.g. that the mean temperature in the conditional density be close to the mean temperature across occurrence locations. The resultant Maxent model has an exponential form. Elith et al. (2011) provided a detailed yet accessible description of the Maxent algorithm. We implemented the algorithm using the Maximum Entropy Species Distribution Modeling Software v3.3.1g using default options and the logistic output. The logistic output format provides habitat suitability values that can be interpreted as the probability of species presence, conditioned on the environmental variables (Phillips & Dudik 2008).

In many applications of Maxent there is 1 data layer for each environmental variable. In such cases, background data (Hirzel et al. 2002) consist of a random sample of points from the study area, with associated values of all the environmental variables. In our study there was 1 data layer for each environmental variable for each 8 d period. Therefore, background data for each model that we trained was taken from 10 000 randomly chosen location–time pairs (latitude, longitude, and time measured in 8 d periods) from the model training dataset (Phillips & Dudik 2008). The quantity 10 000 has been experimentally determined to be a sufficient number of background samples to ensure that the area under the receiver operator characteristic (ROC) curve (AUC; see ‘Model Evaluation’ section below) no longer changes as the number of background samples is increased (Phillips & Dudik 2008), and it is the default number used in Maxent software. Each background sample has the values of each environmental variable at that location and time, as do the right whale presence samples. Background data therefore represent a sample of conditions across space and time that the whales were choosing from. If no whales were sighted during a particular 8 d period, then environmental data from that period were excluded from the data available to be sampled to generate the background dataset.

Experimental design

Environmental and species occurrence data (spanning 1 January to 1 June for the years 2002 to 2006) was temporally partitioned into 5 subsets to facilitate a cross validation procedure in which 4 yr of data were used to train a model and the fifth year (the test year) was used to test the model. For example, the model trained with data from the years 2003 to 2006 was projected onto environmental data from all 8 d

periods in 2002, yielding 1 habitat suitability map for each 8 d period in 2002. Each right whale occurrence from 2002 was then associated with the prediction of habitat suitability for the time (8 d period) and location of the occurrence. Then, performance of the model in the test year was measured (see 'Model Evaluation' section below).

To address H_1 , that right whale habitat preferences are dynamic, we conducted 3 experiments. In the first experiment, we trained and tested models with data from winter (1 January to 21 March), in the second experiment we used data from spring (22 March to 1 June), and in the third experiment we used data from winter–spring (1 January to 1 June). The yearly cross validation scheme was used for each of these experiments, yielding 5 models (1 for each test year) for each experiment. A difference in the relative influence of predictors in winter versus spring experiments would suggest that the habitat preference of right whales changes on a seasonal basis.

To address H_2 , that prey is an important predictor of right whale habitat preferences, we compared the predictive accuracy of models trained with and without modeled *Calanus* for each season outlined above. This doubled the number of experiments (and therefore the number of models). A subscript C, e.g. winter–spring_C, was used to label experiments that included the modeled *Calanus* predictor variable. Statistical significance was determined by estimating confidence intervals using the delete-d jackknife procedure described below. Non-overlapping confidence intervals indicated a statistically significant difference.

Model evaluation

AUC is a standard performance metric for presence-only species distribution models, and for many other classifiers (Hanley & McNeil 1982, Swets 1988, Fawcett 2006). Although AUC is sensitive to sample size and other aspects of model structure (Hernandez et al. 2006, Wisz et al. 2008), it is an important metric for measuring model performance. We calculated 1 AUC score for each seasonal model in each test year, for a total of 30 scores. The ROC curve is a plot of the true positive classification rate (sensitivity) versus the false positive classification rate (1-specificity). The total area of the plot is equal to 1. A ROC curve that is a diagonal line from the lower left to the upper right corner is the theoretical ROC curve for a random model and produces an AUC = 0.5. A higher AUC score indicates better predictive accuracy of the

model. For example, an AUC of 0.8 indicates that there is an 80% chance that the predicted habitat suitability for a randomly drawn species presence will be higher than that of a randomly drawn absence (Fawcett 2006). In a presence-only model there are no species absences, so computing a false positive rate is not possible. Rather than distinguish presence from absence, we distinguished presence from random or background data, also known as pseudo-absence data (Ferrier et al. 2002, Phillips et al. 2006, Phillips & Dudik 2008).

Commonly, a single model is applied to a single set of data layers, and a single AUC score is calculated. In our study, we applied a single seasonal model to many sets of environmental data layers, generating 10, 9, and 19 habitat suitability maps for experiments from the winter, spring, and winter–spring periods, respectively. To deal with this, we concatenated habitat suitability maps from all 8 d periods within each experiment and test year and calculated 1 AUC score. For example, we calculated 1 AUC score for winter 2002. To accomplish this, all habitat suitability maps for 8 d periods in test year 2002, during which right whales were observed, were layered to create 1 array. Time periods with no right whale sightings were excluded to bring the bias of the background data into closer agreement with the bias of the aerial survey data (Phillips et al. 2009). From the layered array, 10 000 coordinate locations, indexed by i , were chosen uniformly at random with replacement. The value of habitat suitability, $S(i) \in [0, 1]$, at each of those locations was stored, and these $S(i)$ were the pseudo-absence background data. For a series of threshold values, $t \in [0, 1]$, $t = 0, 0.05, 0.01, \dots, 1$, 1-specificity was calculated as the percentage of $S(i) \geq t$. The true positive classification rate (TPR or sensitivity) was calculated as the fraction of right whales occurring at locations where the habitat suitability value was $\geq t$. TPR was then plotted as a function of 1-specificity to generate the ROC curve. Intuitively, the quantity 1-specificity can be thought of as, and is nearly the same as (Phillips et al. 2006), the proportion of pixels predicted to be suitable given the threshold value t . The area under the ROC curve was then summed to calculate the AUC score.

We used the delete-d jackknife resampling procedure (Efron & Tibshirani 1994) to generate 90% confidence intervals around each AUC score. The procedure was implemented in the following way: from the total of n occurrence locations for each season / test year combination, a subset of size $n - d$, where $d = \sqrt{n}$ rounded up to the nearest integer, was drawn randomly and without replacement. This procedure was

repeated 100 times for each season and test year. One hundred Maxent models were then fit, and 100 AUC scores were calculated. For consistency with original experiments, pseudo-absence background data were sampled in the manner described above. Confidence intervals were plotted by ranking the AUC scores and plotting the 5th through the 95th greatest values (Fig. 2). Confidence intervals for models fit with and without *Calanus* in each test year were compared to determine whether there was a statistically significant difference in AUC, with non-overlapping intervals indicating a significant difference.

To assess the capacity of seasonal models to generate consistent predictions regardless of test year, we calculated the variance in true positive rate (hereafter TPR variance). TPR variance is measured directly from the data in ROC plots, and it is the variance in TPR across the 5 ROC curves (1 for each year) for each experiment. TPR variance was measured at all threshold values, $t \in [0, 1]$, $t = 0, 0.05, 0.01, \dots, 1$, and was plotted as a function of 1-specificity. The advantage of calculating TPR variance versus variance in AUC scores is that TPR variance provides information on how predictions vary as the threshold for good habitat varies, and that information is lost when an AUC score is calculated. If the TPR variance for a set of models (i.e. spring) was low and the associated AUC scores were high, it means that models trained for that season made consistently good predictions in all years. If the TPR variance was low and the associated AUC scores were low, it means that the models trained for that season made consistently

poor predictions. Thus, low TPR variance among models is associated with low interannual variability in predictive accuracy.

RESULTS

AUC

All AUC scores (Table 1) obtained were above 0.7. Thirty percent were between 0.7 and 0.799, 53.33% were between 0.8 and 0.899, and 16.67% were between 0.9 and 1.0 (Table 1). Mean AUC for the winter period experiments was higher than for the winter–spring period experiments, and the score for the winter–spring period experiments was higher than for the spring period experiments. The mean AUC score for the winter–spring models using *Calanus* (winter–spring_C) was higher than the models without *Calanus* (winter–spring). The mean AUC for the spring models with *Calanus* (spring_C) was also higher than the models without *Calanus* (spring).

In all models trained with data from the winter–spring period, except 2006, the inclusion of *Calanus* significantly improved AUC scores (Fig. 2a). Inclusion of *Calanus* made the AUC score for the year 2006 significantly worse. There was no significant effect of *Calanus* in predictions of models trained with data from the winter period (Fig. 2b). Three out of 5 models trained with data from the spring period were significantly improved by the inclusion of *Calanus* (Fig. 2c). As with the winter–spring period,

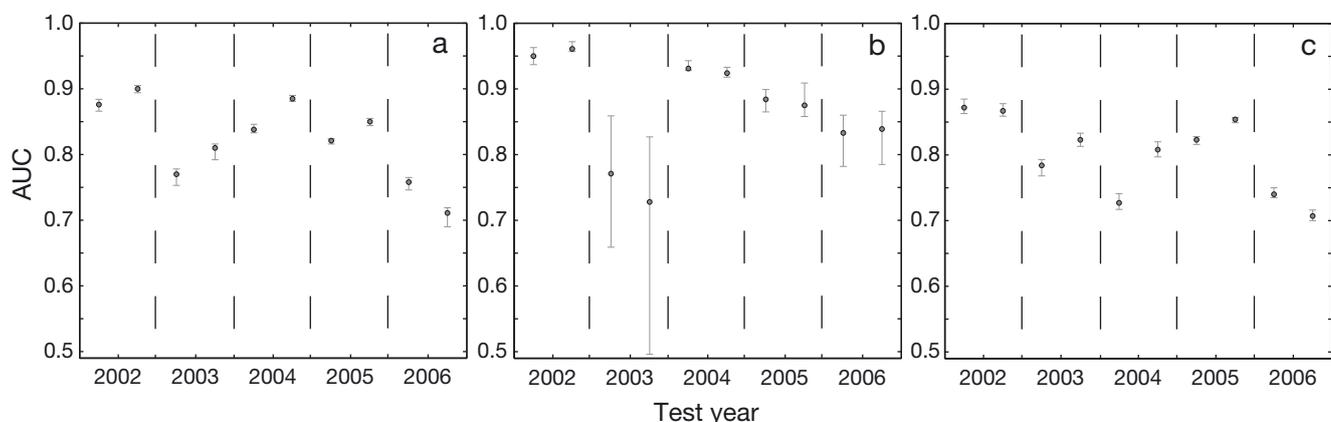


Fig. 2. Confidence intervals (90%) on the area under the receiver operator characteristic (ROC) curve (AUC) scores for each model, generated with the delete-d jackknife resampling procedure. Models are (a) winter–spring and winter–spring_C, (b) winter and winter_C, and (c) spring and spring_C. Subscript C indicates models in which *Calanus* was included as a predictor variable. Years are separated by dashed lines. Dots are original model AUC scores shown in Table 1. Data on the left side of each column (i.e. year) are for the model trained without *Calanus*, and data on the right side of columns show models where *Calanus* was included as a predictor variable. Non-overlapping confidence intervals in each test year indicate that there was a significant difference between models trained with versus without *Calanus*. Note that vertical axes show approximately half of the full range [0 1] for AUC

Table 1. Area under the curve (AUC) scores, calculated from receiver operator characteristic (ROC) curves of habitat suitability maps in each test year for each experiment: winter–spring and winter–spring_C (1 January to 1 June), winter and winter_C and (1 January to 21 March), spring and spring_C (22 March to 1 June). Subscript C indicates models in which *Calanus* was included as a predictor variable. Numbers in parentheses are the number of right whale occurrences used in model testing and training, respectively

Test year	Winter–spring	Winter–spring _C	Winter	Winter _C	Spring	Spring _C
2002	0.876 (133, 856)	0.900 (133, 856)	0.950 (18, 188)	0.961 (18, 188)	0.872 (115, 668)	0.867 (115, 688)
2003	0.770 (143, 846)	0.810 (143, 846)	0.771 (7, 199)	0.728 (7, 199)	0.784 (136, 647)	0.823 (136, 647)
2004	0.838 (227, 762)	0.885 (227, 762)	0.931 (120, 86)	0.924 (120, 86)	0.727 (107, 676)	0.808 (107, 676)
2005	0.821 (271, 718)	0.850 (271, 718)	0.884 (43, 163)	0.875 (43, 163)	0.823 (228, 555)	0.854 (228, 555)
2006	0.758 (215, 774)	0.711 (215, 774)	0.833 (18, 188)	0.839 (18, 188)	0.740 (197, 586)	0.707 (197, 586)
Mean AUC	0.813	0.831	0.874	0.865	0.789	0.812

the spring period model for test year 2006 was made significantly worse by adding *Calanus*. To alleviate concerns over the potential impact of sample size on AUC (Hernandez et al. 2006, Wisz et al. 2008), we performed a linear regression between AUC and the number of occurrence records used in training or testing of each model. No relationship was found.

One of the primary objectives of this study was to produce a model that could make good predictions of habitat suitability, not only within each year, but also from year to year. We were therefore interested in models for which AUC scores (Table 1), and shape of ROC curves, were similar from year to year. As was reported above, AUC scores for individual test years were generally good. The main exceptions were the scores for winter–spring_C and spring_C models for test year 2006 (Table 1), which were low in comparison with others.

Ideally, the modeled relationship between species and environment should provide accurate results even as environmental conditions change from year to year. We assessed the capacity of models to provide consistent performance across test years by plotting TPR variability (Fig. 3). Lower variability corresponds to more consistent model performance across test years. Variability in the winter period experiments was more than 5 times higher than for the winter–spring and spring period experiments. To aid examination of the low variance results, we removed the higher variance results (Fig. 3c,d). Removing the results for 2006 (Fig. 3a,c) allowed for comparison of TPR variance in winter–spring and spring period experiments with and without *Calanus*. The degree of consistency in model performance, from greatest to least after removing the results for 2006, is winter–spring_C, spring_C, winter–spring, and spring. Predictions from models trained with *Calanus* had greater consistency across years than models trained without *Calanus*.

Habitat suitability maps

The results presented in Figs. 2 & 3 are quantitative summaries of 380 habitat suitability maps. We present a small subset of these maps in Fig. 4, which shows hindcasted habitat suitability (from the winter–spring_C experiment) for all 8 d periods in 2002 during which whales were sighted. General patterns observed across all predicted habitat suitability maps are reported below.

A visual examination of right whale habitat suitability maps confirms what is reflected in the AUC scores. Years with high AUC tend to have more sightings in areas of high habitat suitability, and years with relatively low AUC tend to have fewer sightings in areas of high habitat suitability. The transition of whales from Cape Cod Bay to the Great South Channel, a well known phenomenon which has been documented in several studies (CETAP 1982, Winn et al. 1986, Kenney & Wishner 1995, Kenney et al. 2001), was best reflected in maps from the winter–spring_C experiment. Maps from winter period experiments clearly showed that Cape Cod Bay is an area with high habitat suitability from 1 January to 21 March, but they did not capture the transition to the Great South Channel due to their short time span. Maps from the spring period experiments (22 March to 1 June) showed the shift in highly suitable habitat from Cape Cod Bay to the Great South Channel as habitat suitability in Cape Cod Bay declined, but that transition was less pronounced than that seen in maps from the winter–spring period experiments.

Habitat preferences

We used the relative influence of predictor variables (Table 2) in each model as a proxy for habitat preferences of right whales. *Calanus* was the most

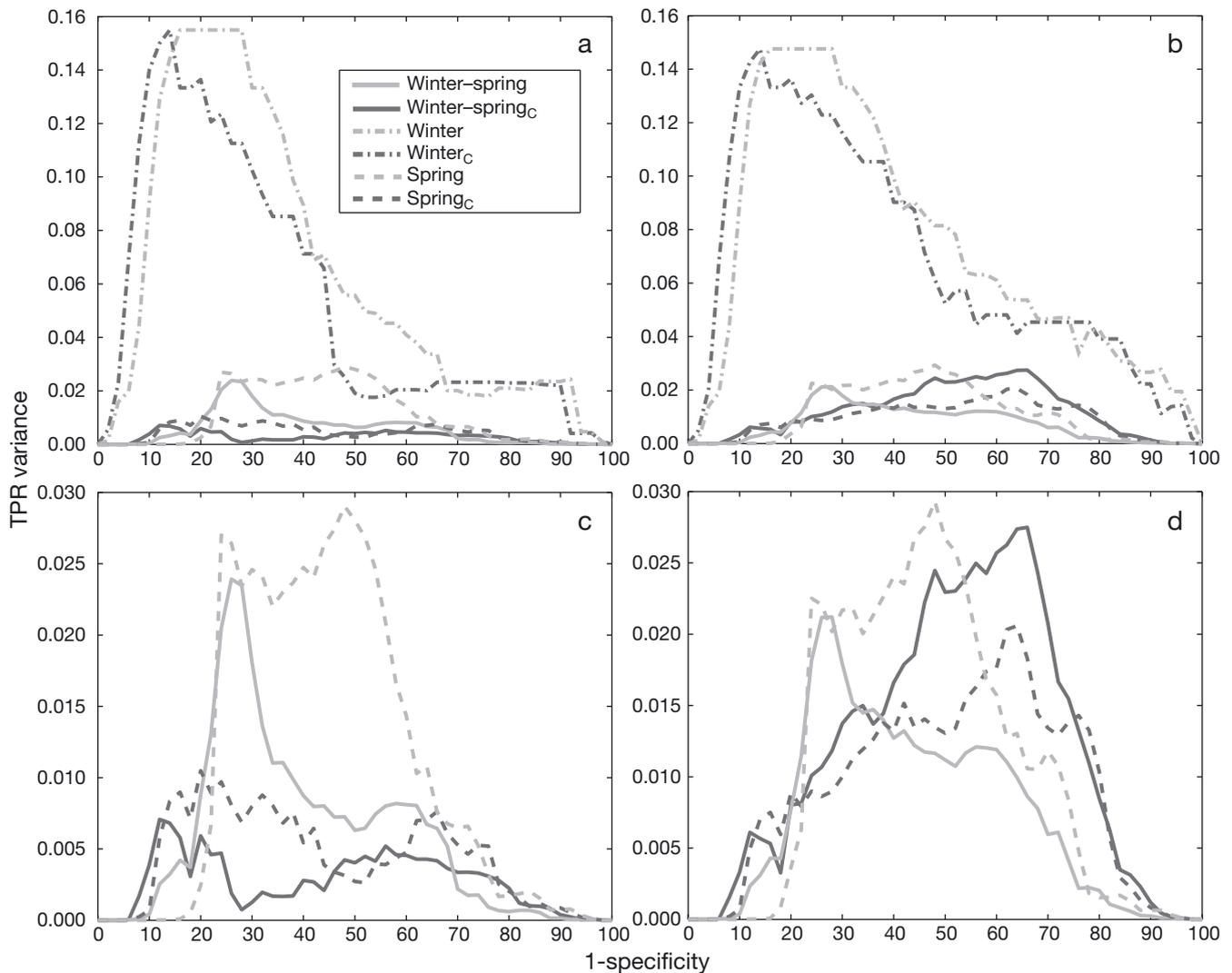


Fig. 3. *Eubalaena glacialis*. Variance in true positive rate (TPR) of right whale classification for each experiment, as a function of 1-specificity (pseudo-absence derived false positive rate), calculated from receiver operator characteristic (ROC) curves. Variance of each model (a,c) without and (b,d) with the result from 2006. TPR variance (a,b) with and (c,d) without winter and winter_C experiments to better compare models with lower variability. Winter–spring and winter–spring_C (1 January to 1 June), winter and winter_C (1 January to 21 March), spring and spring_C (22 March to 1 June). Subscript C indicates models in which *Calanus* was included as a predictor variable. Legend is the same for all subplots

influential predictor in the winter–spring_C experiment followed in importance by bathymetry, chlorophyll, and finally SST. The ranking was the same for the spring_C experiment in all years except 2004. The influence of *Calanus* was greater in the spring_C than in the winter–spring_C experiment. When *Calanus* was not included as a predictor variable, in winter–spring and spring experiments, the ranking of variable importance was bathymetry as the most influential, followed by chlorophyll and SST. The winter period experiments showed the opposite pattern: SST was the most influential predictor, followed by chlorophyll, bathymetry, and *Calanus* when it was

included as a predictor variable. The winter_C 2006 model was an exception to this ranking, with bathymetry (22.9%) slightly more influential than chlorophyll (22.6%).

Predicted habitat suitability at whale occurrences during winter–spring

There were clear differences in the value of environmental variables at the time and place of right whale occurrence. Modeled *Calanus* abundance was $<10 \text{ ind. m}^{-3}$ at 95% of winter right whale

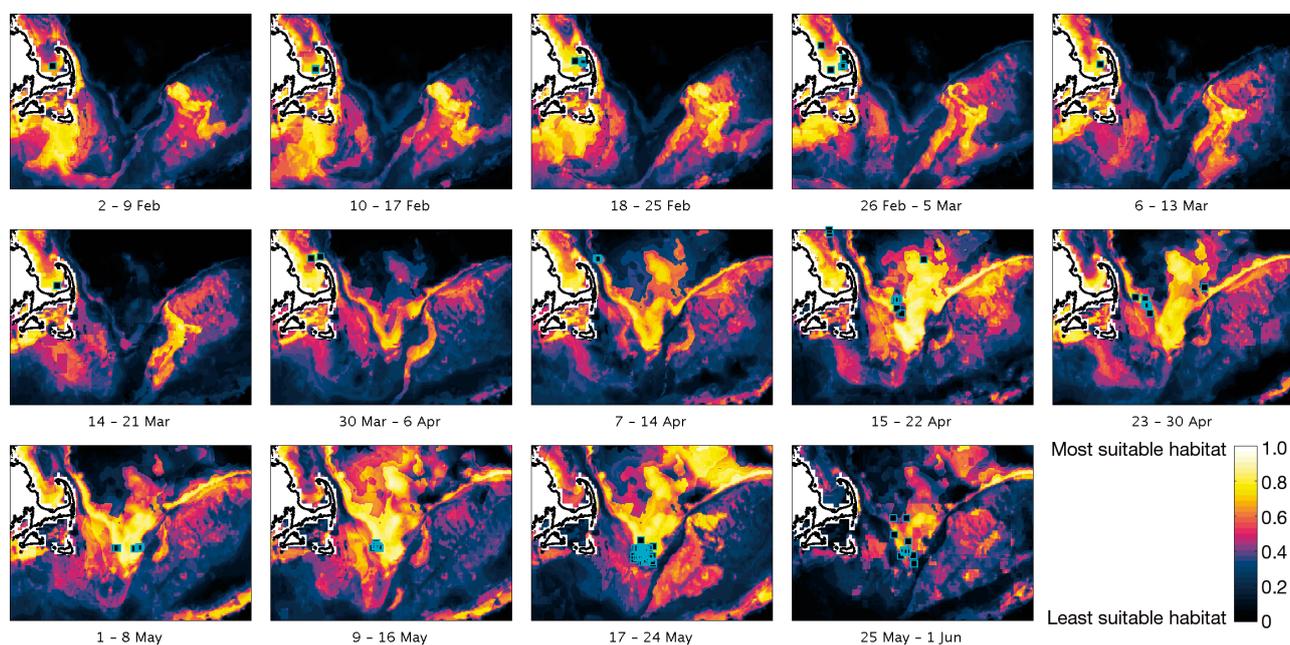


Fig. 4. *Eubalaena glacialis*. Hindcasts of right whale habitat suitability for all 8 d periods in 2002 during which right whales (squares) were sighted. White area at left is Cape Cod, Massachusetts. Per the color bar, black is low and white is high habitat suitability. All hindcasts are from the winter–spring_C experiment

Table 2. Percent variable contribution for all model test years 2002 to 2006, for all experiments: winter–spring and winter–spring_C experiments (1 January to 1 June), winter and winter_C and (1 January to 21 March), spring and spring_C (22 March to 1 June). Subscript C indicates models in which *Calanus* was included as a predictor variable. The greatest contribution to each model is shown in **bold**. SST: sea surface temperature, na: not applicable

Test year	Predictor variable	Winter–spring	Winter–spring _C	Winter	Winter _C	Spring	Spring _C
2002	<i>Calanus</i>	na	35.3	na	10.8	na	45.1
	Bathymetry	42.6	32.5	23.2	20.7	54.3	34.3
	Chlorophyll	33.7	19.6	33.8	26.9	24.0	12.1
	SST	23.7	12.6	43.0	41.7	21.7	8.4
2003	<i>Calanus</i>	na	37.6	na	17.6	na	44.4
	Bathymetry	44.4	28.5	28.2	21.1	57.3	35.6
	Chlorophyll	29.0	18.5	30.3	26.2	21.7	11.2
	SST	26.6	15.4	41.6	35.1	21.0	8.8
2004	<i>Calanus</i>	na	36.9	na	18.7	na	35.7
	Bathymetry	48.4	31.8	26.3	22.7	56.7	38.6
	Chlorophyll	27.7	18.0	30.0	24.6	22.9	14.2
	SST	23.9	13.4	43.7	34.0	20.4	11.4
2005	<i>Calanus</i>	na	33.2	na	15.4	na	42.5
	Bathymetry	45.0	31.6	26.7	20.6	60.6	39.0
	Chlorophyll	35.0	19.2	27.9	24.3	21.4	11.4
	SST	25.0	16.0	45.3	39.6	18.1	7.1
2006	<i>Calanus</i>	na	40.9	na	16.9	na	46.3
	Bathymetry	47.7	31.0	26.5	22.9	61.4	34.9
	Chlorophyll	27.8	15.8	31.4	22.6	20.7	11.6
	SST	24.4	12.3	42.1	37.6	17.9	7.1

occurrences (Fig. 5a). A portion of spring right whale occurrences were in areas of low *Calanus*; however, for a large portion of occurrences (65%) modeled *Calanus* was >10 ind. m^{-3} . There was a positive trend between predicted habitat suitability

and modeled *Calanus* abundance during spring. Chlorophyll at right whale occurrences was generally higher in winter than in spring (Fig. 5b). Almost all of the whale occurrences during winter (75%) were associated with chlorophyll values >0.5 (\log_{10}

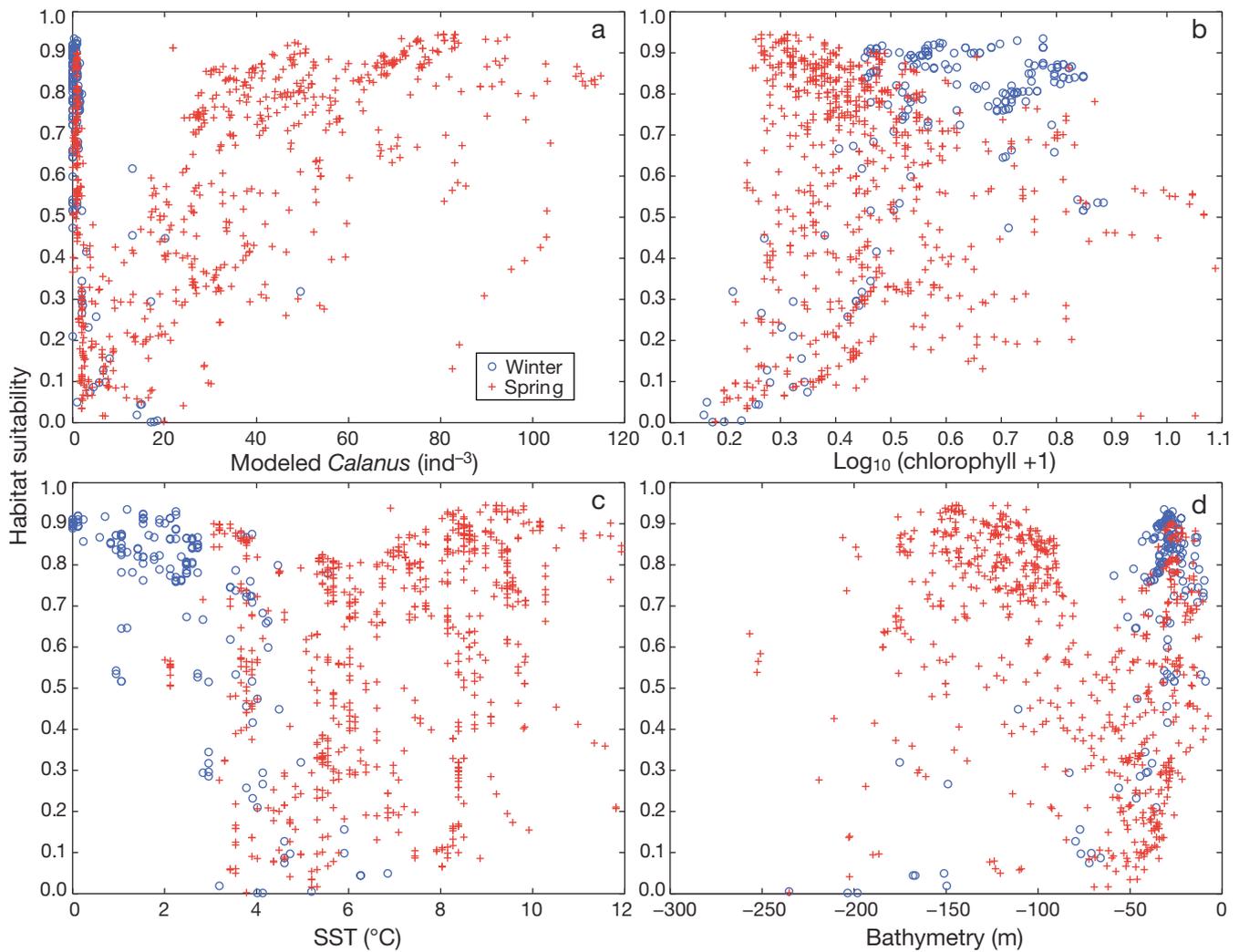


Fig. 5. *Eubalaena glacialis*. Habitat suitability values at each right whale presence location and time from 2002 to 2006 (N = 989) versus: (a) modeled *Calanus*, (b) $\log_{10}(\text{chlorophyll} + 1)$, (c) sea surface temperature (SST) and (d) bathymetry. Circles indicate right whale presence records from the winter period (1 January to 21 March). Crosses indicate right whale presences from the spring period (22 March to 1 June)

[$x + 1$]), while 74% of spring right whale occurrences were associated with chlorophyll < 0.5 . During winter there was a cluster of whale sightings at SSTs less than 3°C and a steep decline thereafter (Fig. 5c). Eighty-eight percent of winter right whale sightings were associated with SST below 4°C . Eighty-five percent of spring right whale sightings were associated with SST between 4 and 12°C . There was no discernible trend in the temperature at whale occurrences during summer. There was a large cluster of winter right whale occurrences at a depth of less than 50 m. Ninety-one percent of winter right whale occurrences were in water less than 63 m depth (the maximum depth of Cape Cod Bay), and 56% of spring right whale occurrences were in water greater than 63 m depth.

DISCUSSION

We found that weekly habitat suitability for North Atlantic right whales, a highly migratory marine mammal, can be hindcasted with reasonable accuracy using a species distribution model. In the vast majority of applications, species distribution models have used multi-year climatologically averaged predictor variables, such as those from the WorldClim database (Hijmans et al. 2005), yielding estimates of the absolute range or habitat suitability of a species. In such predictions, year-to-year variability is averaged out. While there has been some work on seasonal models (Suárez-Seoane et al. 2008), we believe our work to be the first application of a species distribution model on the weekly timescale.

H₁: right whale habitat preferences are dynamic

Differences in the influence of predictor variables on model results for winter/winter_C and spring/spring_C experiments (Table 2) suggest that the habitat preferences of right whales are not static. Results from models of the winter period suggest that the winter distribution of right whales is determined primarily by SST. The significance of ocean temperatures to right whales is not well understood. SST has been documented to be an important predictor of right whale distribution in certain areas. The distribution of SST at right whale occurrences in the southeast US right whale Critical Habitat is non-random (Keller et al. 2006), with right whales being found in waters cooler than 22°C (Good 2008). In the northern habitats such as the Bay of Fundy and Roseway Basin, where feeding is common, the influence of temperature on the distribution of right whales is less clear. Woodley & Gaskin (1996) found significantly higher surface temperatures where right whales were present than where they were absent in the lower Bay of Fundy. Baumgartner et al. (2003b) found some indication that the SST gradient could explain interannual variability in right whale occurrence in Roseway Basin, adjacent to the Bay of Fundy. However, in that same area, Patrician & Kenney (2010) did not find a relationship between SST and right whale abundance. Differing spatial and temporal scales of investigation confound comparison of these studies. The strong influence of temperature in our winter models may be a proxy for an environmental variable not considered in our study. The copepods *Pseudocalanus* spp. are important winter prey for right whales (Mayo et al. 2004), but temperature has not been found to impact the abundance of *Pseudocalanus* spp. (DeLorenzo Costa et al. 2006, Turner et al. 2011). It is also important to note that satellite-derived SST provides the temperature in only the first ~1 mm of the ocean, and right whales spend a considerable amount of time at depth where temperatures are typically different.

Models for the winter–spring and spring periods were most heavily influenced by *Calanus*, with that variable being more important in models of spring than of the winter–spring period. An increased influence of *Calanus* as the year progresses agrees with what we have observed empirically: during winter in Cape Cod Bay, right whales feed on *Pseudocalanus* spp. and possibly *Centropages* spp. (Mayo & Marx 1990), after which they transition to a diet of *Calanus* as that prey taxon becomes more abundant (Pendleton et al. 2009). Our results support the findings of

several other studies that have found or inferred *Calanus* to be a major component of the right whale diet (Murison & Gaskin 1989, Mayo & Marx 1990, Wishner et al. 1995, Beardsley et al. 1996, Woodley & Gaskin 1996, Baumgartner & Mate 2003).

Right whales have been found in a wide range of depths. On the calving grounds in the southeast USA, right whales are commonly found in water 10 to 20 m deep, which is consistent with calving activity in that region (Garrison 2007). Our study found right whales in water with depths of 8 to 257 m in the Gulf of Maine. Aggregations of right whales regularly occur over the deep basins of the lower Bay of Fundy and Roseway Basin in waters 100 to 200 m deep. It is likely that bathymetric features give rise to physical and/or biological processes that are important to right whales.

Relative to other predictor variables, chlorophyll had a moderate influence on model results. Chlorophyll provides a reasonable proxy for phytoplankton abundance, which is directly related to *Calanus* egg production. The time lag between a phytoplankton bloom and increases in *Calanus* abundance mean that weekly chlorophyll data may not be a good indicator of *Calanus*, and thus right whale distributions. Hlista et al. (2009) found that a 2 yr lagged index of chlorophyll concentration (measured when right whales are expected in Cape Cod Bay, the Great South Channel, and the Bay of Fundy feeding habitats) was positively correlated with annual right whale calving numbers. Thus, on longer time scales, chlorophyll is an important variable to consider in right whale habitat models. Temporally lagged or longer-term averages of chlorophyll concentration may have provided a stronger predictor than the weekly averages used in our models.

We modeled the distribution of right whale habitat based upon only 4 predictor variables. It is widely thought that several other factors may influence right whale distributions, including thermal fronts and the bottom mixed layer (Baumgartner et al. 2003b), abundance of other prey species such as *Pseudocalanus* spp. and *Centropages* spp. (Mayo & Marx 1990, Pendleton et al. 2009), and bottom type (Good 2008).

Our training data sets were constructed by associating each whale occurrence with average environmental conditions for the 8 d period in which the whale was observed. Right whales can travel a considerable distance in 1 wk (Mate et al. 1997); therefore, the approximation of environmental conditions at the time of whale occurrence probably introduced some inaccuracy. A more precise model could be

made by associating occurrences of non-transiting right whales with daily values of environmental conditions. Our results are based upon statistical relationships between known right whale presences and the value of predictor variables associated with those presence records. As with all results of correlative studies, our results are not necessarily indicative of physical or biological processes.

***H*₂: prey data improve predictive accuracy**

Torres et al. (2008) found that predictive accuracy was not improved with the inclusion of prey data in a generalized additive model of bottlenose dolphin *Tursiops truncatus* habitat. In the present study, in 2 of the 3 time periods examined (winter–spring and spring), the inclusion of modeled prey abundance improved overall predictive accuracy (Fig. 2, Table 1), sensitivity to interannual environmental variability (Fig. 3), and the ability to discriminate good from bad habitat (i.e. greater range in habitat suitability values makes it easier to delineate differences). Because we are modeling right whale habitat suitability in areas where whales are known to feed much of the time (Mayo & Marx 1990, Wishner et al. 1995), it makes sense that the availability of prey would be an important predictor of the distribution of whales.

Right whales feed on ultra-dense patches of copepods (Watkins & Schevill 1976, Wishner et al. 1988, Mayo & Marx 1990, Beardsley et al. 1996, Baumgartner et al. 2003a), often occurring at the scale of 1 to 10s of meters. It is therefore interesting that our estimates of copepod abundance at 1 km resolution served as an important predictor of the distribution of potential right whale habitat, given that the modeled copepod fields do not contain patches. This, along with the studies of Pendleton et al. (2009) and Pershing et al. (2009b), reinforce the view that high regional-scale mean abundance of copepods increases the likelihood of formation of ultra-dense patches of copepods. Under this view, adding SST, chlorophyll, and bathymetry to the habitat model may indicate the physical conditions that encourage patch formation.

Interannual variability

Interannual variability was well captured in results from the winter–spring_C experiment (Fig. 3). At the other end of the spectrum were the models of the winter period, which performed well in some years and

poorly in other years. The high variability in winter models (Fig. 3a,b) may be an indication that we are missing an important environmental variable, such as *Pseudocalanus* spp., a winter taxon which is thought to be an important resource for right whales during winter (Mayo & Marx 1990, Pendleton et al. 2009).

Another important consideration is the influence of sample size on AUC. Hernandez et al. (2006) and Wisz et al. (2008) found that models trained with fewer presences have a lower AUC. Our models of the winter season were trained on considerably fewer presence locations than models for winter–spring or spring periods. However, the lowest number of presences used in any of our models (N = 86 in winter 2004 and winter_C 2004) was well above the number of presence locations that produced artificially low AUC values in Hernandez et al. (2006) and Wisz et al. (2008). There were relatively few presence locations available for testing winter 2002 (N = 18) and winter 2004 (N = 7) (Table 1). Using a small number of test points to construct a ROC curve will result in a curve with fewer nodes or points to connect. This could result in an over- or underestimate of the AUC that would have been calculated with a large number of test points.

Models fit with *Calanus* from the winter–spring and spring periods for 2006 did not perform well (Table 1). This, along with the fact that models fit with data from winter and tested in 2006 (with and without *Calanus*) were not far from average leads us to believe that the modeled distribution of *Calanus* in the spring was responsible for the poor predictions. Alternatively, right whales may have had a strong response to an environmental covariate that we did not model, such as *Pseudocalanus* spp., or right whales may have been drawn to a more suitable habitat outside of our model domain (and outside of the typical range of right whales in winter and spring). We analyzed the spatial and temporal patterns of sighting data in 2006 versus 2002 to 2005 and found that the distribution of whales in 2006 was not unusual in comparison to 2002 to 2005. An examination of environmental variables revealed that modeled *Calanus* abundances were anomalously high between late March and late May of 2006. This means that the models trained with data from 2002 to 2005 were informed by relatively low springtime abundances of modeled *Calanus* and were tested in a year with relatively high springtime abundances of modeled *Calanus*. This difference in training and testing levels of *Calanus* probably contributed to below average performance of winter–spring_C and spring_C models for 2006.

CONCLUSIONS

We have found that right whale habitat suitability can be estimated on a weekly timescale with only 4 predictor variables. Models that incorporated data from both winter and spring seasons were more accurate and provided more consistent performance than winter-only or spring-only models. Our results confirmed empirical observations that right whale habitat preferences change on a seasonal timescale. The inclusion of prey as a predictor variable improved predictive accuracy of models. The framework presented here is an important step toward the goal of having a near real-time habitat modeling system that could be used to answer ecological questions and assess risk to endangered species by proposed management actions. Our approach could also be extended to other populations of highly mobile animals, both on land and in the ocean.

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