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2	Sea shelf during summers of 1994–2009
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17 Abstract

On the southeastern Bering Sea shelf, mesozooplankton play an important role in 18 19material transfer between primary producers and fisheries resources. The biomass of 20mesozooplankton in this region is known to vary annually, but little is known about annual changes in community structure and species composition. In the present study, 21regional and long-term changes in abundance, biomass and community structure of 2223copepods and chaetognaths on the shelf were evaluated based on NORPAC net samples 24collected during summers of 1994–2009. During the study period, regime shifts occurred from high interannual variability regime (1994-1999) to low interannual 2526variability regime with high temperature (2000-2005), then to a low interannual variability regime with low temperature (2007-2009). A total of 24 calanoid copepod 2728species belonging to 21 genera were identified from samples. Copepod abundance ranged from 150 to 834,486 inds. m⁻², was greatest on the Middle shelf, and was higher 29in cold years, than in warm years. Copepod biomass ranged from 0.013 to 150 g DM 30 m^{-2} , and was also higher in cold years than in warm years. Based on the results of 31 cluster analysis, the copepod community was divided into six groups (A-F). The 3233 regional and interannual distributions of each group were distinct. Interannual changes in abundance of the dominant copepod on the Outer shelf and Middle shelf were highly 34significant (p < 0.0001), and their abundances were negatively correlated with 35temperature and salinity. Interannual changes in copepod community that occurred 36 between cold and warm years are thought to have been caused by differences in the 3738 magnitude and timing of the spring phytoplankton bloom between the two regimes.

Abundance and biomass of the chaetognath *Parasagitta elegans* ranged from 30 to 15,180 inds. m^{-2} and from 11 to 1,559 mg DM m^{-2} , respectively. Chaetognath abundance was significantly correlated with the abundance of the dominant copepods (*p* < 0.0001). Differences in cold and warm years may also affect recruitment of walleye pollock. We conclude that on the southeastern Bering Sea shelf, the magnitude and timing of primary production, which is related to climate change, may significantly affect how it is transferred through the food web.

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47 Keywords: Annual variations; Abundance; Biomass; Zooplankton; Community
48 composition; Species diversity

49 Regional terms: Southeastern Bering Sea, Alaska, USA

50 **1. Introduction**

The southeastern Bering Sea shelf is characterized by high biological productivity, has 5152large amounts of phytoplankton, zooplankton, benthos, fishes, seabirds and marine 53mammals, and is an important fishing ground of walleye pollock (Theragra chalcogramma) (Springer, 1992; Springer et al., 1996). Recently, this region has 54undergone large biological/environmental changes (Jin et al., 2009; Stabeno et al., 55562010). In 1997, a coccolithophore (Emiliania huxleyi) bloom was observed 57(Sukhanova and Flint, 1998; Napp and Hunt, 2001; Stockwell et al., 2001), and mass mortality of short-tailed shearwaters (Puffinus tenuirostris) occurred near the Pribilof 58Islands (Baduini et al., 2001). In 1998, a climate regime shift was reported 59(McFarlane et al., 2000), and the biomass of the jellyfish Chrysaora melanaster was 60 61 observed to peak (Brodeur et al., 2008). To describe the effect of climate regime 62 shifts on lower to higher trophic levels, Hunt et al. (2002) proposed the Oscillation 63 Control Hypothesis (OCH). In 2006, a climate regime shift from a warm period to cold period was reported, and the OCH was revised based on observations during this 64 period (Hunt et al., 2011). 65

In the Bering Sea ecosystem, mesozooplankton are important because they connect primary production to higher trophic level production. In the mesozooplankton fauna of this region, the dominant taxon is Copepoda, and various studies have been conducted on their biology and ecology. They include studies of seasonal variations of community structure during spring and summer (Smith and Vidal, 1986), interannual variations in their total wet-weight biomass (Sugimoto and Tadokoro,

721997, 1998; Napp et al., 2002; Hunt et al., 2008), seasonal and spatial variations in community structure (Kang et al., 2006), and the effects of El Niño and La Niña on 7374biomass and production (Coyle and Pinchuk, 2002). However, most studies of 75long-term variations in mesozooplankton have treated only wet-weight biomass data without examining the species caught or separating sub-regions or domains. In studies 76of species and zooplankton community structure, the study areas have been narrow, and 7778 the study periods have been short. Consequently, information on long-term changes in 79mesozooplankton community and their species composition is extremely scarce.

80 The second most abundant taxonomic group in the Bering Sea mesozooplankton biomass is chaetognaths (Coyle and Pinchuk, 2002). 81 They are 82 carnivores, prey upon small zooplankton and fish larvae, contribute to vertical fluxes to deep water (Brodeur and Terazaki, 1999), and sometimes account for nearly 50% of the 83 prey of walleye pollock larvae (Brodeur and Wilson, 1996). Thus, chaetognaths are 84 important because of their role linking lower to higher trophic levels. However, little 85 is known about their long-term variation in the Bering Sea. 86

In the present study, regional and long-term changes in abundance, biomass and community structure of copepods and chaetognaths are evaluated based on samples collected over the southeastern Bering Sea shelf during the summers of 1994–2009. Long-term data on copepods and chaetognaths are compared with hydrographic data, and we discuss possible mechanisms controlling interannual variations in mesozooplankton community structures and species composition.

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93 2. Material and methods

94 *2-1. Field sampling*

95 Sampling was conducted over the southeastern Bering Sea shelf during 1994–2009 96 between 24 June and 8 August on board the T/S Oshoro-Maru of the Faculty of Fisheries, Hokkaido University. For each year, samples were collected within a two 97 week period. Zooplankton samples were collected by vertical hauls of 98 99 flowmeter-equipped NORPAC nets (45 cm mouth diameter, 0.335 mm mesh; Motoda, 100 1957) from 150 m depth or near the bottom (where the depth was shallower than 150 m) Zooplankton samples were immediately preserved with 5% 101 to the surface. 102formaldehyde-seawater buffered with sodium tetraborate. In addition, temperature and salinity were measured with a CTD (Neil Brown, Mark IIIB during 1994-2001 and 103 104 Seabird SBE-911*plus* during 2002-2009). Water samples from the CTD rosette were 105filtered through GF/F filters, extracted with acetone or DMF and chlorophyll a 106 concentration was measured using a fluorometer (Turner Designs, Inc.).

The number of sampling stations in any one year was 9-49 (total = 96), and the 107 total number of samples was 428 (Fig. 1). Based on bottom depth, the southeastern 108 109 Bering Sea shelf was divided into three regions: Inner shelf (< 50 m), Middle shelf (50-100 m) and Outer shelf (100-200 m) (Coachman and Charnell, 1979; Coachman, 110 111 1986). Areas with depth of 200-1000 m and > 1000 m were classified as Slope and Basin, respectively. Six stations on the Outer shelf and six on the Middle shelf were 112sampled every year of the study (Fig. 1), so only the data from these twelve stations 113114 were used to analyze long-term changes.

115 2-2. Sample analysis

Based on the biomass of the samples, subsamples were created with a Motoda planktonsplitter (Motoda, 1959), and the subsamples were examined under a stereomicroscope.

Copepods were staged and identified to the lowest possible taxonomic level. 118For copepods, the most important genus in this region (Calanus) is reported to be 119 120mixture of C. glacialis and C. marshallae (Nelson et al., 2009). We could not distinguish the two species and treated them as "Calanus spp.". For biomass 121122estimation, copepods were separated into four groups: Calanus spp., Neocalanus spp., 123Eucalanus bungii and other copepods. One hundred individuals in each group were chosen randomly for measurement of total lengths (TL, µm) by ocular micrometer. 124125Dry mass (DM, µg) was estimated using the following equations derived for organisms 126from the Oyashio region (Imao, 2005):

127	Calanus and Neocalanus spp. (less lipids)	$\log_{10} \text{DM} = 2.48 \log_{10} \text{TL-6.00}$	(1)
128	Calanus and Neocalanus spp. (more lipids)	$\log_{10} DM = 3.00 \log_{10} TL\text{-}7.70$	(2)
129	Eucalanus bungii	$\log_{10} \text{DM} = 3.16 \log_{10} \text{TL}-9.16$	(3)

130 Other copepods

 $\log_{10} \text{DM} = 2.62 \log_{10} \text{TL-6.40}$ (4)

For *Calanus* spp. and *Neocalanus* spp., lipid accumulation was separated into two: less lipids (<20% of prosome volume) and more lipids (\geq 20% of prosome volume). Then the equations (1) and (2) were applied for less and more lipid specimens, respectively. The total biomass of copepods (mg DM m⁻²) was estimated by multiplying the average individual mass (mg DM ind.⁻¹) by the abundance (inds. m⁻²).

136	Chaetognaths were identified to species based on the samples collected during
137	1996-2009. The total length (TL, mm) of the dominant chaetognath, Parasagitta
138	<i>elegans</i> , was measured using a ruler for large individuals (TL ≥ 10 mm) and an ocular
139	micrometer for small specimens (TL < 10 mm). Specimens were classified according
140	to McLaren (1969), Sameoto (1973) and Zo (1973): Stage I (juveniles), Stage II
141	(immature) and Stage III (mature). To estimate DM, ash-free dry mass (AFDM, mg)
142	was estimated from the TL using the following equation (Mumm, 1991):
143	$AFDM = 0.0002 \text{ TL}^{2.6924} $ (5)

Dry mass was estimated as DM = AFDM / 0.9 (Båmstedt, 1986). 144

2-3. Data analysis 145

146 Zooplankton samples were collected day and night. Since the depths of most sampling 147stations were < 150 m and the sampling was conducted throughout the water column, day-night differences in abundance and biomass were expected to be small. 148 There 149were no significant differences between day and night abundance or biomass of 150copepods and chaetognaths for all years (p = 0.06-0.93, U-test). Thus, no day-night 151conversions were done for abundance or biomass.

To examine community structure, we conducted cluster analysis and 152non-metric multidimensional scaling (NMDS) ordination. Abundance data (X: inds. 153 m^{-2}) of each species were transformed to $log_{10}(X+1)$ prior to analysis to reduce the bias 154of abundant species. Rare species which occurred only <2% of the whole stations 155were eliminated from the data for analysis (cf. Table 1). Similarities between samples 156

were examined by Bray-Curtis index (Bray and Curtis, 1957) according to the differences in species composition. For grouping the samples, the similarity indices were coupled with hierarchical agglomerative clustering with a complete linkage method. The NMDS ordination was carried out to delineate the sample groups on the two-dimensional map. All of these analyses were carried out using BIOSTAT II software (Sigma Soft).

163To evaluate environmental factors that may have influenced sample groups determined by cluster analysis, nonmetric multi-dimensional scaling (NMDS) 164ordination was carried out to delineate the sample groups in two-dimensional space. 165We then conducted multiple-regression analysis $(Y = aX_1 + bX_2 + c)$ with the 166hydrographic data (Y: latitude, longitude, depth, sea surface and bottom temperature and 167 168 sea surface and bottom salinity) are the dependent variables and two-dimensional NMDS as independent variables (thus, axis 1 as X_1 and axis 2 as X_2) (a, b, c are fitted 169170constants). Shannon's diversity index (H') in each group was calculated using the equation: 171

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$$H' = -\sum n_i / N \times \ln n_i / N \tag{6}$$

where n_i is abundance (inds. m⁻²) of the *i*th species, and *N* is total copepod abundance (inds. m⁻²) in the group (Shannon and Weaver, 1949).

To evaluate factors affecting the variability of copepod abundance, two-way ANOVA with year and station as independent variables was used. To determine which factors controlled annual changes in copepod abundance, correlation analysis was done between the abundance of dominant copepods and three factors: water temperature (throughout whole water column, in upper layer and in lower layer), salinity andchlorophyll *a* (whole water column).

For *P. elegans*, cohort analysis was done based on the TL histogram data from
the Middle shelf using Microsoft Excel Solver (Aizawa and Takiguchi, 1999).

183 To evaluate what factors controlled the interannual changes in chaetognath 184 abundance, we tested for correlations between the chaetognath abundance for the Outer 185 shelf and Middle shelf and four factors: water temperature (whole water column, upper 186 and lower layer), salinity (whole water column), chlorophyll *a* (whole water column) 187 and abundance of the dominant copepods.

188 **3. Results**

189 *3-1. Hydrography*

In all years, a thermocline was present 20–30 m, and the bottom temperature of Middle shelf ranged between 0° and 4°C, but was higher for the Inner and Outer shelves (Fig. 2A). The Middle shelf, bottom temperature was about 4°C during 2001–2005, and lower (0–3°C) during 2007–2009. Owing to Stabeno et al. (2012), there were three regimes in depth-averaged ocean temperature: cold; 1995, 1997, 1999, 2007-2009, average; 1996, 2000, 2006, and warm; 1998, 2001-2005. Our results confirmed this pattern (Fig. 2A).

197 The peak of summer chlorophyll *a* at each station varied between 0.5 and 25.4 198 mg m⁻³, and chlorophyll *a* was high above the thermocline and on the Inner shelf (Fig. 199 2B). The peak of chlorophyll *a* was low (ca. 1 mg m⁻³) during the warm regime

201 *3-2. Copepods*

202 Regional and interannual changes in total copepod abundance and biomass were observed (Fig. 3). Copepod abundance ranged from 150 to 834,486 inds. m⁻², 203204 and was greatest on the Middle shelf (Fig. 3A). Copepod abundance was high during 205cold years (1995, 1997, 1999 and 2007-2009), and low in warm years (2001-2005) (Fig. 3A). Copepod biomass ranged from 0.013 to 150 g DM m^{-2} , and was highest on the 206207 Outer and Middle shelves (Fig. 3B). The biomass peak occurred in more oceanic 208waters than the abundance peak due to the dominance of large-sized oceanic copepods. 209Copepod biomass and abundance were both high in cold years and low in warm years.

Throughout the study period, 24 calanoid copepod species from 21 genera were identified (Table 1). *Pseudocalanus* spp. was the most numerous, and *Calanus* spp. was dominated the copepod biomass. Large-sized oceanic copepods that diapause at depth (i.e., *E. bungii, M. pacifica, N. cristatus, N. flemingeri* and *N. plumchrus*) were collected on the Outer shelf, on the Slope and in the Basin (Table 1).

Based on abundance of each species, cluster analysis divided the copepod community into six groups (A–F). The groups contained 24–127 stations (Fig. 4A). From multiple-regression analysis, environmental factors that were significantly related with the group separation were latitude, longitude, depth, sea surface temperature, sea surface salinity and bottom salinity. The most important factors were latitude and depth; they determined 70–77% of the variability of the copepod community (Fig. 4B).

221Mean copepod abundance was highest in group D and lowest in group E (Fig. 4C). Species diversity was higher in groups A and C (H' = 1.6) than in groups B, D, E and F 222223(H' = 1.0-1.1). These group-specific differences in species diversity were caused by their species composition. For groups A-C, Pseudocalanus spp. and large, oceanic 224copepods (M. pacifica, E. bungii, Neocalanus spp.) were abundant (Fig. 4C). For 225226groups D and E, species diversity was lower, and the species compositions were similar; 227they were comprised of *Pseudocalanus* spp. (55–58%), *Acartia* spp. (24%) and *Calanus* 228spp. (18–19%). In group F, the proportion of Calanus spp. decreased, and the main species were Pseudocalanus spp. (69%), and Acartia spp. (18%). Endemic species, 229230such as Epilabidocera amphitrites, Eurytemora herdmani and Tortanus discaudatus, 231were also observed (Fig. 4C, Table 1).

The regional and interannual distributions of each group were clearly separated (Fig. 5A). Composition of group A was greater in cold years (1995, 1997, 1999 and 2007-2009) on the Outer shelf, and that of group B was greater in warm years (1998, 2001-2005) on the Outer shelf, and group C was seen mainly on the Outer shelf and in the Basin, but also on the Middle shelf in 2006–2008 (Fig. 5B). Group D was mainly observed in cold years on the Middle shelf, group E was seen in warm years on the Middle shelf, and group F was centered on the Inner shelf (Fig. 5A).

Two-way analysis of variance (ANOVA) on annual and geographical differences in abundance of the dominant copepods showed significant interannual changes for *Acartia* spp., *M. pacifica* and *Neocalanus* spp. on the Outer shelf (Table 2). Significant station-to-station differences were detected only for *Calanus* spp. on the Middle shelf. All dominant copepods on the Middle shelf (*Acartia* spp., *Calanus* spp. and *Pseudocalanus* spp.) showed highly significant interannual changes in abundance (p< 0.0001, Table 2).

Correlation analysis was used to evaluate which environmental parameters were related to the interannual changes in copepod abundance. Abundances of the three dominant copepods on the Middle shelf, which all showed high interannual variability, were strongly negatively correlated with temperature and salinity (Table 3).

250 *3-3. Chaetognaths*

Two chaetognath species (*P. elegans* and *Eukrohnia hamata*) were collected. *Parasagitta elegans* dominated the chaetognath fauna, composing 84% of the total chaetognath abundance and 89% of the chaetognath abundance on the Middle shelf. For this reason, *E. hamata* was not included in the subsequent analysis of chaetognaths.

Parasagitta elegans abundance ranged from 30 to 15,180 inds. m^{-2} , and was highest on the Middle shelf (Fig. 6A). Abundance was low during 2000–2004 and high during 1996–1999 and 2005–2009. *Parasagitta elegans* biomass varied from 11 to 1,559 mg DM m⁻² (Fig. 6B). Biomass was low during 2002–2004 and high during 1996–1999 and 2005–2009. Differences in the timing of the abundance and biomass peaks might have been caused by interannual changes in *P. elegans* body-size composition.

In every year, *P. elegans* TL composition on the Middle shelf had two or three cohorts (Fig. 7). The small-sized cohort was dominated by Stage I (juveniles), the

middle-sized cohort by Stage II (immature), and the large-sized cohort by Stage III (mature) individuals. Stage III was not observed in 2003 or 2004 (Fig. 7). The average TL of the small-sized cohort (Stage I) was significantly larger (3.5-8.5 mm) during 1996–2005 than those (2.5-5.5 mm) during 2006–2009 (Fig. 7) (p < 0.05, *U*-test).

Parasagitta elegans abundance on the Outer shelf was positively correlated 269270with dominant copepod abundance (Acartia spp., Calanus spp. and Pseudocalanus spp.), 271but not with environmental factors (Table 4). On the Middle shelf, P. elegans had a 272highly significant positive correlation with dominant copepod abundance (p < 0.0001) 273and significant negative correlations with temperature and salinity. Thus on the 274Middle shelf, *P. elegans* abundance was high when temperature and salinity were low, 275and low when temperature and salinity were high, which is similar to the pattern 276observed in copepod abundance (Table 4).

4. Discussion

The copepod community on the Bering Sea shelf showed large interannual differences across the shelf (Fig. 5A). The major environmental factors that were related to the copepod community were latitude and depth (Fig. 4B), and species diversity was highest on the Outer shelf because of the occurrence of the oceanic copepods in addition to the shelf copepods (Fig. 4C). These findings correspond well with the horizontal distribution of copepod community structure reported by Cooney and Coyle (1982). These cross-shelf differences in copepod species may be related to the occurrence of frontal structures (Coachman and Charnell, 1979; Coachman, 1986). The large number of oceanic species on the Middle shelf in 1998 (Fig. 5C) may be caused by the transportation of oceanic copepods by strong cross-shelf advection (Stabeno et al., 2001) (Fig.1). Stabeno et al. (2012) also reported the transport of oceanic species during warm years.

290 Long-term changes in zooplankton on the southeastern Bering Sea shelf based 291upon wet-weight biomass have been reported (Sugimoto and Tadokoro, 1997, 1998; 292Napp et al., 2002). Recently, Hunt et al. (2008) also reported that the zooplankton wet 293mass was lower in the early 2000s than in the late 1990s. This corresponds with the 294interannual changes in copepod abundance and biomass observed in the present study 295(Fig. 3). In the present study, the copepod community greatly varied between cold years (1995, 1997, 1999 and 2007-2009) and warm years (1998, 2001-2005) on both 296297the Outer and Middle shelves. In warm years, oceanic species decreased on the Outer 298shelf, but showed no change on the Middle shelf, and the abundance of all copepod 299species drastically decreased on both the Outer and Middle shelf (Fig. 5). Thus, these 300 findings clearly indicate that both the wet mass and copepod community structure 301 showed interannual changes on the southeastern Bering Sea shelf (Fig. 8).

Physical and biological changes have occurred in the southeastern Bering Sea shelf over the last two decades (Fig. 8). During the 1997 El Niño/Southern Oscillation (ENSO) event, there were anomalous atmospheric conditions over the southeastern Bering Sea shelf (Overland et al., 2001), a warm surface layer, a bloom of the coccolithophorid *Emiliania huxleyi* (Sukhanova and Flint, 1998) and mass mortality of

307 short-tailed shearwaters (P. tenuirostris) (Baduini et al., 2001). Short-tailed shearwaters prey on large zooplankton, especially adult euphausiids (Hunt et al., 1996), 308 and the starvation of shearwaters that occurred during the summer of 1997 may have 309 310 been due to low densities of adult euphausiids (Stockwell et al., 2001) or difficulty in 311 finding prey patches due to low water transparency caused by the coccolithophore 312blooms (Lovvorn et al., 2001). Thus, the mass mortality of short-tailed shearwaters 313 may not have been related with the pattern of mesozooplankton abundance and biomass 314 described in this study.

315Coccolithophore (*E. huxlevi*) blooms in recent times were first observed in this 316 region in September, 1997 (Vance et al., 1998). Large-scale coccolithophore blooms 317were also observed in June, 1998 and 2000 (Sukhanova and Flint, 1998; Iida et al., 2002, 318 2008, 2012). As an ecological consequence of coccolithophores, negative effect for 319 short-tailed shearwaters and their nutritional values for copepods should be considered. 320 Information in the literature regarding ingestion of coccolithophores by copepods is 321contradictory. For example, Huskin et al. (2000) state that E. huxleyi is difficult to 322digest and has low nutritional value, it is not a preferred food item for *Calanus* species. 323However, Nejstgaard et al. (1997) reported that about 75% of the carbon consumed by Calanus finmarchicus during an E. huxleyi bloom came from E. huxleyi (> 30×10^6 cells 324 L^{-1}), and high abundance of *Calanus* spp. was observed on the Inner shelf during a 325coccolithophore bloom (Coyle and Pinchuk, 2002). The latter observations suggest 326 327 that coccolithophore production may have little effect on *Calanus* spp. In fact, years 328 of coccolithophore blooms (1997, 1998 and 2000) did not correspond with years of low

copepod abundance or biomass (Fig. 8B). This discrepancy between coccolithophore blooms and copepod abundance could be explained if coccolithophorids are ingested by *Calanus* species (Nejstgaard et al., 1997) and have some nutritional value. As the other possible cause, turbid waters of coccolithophore bloom may provide copepods refuge from visual predators (Lovvorn et al., 2001).

During the study period, there were cold (1995, 1997, 1999, 2007-2010), 334 335 average (1996, 2000, 2006) and warm years (1998, 2001-2005) (Fig. 2A). During cold 336 and warm years, the timing and magnitude of the spring bloom differ (Stabeno et al., 2001; Hunt et al., 2002, 2011). During cold years, the sea ice remains during the 337 338 severe winter storms, and ice-edge blooms occur. In warm years, the sea ice melts 339 before the spring storms, and strong wind mixing delays the start of the bloom until 340 stratification occurs when the thermocline develops. From long-term (1960–2005) 341simulation modeling, Jin et al. (2009) reported that in cold years (PDO < -1), 342phytoplankton blooms start early, last long and are large, while in warm years (PDO > 3431), they occur later with short pulses and are small. These differences in bloom timing between cold and warm years have also been recorded in mooring observations in this 344345region (Hunt et al., 2011).

Copepod abundance and biomass on both the Outer shelf and Middle shelf were high during cold years (1995, 1997, 1999 and 2007-2009) and low during warm years except 1998 (2001–2005) (Fig. 3, Table 3). The dominant copepod, *Calanus* spp., has an extended spawning period; it starts well before the spring phytoplankton bloom (February) and ends in May, however the survival rate of early developmental

stages depends on whether they encounter a phytoplankton bloom or not, and 351recruitment of copepodid stages increases if they encounter early ice associated blooms, 352353which are highly productive (Baier and Napp, 2003). Low temperatures in cold years 354may also lead to greater abundance of the small copepods Pseudocalanus spp. and Acartia spp. especially in the Middle shelf (Table 3). Cold temperature may provide 355356the longer growing season and high primary productivity of ice-edge blooms, sufficient 357 food condition for copepods, resulting in increased abundance of both large and small 358copepods.

359 In this study, copepod abundance on both the Outer and Middle shelf was 360 negatively correlated with temperature (Table 3). In contrast, on the Inner shelf, 361positive correlations were reported between spring temperatures and the abundance of 362 three copepods (Calanus spp., Pseudocalanus spp. and Acartia spp.), and between 363 summer temperatures and two copepods (Pseudocalanus spp. and Acartia spp.) (Coyle 364 and Pinchuk, 2002). This suggests that the response of copepods to temperature may vary between the Outer-Middle shelf (negative, this study), and the Inner shelf (positive, 365366 Coyle and Pinchuk, 2002). In this study, differences between cold and warm years on 367 the Middle shelf were observed in copepod abundance, but not in species composition 368 (group D and E in Fig. 4C). This is partly because both large and small copepods were 369 abundant in cold years and less abundant in warm years (Table 3). Coyle et al. (2008) compared the summer zooplankton community on the Middle shelf between a cold 370 371(1999) and warm year (2004), and reported that the dominant species shifted from large 372species in the cold year to small species in the warm year. However, this phenomenon

was not observed during the 16 years of the present study. Our results showed that
both large and small-sized copepods showed a clear negative correlation with habitat
temperature (Table 3).

376 Similar to copepods, the biomass of the jellyfish *Chrysaora melanaster* is high in cold years and low in warm years (Brodeur et al., 2008). The increase in cold years 377378 may be due to improved feeding conditions for polyps (mass sinking of phytoplankton 379 after ice-edge bloom), and subsequent abundant food (copepods) for pelagic medusae 380 may improve survival and growth (Brodeur et al., 2008). In contrast, oceanic blooms 381 in warm years are less productive and provide less food (sinking phytoplankton) for 382polyps, as well as less food (copepods) for planktonic medusae, resulting in low growth 383 and survival of polyps and medusae.

384 Interannual changes in chaetognath abundance and biomass were somewhat 385similar to the changes in copepods (Fig. 8). This may be an example of bottom-up 386 control on predatory chaetognath abundance by prey (copepods) abundance. 387 Chaetognaths in this region change their food items and size at maturity between cold 388 and warm years; in cold years, they prey on large *Calanus* spp. and mature at large size, 389 while in warm years, they feed on small Pseudocalanus spp. and mature at small size 390 (Baier and Terazaki, 2005). The TL of most mature (Stage III) individuals in the 391present study was > 20 mm in cold years (1997, 1999 and 2007-2009), and < 20 mm in warm years (1998, 2001, 2002 and 2005), and no mature individuals were observed in 392393 the warm years 2003 and 2004 (Fig. 7). These facts support the finding of Baier and 394 Terazaki (2005), who reported that sufficient food conditions in cold years result in a

395 large size at maturity, while food limitation in warm years induces small size at 396 maturity.

397 For the early life history stages of walleye pollock, an important fisheries 398 resource in this region, food conditions differ between cold and warm years (Napp et al., 2000; Hunt et al., 2011). In cold years, there may be a mismatch between the 399 production of pollock larvae and their prey (Napp et al., 2000), but an early start of the 400 401 ice-edge bloom and high abundance of lipid-rich prey for age-0 fish (copepods and 402euphausiids) results in high survival rates and low cannibalism (Hunt et al., 2011; 403 Heintz et al., this issue). In warm years, the late start of the open-water bloom results 404 in a low abundance of lipid-rich prey for the age-0 pollock and chaetognaths. Under 405this food-limited condition, growth and survival of walleye pollock juveniles decrease, 406 and the proportion of cannibalism may increase. This results in a decrease in 407recruitment (Hunt et al., 2011, Heintz et al., this volume). Thus, the timing and 408 magnitude of primary production related with climate change affects production through the entire food web structure on the southeastern Bering Sea shelf. 409

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576	Basin.	Limnol. Oceano	ogr. 18, 750–7	56.					

578 Figure captions

- Fig. 1. Location of the sampling stations in the southeastern Bering Sea during the
 summers of 1994–2009 (circles). Solid symbols denote stations where
 sampling was conducted in all years from 1994–2009, providing the data used
 for annual comparisons. Line indicates 166°W transect where the hydrographic
 data were analyzed (cf. Fig. 2).
- Fig. 2. (A) Temperature and (B) chlorophyll *a* stations along 166°W during summers
 of 1994–2009.
- 586 Fig. 3. Regional variations in (A) copepod abundance and (B) biomass in the 587 southeastern Bering Sea during summers of 1994–2009.
- 588Fig. 4. Copepod assemblages: (A) dendrograms from the cluster analysis based on 589copepod abundance. Six groups (A-F) were identified with dissimilarity at 59020% (dashed line). Numbers in the parentheses indicate quantity of stations included in each group. (B) Nonmetric multi-dimensional scaling plots of the 591six groups, with arrows and percentages indicating directions of environmental 592593parameters and coefficient of determinations (r^2) , respectively. Lat: latitude, 594Lon: longitude, SSS: sea surface salinity, SST: sea surface temperature, BS: bottom salinity. (C) Mean abundance with standard deviations and species 595596 diversity (H') (upper) and taxonomic composition of each group (lower).
- Fig. 5. (A) Spatial and temporal distribution of six copepod communities identified
 from Bray-Curtis dissimilarity analysis. Annual distribution of groups on the
 (B) Outer shelf and (C) Middle shelf.

- Fig. 6. Regional variations in *Parasagitta elegans*: (A) abundance and (B) biomass
 during summers of 1996–2009.
- Fig. 7. Annual patterns of *Parasagitta elegans* length frequency on Middle shelf
 during summers of 1996–2009. n: number of measured specimens.
- Fig. 8. (A) Annual changes in biological/environmental conditions, (B) copepod
 abundance and biomass and (C) chaetognath (*Parasagitta elegans*) abundance
 and biomass on Middle shelf of the southeastern Bering Sea during summers of
 1994–2009. Values are means and error bars indicate standard deviations.
 For references in (A), 1: Sukhanova and Flint (1998), Iida et al. (2002, 2008), 2:
 Baduini et al. (2001), 3: McFarlane et al. (2000), 4: Brodeur et al. (2008), Hunt
 et al. (2010), 5: Stabeno et al. (2012).

Table 1. Calanoid copepod species collected in the southeastern Bering Sea during summers of 1994–2009. Major region of occurrence is also shown for each species. ●: Large oceanic copepods that have a diapause phase in deep layer (Miller et al., 1984). Region abbreviations are; I: Inner shelf, M: Middle shelf, O: Outer shelf, S: Slope and B: Basin. *: Rare species (occurred only <2% of the whole stations) omitted from data for cluster analysis (cf. Fig. 4).

Species	Major region of occurrence
Acartia longiremis (Lilljeborg, 1853)	I, M, O
A. tumida Willey, 1920	М, О
* Aetideopsis rostrata Sars, 1903	В
*Aetideus pacificus Brodsky, 1950	В
Calanus spp.	M, O, S, B
Candacia columbiae Campbell, 1929	S, B
Centropages abdominalis Sato, 1913	I, M
Epilabidocera amphitrites (McMurrich, 1916)	Ι
 Eucalanus bungii Giesbrecht, 1892 	O, S, B
Eurytemora herdmani Thompson and Scott, 1897	7 I
Gaetanus intermedius Wolfenden, 1905	S, B
G. simplex Brodsky, 1950	S, B
Heterorhabdus tanneri (Giesbrecht, 1895)	S, B
Metridia pacifica Brodsky, 1950	O, S, B
Microcalanus pygmaeus (Sars, 1900)	М, О
 Neocalanus cristatus (Kröyer, 1848) 	O, S, B
 N. flemingeri Miller, 1988 	O, S, B
 N. plumchrus (Marukawa, 1921) 	O, S, B
Paraeuchaeta elongata (Esterly, 1913)	В
Pleuromamma scutullata Brodsky, 1950	S, B
Pseudocalanus spp.	I, M, O, S, B
*Racovitzanus antarcticus Giesbrecht, 1902	В
Scolecithricella minor (Brady, 1883)	O, S, B
* S. ovata (Farran, 1905)	В
Tortanus discaudatus (Thompson and Scott, 189	7) I
Undinopsis pacificus Brodsky, 1950	O, S

Domain		F -value $_{df_1, df_2}$			
	Species	Year _{15,30}	Station _{2,30}		
Outer shel	f				
	Acartia spp.	2.136*	1.215 ^{ns}		
	Calanus spp.	1.259 ^{ns}	3.027 ^{ns}		
	Eucalanus bungii	1.402 ^{ns}	2.610 ^{ns}		
	Metridia pacifica	2.060*	1.464 ^{ns}		
	Neocalanus spp.	5.228***	1.956 ^{ns}		
	Pseudocalanus spp.	1.190 ^{ns}	0.194 ^{ns}		
/liddle sh	elf				
	Acartia spp.	6.574****	2.798 ^{ns}		
	Calanus spp.	6.026****	3.745*		
	Pseudocalanus spp.	5.791****	0.195 ^{ns}		

Table 2. Variance analysis (two-way ANOVA) on year-to-year and station-to-station differences in abundance (inds. m⁻²) of dominant copepod taxa on Outer and Middle shelf of the southeastern Bering Sea during summers of 1994–2009. *: p < 0.05, **: p < 0.01, ***: p < 0.001, ****: p < 0.0001, ns: not significant.

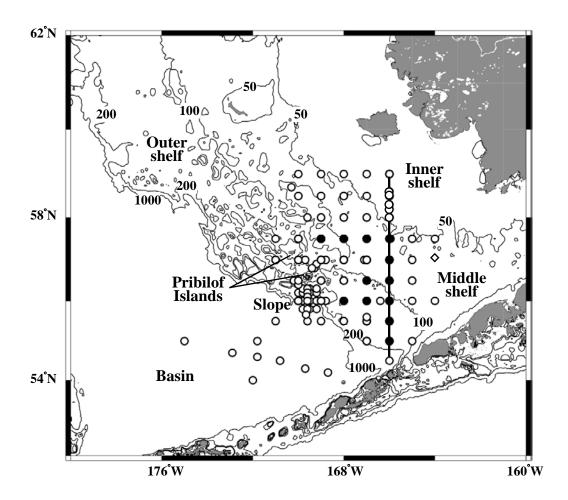
Table 3. Correlation coefficients (r) between abundance of dominant copepods and various environmental parameters in the
southeastern Bering Sea during summers of 1994–2009.

*: p < 0.05, **: p < 0.01, ***: p < 0.001, ****: p < 0.0001.

Domain	Temperature (n= 91)			Salinity (n= 91)	Chlorophyll <i>a</i> (n=45)	
Species	Whole water column	Upper layer	Lower layer	Whole water column	Whole water column	
Outer shelf						
Acartia spp.	0.207 *	0.149	0.186	-0.140	0.067	
Calanus spp.	-0.423 ****	-0.101	-0.406 ***	-0.294 **	0.192	
Eucalanus bungii	0.138	-0.143	0.307 **	0.517 ****	0.071	
Metridia pacifica	-0.231 *	-0.258 *	-0.105	0.076	0.370 *	
Neocalanus spp.	-0.285 **	-0.404 ***	-0.108	0.252 *	0.369 *	
Pseudocalanus sp	р0.159	-0.142	-0.044	-0.028	0.341 *	
Middle shelf	Middle shelf					
Acartia spp.	-0.410 ***	-0.146	-0.431 ****	-0.506 ****	-0.111	
Calanus spp.	-0.448 ****	-0.340 **	-0.457 ****	-0.407 ***	-0.126	
Pseudocalanus sp	pp0.567 ****	-0.292 **	-0.539 ****	-0.484 ****	-0.037	

Parameters	Outer shelf	Middle shelf
Temperature Whole water column	-0.137	-0.519 ****
Upper layer	-0.057	-0.301 **
Lower layer	-0.187	-0.547 ****
Salinity Whole water column	-0.172	-0.513 ****
Copepods Acartia spp.	0.343 **	0.563 ****
Calanus spp.	0.431 ****	0.503 ****
Eucalanus bungii	0.129	_
Metridia pacifica	0.195	_
Neocalanus spp.	0.093	_
Pseudocalanus spp.	0.291 **	0.566 ****

Table 4. Correlation coefficients (*r*) between abundance of chaetognath *Parasagitta elegans* and various parameters (water temperature, salinity and copepod abundance) on the Outer shelf and Middle shelf in the southeastern Bering Sea during summers of 1996–2009. *: p < 0.05, **: p < 0.01, ***: p < 0.001, ****: p < 0.0001. —: not applicable.



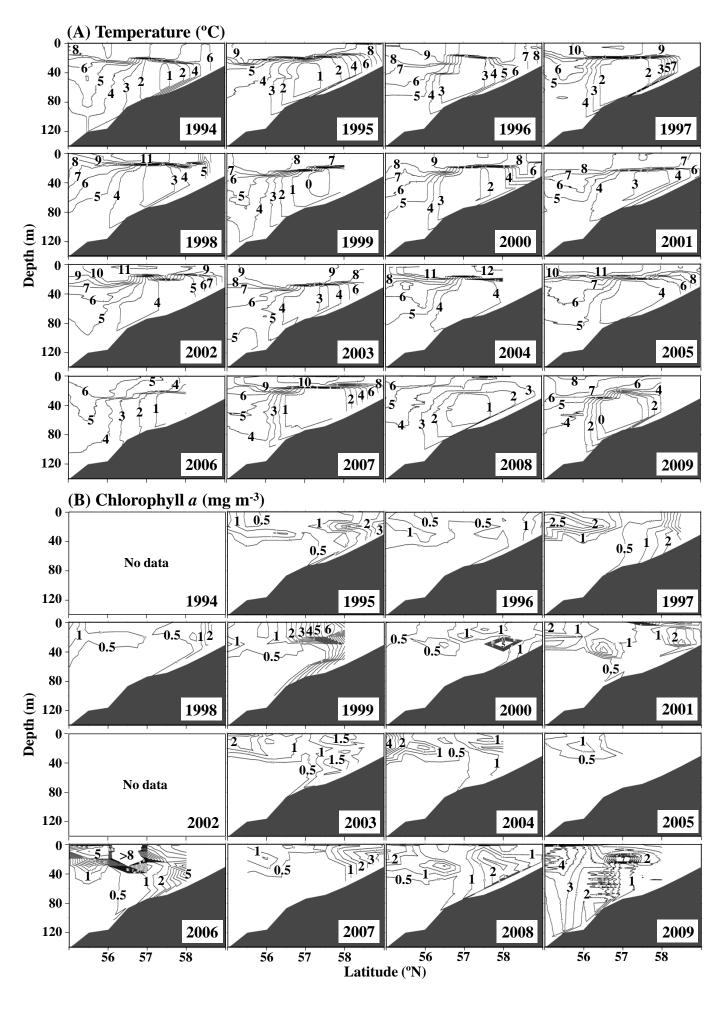


Fig. 2. Ohashi et al.

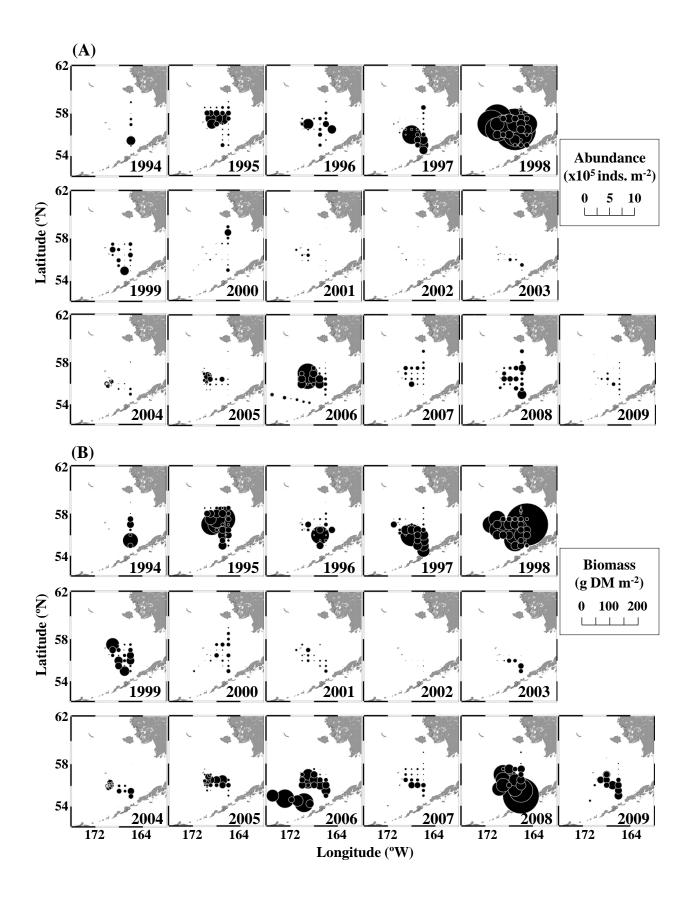


Fig. 3. Ohashi et al.

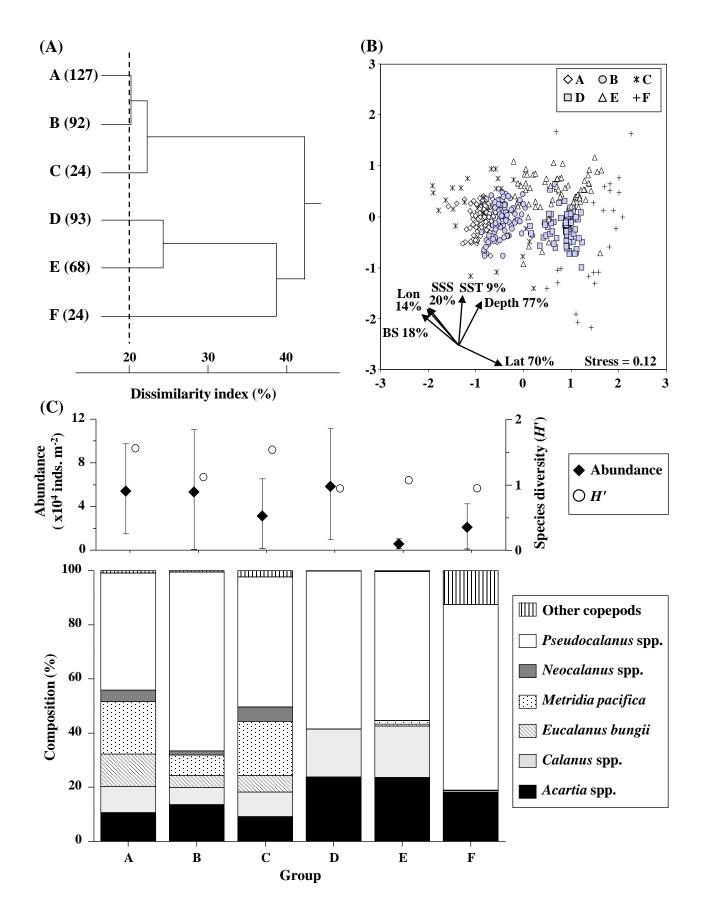


Fig. 4. Ohashi et al.

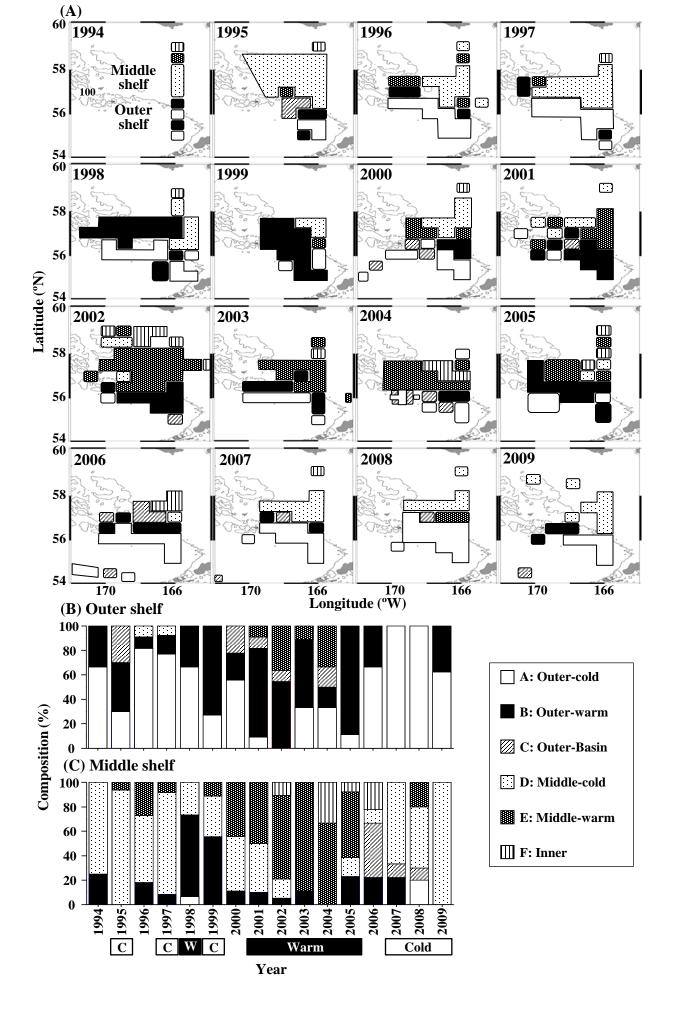


Fig. 5. Ohashi et al.

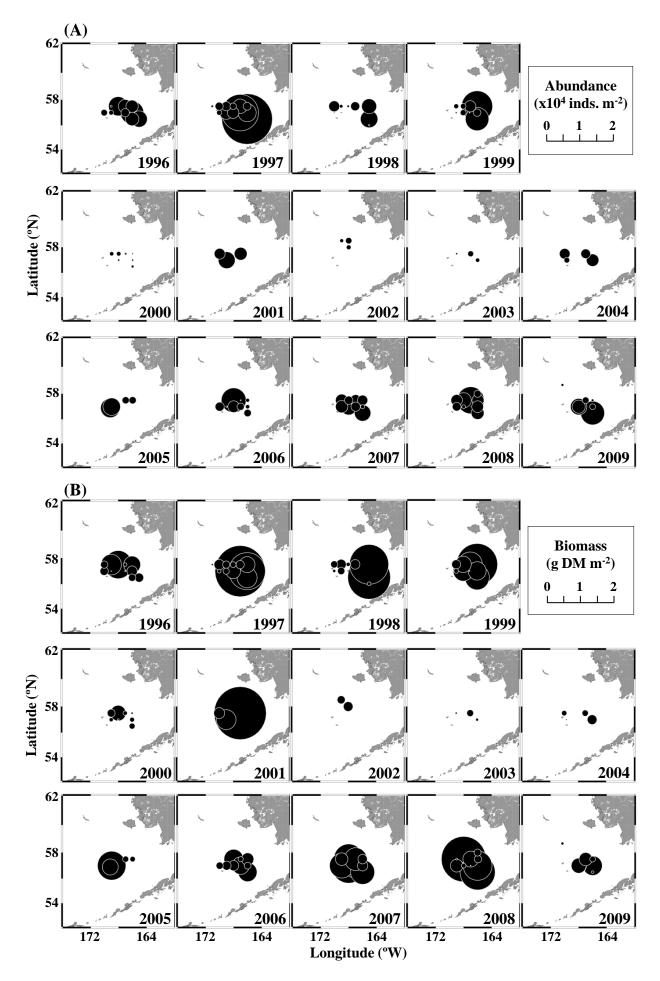


Fig. 6. Ohashi et al.

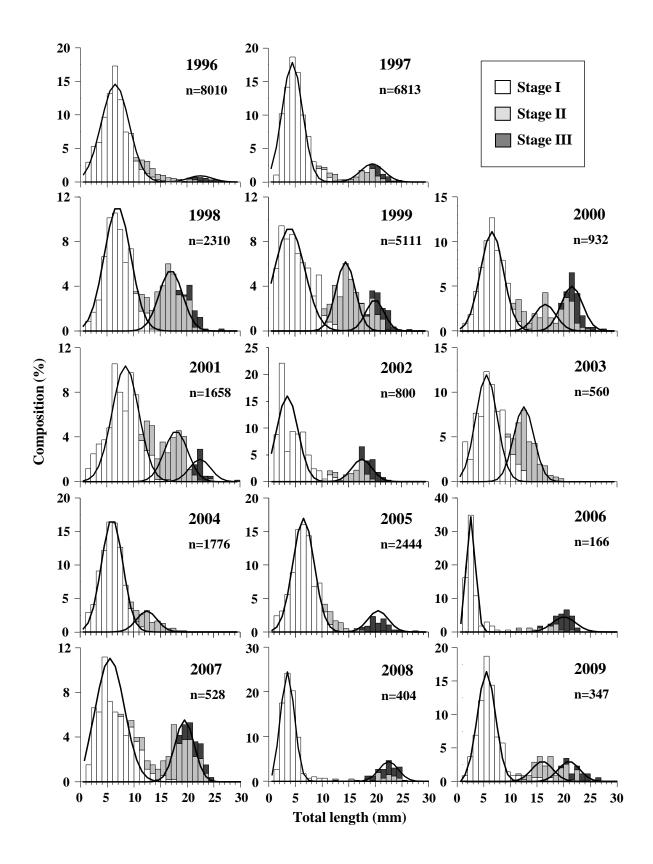


Fig. 7. Ohashi et al.

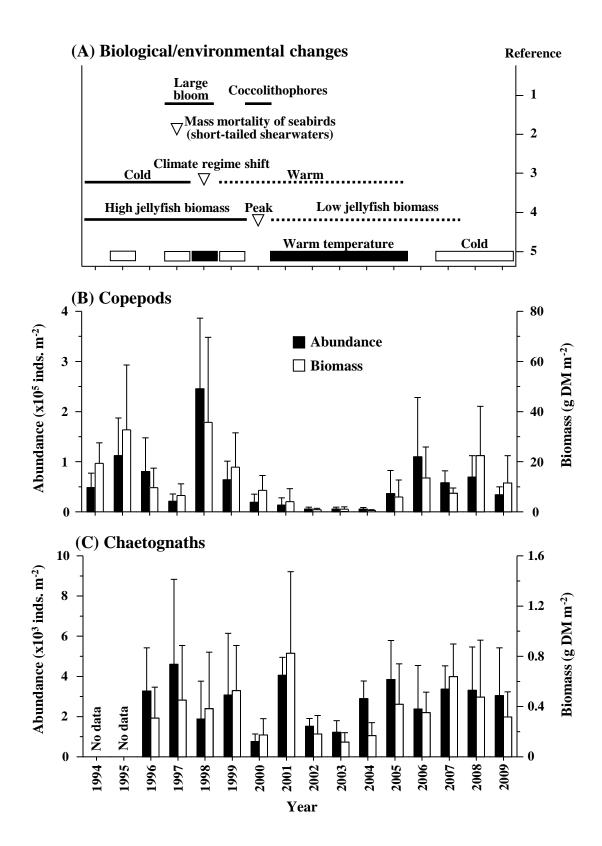


Fig. 8. Ohashi et al.