



# Differing prey associations and habitat use suggest niche partitioning by fin and humpback whales off Kodiak Island

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**ABSTRACT:** Fin *Balaenoptera physalus* and humpback *Megaptera novaeangliae* whales share foraging areas and may compete for the same prey, but little is known about the extent to which they partition prey resources. Visual cetacean surveys and simultaneous acoustic-trawl surveys of prey were conducted around 2 submarine canyons off Kodiak Island, Alaska, in 2004 and 2006. Statistical models were used to examine the associations between sightings of fin and humpback whales and measures of their potential prey and environment. Observations and models indicate that fin whales were disproportionately abundant in areas with the highest observed euphausiid concentrations, while humpback whales were abundant at lower euphausiid concentrations and in areas where juvenile walleye pollock were abundant. Fin whales were abundant in the areas where euphausiid biomass was deepest and in the deepest areas surveyed (>150 m depth). In contrast, humpback whales primarily occurred in shallower areas and near more shallowly distributed euphausiids. The different depth and prey affinities of fin and humpback whales suggest niche and habitat partitioning between these 2 co-occurring species. Abundance models built using acoustic estimates of prey density are a useful tool to further understanding of the abundance, distribution, and behavior of these animals.

**KEY WORDS:** Fisheries acoustics · Cetacean habitat · Niche partitioning · Fin whales · Humpback whales · Euphausiids

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## 1. INTRODUCTION

Worldwide, both fin *Balaenoptera physalus* and humpback *Megaptera novaeangliae* whale populations aggregate in highly productive coastal environments such as those in the North Pacific off northern California (Fleming et al. 2016), British Columbia (Darlings et al. 1996, Dalla Rosa et al. 2012), and Kodiak Island (Waite et al. 1999); and in the Atlantic, including the Bay of Fundy (Hazen et al. 2009), the Gulf of Maine (Clapham & Seipt 1991, Clapham et al. 2003), the Gulf of St. Lawrence (Simard & Lavoie

1999), Newfoundland (Whitehead & Carscadden 1985), and the Barents Sea (Mizroch et al. 2009, Skern-Mauritzen et al. 2011, Ressler et al. 2015). High productivity in these regions during summer months creates foraging hotspots for multiple species, including fin and humpback whales (Mizroch et al. 2009, Block et al. 2011). The energy-rich prey they consume in high latitude feeding grounds fuels their long distance migrations to breed, and increases survival rates of calves (Baker et al. 1985, Calambokidis et al. 2001, Goldbogen et al. 2007, 2008, Sigler et al. 2012, Williams et al. 2013). Because both species

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often forage simultaneously in the same productive coastal habitats, there is potential for competitive interactions.

A persistent question in ecology is how 2 organisms with overlapping diets partition resources so they can coexist in the same niche or ecological space (Werner & Gilliam 1984, Warren et al. 2008). Physiological and behavioral adaptations to facilitate this partitioning result from competitive pressure and demonstrably occur in plants (Kelly et al. 2008, Leibold 2008) and terrestrial and aquatic animals (Dumas 1956, Siemers & Schnitzler 2004). Species with similar environmental requirements can minimize direct competition through spatial, temporal, and trophic partitioning (Hutchinson 1957, Porter 1976, Fox & Bellwood 2013). For example, terrestrial animals may spatially partition a niche and modulate sun exposure through movement up or down a tree trunk (Dumas 1956), whereas aquatic organisms may spatially partition by inhabiting different depths (Porter 1976, Cullen & MacIntyre 1998). Temporal partitioning can be mechanistic and direct, as with terrestrial animals shifting their niches diurnally or nocturnally and thus avoiding direct competition (Kronfeld-Schor & Dayan 1999, Albrecht & Gotelli 2001), or it can be more dynamic and occur through sensitivity to seasonal fluctuations and long-term climate conditions (Engelbrecht et al. 2007, Kelly et al. 2008). Trophic partitioning occurs when organisms consume different prey assemblages (Ross 1986), and often occurs in tandem with some form of fine-scale spatial partitioning, especially in the marine environment (Fox & Bellwood 2013, Hinke et al. 2015).

Large cetaceans exhibit multiple forms of niche partitioning. Both trophic and spatial partitioning occur between minke *B. acutorostrata* and humpback whales (Friedlaender et al. 2006, 2009b) and between fin and humpback whales in highly productive cold water feeding areas (Witteveen et al. 2015, Herr et al. 2016, Witteveen & Wynne 2016). Trophic partitioning occurs between blue *B. musculus*, fin, humpback, and minke whales in the Gulf of St. Lawrence (Gavrillchuk et al. 2014), as does a combination of spatial and temporal partitioning in relation to dynamic oceanographic features (Doniol-Valcroze et al. 2007). In the Bering Sea, fin and humpback whale distributions were associated with higher euphausiid (krill) biomass, and minke whale distributions were associated with higher juvenile walleye pollock *Gadus chalcogrammus* (hereafter pollock) biomass (Zerbini et al. 2016). In the same study, fin whales aggregated near the shelf edge and hump-

backs aggregated on the mid-shelf, suggesting both trophic niche partitioning and spatial partitioning (Zerbini et al. 2016). Blue whales and humpback whales exhibit resource partitioning in the California Current through spatial, temporal, and trophic partitioning, with blue whales foraging exclusively on euphausiids during the seasonal peak on the shelf break, and humpbacks foraging on a wider variety of prey both on the shelf break and inshore waters in both summer and fall (Fossette et al. 2017). In some cases, niche partitioning is less clear, with 2 or more species appearing to target the same prey simultaneously. In the Barents Sea, both fin and humpback whale occurrences were correlated with euphausiid aggregations, suggesting possible direct competition for prey (Skern-Mauritzen et al. 2011, Ressler et al. 2015). Understanding the mechanisms by which sympatric fin and humpback whales niche-partition helps us to better understand the potential impacts they may have on prey populations and on prey availability for other top predators.

The region near Kodiak Island, Alaska, is a foraging area for both fin and humpback whales during summer months (Waite et al. 1999, Baraff 2006, Zerbini et al. 2006, Witteveen & Quinn 2007, Mizroch et al. 2009, Witteveen et al. 2015, Zerbini et al. 2016). The potential for competition exists as both species have been recorded foraging on the same prey items (e.g. euphausiids, copepods, capelin *Mallotus catervarius*, mackerel *Scomber japonicus*, pollock) in this region and in other foraging hotspots (Nemoto 1959, Dolphin 1988, Clapham & Meade 1999, Flinn et al. 2002). While population levels of both fin and humpback whales are increasing in the Gulf of Alaska (GOA), including in the Kodiak Island region, they have not yet risen to pre-whaling levels (Zerbini et al. 2006, Witteveen & Quinn 2007, Mizroch et al. 2009, Allen & Angliss 2015). With low (but increasing) population sizes, we would not expect to see direct competition driven by large-scale prey depletion in this area (Clapham & Brownell 1996, Friedlaender et al. 2006). But, given the high energy costs for foraging dives (Croll et al. 2001, Acevedo-Gutiérrez et al. 2002, Hazen et al. 2015) and the need to target the highest density patches of prey to maximize foraging efficiency and energy gain (Hazen et al. 2009, Burrows et al. 2016, Fossette et al. 2017), there is potential for localized depletion of prey in this region (Witteveen et al. 2006, Straley et al. 2018), which could be a factor in influencing behavior of these 2 species (Clapham & Brownell 1996).

Recent tagging and stable isotope studies in coastal bays around Kodiak Island have considered

how fin and humpback whales partition their foraging environment (Witteveen et al. 2008, 2015, Witteveen & Wynne 2016). The tagging study (Witteveen et al. 2015) focused on the nearshore and inner bays around Kodiak and measured individual fin and humpback dive behavior for brief periods of time, concurrent with acoustically quantified prey abundance generalized to trophic level. The isotope study (Witteveen & Wynne 2016) examined the relative contributions of zooplankton and forage fishes to fin and humpback whale diets in coastal bays in the Kodiak and Shumagins regions. These studies suggest that in the Kodiak region in nearshore waters, fin whales are functionally planktivores, while humpback whales switch from consuming zooplankton to forage fishes, depending on the relative abundance of these prey (Anderson & Piatt 1999, Witteveen & Wynne 2016). Additional research suggests that spatial segregation of fin and humpback whales may occur by depth (Wynne et al. 2005). Evidence from other ecosystems also suggests that humpback and fin whales may undergo prey switching between zooplankton and forage fishes (Skern-Mauritzen et al. 2011, Ressler et al. 2015, Fleming et al. 2016), and that fin whales may forage in areas with lower prey densities than humpbacks (Baraff 2006, Witteveen et al. 2015, Witteveen & Wynne 2016). Alternately, it has been suggested that it may be critical for larger rorquals, such as fin whales, to exploit the densest prey patches (Acevedo-Gutiérrez et al. 2002, Goldbogen et al. 2007, 2012). The 2 species' foraging behaviors and prey preferences remain a research need in the GOA and particularly in the offshore areas around Kodiak Island (Witteveen et al. 2015, Witteveen & Wynne 2016, Wright et al. 2016).

In 2004 and 2006, the Alaska Fisheries Science Center conducted acoustic-trawl surveys northeast of Kodiak Island, focused on capelin and pollock (Wilson et al. 2003, Hollowed et al. 2007, Logerwell et al. 2007, Walline et al. 2012), that also included opportunistic visual observations of marine mammals. Together, these data allowed for concurrent measurements of the distribution of large cetaceans, the oceanographic features of their habitat, characterization of the available prey fields using acoustic and midwater trawl sampling, and an evaluation of prey partitioning in the fin and humpback foraging grounds. This single-platform, concurrent sampling approach complements short-term focal follows and stable isotope work in previous studies in the area by providing detailed data on prey availability and cetacean behavior simultaneously over several months in more offshore environments. A

better understanding of the environmental and prey conditions that influence niche partitioning of these whales can potentially inform management by ensuring that marine protected areas are designed to include suitable habitat and by incorporating prey requirements into fisheries management processes (Trites et al. 1997, Gregr & Trites 2001, Bailey & Thompson 2009).

## 2. MATERIALS AND METHODS

### 2.1. Study site

Barnabas and Chiniak are 2 adjacent troughs on the east side of Kodiak Island in the GOA, USA, which support high primary production from April–November (Hollowed et al. 2007). This region is a summer feeding ground for humpback and fin whales (Waite et al. 1999, Zerbini et al. 2006, Witteveen & Quinn 2007). The 2 troughs have similar bathymetric features, with depths in the surveyed area ranging from 50–250 m, and an average depth of approximately 135 m.

The Alaska Fisheries Science Center conducted acoustic-trawl surveys of Barnabas and Chiniak Troughs with the NOAA ship 'Miller Freeman' in 2004 and 2006 with the goal of understanding the drivers of distribution of capelin and pollock and their relationship to commercial fishing in these regions (see Fig. 1; Logerwell et al. 2007, Walline et al. 2012). Opportunistic visual observations of marine mammals were collected during the acoustic-trawl survey. Three replicate survey passes were completed in Barnabas and 4 in Chiniak from 13 August through 6 September 2004. Five survey passes were completed in each trough from 11 August through 5 September 2006. Chiniak and Barnabas troughs (see Figs. 1 & 2) were surveyed using parallel transects spaced 5.5 km apart, with transect start and end points located at depths of 75–100 m past the trough edges. Vessel speed averaged  $6.0 \text{ m s}^{-1}$  (11.7 knots) during acoustic data collection. The acoustic-trawl surveys took place during daylight hours (approximately 14–15 h  $\text{d}^{-1}$ ).

### 2.2. Cetacean survey

A single observer made visual cetacean observations during daylight hours of the acoustic-trawl survey. Line-transect data were collected using standard cetacean survey protocols (e.g. Zerbini et al.

2006). As there was only a single observer, most scanning for whales was done by eye, but a set of pedestal-mounted 20× power ‘big-eye’ binoculars were used to estimate the angle from the trackline to the sighting, radial distance to the sighting (from reticles in the ocular), and to confirm species identification. All sighting and environmental data were entered into the program WINCRUZ (available for download at the SWFSC PRD software portal; Friday et al. 2013), a computer database interfaced to a GPS unit to gather positional and navigational information. Only sightings of whales obtained while the observer was ‘on-effort’, when the Beaufort sea state was 5 or lower, and when visibility was >3 km were used. Radial distance to each sighting was calculated using ‘approximation 2’ of Lerczak & Hobbs (1998) from the binocular reticule measurements and platform height. Perpendicular distance was calculated by multiplying the radial distance by the sine of the angle to the sighting. The latitude and longitude of each whale sighting was calculated from the GPS position of the ship and the angle and radial distance to the sighting. The observer recorded species, group size, and location of sightings. As tracklines were 5.5 km apart, only sightings within 2.75 km of the trackline were compiled and used in the final analysis. As there was only a single observer, the cetacean observations were not continuous: observations were made along 45% of the survey transects in 2004 and 65% in 2006.

### 2.3. Acoustic-trawl survey

Acoustic-trawl methods (Simmonds & MacLennan 2005) were used in the collection and analysis of acoustic and trawl data to determine the abundance of potential prey. Acoustic backscatter was measured continuously during daylight hours along the cruise track with centerboard-mounted Simrad EK500 (38 kHz) and EK60 (18, 120, 200 kHz) echosounders, which were synchronized to transmit simultaneously. In 2006, a Simrad EK60 echosounder was used at the 4 frequencies described above. The echosounders were calibrated using the standard sphere method (Foote et al. 1987) at the beginning and end of each cruise. Acoustic data were processed with Sonardata echoview software (version 4.9).

Trawl hauls were conducted to identify the size and species composition of the primary sound-scattering organisms detected with the echosounder. Trawl hauls were targeted in areas of high backscatter, and the catch was sampled to determine the species and size compositions of the sound-scattering organisms (Wilson et al. 2003). An Aleutian Wing 30/26 pelagic trawl fitted with a 13 mm codend liner (Honkalehto et al. 2009) was used to sample midwater fish aggregations. A poly Nor’eastern bottom trawl with roller gear was used to sample demersal fishes (Wilson et al. 2003). A Methot trawl with a mouth opening of 5.2 m<sup>2</sup>, constructed with 2 × 3 mm oval mesh with a 1 mm mesh codend was used to sample suspected euphausiid aggregations (Hollowed et al. 2007). A total of 59 midwater trawls were conducted (28 in Chiniak and 31 in Barnabas) in 2004. Five bottom trawl hauls and 6 Methot hauls were also conducted in each trough during the study period. A total of 47 midwater trawl hauls were conducted in 2006 (23 in Chiniak and 24 in Barnabas). Fourteen bottom trawl hauls (11 in Barnabas, 3 in Chiniak) and 4 Methot tows (2 trough<sup>-1</sup>) were conducted during the study period (Fig. 1). Each midwater and bottom haul catch was weighed and enumerated, a subsample of ~300 pollock >10 cm standard lengths (SL) were measured, and ~50 were weighed.

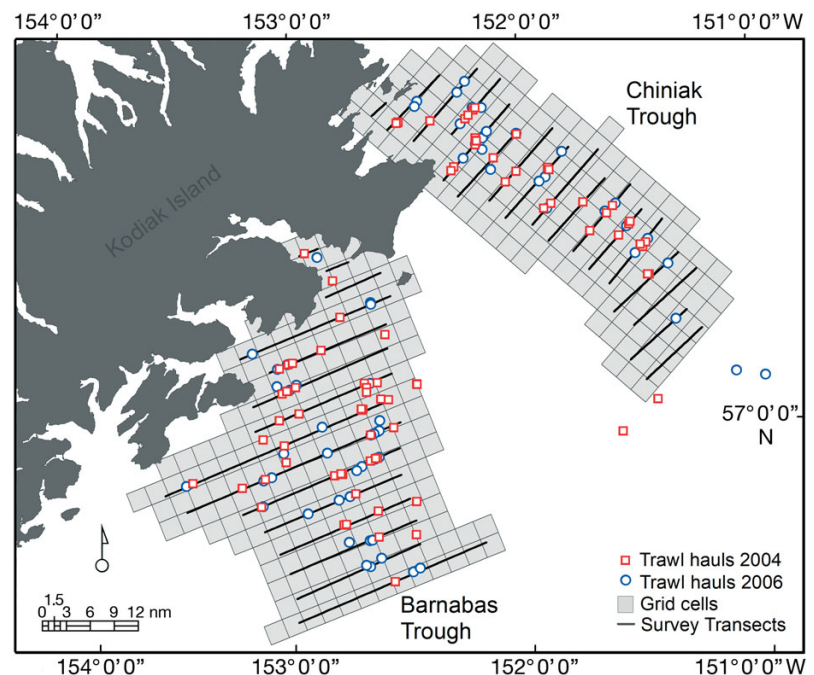


Fig. 1. Survey regions (Barnabas and Chiniak Troughs) for summer 2004 and 2006 cruises aboard the NOAA Ship ‘Miller Freeman’. Red boxes: trawl hauls performed in 2004; blue circles: hauls in 2006. Survey transects were similar for all passes and both years

When age-0 pollock ( $\leq 10$  cm) or capelin (10–17 cm) were captured, approximately 50 length and 10 length–weight measurements were collected. Methot catches were processed by first removing large medusae and then preserving a subsample of the remaining catch in 3.7% buffered formaldehyde solution. The organisms in the Methot samples were later sorted and identified in the laboratory using a binocular microscope.

#### 2.4. Acoustic data analysis

Acoustic backscatter was attributed to several species groups (Table 1), based on interpretation of the depth distribution and aggregation patterns observed in the echosounder data and the results of targeted trawl sampling (Walline et al. 2012). The primary sound scatterers were large pollock (95% were  $>35$  cm in both troughs), euphausiids, and a mixture of capelin and age-0 walleye pollock. Backscatter from euphausiids in the 18–200 kHz range is strongly frequency-dependent (Demer & Conti 2005, McKelvey & Wilson 2006). An index of euphausiid abundance was generated using a multifrequency technique based on observations from trawl ground-truthed euphausiid aggregations in Alaska (De Robertis et al. 2010) and applied to these data. Volume backscatter was averaged into 5 ping  $\times$  5 m cells, and volume backscattering 12–32 dB higher at 120 kHz relative to 18 kHz (i.e. 15.8–158.5 fold higher at 120 kHz) was classified

as euphausiids. The nautical area scattering coefficient ( $s_A$ ;  $m^2 \text{ nmi}^{-2}$ ) at 38 kHz for fishes and near-surface backscatter, and at 120 kHz for euphausiids, was calculated in 10 m vertical layers (14 m below surface to 0.5 m off bottom) in 0.5 nmi horizontal intervals along the survey track. The mean-weighted depth (MWD) was calculated for each of the species groups.

The echo integral from the age-0 pollock and capelin mixture was partitioned into backscatter from each species based on the proportion of the species in the trawl catches, the size distribution, and the target strength (TS) of each species (De Robertis et al. 2017, their Eq. 4). We used the TS to fork length (FL, cm) relationship for pollock ( $TS = 20\log[FL] - 66.0$ ; Traynor 1996) and a TS to total length (TL, cm) relationship to capelin, derived from *in situ* measurements in the GOA ( $TS = 20\log[TL] - 70.3$ ; Guttorf & Wilson 2009). Given that conversion of euphausiid backscatter to abundance remains uncertain (Hunt et al. 2016), we did not calculate euphausiid biomass and abundance from the backscatter measurements. Euphausiid backscatter is a reasonable crude index of abundance and an effective proxy for biomass in this region (Ressler et al. 2012, Simonsen et al. 2016). Acoustic backscatter can be used as proxy for animal abundance as it is proportional to abundance if scattering strength is constant (Foote 1983) and was used as a proxy for biomass for all potential prey species for consistency. Acoustic analyses were performed using Echoview software version 8.0.

Table 1. Predictor variables for surveys off Kodiak Island in summer 2004 and 2006

| Predictor variable        | Description   |
|---------------------------|---|
| Krill                     | Euphausiid backscatter, mean nautical area scattering coefficient in ( $m^2 \text{ nmi}^{-2}$ ) per grid cell                               |
| Adult pollock             | <i>Gadus chalcogrammus</i> backscatter, mean nautical area scattering coefficient in ( $m^2 \text{ nmi}^{-2}$ ) per grid cell               |
| Age-0 pollock             | <i>Gadus chalcogrammus</i> young-of-year backscatter, mean nautical area scattering coefficient in ( $m^2 \text{ nmi}^{-2}$ ) per grid cell |
| Capelin                   | <i>Mallotus villosus</i> backscatter, mean nautical area scattering coefficient in ( $m^2 \text{ nmi}^{-2}$ ) per grid cell                 |
| Chlorophyll               | Chlorophyll ( $\mu\text{g m}^{-2}$ )  |
| SST                       | Sea surface temperature ( $^{\circ}\text{C}$ ) from shipboard sensors   |
| Depth                     | Water column depth (m)  |
| Trough                    | Chiniak or Barnabas troughs east of Kodiak Island   |
| Effort                    | Distance (nmi) per grid cell during which observers were manning 'big eyes' and whale search effort was 'on'                                |
| Mean-weighted depth (MWD) | Water-column depth (m) at which the majority of prey backscatter occurs   |

## 2.5. Oceanographic data

Underway sea surface temperature (SST), salinity from a Sea-Bird Electronics SBE-21 probe, and raw voltage from a Turner Designs 10AU fluorometer were measured through the survey from water obtained from a ship intake on the hull that was located 5 m below the surface of the water. Temperature and salinity profiles were collected at trawl sites from the downcast of a Sea-Bird CTD system, and discrete chlorophyll samples were collected at 10 m intervals from 50 m depth to the surface (Hollowed et al. 2007, Logerwell et al. 2007). Surface chlorophyll samples were used to calibrate the underway fluorometry data and calculate chlorophyll values in  $\mu\text{g m}^{-2}$  for the survey area. Oceanographic data were processed using Matlab version R2015b.

## 2.6. Statistical analyses

Acoustic data, cetacean sighting data, and oceanographic data were collected at different spatial and temporal resolutions (Table 1, Fig. 2). Thus, it was necessary to summarize them on a common spatial scale. We computed semi-variograms of the acoustic backscatter and cetacean count data to estimate the appropriate grid cell size to minimize spatial autocorrelation. These analyses (not shown) revealed that scaling the data to a 4.0 km grid cell would eliminate most of the spatial autocorrelation in both the measurements of acoustic backscatter and the whale sighting data. A 4.0 km grid-cell was used to minimize spatial autocorrelation for both data types and include all of the data points from the parallel transects (Fig. 1). Effort was defined as the distance along a transect within a given grid cell where whale observations were taken. Cells with missing acoustic or environmental data or where

<2.3 km (1.25 nmi, half of the distance between transects) of whale observations were taken were eliminated from further analysis. After this process, 72% of the originally surveyed cells remained. The whale sighting effort coverage in the cells retained in the analysis was 90%.

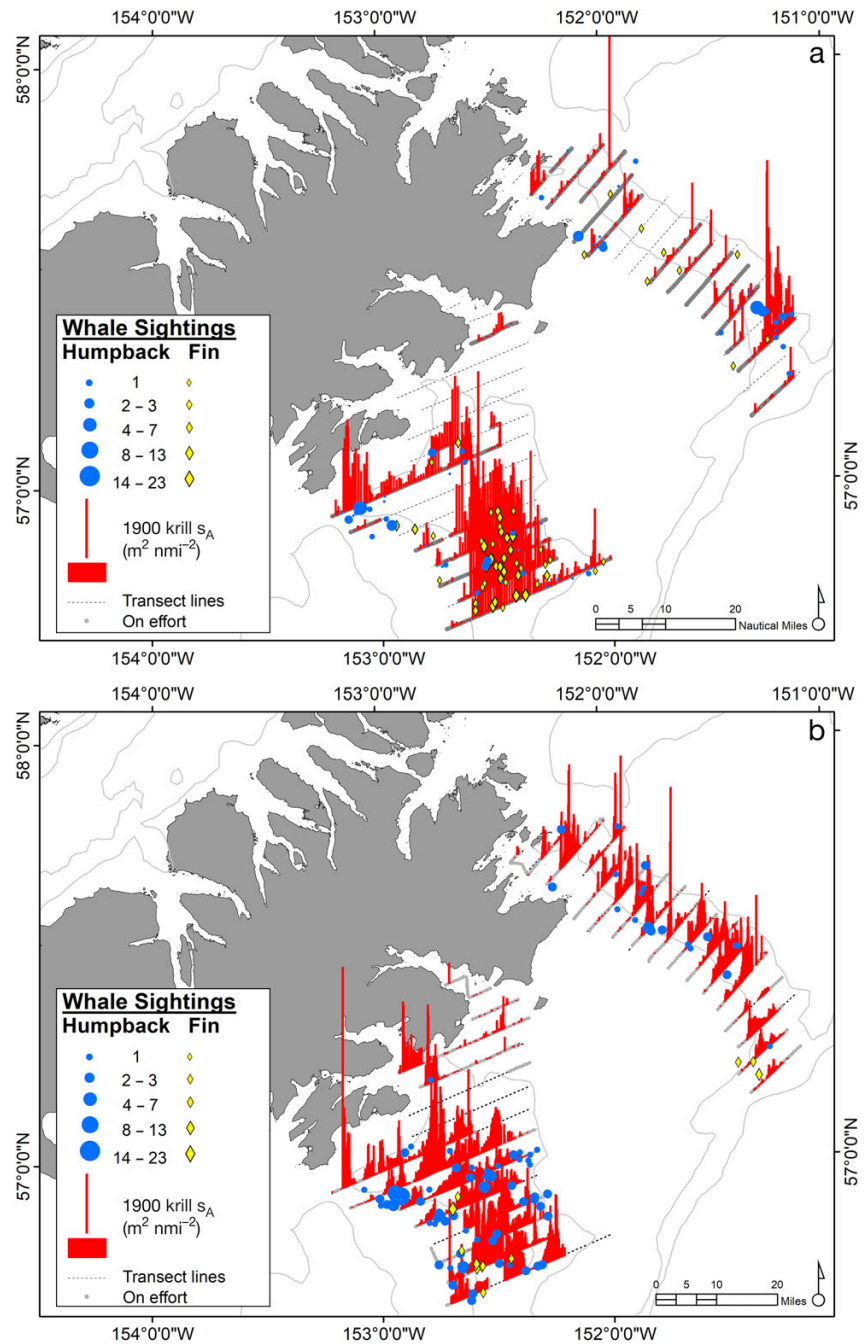


Fig. 2. Euphausiid (krill) acoustic backscatter at 120 kHz ( $S_A$ ;  $\text{m}^2 \text{nmi}^{-2}$ ) and fin and humpback whale sightings by group size during summer (a) 2004 survey passes 7 and 8 and (b) 2006 survey passes 9 and 10 aboard the NOAA Ship 'Miller Freeman'. Pale grey dots: whale observer 'on effort'; dashed lines: transects; light gray lines: 100 and 200 m depth contours

### 2.6.1. Data exploration

We conducted a univariate data exploration to visualize the environmental conditions and prey density under which fin and humpback whales were likely to be found. This visualization allowed us to compare the prey conditions with which whales were associated to those available to the whales within the survey area. Mean environmental conditions, prey density, and presence/absence of fin and humpback whales by grid cell were used in this analysis. To evaluate whether whales were disproportionately present in grid cells with high concentrations of fishes or zooplankton or specific environmental conditions, we compared the cumulative distribution of each of the measured environmental metrics weighted by whale presence to the distribution of these metrics in the entire surveyed area. This produced an index of the minimum level of environmental conditions (e.g. temperature or prey density) in the grid cells at which a given proportion of fin and humpback whale sightings occurred for each of the acoustically measured potential prey fields and each of the environmental variables.

We computed the value of the minimum prey backscatter (or oceanographic variable) that occurred at a given percentile of cells with positive sightings for both fin and humpback whales (De Robertis & Cokelet 2012). To do this, we sorted the whale sightings ( $W$ ), which were paired with prey backscatter or oceanographic measurements ( $E$ ), in ascending order of  $E$ , and computed the minimum values of  $E$  with a given percentile,  $P$  (i.e. 5<sup>th</sup>, 25<sup>th</sup>, 50<sup>th</sup>, 75<sup>th</sup>, 95<sup>th</sup>) of the cells where whales were observed. This can be expressed as the lowest value of the environmental or prey variable in a cell,  $E_j$  that fulfills the following expression:

$$\sum_{i=1}^j W \geq \frac{P}{100} \left( \sum_{i=1}^n W, i \right) \quad (1)$$

where  $i$  and  $j$  are indices into the sorted vector of  $E$ , and  $n$  is the total number of grid cells. We performed an equivalent analysis for each of the prey variables, where  $E$  = prey backscatter (i.e.  $s_A$ ), and for each of the oceanographic measurements, where  $E$  = temperature (°C), chlorophyll ( $\mu\text{g m}^{-2}$ ), or depth (m), and we paired these variables with both fin and humpback whale counts. This allowed a comparison of the prey and ocean conditions in the survey region with the conditions at the locations where whales were observed. Box plots were used to summarize the cumulative distribution of cetaceans in relation to prey backscatter (krill, capelin, and juvenile and

adult pollock), prey depth in the subset of cells where the prey was found, and habitat (oceanographic variables; see Fig. 3). Comparing the 10<sup>th</sup>, 25<sup>th</sup>, 50<sup>th</sup>, 75<sup>th</sup>, and 90<sup>th</sup> percentiles of the environmental measurements in the study area with the conditions under which whales were sighted allows one to visualize whether the whales are disproportionately abundant under a given range of environmental conditions (e.g. are whales associated with high prey abundances?). This portion of the statistical analysis was done using Matlab version R2015b.

### 2.6.2. Multivariable analysis

We developed a multivariable predictive model of whale abundance that simultaneously considered the different environmental and potential prey variables in relation to cetacean sightings. Preliminary analysis indicated that whale count data were overdispersed with respect to the binomial distribution (i.e. additional variance in the response data; see Fig. 3 in Hinde & Demétrio 1998) and were zero-inflated (ZI; Zuur et al. 2009). Initial investigation of the grid-cell-aggregated count data indicated that both fin and humpback whales exhibited distributions that were better characterized by a negative binomial (NB) distribution than a Poisson distribution as the variance was 5 or more times the mean (see Table 2, Fig. 3; Zuur et al. 2009, Preisser et al. 2012). Within the negative binomial family, we considered 2 generalized linear models: NB and zero-inflated negative binomial (ZINB). We performed Vuong's non-nested hypothesis test (Vuong 1989), which indicated that due to the zero-inflation, the ZINB was a better fit for the data.

The ZINB regression model consists of a NB generalized linear model (GLM) used to describe the expected cetacean count and a logit regression used to model the zero observations (Ver Hoef & Boveng 2007, Zuur et al. 2009, 2012). The probability that we measure a certain number of cetaceans in a given grid cell is  $\Pr(Y_i)$ ,  $\Pr(Y_i) \sim \text{ZINB}(\mu_i, \pi_i, k)$ . The variance structure for the ZINB model is  $\text{var}(Y_i) = (1 - \pi_i) \times \mu_i \times (1 + \pi_i \times \mu_i + \mu_i / k)$ , where  $k$  is the dispersion parameter of the negative binomial distribution; and the equation for the mean is  $E(Y_i) = \mu_i \times (1 - \pi_i)$ . The count portion (NB distributed) of the model represented by  $\mu_i$  (Eq. 2) predicts the expected cetacean observed count, and the zero portion of the model with probability  $\pi_i$  (Eq. 3) predicts only the excess zeros (Zeileis et al. 2008, Zuur et al. 2009, 2012). The predictor variables are included in both portions of the model as shown:

$$\mu_i = e^{\alpha + \beta_1 \times X_{i1} + \dots + \beta_q \times X_{iq}} \tag{2}$$

where  $X$  represents a matrix of the prey and environmental variables (i:q) included in Table 1,  $\alpha$  is the population intercept, and  $\beta$  is vector of slopes. In the zero portion of the model,  $\nu$  is the intercept and  $\gamma$  is a vector of slopes:

$$\pi_i = \frac{e^{\nu + \gamma_1 \times X_{i1} + \dots + \gamma_q \times X_{iq}}}{1 + e^{\nu + \gamma_1 \times X_{i1} + \dots + \gamma_q \times X_{iq}}} \tag{3}$$

The combined model prediction, the mean of the ZI and NB components presented together, represents the expected whale count observed in a given cell (Lambert 1992, Martin et al. 2005, Zuur et al. 2009, Otero et al. 2016). The predictor variables in the full model include potential prey and environmental predictors (Table 1), with MWD and acoustic proxies for abundance included for each of the prey variables. Basin (i.e. Chiniak or Barnabas) was included as a factor variable to account for differences in whale abundance between basins (see Table 2). The relationships for other covariates and whale abundance were similar when models were fit to each basin separately, which indicates a lack of strong interactions with basin, although confidence intervals were broader. MWD for the potential prey species can only be included in the model where the prey are present, and cells where a given prey category was not present (11% of cells) were excluded. We used stepwise backward selection based on Akaike’s information criterion (AIC; Akaike 1987) to obtain a final model. AIC was calculated for the full model with the 2 whale species as response variables. Second, reduced models were run by removing one predictor variable at a time. If the AIC of the reduced model was lower than the AIC of all other reduced models and the full model, that variable was removed. This

process continued until no further reduction in the AIC was achieved.

The predictor effects from final models are presented as partial dependence plots highlighting individual variables (i.e. plots where all other predictors are kept constant at their means, Minami et al. 2007; Hastie et al. 2009). These partial dependence plots represent the dependence of predicted whale count on individual variables which were significant in the final model. The predicted effects are shown over the observed range of values of the predictors during the surveys. Analyses, treatment, and visualization of data for this portion of the analysis were performed in R v.3.4.9 using packages ‘pscl’ (v.1.4.9; Zeileis et al. 2008), ‘MASS’ (v.7.3-49; Venables & Ripley 2013), ‘AER’ (v.1.2-7; Kleiber et al. 2020), and ‘ggplot2’ (v.3.2.0; Wickham et al. 2016).

### 3. RESULTS

#### 3.1. Biological summary

Overall, sightings of whales were high for the entire survey, and there were more than 4 times as many humpback whales ( $n = 1055$ ) as fin whales ( $n = 247$ ) in the area around Kodiak Island. In 2004, humpback whales were sighted at a rate of 0.66 animals  $\text{nmi}^{-1}$  of search effort, and fin whales at a rate of 0.20 animals  $\text{nmi}^{-1}$ . In 2006, humpback whales were sighted at a rate of 0.38 animals  $\text{nmi}^{-1}$  of search effort, and fin whales at a rate of 0.05 animals  $\text{nmi}^{-1}$  (Table 2). Humpback and fin whales were more abundant in Chiniak, where over both years of the study, humpback whales were sighted at a rate of 0.61  $\text{nmi}^{-1}$ ,

Table 2. Basic statistics (mean [SD]) for humpback and fin whale counts, krill and fish acoustic backscatter ( $s_A$ ;  $\text{m}^2 \text{nmi}^{-2}$ ), and oceanographic variables for surveys off Kodiak Island in Barnabas and Chiniak Troughs in summer 2004 and 2006. Backscatter cannot be summed over the survey area by year

|   | Barnabas 2004   | Chiniak 2004    | Total | Barnabas 2006   | Chiniak 2006    | Total |
|---|-----------------|-----------------|-------|-----------------|-----------------|-------|
| Humpback whales (sightings per grid cell) | 1.170 (3.33)    | 1.780 (4.45)    | 566   | 0.410 (1.38)    | 1.180 (5.3)     | 489   |
| Fin whales (sightings per grid cell)      | 0.050 (0.22)    | 0.710 (1.91)    | 171   | 0.020 (0.17)    | 0.210 (1.39)    | 76    |
| Effort (km)                               | 2.200 (0.42)    | 2.320 (0.32)    | 1530  | 2.320 (0.45)    | 2.340 (0.37)    | 2454  |
| Krill                                     | 102.680 (137.1) | 284.230 (328.9) | –     | 85.180 (175.8)  | 168.220 (210.1) | –     |
| Adult pollock                             | 110.420 (212.9) | 179.000 (401.7) | –     | 63.610 (124.7)  | 52.490 (152.1)  | –     |
| Age-0 pollock                             | 215.800 (439.4) | 237.560 (553)   | –     | 603.920 (902.8) | 310.600 (787.1) | –     |
| Capelin                                   | 98.280 (290.9)  | 21.100 (81.08)  | –     | 315.250 (596.7) | 63.280 (295.4)  | –     |
| Chlorophyll ( $\mu\text{g m}^{-2}$ )      | 12.550 (7.29)   | 8.980 (3.45)    | –     | 9.490 (5.64)    | 8.050 (3.08)    | –     |
| SST ( $^{\circ}\text{C}$ )                | 10.310 (0.43)   | 10.750 (0.4)    | –     | 10.460 (0.5)    | 10.820 (0.41)   | –     |



and fin whales were sighted at a rate of  $0.17 \text{ nmi}^{-1}$ . In Barnabas, humpback whales were sighted at a rate of  $0.30 \text{ nmi}^{-1}$  and fin whales were sighted at a rate of  $0.02 \text{ nmi}^{-1}$ .

In the context of population assessments of humpback and fin whale populations in the North Pacific (Muto et al. 2018), our sighting rates are relatively high, suggesting high overall population numbers in the study area during our study period. In a survey of the southeastern Bering Sea in 2000, humpback whales were sighted at a rate of  $0.0033 \text{ nmi}^{-1}$ , and fin whales were sighted at a rate of  $0.024 \text{ nmi}^{-1}$  (Moore et al. 2002). Between 2001 and 2003, a survey of the Aleutian Islands and GOA found  $0.035 \text{ fin whales km}^{-2}$  ( $0.12 \text{ nmi}^{-2}$ ) and  $0.054 \text{ humpback whales km}^{-2}$  ( $0.18 \text{ nmi}^{-2}$ ) in the Kodiak Island region (Zerbini et al. 2006).

The composition of potential prey items differed between survey years. Higher backscatter from adult pollock and euphausiids was observed in 2004 than in 2006 (Table 2), but higher backscatter from capelin and age-0 pollock was observed in 2006 (Table 2). During the 2004 survey, adult pollock was the predominant species caught by weight in midwater trawls (95.3%) and age-0 pollock were the most abundant by numbers (32.0%), followed by adult pollock (23.9%), euphausiids (20.7%), and capelin (14.0%). Euphausiids were the predominant catch in Methot trawl hauls, both by weight (68.4%) and numbers (99.2%). The remainder of the Methot catch by weight consisted of large jellyfish (Scyphozoa; 31%) unlikely to contribute to the acoustic observations (De Robertis & Taylor 2014). During the 2006 survey, adult pollock made up more than half of the catch in midwater trawls by weight (51.0%), while age-0 pollock were the most abundant by numbers (31.0%), followed by age-1 pollock (22.0%), capelin (20.2%), euphausiids (10.5%), and adult pollock (4.6%). Euphausiids again were the predominant catch in Methot trawl hauls, both by weight (72.1%) and numbers (99.9%). Jellyfish (Scyphozoa) composed the majority of the remaining Methot catch (27%).

A subset of Methot hauls were assessed for species composition, 6 of 15 methot tows in 2004 and 2 of 4 methot tows in 2006. *Thysanoessa inermis* and *T. spinifera* dominated these euphausiid catches in both 2004 and 2006. In 2004, *T. inermis* (mean length 21 mm) accounted for 67% of the catch and *T. spinifera* (mean length 16 mm) accounted for 31% of the catch. In 2006, *T. inermis* (mean length 17 mm) accounted for 95% of the catch and *T. spinifera* (mean length 26 mm) accounted for 4%, which was similar to

Methot samples from the same region in 2011 and 2013 (Simonsen et al. 2016).

In both years, most adult pollock were detected as dense near-bottom aggregations. Age-0 pollock (5–10 cm SL) were often spatially co-located with capelin. These mixed aggregations of age-0 pollock and capelin typically occurred higher in the water column than adult pollock and were broadly distributed throughout Chiniak Trough and predominantly in the northern portion of Barnabas Trough.

### 3.2. Exploratory analysis

Fin whale sightings were restricted largely to the areas with the highest euphausiid backscatter in the survey area, while humpback whales were found over a wider range of available prey concentrations (Fig. 3). Fin whales were consistently observed in areas with the deepest euphausiid aggregations (Fig. 3c,d). Humpback whales tended to occur in areas with higher euphausiid concentrations, though to a lesser degree than fin whales, and humpbacks were also more frequently sighted in areas with higher age-0 pollock concentrations than those in the surveyed area (Fig. 3a,b).

Fin whale sightings were almost entirely restricted to the top 50% of euphausiid backscatter available in grid cells in both 2004 (Fig. 3a) and in 2006 (Fig. 3b). Fin whale sightings occurred more often in the deepest 65% of the euphausiid depth distribution by cell in 2004 and in the deepest 60% of the euphausiid depth distributions in 2006 (Fig. 3c,d). Fin whales were primarily distributed in deep water. In 2004, 90% of the fin whale sightings occurred in water >150 m depth, which accounts for 35% of the survey area (Fig. 3g); in 2006 the same trend continued, with 50% of fin whale sightings occurring in water deeper than 150 m (Fig. 3j). Although the total environmental variation in SST was only about  $1.5^\circ\text{C}$ , fin whales were more likely to occur in warmer water in both years (Figs. 3e,h). Fin whales were also seen more often in the deepest part of the adult pollock depth distribution in 2006 (Fig. 3b), although this is likely due to fin whales targeting deep euphausiid aggregations, which co-occur with adult pollock.

Humpback whale sightings in both 2004 and 2006 occurred more often in the highest 50% of the euphausiid backscatter in the survey area; however, this pattern in distribution was less pronounced than it was for fin whales (Fig. 3a,b). Humpback whales were observed more often in the upper 50% of the age-0 pollock backscatter in the surveyed area in

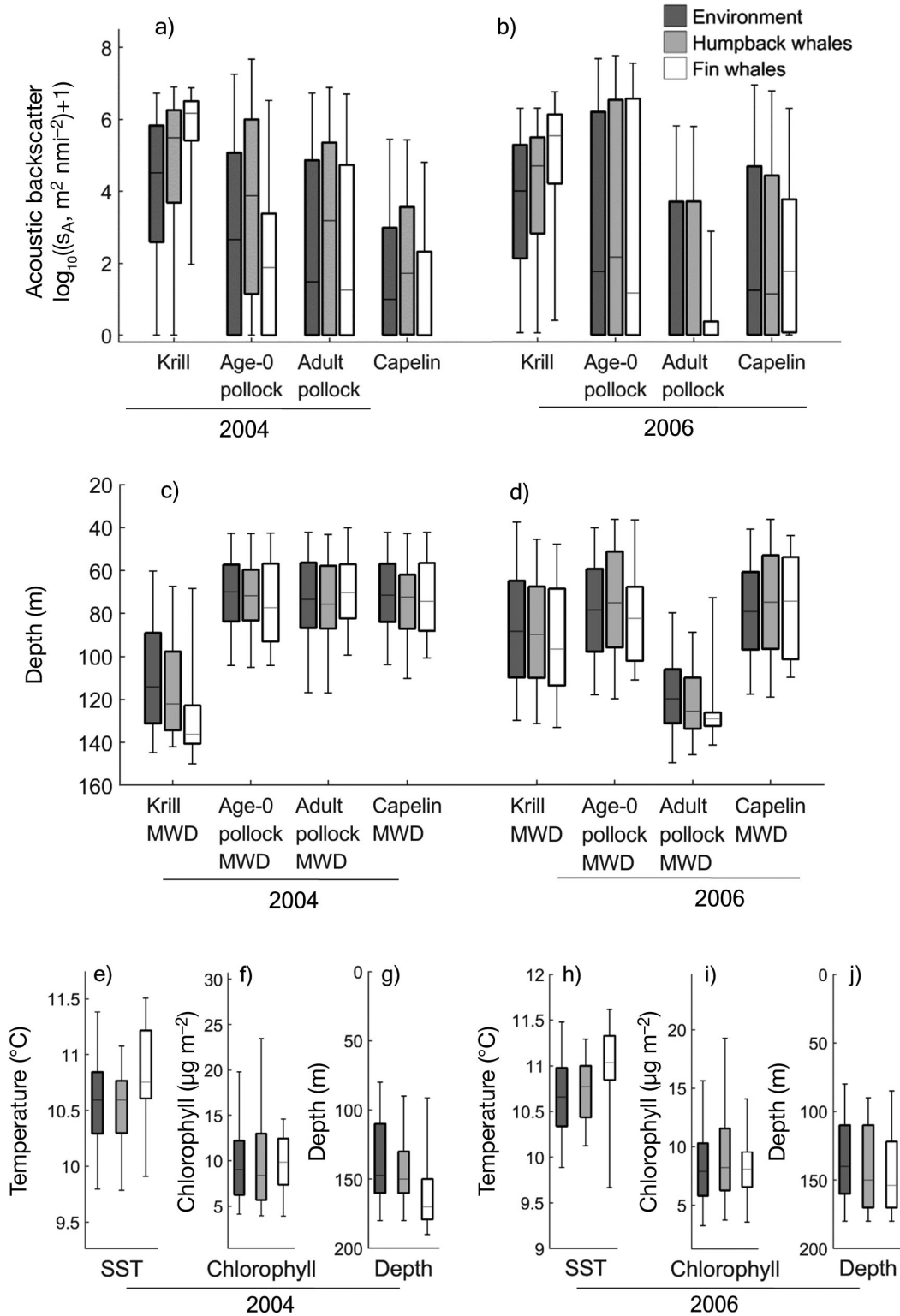


Fig. 3. Conditions in the environment sampled and in areas where fin and humpback whales were observed. Box plots show the environmental features at which the 5<sup>th</sup>, 25<sup>th</sup>, 50<sup>th</sup>, 75<sup>th</sup>, and 95<sup>th</sup> percentiles of total whale sightings occurred for prey backscatter in (a) 2004 and (b) 2006; prey depth in (c) 2004 and (d) 2006; and sea surface temperature (SST), chlorophyll concentration, and depth in (e–g) 2004 and (h–j) 2006, respectively. ‘Environment’ box plots represent the same percentiles for the entire survey area. Cases where the distributions in the sampled environment differ from those where whales were detected indicate that the whales are disproportionately present under a subset of environmental characteristics

2004 (Fig. 3a) and tended to be observed more in areas with higher age-0 pollock concentrations in 2006 (Fig. 3b). Humpback whales were observed in a wider range of depths than fin whales in both years of the survey (Fig. 3g,j), and did not show noticeable trends in distribution relative to SST. Neither species of whale was associated with increased chlorophyll concentration (Fig. 3f,g).

### 3.3. Multivariable analysis

The final model for prediction of whale sightings for both fin and humpback whales included the following predictors: krill density, krill depth, age-0 pollock density, water column depth, trough (i.e. Barnabas vs. Chiniak), search effort, and year in multivariable predictive models with cetacean count as the response. Final model results are presented in Table 3. The structure of the 2 final models predicting humpback (Eq. 4) and fin (Eq. 5) whale distributions are below. The terms pertaining to the count portion of the model are given in black, and those in the zero-inflated portion in grey. In both cases, search effort had little impact on AIC but was retained in the model:

$$Hcount \sim krill + krill\_mwd + pk0 + factor(trough) + factor(year) | krill + pk0 + effort \tag{4}$$

$$Fcount \sim krill\_mwd + factor(trough) | krill + factor(year) + Beaufort + effort \tag{5}$$

For fin and humpback whales, trough was significant in predicting the whale count per grid cell, with fewer whales of either species in Barnabas and a lower total count in 2006 than in 2004 (Tables 2 & 3, see Figs. 4–8). Models for both fin and humpback whales included year as a factor, and residuals plotted relative to predictor values for each of the 2 years were similar, suggesting that the magnitude and trend of predictors of whale presence were similar in both years. Predicted sightings of both fin (Fig. 4) and humpback (Fig. 5) whales increased with euphausiid backscatter, although the relationships differed. In the case of fin whales in Chiniak, for example, a steep increase in predicted sightings occurred between euphausiid  $s_A$  of 500 and 1000  $m^2 nmi^{-2}$  (Fig. 4), whereas predicted sightings for humpbacks increased gradually and steadily with greater euphausiid concentrations in both troughs (Fig. 5). While bootstrapped 95% CIs are wide for these models, trends persist throughout those CIs. Model

Table 3. Model components and model fit for zero-inflated negative binomial models of humpback and fin whale counts. MWD: mean-weighted depth; \*\*\*p < 0.0001; \*\*p < 0.001; \*p < 0.05;  $\dot{p}$  < 0.1

| Parameter  | Estimate | 95 % CI |        | z-value | Pr(> z )       |
|--|----------|---------|--------|---------|----------------|
| <b>Humpback whale model</b>                                  |          |         |        |         |                |
| Count model coefficients (negative binomial with log link)   |          |         |        |         |                |
| (Intercept)  | 1.646    | 0.485   | 2.808  | 2.778   | 0.005**        |
| Krill  | 0.001    | 0.000   | 0.001  | 2.171   | 0.030*         |
| Krill MWD  | -0.013   | -0.022  | -0.003 | -2.681  | 0.007**        |
| Age-0 pollock  | 0.000    | 0.000   | 0.000  | 1.799   | 0.072 $\dot{}$ |
| Trough (factor)  | 0.632    | 0.190   | 1.074  | 2.800   | 0.005**        |
| Year (factor)  | -0.562   | -1.018  | -0.107 | -2.421  | 0.015*         |
| Zero-inflation model coefficients (binomial with logit link) |          |         |        |         |                |
| (Intercept)  | 2.573    | 0.516   | 4.630  | 2.452   | 0.014*         |
| Krill  | -0.007   | -0.013  | 0.000  | -1.930  | 0.054 $\dot{}$ |
| Age-0 pollock  | -0.001   | -0.001  | 0.000  | -2.505  | 0.012*         |
| Effort   | -0.845   | -1.753  | 0.062  | -1.825  | 0.068 $\dot{}$ |
| <b>Fin whale model</b>                                       |          |         |        |         |                |
| Count model coefficients (negative binomial with log link)   |          |         |        |         |                |
| (Intercept)  | -4.311   | -6.016  | -2.585 | -4.899  | 0.000***       |
| Krill MWD  | 0.023    | 0.011   | 0.035  | 3.839   | 0.000***       |
| Trough (factor)  | 1.866    | 1.007   | 2.725  | 4.259   | 0.000***       |
| Zero-inflation model coefficients (binomial with logit link) |          |         |        |         |                |
| (Intercept)  | 4.072    | 0.929   | 7.215  | 2.539   | 0.011*         |
| Krill  | -0.004   | -0.006  | -0.002 | -3.771  | 0.000***       |
| Year (factor)  | 1.283    | 0.507   | 2.058  | 3.242   | 0.001**        |
| Beaufort   | -0.314   | -0.597  | -0.032 | -2.184  | 0.029*         |
| Effort   | -0.675   | -1.909  | 0.559  | -1.072  | 0.284          |

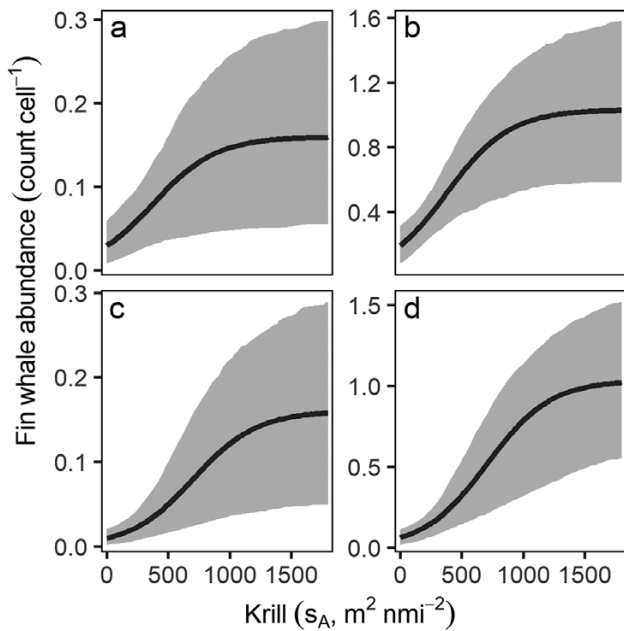


Fig. 4. Predicted fin whale sightings with krill backscatter ( $s_A$ ) varied and other model components held constant. Shading represents bootstrapped 95% CIs. (a) Barnabas and (b) Chiniak basins in 2004; (c) Barnabas and (d) Chiniak basins in 2006

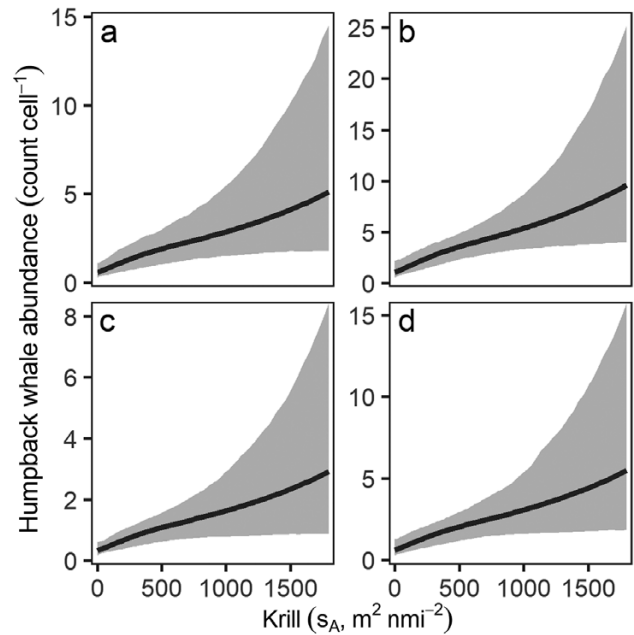


Fig. 5. Predicted humpback whale sightings with krill backscatter ( $s_A$ ) varied and other model components held constant. Shading represents bootstrapped 95% CIs. (a) Barnabas and (b) Chiniak basins in 2004; (c) Barnabas and (d) Chiniak basins in 2006

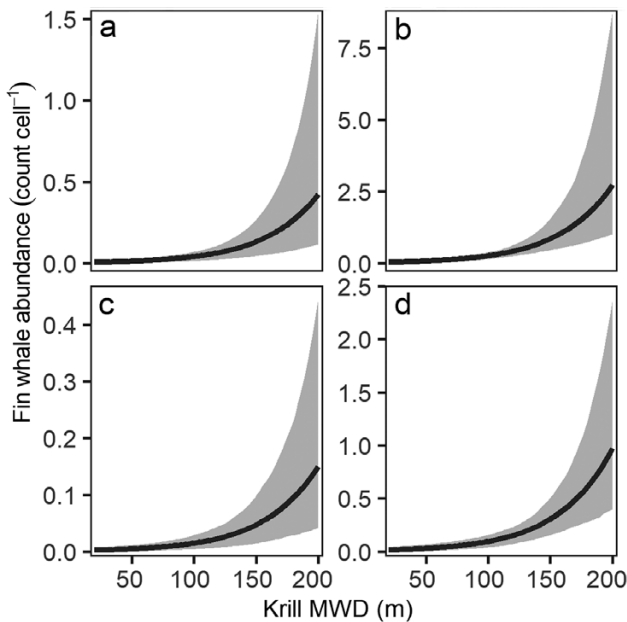


Fig. 6. Predicted fin whale sightings with krill mean-weighted depth (MWD) varied and other model components held constant. Shading represents bootstrapped 95% CIs. (a) Barnabas and (b) Chiniak basins in 2004; (c) Barnabas and (d) Chiniak basins in 2006

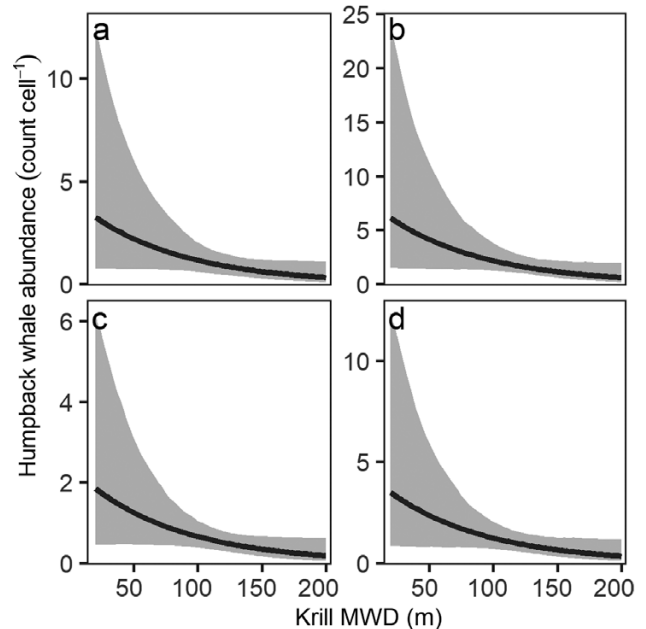


Fig. 7. Predicted humpback whale sightings with krill mean-weighted depth (MWD) varied and other model components held constant. Shading represents bootstrapped 95% CIs. (a) Barnabas and (b) Chiniak basins in 2004; (c) Barnabas and (d) Chiniak basins in 2006

effects for the predicted whale sightings differed with respect to euphausiid depth (Figs. 6 & 7). That is, increased euphausiid MWD was associated

with increased fin whale sightings, but decreased humpback sightings. Finally, humpback whales were associated with higher age-0 pollock densities

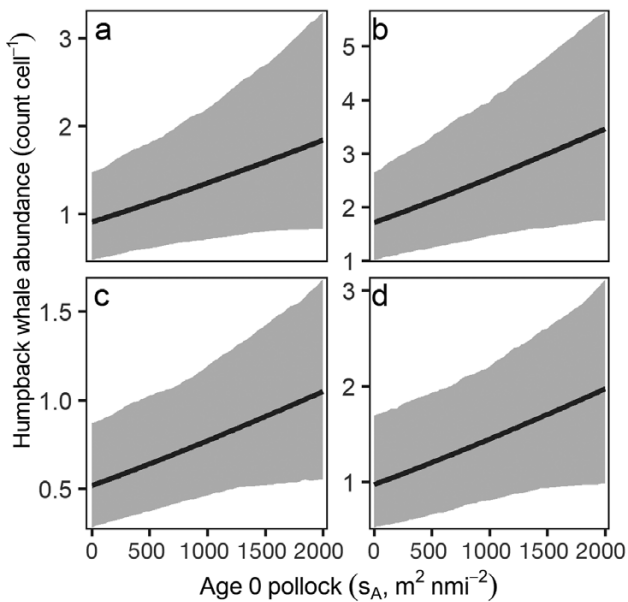


Fig. 8. Predicted humpback whale sightings with age-0 pollock  $s_A$  varied and other model components held constant. Shading represents bootstrapped 95% CIs. (a) Barnabas and (b) Chiniak basins in 2004; (c) Barnabas and (d) Chiniak basins in 2006

(Fig. 8), whereas age-0 pollock was dropped from the fin whale model (Table 3).

#### 4. DISCUSSION

Concurrent observations of the distribution of whales and their prey suggest that fin and humpback whales engage in spatial and trophic niche partitioning at their feeding grounds near Kodiak, Alaska. The most notable contrast in how fin and humpback whales shared available habitat in the study area was seen in exploratory analyses suggesting that fin whales targeted the deepest, densest patches of euphausiids available, while humpback whales were associated with shallower and lower-density aggregations of euphausiids. Building on the exploratory analyses, multivariable predictive models suggested that euphausiid depth was an important predictor of fin whale sightings. Combined, the exploratory and multivariable approaches provide evidence that fin whale habitat use in this region differs from that of humpbacks. This may be due to the ability of fin whales to more efficiently target deeper prey patches due to their larger body size compared to humpback whales. Recent work examining diving behavior relative to prey patch depth suggests that whale foraging behavior is driven in part by species-specific

morphological differences (Friedlaender et al. 2015, Keen & Qualls 2018).

Sightings of both fin and humpback whales were related to euphausiid abundance near Kodiak, but analyses suggested that the whales may have different foraging thresholds. Fin whales were associated with the highest euphausiid abundances in the area, and predicted fin whale sightings increased steeply at intermediate levels of euphausiid backscatter (500–1000  $\text{m}^2 \text{nmi}^{-2}$ ), which suggests that they exhibited threshold foraging on euphausiid patches. The relationship between euphausiid backscatter and predicted humpback whale sightings was linear and more gradual than it was for fin whales, suggesting a lower foraging threshold. This contrast could be related to species-specific differences in energetic requirements and ability to efficiently forage at lower prey concentrations. Larger body size means that fin whales require more energy than humpback whales, so they must target the densest available prey patches to be efficient (Croll et al. 2001, Goldbogen et al. 2007, 2012). Comparatively, humpbacks generally exploit lower density prey aggregations with lower search costs rather than seeking out deeper, higher density prey aggregations (Friedlaender et al. 2009b, Tyson et al. 2016, Fossette et al. 2017, Keen & Qualls 2018). Recent work also suggests that fin whales and other large cetaceans are optimized for deeper diving to target dense prey (Hazen et al. 2015, Keen 2017, Keen & Qualls 2018). The combined factors of higher energetic requirements and more efficient foraging at depth may account for differences in associations seen in the present study.

The concentration of forage fishes near Kodiak had little effect on the presence of fin whales, but humpback whales were more likely to be found in areas where age-0 pollock were abundant. Although fin whales consume fishes in other areas (Nemoto 1959, Pauly et al. 1998, Sigler et al. 2012), we detected no relationship between fin whales and forage fishes in our study area. Given that fin whales were strongly associated with dense euphausiid patches, but not with forage fish, suggests prey partitioning by fin whales. In contrast, we observed that humpback whales were spatially associated with both euphausiids and small fishes. Differences in the prey partitioning strategies of fin and humpback whales are supported by prior work in the Kodiak region (Witteveen et al. 2015, Witteveen & Wynne 2016, Wright et al. 2016). Humpback whales also demonstrate an ability to switch target prey with changing prey availability (Friedlaender et al. 2009a, Hazen et al. 2009, Fleming et al. 2016), and this may account for findings in this

study. Historically, there have been reports of humpback whale predation on juvenile pollock (15–30 cm SL; Nemoto 1959, Kawamura 1980), but we are aware of no published work that addressed patterns in their distribution in relation to age-0 pollock (SL  $\leq$  10 cm). While other studies have suggested that humpback and fin whales may also exploit oily fish like capelin (Krieger & Wing 1984, Piatt & Methven 1992, Witteveen et al. 2015, Fleming et al. 2016, Wright et al. 2016), this was not observed in our study despite some seasonal overlap. There may be several explanations for why euphausiids, and not capelin, appear to be a driver of whale distributions near Kodiak. First, euphausiids are relatively abundant near Kodiak (Simonsen et al. 2016), and this may attract whales to the area. Second, while capelin consistently occur on the Kodiak shelf, they exhibit high spatial, temporal, and interannual variability (Jones et al. 2014, McGowan et al. 2019, 2020), and therefore they may be more costly to exploit, in spite of their high caloric content. Finally, it is possible that the capelin densities, even those in patches within the study area, were insufficient to make foraging for them energetically viable. Future studies of the predator–prey interactions of these species could be helpful in elucidating these patterns.

While prey abundance was an important predictor of fin and humpback whale presence, environmental variables were not. In this study, SST and chlorophyll were not related to whale distribution, but have been useful predictors in larger scale, basin-wide studies (Gregar & Trites 2001, Moore et al. 2002, Zerbini et al. 2016). Within our survey area, Chiniak Trough consistently had higher numbers of whales of both species, possibly due to higher fishing pressure in Barnabas. Despite the difference in magnitude, the predictive trends in the models persist between the 2 troughs. Given that prey availability changes within this survey area over both seasonal (Hollowed et al. 2007, Logerwell et al. 2007) and longer time scales (Anderson & Piatt 1999), and that environmental conditions change over short and long time scales, there is potential that the relationships shown by this work have shifted over time.

This study provides evidence that sympatric fin and humpback whales near Kodiak Island, Alaska, exhibit spatial and trophic prey partitioning through different foraging strategies and efficiencies. This work demonstrated the importance of simultaneously collecting acoustic-based estimates of prey and visual estimates of cetaceans, and future applications of this approach will likely lead to improvements of the predictive models needed for conservation and

management. Evidence for niche partitioning observed here, together with previous observations (Friedlaender et al. 2006, Witteveen et al. 2015), suggests a mechanism that reduces or eliminates interspecific competition between these frequently co-occurring Balaenopteridae. More work is needed to determine whether foraging strategies that reduced competition in this study are used by fin and humpback whales elsewhere and also to determine short- and long-term temporal trends in niche partitioning by these 2 species in the Kodiak Island region.

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