- 1 Decreased lipid storage in juvenile Bering Sea crabs (*Chionoecetes* spp.) in a warm (2014)
- 2 compared to a cold (2012) year on the southeastern Bering Sea.

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Abstract

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The decline of eastern Bering Sea snow (Chionoecetes opilio) and Tanner (Chionoecetes bairdi) crab has coincided with loss of spring sea ice extent and northward contraction of the 'cold pool', a layer of cold (< 2 °C) summer bottom water. We measured temperature-associated growth and lipid storage of lab-reared Tanner crab, as well as the fatty acid content of fieldcollected juvenile *Chionoecetes* spp. Field collections occurred during a cold, high sea ice year (2012) and a warm, low ice year (2014), representative of cold and warm climate stanzas in the southeastern Bering Sea. Lab-reared Tanner crab maintained the lowest growth rates but highest lipids under cold conditions (2 °C). In the field, crabs contained higher fatty acids per wet weight (mg.g⁻¹) during 2012 than during 2014. Water column integrated chlorophyll a (chla, an indicator of phytoplankton biomass) from large particles (>10 µm) such as diatoms, was elevated in the colder year particularly over the central middle shelf. During the cold-year, crab storage of diatom-sourced fatty acids (16:1n-7 to 16:0, r²=0.72) as well as a station-specific relationship between large size-fraction integrated chla and crab total fatty acids ($r^2=0.5$) points to the potential importance of diatoms to juvenile crab nutrition. Our results suggest that continued warming and loss of sea ice across the Bering Sea may reduce juvenile crab lipid storage through both direct thermal effects as well as through the reduction of large size phytoplankton delivered to the benthos.

Key Words

39 Lipid, fatty acids, diatom, chlorophyll a, sea ice

Introduction

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Stocks of most major crab species in the North Pacific (red king crab, *Paralithodes* camtschaticus; blue king crab, P. platypus; snow crab, Chionoecetes. opilio, and Tanner crab, Chionoecetes bairdi) have been in decline for the last three decades (Armstrong et al. 1998; Orensanz et al. 1998; Woodby et al. 2005). Declining snow crab stocks have resulted in major fisheries reductions with a 50% drop in the total allowable catch for 2017, making this the lowest value for snow crab since 1982 (NPFMC 2018). The reason for Alaskan crab declines is poorly understood, but is generally thought to be due to both over-fishing and a changing climate (Szuwalski et al. 2020; Zheng and Kruse 2000; Zheng and Kruse 2006). On the southeastern Bering Sea (SEBS) shelf, south of 60N, two commercially important congener species of Chionoecetes overlap in their distributions. Snow crab is an Arctic species that occupies deep waters in the North Atlantic, North Pacific, and throughout the Arctic region; Tanner crab is a boreal species that ranges from Puget Sound, WA to the SEBS in the north. Recent climate warming (1982 to 2017) has resulted in a shift in the distribution of these species with adult snow crab moving northward and Tanner crab moving to deeper habitats to avoid rapidly warming bottom water and increased predation from northward moving ground fish (Landeira et al. 2018; Murphy 2020; Orensanz et al. 2004).

Temperature is the most important environmental variable controlling vital rates in ectotherms and has previously been shown to dramatically affect growth rates in many juvenile Alaskan crab species (Ryer et al. 2016; Stoner et al. 2013; Stoner et al. 2010). It has been suggested that juvenile snow crab have very narrow temperature tolerances, being generally restricted to bottom waters < 3 °C (Dionne et al. 2003), while Tanner crab juveniles prefer warmer conditions and avoid temperatures < 2 °C (Ryer et al. 2016), making both species

susceptible to climate-mediated changes in bottom water temperatures. The SEBS has experienced a number of warm (2002-2005 and 2014-2019) and cold stanzas (2006-2013) during the recent period of environmental monitoring (Duffy-Anderson et al. 2019; Eisner et al. 2016; Mueter and Litzow 2008; Stabeno and Bell 2019; Stabeno et al. 2012; Stevenson and Lauth 2019). The changing climate in the SEBS also influences sea ice dynamics and the resulting formation of the 'cold pool', a layer of cold (<2 °C) subsurface water (below the pycnocline) that forms over the middle shelf (50 to 100 m depth) during sea ice retreat in the spring (Wyllie-Echeverria and Wooster 1998). In cold years with extensive sea ice cover, the cold-pool can extend south to ~56 °N while during warm periods with reduced sea ice cover the cold pool retreats northward to >60 °N. In recent years (2018 & 2019), this retraction has been so extreme that the cold-pool has been completely absent from the SEBS and most of the north Bering Sea benthic system (Stabeno and Bell 2019; Stevenson and Lauth 2019).

The relative importance of climate-influenced, bottom up trophic cascades to juvenile crab condition and recruitment remains largely unknown. The Oscillating Control Hypothesis provides a framework to examine the complex, indirect food web effects on marine species dynamics in the Bering Sea, but focusses on pelagic food web dynamics. This theory links climate oscillation and variable sea ice extent to the timing of the spring bloom and subsequent energetic condition and recruitment of juvenile walleye pollock (*Gadus chalcogramma*) and Pacific cod (*Gadus macrocephalus*) (Coyle et al. 2011; Eisner et al. 2020; Farley et al. 2016; Heintz et al. 2013; Hunt et al. 2011; Hunt et al. 2002; Hunt et al. 2008; Sigler et al. 2014). However, we have less understanding of how these environmental processes affect early life history stages of juvenile crabs. This is because annual assessments of snow and Tanner crab over the SEBS focus on larger crabs with a mean carapace width of 48 mm and 4 to 7 years post

settlement age (Murphy 2020). Larger crabs likely have a nutritional condition that is integrated over multiple seasons and is less sensitive to short-term pulses in phytoplankton production.

Juvenile stages of fish and crabs are characterized by lower energy storage and faster metabolic rates which makes them relatively more sensitive to changes in food quantity/quality and more vulnerable to starvation (Beder 2015; Copeman et al. 2014; Houde 2008; Hurst 2007).

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Increased warming and loss of sea ice is forecast to decrease both food quality and quantity supplied to the benthos through decreased flux from ice-edge and pelagic phytoplankton blooms (Grebmeier 2012; Grebmeier et al. 2015; Renaud et al. 2015; Wassmann and Reigstad 2011). Benthic-pelagic coupling represents an essential source of energy, lipids and more specifically essential fatty acids for benthic fauna (Copeman and Parrish 2004; Ramos et al. 2003; Richoux et al. 2004). Characterization of fatty acid (FA) pools in benthic organisms has become increasingly important in ecological studies because FAs can provide time-integrated information about dietary utilization and trophic relationships (Budge et al. 2006; Dalsgaard et al. 2003; Parrish 2013). FA trophic markers are synthesized at lower trophic levels and can link consumers with various sources of primary production such as diatoms, dinoflagellates, terrestrial runoff, calanoid copepods and bacterial sourced organic matter (Budge and Parrish 1998; Copeman et al. 2009; Dalsgaard et al. 2003; Parrish 2013). Transfer of FAs from lower to higher trophic levels is generally conservative and therefore FAs are used to indicate dietary sources for invertebrates (Galloway et al. 2014; Kelly and Scheibling 2012; Spilmont et al. 2009), fish (Copeman et al. 2016; St John and Lund 1996), and mammals (Budge et al. 2008; Tucker et al. 2009). Sinking phytoplankton contain high proportions of polyunsaturated fatty acids (PUFA) that are essential to juvenile stages of many different benthic fauna (Kelly and Scheibling 2012; Richoux et al. 2005).

Recent laboratory and field observations have demonstrated the significance of both food quantity and food quality on the growth and lipid storage of newly settled Tanner crabs from shallow nearshore embayments in the Gulf of Alaska (Copeman et al. 2018). Here we provide the first data that examines trophic dynamics of juvenile *Chionoecetes* spp. across the SEBS. Our objectives were to examine crab lipid storage across offshore sites on the SEBS during two environmentally disparate years, 2012 (cold) and 2014 (warm). Specifically, we aimed to 1) determine the direct effect of temperature on lipid concentration and growth in newly settled juvenile Tanner crab in the lab, 2) determine spatial and annual differences in phytoplankton biomass (total integrated chla) and large particle phytoplankton biomass (>10 µm integrated chla) which is often indicative of diatoms, 3) determine spatial and annual differences in crab total fatty acid concentrations (mg.g⁻¹) and individual fatty acid proportions in relation to late summer phytoplankton biomass, and 4) discuss crab trophic dynamics and lipid storage in the context of continued warming across the sub-Arctic Bering Sea.

Methods

Live crab collection and laboratory temperature-growth experiment

Details of crab collection protocols can be found in Ryer et al. (2015). Briefly, Tanner crabs (4.2 to 5.6 mm carapace width) were collected from shallow bays surrounding Kodiak Island, Gulf of Alaska, AK, USA using an epi-benthic sled in July of 2013. The codend of the sled was made of 3 mm mesh seine fabric, which retained C2 (molt stage 2) and larger juvenile Tanner crabs. The sled was towed along the seafloor at depths ranging from 10 to 30 m, parallel to shore, at a speed 0.5 m.s⁻¹ for ~30 m. Tanner crabs (~5 mm carapace width) were retained alive and shipped overnight in insulated containers to the Hatfield Marine Science Center (HMSC) in Newport, OR, USA.

At the laboratory, crabs were placed into individual growth cells immersed in temperature-controlled, flow-through seawater tanks as detailed in Copeman et al. (2018) and Ryer et al. (2016). Individual C2 crabs were gradually acclimated to 9 °C and 2 °C and fed a marine gelatinized food (diet composition as detailed in Copeman et al. (2018)). Crabs were fed a quantity of food that was more than their 24-hour satiation ration (based upon previous experiments). Three times per week, remaining food was removed and new food was added to growth chambers. Upon molting (C3) we measured crab carapace width using digital photography and this was considered the beginning of the temperature-growth experiment. Ten replicate crabs were maintained at 9 °C and 2 °C and the experiment was continued until the midmolt point between the C4 and C5 stages. Mid-molt was estimated based upon prior data on the relative increase in intermolt periods from one stage to the next (Ryer et al. 2016). Growth cells, and hence crab placement within temperature controlled water baths were re-randomized on a weekly basis to preclude tank effects. All water baths were provided with continuous flowthrough seawater (salinity ~33) and crabs were checked daily for molting activity. Crabs were re-measured 2 to 5 d after each molt to help insure that crab exoskeletons were sufficiently hard to resist damage during handling. At mid molt of the C4 stage, a randomly chosen subsample of 6 crabs from each temperature treatment was measured, weighed and placed in a vial with 2 mL of chloroform under a layer of nitrogen and frozen (-20 °C) for later lipid extraction (<6 months). Intermolt period was determined in each temperature treatment using the days from C3 to C4. Molt increment, i.e. percent increase in carapace width from one stage to the next, was also analyzed across this same developmental stages. Intermolt period was analyzed to test for temperature effects using a Kruskal-Wallis one-way nonparametric ANOVA while, temperature

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effects on molt increment and total lipids (\sum lipid, mg.g⁻¹) were analyzed using a one-way ANOVA with a significance of $\alpha > 0.05$.

Crab sampling over southeastern Bering Sea 2012 and 2014

Small mesh beam trawling was conducted over the SEBS shelf in 2012 and 2014 as detailed in Hurst et al. (2015) and Ryer et al. (2016). This sampling was conducted to target juvenile crabs on an exploratory basis as part of the larger NOAA Bering-Arctic-Subarctic Integrated Survey (BASIS). Crabs were sampled on the shelf (< 200m depth) from the Alaska Peninsula to 60 °N in August to early October of 2012 and August to September of 2014. A 3-m beam trawl, with 7-mm mesh and a 4-mm mesh codend liner were used to make benthic tows. Global positioning system (GPS) coordinates were used to determine the length of each tow and were paired with the effective fishing width of the net (2.26 m) to calculate catch per unit effort (CPUE, crabs.100 m⁻²). The speed and duration of the trawl was 1.3 m.sec⁻¹ for 5 to 10 minutes. *Chionoecetes* crabs < ~15 mm carapace width (C6 stages and smaller) were sorted, counted and then immediately frozen on-board.

To compare crabs across years but from within similar geographical regions, we divided crabs by latitude and into those from middle and outer shelf habitats. These regions are commonly used due to their distinct physical oceanography and ecosystem dynamics (Ortiz et al. 2016). The four crab sampling regions on the SEBS shelf included: 1) central middle (CM) between the 50 and 100 m bottom depth contours and from between 57 and 60 °N, 2) south middle (SM) between the 50 and 100 m depth contours and from 57 °N south to the Alaskan peninsula, 3) the central outer (CO) between the 100 and 200 m depth contours and offshore from 57 to 56 °N, and 4) south outer (SO) between 100 and 200 m depth contours and from 56

°N south to Unimak Pass. Sample sizes were based on availability and ranged from 12 to 37 crabs per year in a given sampling region (Table 1).

Crab morphometrics and lipid analyses

In 2012 and 2014, crab were sorted from the small mesh beam trawl and were frozen in a -20 °C walk-in freezer on board the research cruise. Following the cruises, crabs were shipped frozen overnight to the Hatfield Marine Science Center, where they were stored in a -80 °C freezer until they were identified, measure and weighed. All crabs were sampled within 6 months of storage at -80 °C. They were placed directly into 2 mL of chloroform under a layer of nitrogen and returned to a -20 °C freezer. All crabs were extracted for lipids within the following 6 months (see below).

For the lipid laboratory analyses, crabs were first identified using the scores of carapace characteristics developed by Urban et al. (2002) to differentiate Tanner crabs (score = 1), snow crabs (score = 5) and possible range of intermediate morphology that represented hybrids (score = 2 to 4). However, especially for the smaller juveniles, we found that typical carapace shape was not sufficiently developed to be reliable, and eye color was not distinguishable, especially after freezing. Therefore, we relied on epistome margin as the diagnostic character. Due to this difficulty in distinguishing species from field sampled juveniles, we grouped all crabs as *Chionoecetes* spp. for most statistical analyses of the effect of region and year on crab condition and biomarker composition. The one exception, is a comparison of crab total fatty acid storage on the central middle shelf, where we examined the differences between years as a function of hybrid status (1- Tanner, 2-Tanner Hybrid, 3- snow hybrid). Here we had a high percentage of stage 3 snow crab hybrids (Urban et al. 2002) which allowed us to look at annual differences as a function of morphologically defined hybrid status. Carapace widths were measured with calipers

to the nearest 0.1 mm. The minimum carapace width in the 2012 samples was 4.33 mm (C3 stage) and 6.08 mm (C4 stage) in 2014. For the juvenile data analysis, we only included crabs <15.3 mm (C6 or under).

Once crabs were weighed and measured (both field samples and laboratory experiments) they were processed for lipid content in the Marine Lipid Ecology Laboratory at Oregon State University, Hatfield Marine Science Center. We did not have excess crabs to process for dry weights (DWT) or for ash-free DWT (AFDWT). The small size of C5 and C6 molt stages precluded splitting crab tissue between WWT, DWT and AFDWT measures. Therefore, we used conversion factors from a previous experiment on Alaskan Tanner crab to express crab total fatty acids per WWT (measured) into crab total fatty acids per DWT and AFDWT (estimated with conversion factors from Copeman et al. (2018), see Table 1). Lipids from whole individual crabs were extracted in chloroform and methanol according to (Parrish 1987) using a modified Folch procedure (Folch et al. 1956). Lipid class analyses were only completed on laboratory temperature experiment crabs while total fatty acids were measured on field collected crabs. The analyses of total fatty acids is faster and more efficient, and there is a strong positive relationship between total lipids and total fatty acids, as previously demonstrated in Tanner crabs collected around Kodiak Island, Alaska (Copeman et al. 2018).

Total lipids were determined using thin layer chromatography with flame ionization detection (TLC/FID) with a MARK V Iatroscan (Iatron Laboratories, Tokyo, Japan) as described by Lu et al. (2008) and Copeman et al. (2017). Extracts were spotted on duplicate silica-gel-coated Chromarods, and a three-stage development system was used to separate wax/steryl esters, triacylglycerols, free fatty acids, sterols and polar lipids. Polar lipid is mostly comprised of phospholipids with minor amounts of other acetone mobile polar lipids. The first rod

development was in a chloroform: methanol: water solution (5:4:1 by volume) until the leading edge of the solvent phase reached 1 cm above the spotting origin. The rods were then developed in hexane: diethyl ether: formic acid solution (99:1:0.05) for 48 min, and finally rods were developed in a hexane: diethyl ether: formic acid solution (80:20:0.1) for 38 min. After each solvent development, rods were dried (5 min) and conditioned (5 min) in a constant humidity chamber (~32%) that was saturated with aqueous CaCl₂. Following the last development, rods were scanned using Peak Simple software (ver. 3.67, SRI Inc.) and the signal detected in millivolts was quantified with calibration curves using the following commercial standards from Sigma (St Louis, MO, USA): cholesteryl stearate (wax ester), glyceryl tripalmitate (triacylglycerols), palmitic acid (free fatty acids), cholesterol (sterols), L-alphaphosphatidylcholine (polar lipids). Calibrated relationships between lipid class areas and standard lipid amounts (μg) had correlations with an r² >0.98 for all classes. Lipid class data did not show significant proportional differences (Appendix 1) so only total lipids (ΣIatroscan total lipids) are shown as absolute amounts per wet weight (mg.g-1).

Prior to derivatization of field-collected crabs, an internal standard, tricosanoic acid methyl ester (23:0), was added to lipid extracts in an amount that was ~10% of the total fatty acids. Total lipid extracts were derivatized to form fatty acid methyl esters (FAME) using acid transesterification in H₂SO₄ in MeOH (Budge et al. 2006). An HP 7890 GC FID equipped with an autosampler and a DB wax GC column (Agilent Technologies, Inc., USA) was used to analyze FAME samples. The DB wax column was 30 m in length, with an internal diameter of 0.25 mm and a film thickness of 0.25 μm. The column temperature began at 65°C, held for 0.5 min, and then raised to 195°C (40°C.min⁻¹), held for 15 min and then increased again (2 °C.min⁻¹) to a final temperature of 220°C and held for 1 min. The hydrogen carrier gas was flowing at a

rate of 2 mL.min⁻¹ and the injector temperature was set at 250°C, while the detector temperature was held constant at 250°C throughout the 31-min run. Peaks were identified using retention times based on standards purchased from Supelco (37 component FAME, BAME, PUFA 1, PUFA 3). Nu-Check Prep GLC 487 (Elysia, MN, USA) quantitative FA mixed standard was used to check column function and chromatograms were integrated using Chem Station (version A.01.02, Agilent).

Statistical differences by year and region of collection were examined for crab total fatty acids per wet weight (WWT, mg.g⁻¹), crab storage of diatom indicator fatty acids (ratio of 16:1n-7 to 16:0, Viso and Marty (1993)) and crab storage of bacterial fatty acids per wet weight (mg.g ¹, \sum odd and branched chains, Kaneda (1991)) using 2-way ANOVAs with Tukey's pairwise comparisons (p<0.05, Minitab 19). We included individual fatty acid proportions > 0.5% in at least one region (Table 1) as well as total fatty acids per WWT in a multivariate analyses using PRIMER v.7 (Primer-E) with a Permutational ANOVA, PERMANOVA add-on package (Primer-E Ltd). We included Σ FA (mg.g⁻¹) in our multivariate analyses of crabs in order to demonstrate fatty acids associated with increased total lipid storage. Fatty acids were squareroot transformed prior to analyses and were then used to calculate a triangular matrix of similarities (Bray-Curtis similarity) between each pair of samples. Non-metric multidimensional scaling (nMDS), an iterative process that uses ranks of similarities, was utilized to explore the effect of region and year of collection on the fatty acid composition of juvenile *Chionoecetes* spp.crabs. We performed a 2-way PERMANOVA with carapace width as a covariate to control for size and examined the significance of region of collection (CM, SM, CO, SO) and year (2012 or 2014) on the fatty acid composition of crabs.

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To compare crab condition and trophic biomarkers to an index of phytoplankton biomass 270 we used water column integrated chla (mg·m⁻²) values collected in 2012 and 2014 from the same 271 272 biennial BASIS survey that hosted crab collections. The BASIS fisheries oceanography surveys (2003 to present) provide valuable data on oceanography, and primary and secondary production 273 274 across the Eastern Bering Sea from mid-August to October (details as in Eisner et al. (2016)). Variations in chla concentrations were used to evaluate spatial and inter-annual differences in 275 total phytoplankton biomass. In addition, the ratio of large size-fraction to total phytoplankton 276 277 biomass (>10 µm chla / total chla) was estimated from discrete water samples filtered through polycarbonate 10 µm and GF/F (nominal pore size 0.8 µm) filters, and analyzed with standard 278 fluorometric methods (Parsons et al. 1984). Increases in the ratio of large sized phytoplankton 279 280 can indicate higher proportions of diatoms and thus may be indicative of diet quality. Total integrated chla (mg.m⁻², over the top 50 m of the water column or to the bottom for shallower 281 stations) was estimated from CTD fluorescence profiles, calibrated with discrete total chla 282 (GF/F) samples. At each station, integrated chla > 10 μm was estimated by multiplying the mean 283 large size fraction ratio (>10 µm chla/ total chla) from discrete samples by the total integrated 284 285 chla value. Total and >10 µm integrated chla at each station were averaged over each of the four regions used to compare juvenile crabs (CM, SM, CO, SO) in both 2012 and 2014 (Table 1). 286 Additionally, we looked at the station-specific relationship between late summer large size 287 288 phytoplankton biomass and crab condition over the shelf in both years and just in 2012 alone. Specifically, we compared the integrated >10 μ m chla to station-specific averages for crab Σ FA 289 (mg.g⁻¹) and the fatty acid diatom indicator (ratio of 16:1n-7 to 16:0) in crab tissue using linear 290 291 regressions (SigmaPlot version 14, p<0.05).

Spatial patterns in bottom temperatures and chla biomass data from survey sampling stations was visualized using Empirical Bayesian Kriging (Krivoruchko and Gribov 2019) using ArcGIS Desktop 10.7. Bottom temperature for the BASIS juvenile crab distribution maps was obtained from CTD casts conducted at survey stations as described in Eisner et al. (2016). Monthly average sea ice extents were obtained from the National Snow and Ice Data Center's Sea Ice Index version 3 (Fetterer et al. 2017 updated daily).

Results

Our laboratory experiments showed that temperature influenced the growth and lipid storage of juvenile Tanner crabs. Temperature increased the rate at which crabs molted with an average day to molt (C3 to C4 stages) of only 37 ± 1 (n = 9) days at 9 °C compared to 110 ± 5 (n = 10) days at 2 °C (C3: Kruskal-Wallis, $H_1 = 13.58$, p = 0.0002, Fig. 1a). The effect of temperature on the molt increment was also pronounced with a higher increase in size from C3 to C4 at 9 °C (38.5 ± 3.7 , n = 9) than at 2 °C (26.5 ± 3.9 , n = 10, C3: ANOVA, $F_{1,16}$, = 4.82, p=0.043, Fig. 1b). Although growth was faster at warmer temperatures, the effect on lipid storage was the opposite: C4 crabs cultured at 2 °C had significantly higher lipid densities (18.7 ± 3.1 mg.g⁻¹, n = 5) than those cultured at 9 °C (9.4 ± 3.0 mg.g⁻¹, n = 6, ANOVA, $F_{1,9}$ =4.7, p = 0.050, Fig. 1c). Difference in the lipid class proportions were not significant although crabs at colder temperatures trended towards higher triacylglycerols and lower proportions of sterols than those reared at 9°C (Table A1).

Throughout all four regions of the SEBS, bottom temperatures were warmer in 2014 than 2012 (Fig. 2). The average region-specific increase in bottom temperature was 2.1 ± 0.6 °C (n = 4) with the largest increase in temperature across the two central shelf regions (~2.5 °C) and a

smaller increase in bottom temperature along the outer deeper shelf regions (~1.5 °C, Table 1). In both years, juvenile Tanner crabs (species rank 1 & 2) were concentrated in the warmer deeper bottom waters of the outer shelf (100 to 200 m). This data from both years are in agreement with a previous reports for just 2012 where Ryer et al. (2016) noted the distribution of juvenile Tanner crab was concentrated along the outer shelf in avoidance of the cold-pool over the middle shelf. A similar distribution of Tanner crab was found in 2014, although generally CPUE was lower in the southern regions. Juvenile snow-hybrids (rank-3 as defined in Urban et al. (2002)) were only captured on the central middle shelf (Fig. 2), which historically is covered by the cold-pool (<2 °C water) during cold and moderate temperature years.

There was substantial spatial and inter-annual variation in chla across the SEBS during 2012 and 2014 (Fig. 3). The region-specific total integrated chla was higher in 2012 than in 2014 with an average region-specific difference of 42 mg.m⁻² (n = 4 regions, Fig. 3a & 3c) while chla in the large size fraction was 32 mg.m⁻² higher in 2012 than 2014 (Fig. 3b & 3d). Mean annual values for each regions showed the highest integrated chla in 2012 on the south outer (SO) shelf $(143.7 \pm 79.2 \text{ mg.m}^{-2}, n = 4)$ and the lowest mean values in 2014 on the south middle (SM) shelf $(27.7 \pm 7.1 \text{ mg.m}^{-2}, n = 8)$ (Table 1). The regional-specific decrease in integrated chla was highest over the SO shelf with a drop from 2012 to 2014 of 104 mg.m⁻². The least change in total integrated chla was measured across the central middle (CM) shelf where values decreased by 18 mg.m⁻² from 2012 to 2014. Region-specific large particle (>10 μ m) chla ranged from a high of 28 mg.m⁻² on the CM shelf in 2012 to a low of 2 mg.m⁻² in 2014 on the SO shelf (Fig. 3b & 3d). The largest regional differences in chla from large particles was measured on the CM shelf between 2012 and 2014, where values in 2014 were only 21% on average of those measured in 2012 (a reduction from 28 mg.m⁻² down to 6 mg.m⁻², Table 1, Fig. 3).

There was a significant interaction between year and region (general linear model (GLM), $F_{3,178} = 13.90$, p<0.001) on the total fatty acid concentration of juvenile crabs. Crabs had the highest fatty acids per weight on the CM shelf in 2012 and the lowest fatty acids on the SO shelf in 2014. This region-specific difference in fatty acid tissue storage between 2012 and 2014 was the most significant on the CM shelf compared to other survey regions (Fig. 4a). There were significantly higher densities of bacterial fatty acids in crab tissues in 2012 than in 2014 across all regions (GLM, $F_{1,178} = 5.24$, p<0.02). Region of collection also had a significant effect on crab bacterial markers (GLM, $F_{3,178} = 28.04$, p<0.001, Fig. 4b) with crabs collected on the CM shelf having significantly higher levels than crabs from all other regions. Diatom markers in crab tissues were interactively affected by region and year of collection (GLM, $F_{3,178}$ = 19.95, p<0.001) with crabs collected in 2012 from the CM region having the highest ratio of diatom indicator fatty acids while crabs collected on the outer shelf (SO & central outer (CO)) had significantly lower and less variable diatom indicators in both years. A significant decrease in the diatom indicator was measured in all regions from 2012 to 2014 except on the SO where levels did not significantly differ between years (Fig. 4c).

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Correlations between the integrated chla in >10 μ m particles and mean crab fatty acid storage (mg.g⁻¹ WWT) as well as crab storage of diatom indicator fatty acids (ratio of 16:1n-7 to 16:0) were evaluated across stations where crab were captured over the middle shelf (CM and SM) in 2012 and 2014. The outer shelf stations were excluded due to the lower numbers of stations sampled. Comparisons over both years showed a week relationship between >10 μ m chla in late summer and crab lipid storage ($F_{1,22} = 6.59$, p = 0.018, r² = 0.2) as well as crab tissue storage of diatom markers ($F_{1,22} = 11.03$, p = 0.003, r² = 0.3, Fig. 5a & 5b). However, when this relationship was examined only in the cold year (2012) we saw a positive relationship between

station-specific >10 μ m chla and crab total fatty acids ($F_{1,9}$ = 11.06, p = 0.001, r² = 0.50) and a highly positive relationship with the diatom fatty acid indicators in crab tissue ($F_{1,9}$ = 26.75, p = 0.009, r² = 0.72). This relationship points to the importance of large size phytoplankton biomass (conceivably diatoms) to juvenile crab lipid storage.

Juvenile *Chionoecetes* spp. from all four regions across the SEBS in two years were distinguished based on individual fatty acids >0.5% and crab total tissue fatty acids $(mg.g^{-1})$ using nMDS (non-metric multidimensional scaling, Fig. 6). Crabs from the CM shelf were characterized by elevated fatty acids characteristic of diatoms (16:1n-7 and C₁₆ PUFA) and elevated total fatty acids per WWT, compared to crabs from outer shelf regions which had lower fatty acid concentrations. In particular, crabs from the CO shelf had higher proportions of long chain C₂₀₊₂₂ monounsaturated fatty acids (MUFA) typical of calanoid copepods (i.e. 20:1 n-9 and 22:1 n-11) which could indicate that these lower lipid crabs were feeding at a higher trophic level. There was a proportional increased in importance of bacterial fatty acids (i.e. ai 17:0 and ai 15:0) in crabs from the CM and SM shelf regions in 2014 compared to 2012, likely due to the relative decrease in diatom-sourced fatty acid in 2014. There was a significant interaction between year and collection region on crab lipid biomarker composition (PERMANOVA, pseudo $F_{3,177} = 9.18$, p <0.001) with larger annual differences in crabs collected on the CM shelf then those from the SO shelf.

Tanner crab (ranks 1), Tanner hybrid (rank 2) and snow hybrid (rank 3, as defined in Urban et al. (2002)) only overlapped geographically on the CM shelf (Fig. 2). Therefore, we only analyzed crab lipid composition as a function of hybridization stage in the CM region while for all other analyses we pooled data for *Chionoecetes* spp. In the CM region, we found a significant interaction between year and hybrid status on total fatty acids per WWT (mg.g⁻¹) in

juvenile crabs (GLM, $F_{2,56} = 5.16$, p = 0.009). The decrease in total fatty acids from a cold (2012) to a warm (2014) year was more significant for snow hybrids (rank 3) than it was for Tanner crabs (rank 1, Fig. 7). Although sample sizes were small, this indicates potentially a more dramatic effect of environmental warming and changes to food quality on snow crab lipid storage then measured for Tanner crabs.

Discussion

Our combined laboratory and field-based analyses on the early juvenile stages of the *Chionoecetes* spp. complex from across the SEBS show both direct thermal and indirect dietary impacts of environmental change on juvenile crab lipid storage. Crabs in the warm year (2014) had lower total fatty acids per weight, and had lower diatom fatty acid biomarkers than crabs from the cold year (2012: one of the last major SEBS cold pool years that decade). These findings give us a mechanistic framework through which to view changes in crab abundance and energetic status during on-going environmental transformation and northward retraction of the Bering Sea cold pool.

Direct thermal effects on juvenile Chionoecetes spp.

Temperature is the most important environmental factor controlling vital rates in ectotherms (Hartnoll 2001) and consequently our finding of faster crab growth with warm (9 °C) compared to cold temperatures (2 °C) is expected (Ryer et al. 2016; Stoner et al. 2013; Stoner et al. 2010). Ryer et al. (2016) noted that slow growth rates of Tanner crabs at cold temperatures (C3 to C4 stage, ~100 days at 2 °C) likely explained their avoidance of the Bering Sea cold pool with higher crab abundance found in warmer deeper outer shelf waters (C3 to C4 stage, ~50 days at 5°C). Temperature-dependent growth rates of *Chionoecetes* spp. (Ryer et al. 2016; Yamamoto

et al. 2015) indicate that crabs in our study were in their second year on the bottom after setting, despite their comparative small size in relation to age-0 crabs from warmer nursery sites in the Gulf of Alaska (Copeman et al. 2018; Ryer et al. 2015). Warmer temperatures in 2014 may explain the general predominance of larger juvenile crabs in our collections from 2014 compared to 2012 (Table 1), but this variance was not extreme, and was within one molt stage (C5 to C6, estimated from the temperature-dependent growth relationship in Ryer et al. (2016)). Previous laboratory studies on juvenile Alaskan crab species have shown variable lipid densities with ontogeny during pelagic stages (Beder 2015; Copeman et al. 2014), but relatively stable lipids per weight across juvenile benthic stages (Copeman et al. 2012). Therefore, annual difference (2012 vs. 2014) in crab lipid storage here are likely not attributable to the minor annual differences in crab molt stages.

Understanding the mechanisms through which thermal effects act on vital rates (oxygen consumption, growth, lipid metabolism) can be improved by utilizing controlled laboratory studies with multiple physiological end points (Pörtner and Knust 2007). We found increased temperature caused direct, but opposing effects on juvenile Tanner crab growth and lipid storage. Vital rates of crustaceans, and ectotherms in general, have a dome-shaped relationship with temperature (Laurel et al. 2016; Schiffer et al. 2014; Storch et al. 2011). Ryer et al. (2016) noted a relatively eurythermal growth response in juvenile Tanner crabs as growth is positively correlated with a wide range of temperatures (ranging from 2 °C up to 12 °C). In contrast, snow crab exhibit a colder and more stenothermal response with higher growth up to 4-5 °C, but decreased growth and survival at warmer temperatures (8 °C, Yamamoto et al. (2015)). However, neither study concurrently measured the variation in energy storage as a function of temperature. While temperature-dependent growth is quite well defined for many species of fish

and crabs, less is known about the tradeoffs between growth and lipid storage with thermal variation. In temperature-controlled experiments on larval and juvenile cod species, the thermal maximum for lipid storage has been found to be lower than that observed for growth (Copeman et al. 2017; Koenker et al. 2018). Similarly, we found reduction in lipid with increased growth in Tanner crabs at 9 °C compared to 2 °C, results that suggests a lower thermal optimum for lipid storage than for growth. Further detailed species-specific experiments with more temperature treatments are required to parameterize the temperature-dependent trade-offs between lipid storage and growth (protein synthesis) in juvenile *Chionoecetes* crabs. Knowledge of these temperature-dependent vital rates could allow more accurate predictions of future effects of warming on crab growth, energetics, and ultimately recruitment potential.

Indirect food web effects

Expected indirect effects of warming and loss of sea ice for benthic invertebrates include diminished quantity and quality of resources fluxing to the benthos (Griffiths et al. 2017; Harada 2016; Hobson et al. 1995; Hunt et al. 2011). Particularly, a reduction in sea ice and associated ice-algae blooms (often diatoms) may substantially reduce the energy deposition to the seafloor. The mechanisms through which environmental change affect benthic-pelagic coupling have been more rigorously studied in the northeastern Bering and Chukchi Seas than in the SEBS likely because northern seas are more ice-dominated and have a greater benthic biomass (Grebmeier 2012; Lovvorn et al. 2005) than the more pelagic-dominated SEBS.

Across all four of our study regions and through both warm and cold years, we found that diatom fatty acid biomarkers in crab tissues were positively associated with increased total fatty acids (mg.g⁻¹ WWT) in juvenile crabs. Therefore, much of the excess lipid storage in juvenile crab tissues is of diatom origin and points to the importance of this primary producer for juvenile

crab metabolism and growth. We did not have samples of food items such as organic "fluff" material or small benthic prey to analyze for lipid biomarkers to directly link crab lipid compositions to their diet. Although we do not have explicit species identification of phytoplankton in our chla samples from 2012 and 2014, evidence from FLOWCAM imaging of phytoplankton undertaken in the Bering and Chukchi seas during summer in the same years noted a predominance of diatoms in 2012, that was much reduced in 2014 (Goes et al. 2014). Further, larger diatoms were more prevalent in late spring of 2012 than in 2011, a warmer year with less extensive sea ice (Stauffer et al. 2014). Our results for these SEBS crabs are also in agreement with previous work on nearshore Tanner crabs in the central Gulf of Alaska (Copeman et al. 2018). In four isothermal nearshore embayments, increased crab growth and lipid storage was positively related to the organic content and in particular to the concentration of bacterial- and diatom-sourced lipids in the top sedimentary layer (Copeman et al. 2018). However, unlike Tanner crabs from the Gulf of Alaska, crabs from the SEBS in cold years also likely store diatom lipids from early spring ice-associated blooms.

In 2012, extensive sea ice covered most of the SEBS throughout early spring with only the SO shelf remaining ice-free. This is in contrast to 2014, where March ice was absent from all regions except for partial coverage of the CM shelf (Fig. 3). The SO shelf was the only region that was not covered by March sea ice in either year and crabs from this region were the only group that did not show a significant difference in their diatom indicator fatty acids between 2012 to 2014 (Fig. 4c). This indicates that ice-associated, diatom production likely contributed significantly to higher condition of 2012 crabs across the middle and CO shelf regions. A potential proxy for late summer diatom production (>10µm chla mg.m⁻², discussed above) over the shelf in 2012 (cold year) was positively correlated with the diatom indicator (16:1n-7 to 16:0)

and total fatty acids (mg.g⁻¹) in crab tissues (Fig. 5d). This suggests that in cold years, both spring ice-edge and summer-fall pelagic diatom production may play significant roles in juvenile crab nutrition over the SEBS shelf. Future use of ice algae specific trophic markers such as compound specific isotopes of diatom fatty acids, sterols and isoprenoid 25 (Brown et al. 2017; Koch et al. 2020; Leu et al. 2020) could help to define the role of spring ice-associated versus pelagic diatom production in juvenile crab nutrition in the Alaskan Arctic.

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There is a very little understanding of how crab lipid metabolism (i.e. fatty acid assimilation rates and tissue wash-out rates) can impact the lipid biomarker signatures in coldwater crab tissues (Copeman et al. 2012; Stoner et al. 2010). Generally, assimilation time for small crabs occurs quickly as exemplified by a recent dietary studies on juvenile Dungeness crabs (Metacarcinus magister) that showed tissue fatty acid assimilation and discrimination after only six weeks (Thomas et al. 2020). Alaskan red king crab (Paralithodes camtschaticus) fed live-food with different fatty acid compositions displayed significant differences in their fatty acid profiles after less than one month of feeding (Beder 2015), while differences in the lipids of juvenile Tanner crabs fed low, medium, and high lipid diets were evident after only one molt stage (C4 to C5, Copeman et al. (2018)). However, even less is known about the effect of reduced food availability or starvation on the measured fatty acid biomarkers in crab tissue. It is likely that crabs store excess lipids and retain these for long times at cold temperatures following the spring sympagic- and pelagic-blooms. The signature of ice-associated diatom production is probably retained throughout the summer months. Additionally, it is likely that spring diatom production sinks to the benthos and that this material serves as a 'food bank' which supplies the benthos well past the short spring production season, as previously described for Arctic (Schollmeier et al. 2018; Weems et al. 2012) and Antarctic (Mincks et al. 2005) benthic food

webs. The unknown crab tissue assimilation rates and turnover times for fatty acids make it directly difficult to partition ice-associated from pelagic-associated diatom dietary input. However, with continued warming of the entire water column, contraction of spring sea ice (2019, Fig. 3), and consequent northward shrinkage of the cold pool over the Bering Sea, it is conceivable that the persistence of this spring-deposited food bank may be decreasing both by reduced benthic-pelagic coupling but also internally by increased secondary consumer metabolic rates on the sea floor.

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The late summer biomass of large sized chla (>10 µm) was elevated across the SEBS in 2012 compared to 2014, but there were also changes in the biomass of small sized chla (< 10 um) between the two years. A large late summer coccolithophore bloom of the prymnesiophyte, Emiliania huxleyi (small cells, ~ 5 µm) formed over the middle shelf in 2014 (60,658 km²) that was practically absent in 2012 (273 km²)(Ladd et al. 2018). The presence of this extensive coccolithophore bloom in 2014 partially explains why total values for late summer chla (mg.m⁻²) remained fairly constant across the two years (CM region, 72 mg.m⁻² to 55 mg.m⁻²), but values for chla in large particles were dramatically lower (28 v. 6 mg.m⁻²) in 2014 as was the ratio of biomass from large particles to total biomass (Table 1). As opposed to diatoms that have relatively large particle sizes, sink rapidly and are common during spring in high nutrient waters, small coccolithophore cells contribute little to the benthos and form blooms often in late summer nutrient-depleted waters (Eisner et al. 2016; Grebmeier et al. 2006; Ladd et al. 2018; Olson and Strom 2002). Coccolithophores likely contribute little to the diet of larger zooplankton and are not necessarily a preferred dietary source for microzooplankton in the SEBS, even given their small size (Olson & Strom 2002). As such, little of this production is likely transferred from

phytoplankton to secondary benthic consumers or to upper trophic level pelagic organisms such as fish, seabirds and marine mammals (Hunt et al. 1999; Olson and Strom 2002).

Crabs collected from the CO shelf were distinguished in a multivariate analysis of their fatty acid composition compared to crabs from other regions, particularly in 2014 (Fig. 6). This differentiation was defined by generally low fatty acid storage (mg.g⁻¹) and higher concentrations of calanoid copepod fatty acids. High proportions of C₂₀₊₂₂ chain length MUFA (16 to 22%) in predators are characteristic of feeding on calanoid copepods with high wax ester lipid storage (Dalsgaard et al. 2003; Lee et al. 2006; Stevens et al. 2004), but these fatty acids are also found in high proportions in some other benthic macroinvertebrates, such as cold-water echinoderms (Copeman and Parrish 2003). Given the small size of these juvenile crabs, the most logical explanation for this elevated C₂₀₊₂₂ MUFA is from the direct consumption of copepod carcasses or their fecal pellets. Large oceanic copepods on the SEBS outer shelf such as *Neocalanus* spp. develop in spring and begin descending to depth to diapause as lipid-rich C5 stages in late May (Miller 1993; Vidal and Smith 1986). Neocalanus cristatus collected in diapause from offshore waters of southeast Hokkaido, Japan were found to contain approximately 80% of their lipids as wax esters with a fatty alcohol composition that was $\sim 81\%$ C₂₀₊₂₂ MUFA (Yamada et al. 2016). It is possible that *Neocalanus* on the outer shelf of the Bering Sea could reach the shallow shelf bottom and be consumed by juvenile crabs during their attempt to reach diapause habitat at greater depths (Miller 1993). However, it is noteworthy that these crabs which are potentially feeding at a higher trophic level were in lower lipid-based condition than crabs from the shallower shelf regions that had higher diatom indicator fatty acids.

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Caveats and future research

Crab total fatty acid values here are expressed relative to WWT because we did not have enough animals to analyze individuals for dry weight (DWT) and ash-free DWT (AFDWT, organic mass). Expression of fatty acids per WWT is not ideal as it is impossible to distinguish differences in lipid content per organic weight from difference in lipid content due to variable shell weight. This can be an issue when comparing crabs that are of disparate size or ontogenetic stage, however, the crabs in our study only varied by one molt stage (Table 1). Previous studies on Tanner and red king crab from Alaska found no significant difference in the proportion of moisture or AFDWT in crab tissues as a function of their molt stage (C2 to C4) (Copeman et al. 2018; Stoner et al. 2010).

Crustaceans can also vary in the proportion of AFDWT as a function of their intramolt cycle and changes in association with ecdysis. (Copeman et al. 2012; Ouellet et al. 1992). Generally, there are 3 biochemical phases during the intramolt period which are typified by a short post molt period, a longer intramolt period, and another short non-feeding premolt period. Studies on juvenile crustaceans have documented a general energetic pattern of no increase in lipids during the postmolt period, a rapid accumulation of lipid during the feeding intramolt stage, and lastly, a small decrease in lipids during a nonfeeding premolt stage (Copeman et al. 2012; Ouellet et al. 1992; Sánchez-Paz et al. 2006; Zhou et al. 1998). We controlled for this intramolt energetic variability in our lab study by sampling at a set and temperature-adjusted mid-molt period. It is not possible to control for energetic variation due to intramolt stage in field collected crabs. Nevertheless, lab studies have shown that there is a large degree of variation in intramolt duration in juvenile snow crabs (i.e. C4 stages at 4 °C ranged from 59-192

d, Yamamoto et al. (2015)) which leads us to conclude that intramolt stage is unlikely to be uniformly different as a function of geographical region across the SEBS.

Through the use of both laboratory and field approaches, we have provided some of the first evidence linking both direct metabolic and indirect food web effects of warming to reduced lipid storage of juvenile *Chionoecetes* spp. Our understanding of factors affecting crab recruitment on the SEBS, could be greatly enhanced by definitive genetic differentiation of Tanner, snow and hybrids (Smith et al. 2005). Snow crabs were only collected on the central middle shelf but we noted that warming conditions appeared to have a more significant impact on their energetic condition than for Tanner crabs. Previous studies on the SEBS have shown that environmentally driven impacts on pelagic food webs are not identical in all warm stanzas (Duffy-Anderson et al. 2017; Eisner et al. 2019; Eisner et al. 2016) and the same may be true for cold stanzas. The addition of more years of crab collection and lipid analysis, with particular urgency focused on sampling during rapidly disappearing SEBS "cold" years, will help us better define the mechanisms driving variability in crab condition. Northward expansion of this type of annual ecosystem study with a focus on small juvenile crabs will provide greater insight into how environmental conditions are impacting the distribution, condition and resultant recruitment variability of *Chionoecetes* spp. across the northern Bering Sea and into the Chukchi Sea.

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Acknowledgements

We would like to thank Janet Duffy-Anderson and Daniel Cooper and the crew of the 2012 and 2014 BASIS NOAA Ecosystem Survey for the collection of juvenile crab, chla and temperature data. We would like to thank Kayln Hubbard for technical assistance with lipid analyses at the Marine Lipid Ecology Lab at Oregon State University. Further, we thank Scott Haines for

assistance with laboratory experiments and help with crab hybrid identification. Benjamin Laurel, Jeffrey Napp, Thomas Hurst and Erin Fedewa provided helpful editorial reviews on earlier versions of this manuscript. Thanks also to Cynthia Sweitzer for providing help with procurement of permits for crab collections. We would like to thank two anonymous peer-reviewers as well as Dr. Patrick Mayzaud for helpful suggestions that greatly improved our manuscript.

Author's Contributions

CR was the principle investigator on this grant. CR and LC contributed to the study conception and design. Data collection was performed by MO, CR, LC, MS, PI and LE. Data analysis was performed by LC. The first draft of the manuscript was written by LC and all authors commented on previous versions of the manuscript. All authors read and provided critical review to the final manuscript.

Funding

This research was funded in part by an Essential Fish Habitat grant awarded to Dr. Clifford Ryer from the NMFS Alaska Region's Habitat Conservation Division.

Declarations

The authors have no conflicts of interest to declare that are relevant to the content of this article.

The findings and conclusions in this paper are those of the authors and do not necessarily

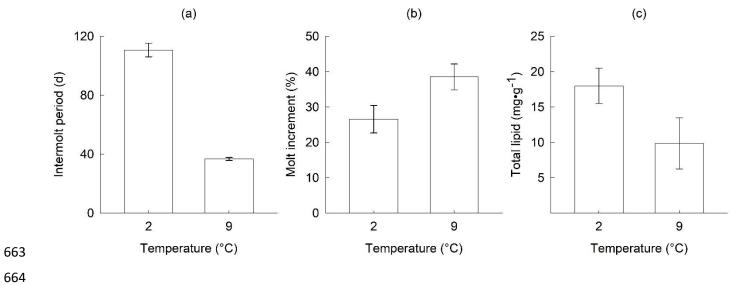
represent the views of the National Marine Fisheries Service.

- 607 Figure Legends
- Fig. 1 The effect of temperature on mean (\pm SE) (a) inter-molt period in days from C2 to C3; (b)
- 609 % intermolt increment from C2 to C3 based on carapace width; and (c) total body lipid by wet
- weight for C3 (2 degrees) and C4 (9 degrees) crabs, midway through their intermolt period.
- Tanner crabs (*Chionoecetes bairdi*) were fed to satiation 3X per week on a marine gelatinized
- 612 diet.
- Fig. 2 Distribution of juvenile Tanner crab (*Chionoecetes bairdi*) and hybrid snow crabs
- 614 (*Chionoecetes opilio*) molt stages C3 to C6 from the southeastern Bering Sea during September
- 2012 and 2014. CPUE are overlain on bottom temperatures collected from CTD casts conducted
- at the sampling sites (open circles). Tan symbols indicate Tanner and Tanner hybrid ranks (1-2)
- while red indicates snow crab and snow crab hybrids (3+, Urban et al. (2002)). Snow crab
- 618 hybrids are found only in the colder central middle (CM) shelf while Tanners and Tanner hybrids
- are predominantly found in the warmer outer and southern regions. Shelf regions include: 1)
- 620 central middle (CM), 2) south middle (SM), 3) the central outer (CO), and 4) south outer (SO).
- 621 Ellipses include sampling sites where crabs were caught within the larger sampling region based
- 622 on Ortiz et al. (2016).
- Fig. 3 Contours of total integrated chla in (a) 2012 and (b) 2014, and integrated chla in particles
- >10 μm in (c) 2012 and (d) 2014. Data are integrated over the top 50 m (methods as in (Eisner
- et al. 2016). Shelf regions include: 1) central middle (CM), 2) south middle (SM), 3) the central
- outer (CO), and 4) south outer (SO). For color bars, interpolated data values were stretched
- along a color ramp using minimum-maximum stretch type, such that high/low values could be
- edited to match between years. Monthly average sea ice extents are shown for 2012, 2014 as
- well as 2019, as an example of recent extremely low sea ice extent (Fetterer et al. 2017 updated
- daily). Ellipses include sampling sites where crabs were caught within the larger sampling
- region based on Ortiz et al. (2016).
- **Fig. 4** Annual and regional differences in the total fatty acids (mg.g⁻¹), total bacterial fatty acids
- (mg.g⁻¹) as well as diatom fatty acid indicator in juvenile crabs (*Chionoecetes* spp.) from four
- 634 geographical regions in the southeastern Bering Sea (2012-2014). Multiple comparisions of
- year*region, different letters indicate a significant difference p<0.05. Shelf regions include: 1)
- central middle (CM), 2) south middle (SM), 3) the central outer (CO), and 4) south outer (SO).
- 537 ΣBacterial markers per WWT include all odd and branched chained fatty acids (Table 1).
- Fig 5 Station-specific relationship between integrated Chla in particles > 10 μm across the SEBS
- shelf in late summer/fall and (a) crab total fatty acids per WWT in 2012 and 2014, (b) diatom
- indicator in crab tissues in 2012 and 2014, (c) crab total lipids per WWT in crabs from 2012
- only, and (d) diatom indicator in crabs from 2012.
- **Fig. 6** Nonmetric multidimensional scaling (nMDS) of individual juvenile crabs (*Chionoecetes*
- spp.) based on their fatty acid composition. Crabs were sampled in 2012 and 2014 from four
- regions across the SEBS: 1) central middle (CM), 2) south middle (SM), 3) the central outer
- 645 (CO), and 4) south outer (SO). Fatty acids used for nMDS include all individual fatty acids
- >0.5% as shown in table 1 as well as total fatty acids per WWT (mg.g⁻¹). Vectors are shown for
- 647 individual fatty acids that had a correlation > 0.5. PERMANOVA for the effects of region and
- year showed a significant interactive effect p<0.001 as illustrated by the larger annual effect
- 649 (2012 versus 2014) in fatty acid composition of crabs on the CM shelf compared to overlapping

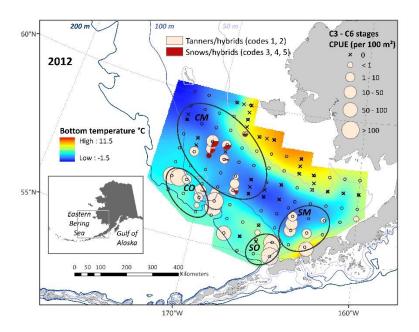
650 composition of crabs from SO region between the 2 years. Fatty acids indicative of diatoms includes C₁₆ PUFA and 16:1n-7, fatty acids indicative of bacteria include odd and branched 651 chains and 18:1n-7, and fatty acids indicative of *Calanoid* copepods include C₂₀₊₂₂ MUFA 652 653 (Dalsgaard et al. 2003; Kelly and Scheibling 2012; Parrish 2013). Fig. 7 Annual and species differences in the total fatty acids (mg.g⁻¹) in juvenile crabs from the 654 central middle (CM) shelf of the SEBS (2012-2014). Samples sizes were Tanner crab 655 (Chionoecetes bairdi.) 2012 n = 6, 2014 n = 7, Tanner hybrid crabs 2012 n = 8, 2014 n = 11, and 656 snow crab (Chionoecetes opilio) hybrids 2012 n = 11, 2014 n = 19. Multiple comparisions of 657

year*region, different letters indicate a significant difference, p<0.05.

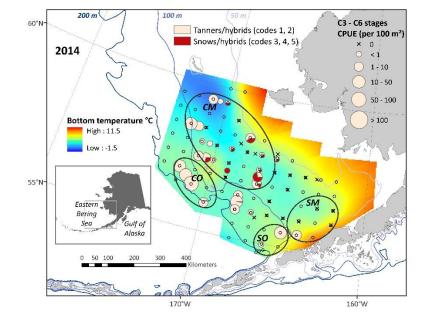
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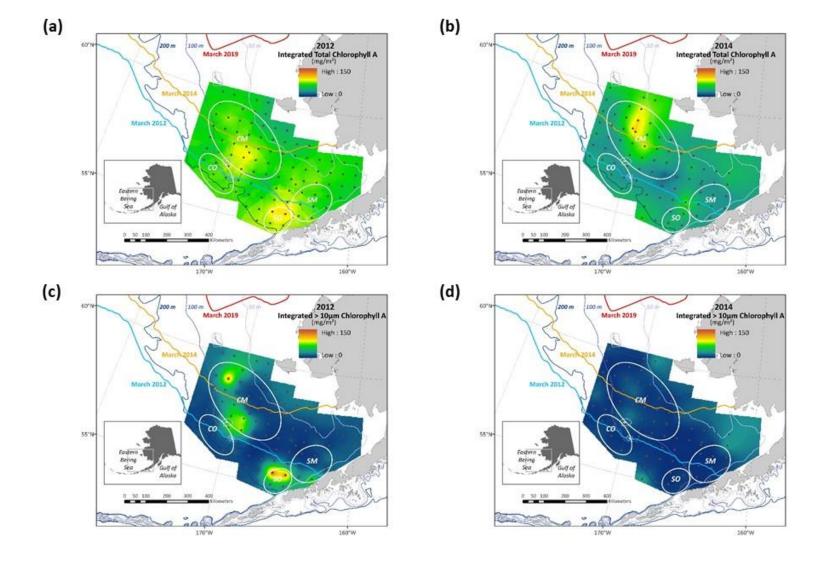


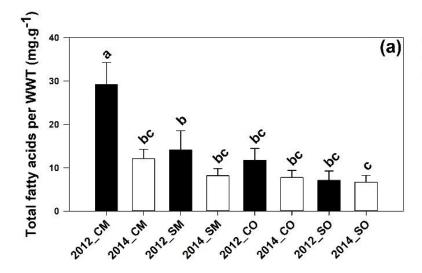




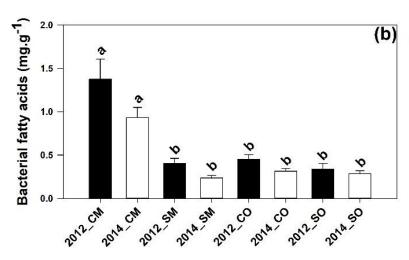




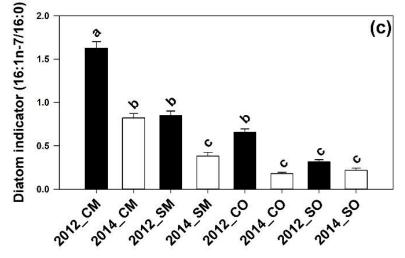




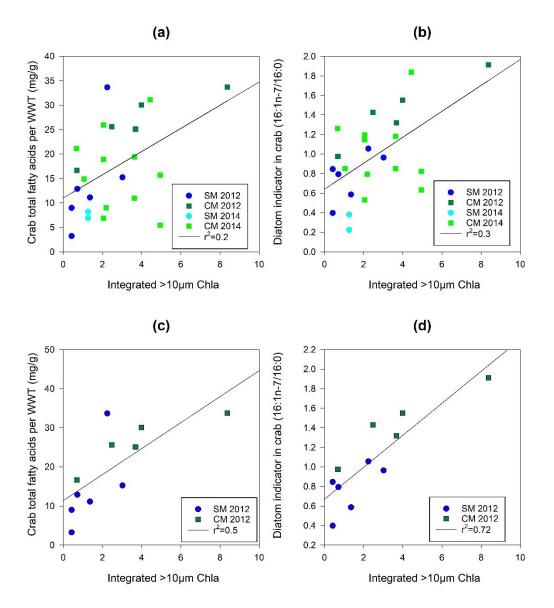
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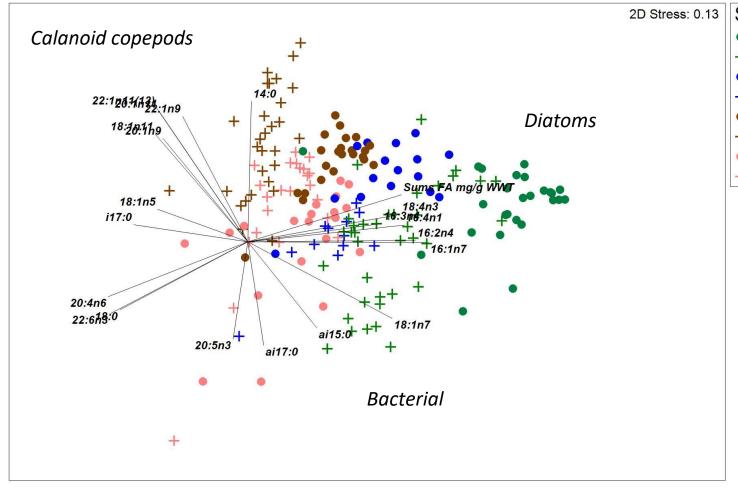


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Transform: Square root

Resemblance: S17 Bray-Curtis similarity



Shelf RegionYear

- CM2012
- + CM2014
- SM2012
- + SM2014
- CO2012
- + CO2014
- SO2012
- + SO2014

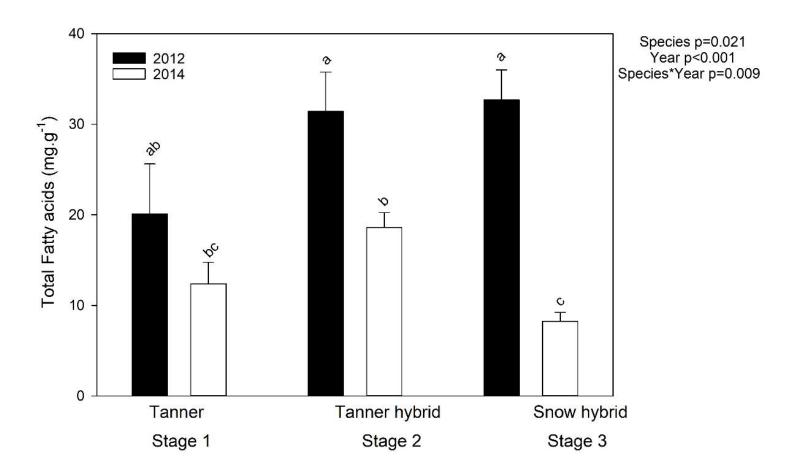


Table 1: Annual and spatial trends in near bottom temperature, mean integrated total and large size (>10 μ m) chla, as well as total fatty acids per WWT and percentage fatty acids (>0.5%) in juvenile crabs (*Chionoecetes* spp.) from the southeastern Bering Sea.

Shelf Region	Central middle (CM)		Central Outer (CO)		South middle (SM)		South outer (SO)	
Year	2012	2014	2012	2014	2012	2014	2012	2014
Station number oceanography	27	25	6	6	8	8	4	6
Bottom temperature (August-September) BASIS °C	1.55±1.78	3.26±1.44	3.02±0.48	4.12±0.19	3.03±2.62	5.94±2.04	4.49±0.54	5.10±0.76
Total integrated chla mg.m ⁻²	72.86±40.05	55.16±37.43	54.35±21.08	33.97±10.8 6	54.07±23.81	27.66±7.07	143.73±79.21	39.05±13.47
Integrated chla in particles >10 μm, mg.m ⁻²	27.93±38.8	5.63 ± 4.99	5.94±4.31	4.28 ± 3.11	15.37±25.54	2.06 ± 1.55	15.37 ± 25.54	3.18 ± 2.18
Ratio of >10µm /total chla	0.29±0.27	0.13±0.11	0.11±0.06	0.11±0.07	0.23±0.28	0.08±0.07	0.56±0.19	0.08±0.04
Sample size crabs	25	37	25	29	16	12	20	23
Carapace Width (mm)	9.0±0.4	11.5 ± 0.4	12.5 ± 0.6	10.9 ± 0.5	9.0 ± 0.2	11.8 ± 1.1	9.0 ± 0.2	9.9 ± 0.4
Molt Stage ¹	C5	C6	C6	C5-C6	C5	C6	C5	C5
Total FA per WWT (mg.g ⁻¹)	29.3±12.7	12.1±6.7	11.8±6.5	7.8±4.3	14.1±8.9	8.2±2.9	7.1±4.8	6.7±3.7
FA per DWT (mg.g ⁻¹) ^d	76.7	31.7	30.9	20.4	36.9	21.5	18.6	17.6
FA per AFDWT (mg.g ⁻¹) ^d	174.1	72.0	70.2	46.3	83.9	48.7	42.3	40.0
% of total FA								
14:0	2.6±0.5	1.5±0.7	3.0±0.9	3.1±1.5	1.8±0.5	1.6 ± 0.5	2.4±0.7	2.3±0.9
ai 15:0	2.3±2.4	3.4 ± 3.4	0.1 ± 0.0	0.1±0.0	0.1±0.2	0.2 ± 0.1	0.2±0.2	0.1±0.1
15:0	0.5±0.1	0.6 ± 0.2	0.5 ± 0.1	0.5±0.2	0.4±0.1	0.4 ± 0.1	0.6±0.1	0.4±0.1
i 16:0	0.2±0.9	1.6±1.4	0.4 ± 0.1	0.4 ± 0.1	0.2±0.1	0.4 ± 0.1	0.6±0.3	0.5±0.2
16:0	12.3±1.2	10.6±1.9	11.1±1.3	10.9±1.8	12.1±0.8	13.6±1.4	15.9±1.8	13.6±1.6
i 17:0	0.2±0.0	0.4 ± 0.1	0.5±0.2	0.7±0.1	0.4±0.1	0.1±0.2	0.6±0.2	0.6±0.3
ai 17:0	0.5±0.4	1.1±0.7	0.4 ± 0.1	0.4±0.1	0.4±0.2	0.4 ± 0.1	0.7±0.1	0.5±0.2
17:0	0.2±0.1	0.4 ± 0.2	0.4 ± 0.2	0.6±0.2	0.3±0.2	0.0 ± 0.2	0.5±0.3	0.4±0.2

18:0	2.3±1.4	3.1±0.6	3.3±0.7	3.8±1.1	3.4±0.6	3.5±0.4	4.1±0.9	4.6±1.1
∑SFA ^a	17.9±2.1	16.7±2.2	18.5±1.9	19.3±2.2	18.1±1.2	19.3±1.5	24.9±5.8	21.5±2.2
16:1n-7	19.8±3.8	8.7±3.8	7.0±1.6	1.9±0.9	10.3±2.4	5.0±1.7	4.9±1.6	2.9±1.4
17:1	0.2±0.4	0.6±0.4	0.5±0.2	0.5±0.2	0.2±0.3	0.6±0.3	0.5±0.2	0.6±0.1
18:1n-11	0.1±0.0	0.2±0.2	0.8±0.3	0.9±0.1	0.3±0.1	0.2±0.1	0.1±0.2	0.5±0.2
18:1n-9	9.8±3.5	8.9±1.9	10.3±3.5	8.6±1.2	10.0±3.1	11.6±1.2	10.9±5.3	10.7±2.2
18:1n-7	11.6±1.0	9.3±0.9	8.2±1.4	4.9±1.2	9.4±1.1	7.7±0.5	7.1±1.1	6.7±1.1
18:1n-5	0.3±0.1	0.5±0.2	0.6±0.2	0.7±0.4	0.8 ± 0.2	0.5±0.2	0.8±0.1	0.7±0.2
20:1n-11	0.9±0.3	1.5±0.9	5.2±1.5	8.4±4.0	2.5±0.8	1.3±0.5	2.0±1.2	3.4±1.5
20:1n-9	0.9±0.4	1.6±0.6	3.4±1.3	4.4±2.0	4.5±2.2	1.9±0.6	2.6±1.9	2.3±0.8
20:1n-7	1.2±0.3	2.0±0.9	2.4±0.6	1.3±1.0	3.2±1.1	1.1±0.4	1.7±0.9	1.3±0.5
22:1n-11	0.1±0.2	0.5±0.6	3.7±1.6	7.4±4.3	2.5±1.6	1.0±0.6	1.5±1.3	2.6±1.6
22:1n-9	0.1±0.1	0.2±0.3	1.0±0.5	1.0±0.5	0.2±0.3	0.2±0.3	0.1±0.2	0.9±0.7
∑MUFAb	46.0±3.5	35.2±4.6	43.6±3.5	40.8±8.2	44.4±3.9	31.9±3.3	32.8±5.9	33.6±4.5
16:2n-4	1.6±0.2	1.1±0.5	0.9 ± 0.2	0.4 ± 0.1	1.1±0.3	0.9±0.3	0.7±0.4	0.6 ± 0.2
16:3n-4	0.6±0.4	0.1±0.3	0.0±0.2	0.0 ± 0.0	0.4 ± 0.3	0.1±0.2	0.0±0.2	0.0 ± 0.0
16:4n-1	0.7±0.3	0.3±0.4	0.2±0.1	0.0 ± 0.0	0.5 ± 0.2	0.2±0.1	0.2±0.2	0.0 ± 0.0
18:2n-6	0.6±0.1	0.8±0.3	0.8±0.2	0.7±0.0	0.6 ± 0.1	1.2±0.2	0.1±0.3	0.9±0.3
18:3n-3	0.1±0.1	0.1±0.1	0.1±0.1	0.2±0.1	0.1 ± 0.1	0.6±0.2	0.3±0.3	0.4 ± 0.2
18:4n-3	1.9±0.4	0.9±0.9	0.8 ± 0.3	0.2±0.2	1.3±0.7	0.8±0.3	0.7±0.5	0.6 ± 0.3
20:2n-6	0.2±0.2	0.8±0.4	0.6±0.5	0.8±0.5	0.6 ± 0.4	1.0±0.2	1.0±0.08	1.0±0.9
20:4n-6	1.4±0.4	2.8±0.7	4.0±1.6	4.2±1.4	2.6±0.9	4.1±0.5	5.2±1.5	5.9±2.6
20:5n-3	17.5.1.2	19.9±2.3	17.2±1.8	17.7±4.2	17.5±3.2	21.0±1.4	17.0±2.7	17.6±2.4
20.311-3	17.5±1.3	19.9±2.3	17.2-1.0	17.7 = 1.2	- , , , , , , , , , , , , , , , , , , ,			
20:5n-3	17.5±1.3 0.4±0.1	19.9±2.3 1.0±0.3	1.3±0.5	1.0±0.3	1.5±0.5	1.2±0.7	0.8±0.7	1.2±0.6
							0.8±0.7 12.3±2.1	1.2±0.6 13.3±1.9

¹Molt stage estimate based on Ryer et al. (2016) and Ryer et al. (2015).

 $^{^{}a}$ ∑SFA, saturated fatty acids contains < 0.5% of *i*15:0, 20:0, 22:0, 24:0,

b∑MUFA, monounsaturated fatty acids contains<0.5% 14:1, 15:1, 16:1n-11, 16:1n-9,16:1n-5, 18:1n-5, 22:1n-7

^c∑PUFA, polyunsaturated fatty acids contains<0.5% 16:4n-3,18:2n-4,18:3n-6, 18:3n-4, 20:3n-6, 20:3n-3, 20:4n-3, 21:5n-3, 22:4n-6, 22:5n-6, 22:4n-3

[&]quot;i" refers to iso branched chains and "ai" refers to anti-iso branched chains.

^d Average conversion factors were calculated for WWT to DWT = 2.63 ± 3.49 (n=107) and WWT to AFDWT = 5.95 ± 8.64 (n=107) using values from wild C3 and C4 Tanner crabs collected near Kodiak Island in the Gulf of Alaska, USA as described in Copeman et al. (2018). Values are to be considered approximate and are given to allow comparison of total fatty acids per weight with published crab studies where fatty acid values are expressed per DWT or AFDWT

Appendix 1: Iatroscan-determined total lipids and lipid classes of juvenile Tanner crabs (*Chionoecetes bairdi*.) reared at 2 °C and 9 °C in individual growth cells in a controlled laboratory growth experiment.

	2°C	9°C
	n=5	n=6
Iatroscan-determined total lipids (mg.g ⁻¹)	18.7 ± 3.1	9.4 ± 3.0
Steryl/Wax Esters	0.2 ± 0.0	-
Triacylglycerols	48.1 ± 5.7	43.7 ± 11.0
Free Fatty acids	1.9 ± 0.3	2.3 ± 0.7
Sterols	6.1 ± 1.2	11.2 ± 4.8
Polar Lipids	43.6 ± 4.5	42.8 ± 6.8

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