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Trophic niche of a neustonic copepod community sustained by local production in high- and low-salinity summer water masses in the northeastern East China Sea

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We investigated the trophic dynamics between neustonic copepod communities and particulate organic matter (POM) sourced from contrasting water masses, i.e., high- and low-salinity areas, within the northeastern East China Sea. The sampling of neustonic copepod communities occurred three times in the summer (June, August, and September) of 2021, and considered the influx of the Yangtze River Diluted Water (YRDW) and the extension of the Tsushima Warm Current (TWC). To examine the relationship between the neustonic copepods and surface POM, we employed stable isotope techniques. Surprisingly, the neustonic copepod community exhibited a reluctance to feed on surface POM. The neustonic copepod community in the YRDW-dominated area in August showed higher $\delta^{15}\text{C}$ values than those in the TWC-dominated area. The trophic groups within the neustonic copepod population appeared to consume selectively depending on the individual preferred prey sizes and composition. Using the POM from the water masses that affected the food sources consumed by neustonic copepods, we found a strong dependence on prey items originating from the TWC. This result suggests that the marine neustonic copepod community does not rely significantly on food sources derived from riverine freshwater.

KEYWORDS

neustonic copepods, Tsushima warm current, Yangtze River Diluted Water, stable isotopes, trophic niche

1 Introduction

Neuston, as defined by Naumann (1917), are organisms that float near the water surface. They are classified into the following three ecological categories: euneuston, which permanently exist on the surface; facultative neuston, which move to the surface by diel vertical migration; and pseudoneuston, which live in the deeper layers but sporadically occur on the surface (Zaitsev, 1971). Facultative neuston and pseudoneuston tend to be commonly regarded as temporary neuston (Rawlinson et al., 2005). Neuston are important components of the marine ecosystem because they connect the surface and deeper layers of the ocean, providing food and energy for many organisms (Albuquerque et al., 2021; Dipper, 2022). Among them, neustonic copepods are a dominant component of marine zooplankton, and their feeding habits fall into two categories: particulate feeders (detritivores and omnivores) and carnivores (Hirabayasi et al., 2016; Kaji et al., 2019), as determined based on their mouthpart morphology, rather than on food type (Ohtsuka and Onbé, 1991; Schmidt et al., 2004; Michels and Stanislav, 2015). This diversity adds complexity to their role in the ecosystem. Suspended particulate organic matter (POM) serves as a significant food source for neustonic copepods (Chen et al., 2018; Im and Suh, 2019), because it consists of various components, such as zooplankton and fish fecal pellets, marine aggregates (marine snow), and phytodetritus from sinking phytoplankton (Turner, 2015). Considering the diel vertical migration of neustonic copepods, there is a potential for them to ingest POM within their vertical migration range. Neustonic copepods exhibit selectivity regarding the ingestion of POM within the pico- to microsize range (Im et al., 2015; Im and Suh, 2023). Understanding their feeding strategies from a morphological perspective is essential for comprehending their nutritional dynamics and trophic interactions within pelagic food webs, particularly in the context of their diel vertical migration.

The northeastern East China Sea (nECS) features an expansive continental shelf with water depths of less than 200 m along the southern coast of the Korean peninsula. The Tsushima Warm Current (TWC), which is a branch of the Kuroshio, flows northward from the Okinawa Trench, passing through the eastern region of Jeju Island and the Korea Strait before entering the East/Japan Sea (Lie and Cho, 2016; Park et al., 2017). During the summer monsoon (late June to August), Yangtze River Diluted Water (YRDW) travels eastward from mainland China and meets the TWC, creating a salinity front (Park et al., 2017). The YRDW also extends into the northwestern waters around Jeju Island, establishing a front through tidal mixing with the South Korea Coastal Water (SKCW) (Meng et al., 2020). These fronts significantly influence the temporal and spatial fluctuations in water masses, which, in turn, affect the composition and distribution of neustonic copepod species (Jeong et al., 2014; Choi et al., 2020). Variations in size-based phytoplankton compositions, such as picosized cyanobacteria in offshore warm waters vs. micro-sized diatoms in coastal cold waters, correspond to the distribution of these water masses (Yoon et al., 2020; Kang et al., 2022). In the nECS, the feeding strategies of *Euchaeta* sp. and *Pleuromamma* spp. have been found to vary according to latitude

using fatty acid biomarkers (Ju et al., 2011). Although there are reports of the trophic levels and energy flow within the zooplankton community (Chi et al., 2020; Kim et al., 2022), and of feeding relationships with apex predators (such as jellyfish and fish) in this shelf sea (Wang et al., 2020; Bai et al., 2021), the precise feeding relationships at lower trophic levels, particularly regarding the different copepod and phytoplankton species compositions in contrasting water masses, remain unknown.

Gut-content analyses serve to confirm the dietary items consumed by the copepods within a brief time frame (Ohtsuka and Onbé, 1991; Nishida and Ohtsuka, 1996; Wu et al., 2004); however, it is significantly affected by the extent of digestion, contingent upon food selection (Baker et al., 2014). In instances where no hard parts are present, the identification of food items becomes challenging because of rapid digestion. In contrast, stable isotope analysis, which is a relatively more recent technique, offers a more precise depiction of dietary assimilation over an extended temporal scale (Chen et al., 2018; Im and Suh, 2019; Pizarro et al., 2019). Consequently, carbon and nitrogen stable isotopes possess an advantage in providing longer-term cumulative insights into the dietary preferences of consumers, thus offering a valuable means to interpret trophic levels within marine ecosystems (Minagawa and Wada, 1984; Kang et al., 2021). Carbon stable isotope ratios ($\delta^{13}\text{C}$) are particularly suited for identifying food sources because of their minimal fractionation (0‰–1‰) between diets and consumers. Therefore, $\delta^{13}\text{C}$ values have found application as useful indicators of the original carbon source in marine food webs. Nitrogen stable isotope ratios ($\delta^{15}\text{N}$) are well suited for comprehending the trophic levels of organisms because of their relatively substantial and consistent increments of around 3.4‰ per trophic step (i.e., between prey and consumers) (Minagawa and Wada, 1984; Post, 2002).

In this research, we analyzed the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of neustonic copepods and size-fractionated POM to investigate the variations in trophic niches within the neustonic copepod community in contrasting water masses. We anticipated that the food sources of neustonic copepods would exhibit differences because of the influence of the distinct water masses (with unique characteristics) of the nECS. Furthermore, we expected the trophic position of neustonic copepods to vary according to their respective food sources. Based on these expectations, we hypothesized that, if the feeding habits of neustonic copepods (i.e., carnivorous or particle feeders) resulted in distinct trophic relationships, their $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values would exhibit variations between the different water masses. Consequently, this would lead to a spatial shift in the isotopic niche of the entire copepod community as a response to the differing compositions of copepod and phytoplankton species within these contrasting water masses. Specifically, because the YRDW plays a crucial role in shaping the spatial distribution of the surface water masses in the nECS, our research aimed to elucidate the impact of YRDW intrusion on the complex trophic interactions that underlie the neustonic copepod food web in this region during the summer monsoon. To accomplish this, we performed a comparative analysis of food web characteristics—focusing on community-wide metrics—regarding the temporal features of the spatial distribution of surface water masses.

2 Material and methods

2.1 Site description

The study area included areas with other physicochemical characteristics, such as YRDW outflow from China during the summer rainy period, TWC expansion with high water temperature, salinity, and inflow of SKCW into Korea. The discharge of YRDW varies from year to year. According to the YRDW discharge provided by the ocean and fisheries research institute in the Jeju special self-governed province (<https://www.jeju.go.kr/jori/monitor/imap.htm>), YRDW was continuously discharged at about 60,000 tons in June 2021, which was the largest value of the year. It gradually decreased from August to September. In contrast, TWC has increased relatively. The inflow of YRDW with low salinity and density characteristics can characterize the fluctuations in water masses across time and space. The variation of heterogeneous water masses affects the copepod community structure and, thus, the complexity of the food web. We performed three onboard observations considering the discharge of YRDW and the expansion of the TWC. We collected POM and neustonic copepod specimens in June (western Jeju Island), August (southern Jeju Island), and September (northeastern Jeju Island) 2021 using a research vessel (Saedongbaek) from the Chonnam National University in South Korea (Figure 1 and Supplementary Table 1). We utilized the period-specific geographical areas identified by Choi et al. (unpublished) based on the neustonic copepod community structure: areas A (affected by the YRDW) and B (remaining shelf water that had a relatively high salinity) in June; areas A (affected by the YRDW) and B (affected by the TWC) in August; and areas A (broad frontal area formed over the coast of Korea) and B (strongly influenced by the TWC) in September. The

range of salinity by areas within month is as follows: areas A (25.7 to 31.3) and B (28.8 to 32.0) in June; areas A (27.9 to 29.7) and B (27.9 to 30.9) in August; and areas A (31.1 to 31.6) and B (31.2 to 32.3) in September.

2.2 Sampling methods

The seawater samples (45 L) of the surface layer were used to treat pico- to microsize POM samples. All filter paper used for treatment was prewashed and precombusted (450 °C for 4 h), then filtered in triplicate. The seawater samples were continuously prefiltered through 200 μm and 20 μm sieves, and particles with a size above 20 μm were collected in \varnothing 47 mm GF/F filter paper (0.7 μm pore size; Whatman, Merck KGaA, Darmstadt, Germany), to obtain microsize POM (mPOM). Nanosized POM (nPOM) was collected by filtering seawater through a 20 μm sieve on \varnothing 47 mm GF/D filter paper (2.7 μm pore size; Whatman), whereas picosized POM (pPOM) was collected by filtering water through GF/D filter paper with GF/F filter paper. All filters were immediately frozen and stored at -80°C until the stable isotope analysis was performed. In the laboratory, all filter samples were completely dried (60 °C for 48 h) in an oven (ThermoStable SON-155, Daihan Scientific, Seoul, Korea). For stable isotope analysis, the filter samples were cut in half and, to remove inorganic carbon and obtain accurate $\delta^{13}\text{C}$ values, acid treatment was performed by fuming it within the acid-containing desiccator overnight. The remaining filter samples were used to obtain $\delta^{15}\text{N}$ values. All filter samples were folded and packed into a tin disk.

All copepods collected with neuston net were considered neustonic copepods because of the difficulty in distinguishing euneuston, facultative neuston and pseudoneuston in this study.

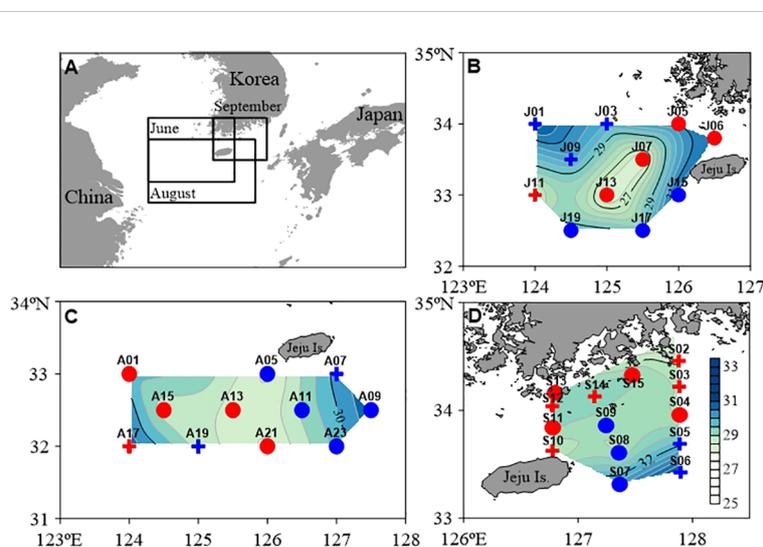


FIGURE 1

Map showing the distribution of surface salinity at the sampling station [(A), study area; (B), June (J); (C), August (A); (D), September (S)] in the northeastern East China Sea. For each sample, the study area was divided into (A) (red, low-salinity) and (B) (blue, high-salinity) according to the non-metric multidimensional scaling (nMDS) model via a Bray–Curtis similarity index correction based on the distribution of neustonic copepods at the species level. The areas used for stable isotope analysis were selected among the two study areas (circles). Summary of the study area is provided in Supplementary Table 1.

A neuston net (70 cm width; 40 cm height; mesh size 333 μm) used for neustonic copepod sampling was towed horizontally at a ship speed of 2.5 to 4 knots at a distance of 2–5 m from the shipside. A flowmeter (model 438 115, Hydro-Bios Co., Altenholz, Germany) was attached to the mouth of the net, to estimate the volume of the filtered seawater. The collected neustonic copepod samples were divided on board using a Folsom-type plankton splitter, for species identification and stable isotope analysis. The samples used for species identification were preserved in 5%–8% neutralized formalin–seawater solution. Subsamples were frozen (-80°C) on board the ship and stored frozen in the laboratory for subsequent stable isotope analysis. After transportation to the laboratory, preserved samples were once more divided using a Folsom plankton splitter before estimating abundance. Neustonic copepod species were identified and counted using a dissecting stereo microscope (SMZ645, Nikon, Tokyo, Japan). When detailed observation was required for species identification, the appendages and any other parts required for this were dissected and observed on a slide glass by mounting in 70% glycerin under high magnification of an optical microscope (ECLIPSE E200, Nikon, Tokyo, Japan). The classification system was devised based on the World Register of Marine Species (WoRMS, <https://www.marinespecies.org>). The density of neustonic copepod was measured and converted into abundance per unit volume (ind. m^{-3}). Log ($x + 1$) transformation of data was performed to avoid bias due to differences in population abundance between survey areas and species and normalize data distribution. To measure the similarity between neustonic copepod stations, hierarchical cluster analysis was performed based on the Bray–Curtis similarity index (Shannon and Weaver, 1963) using the PRIMER software (version 6.1.6., Auckland, New Zealand; Clarke and Gorley, 2006). Subsequently, the cluster was arranged using the non-metric multidimensional scaling (nMDS) model (Figure 1). Based on the nMDS, stations with high abundance of neustonic copepods were selected, and the sampling area for stable isotope analysis was reorganized into areas A and B for each month. The neustonic copepod species in the reconstructed areas were converted to percentages to select dominant copepod species.

The chosen dominant copepod species were as follows (Table 1): areas A (*Calanus sinicus* and *Labidocera rotunda*) and B (*Paracalanus parvus* s. l. and *L. rotunda*) in June; areas A (*Acartia pacifica*, *Canthocalanus pauper*, *L. rotunda*, *Centropages furcatus*, *Temora discaudata* and *T. turbinata*) and B (*A. pacifica*, *L. acuta*, *C. furcatus*, *T. discaudata* and *C. pauper*) in August; and areas A (*C. pauper*, *C. furcatus*, *L. acuta*, *Subeucalanus crassus*, *T. turbinata* and *T. discaudata*) and B (*Undinula vulgaris*, *L. acuta* and *C. pauper*) in September. In addition to the dominant species, the minor species were selected based on their feeding relationships (area A in June: *Pontellopsis yamadae*; and area B in September: *T. discaudata*, *S. crassus* and *C. furcatus*). The feeding type of the selected neustonic copepod species was classified as either particle feeder or carnivorous based on the literature (Anraku and Omori, 1963; Acros and Fleminger, 1986; Price and Paffenhöfer, 1986; Turner, 1986; Ohtsuka and Onbé, 1991; Ohman and Runge, 1994; Hu et al., 2014; Al-Hanoun and Mayya, 2020).

Sufficient individuals of each species (0.3–0.5 mg) selected for stable isotope analysis were collected onto precombusted GF/F (\emptyset

47 mm), dried in an oven (60°C for 48 h), and packed (triplicated) into a tin capsule. Moreover, radiolarians, which were predominant in area B in June, were considered as a food source for the copepods and were selected for analysis (Supplementary Figure 1). Radiolarians have silica-based skeletons, and CaCO_3 content is minimal or negligible (Anderson, 2001; Matsuzaki et al., 2016). Single-celled species range in size from less than 100 μm to very large species reaching 1 to 2 mm in diameter (Anderson, 2001), but radiolarians from the northern East China Sea are reported to be small case (Matsuzaki et al., 2016). Additionally, the radiolarians in this study are relatively very small compared to *L. rotunda* (body length: >1.0 mm; Razouls et al., 2005–2024a). Thus, it was determined that there was a risk of loss during acid treatment; and, after drying, the samples were packed into a tin capsule with a weight of 0.3–0.5 mg.

2.3 Stable carbon and nitrogen isotope analysis

Stable isotope data for all packed POM, copepod, and radiolarian samples were obtained using continuous-flow isotope ratio mass spectrometry (CF-IRMS). Samples were oxidized at a high temperature (980°C) in an Elemental Analyzer (Flash 2000, Thermo Scientific, Waltham, MA), and the resultant CO_2 and N_2 were swept by Helium carrier gas in continuous flow mode and analyzed for stable isotope ratio using a ConFlo IV (Thermo Scientific) interfaced with an isotope ratio mass spectrometer (Delta V^{Plus}, Thermo Scientific). Analyses were conducted by the Alaska Stable Isotope Facility (ASIF) at the University of Alaska Fairbanks. The abundance of each stable isotope is expressed in δ notation as parts per thousand (‰) deviation from the international standards Vienna Pee Dee Belemnite for carbon and Air for nitrogen according to the following formula:

$$\delta X = \left[\frac{R_{\text{sample}}}{R_{\text{standard}}} - 1 \right] \times 10^3$$

where X is ^{13}C or ^{15}N , and R_{sample} and R_{standard} are the ratios of heavy to light isotopes in the samples and standards, respectively. All stable isotope results were subjected to two-point calibrations. Stable isotope ratios were calibrated using international standards of sucrose (ANU $\text{C}_{12}\text{H}_{22}\text{O}_{11}$; National Institute of Standards and Technology (NIST), Gaithersburg, MD, United States) for carbon and of ammonium sulfate ($[\text{NH}_4]_2\text{SO}_4$; NIST) for nitrogen. Using this calibration, the average $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ obtained for the ASIF calibration verification standard (ref/chk/peptone) were $-15.8\text{‰} \pm 0.3\text{‰}$ ($n = 10$) and $7.1\text{‰} \pm 0.1\text{‰}$ ($n = 10$), respectively, which compare well with their accepted values of -15.8‰ and 7.0‰ , respectively. The instrument precision was approximately 0.3‰ and 0.1‰ for $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$, respectively. Lipid extraction was not performed for each copepod sample because it may affect the value of $\delta^{15}\text{N}$ (Bodin et al., 2007; Mintenbeck et al., 2008); rather, the $\delta^{13}\text{C}$ of the copepod samples was calculated according to Smyntek et al. (2007), as follows:

$$\delta^{13}\text{C}_{\text{ex}} = \delta^{13}\text{C}_{\text{bulk}} + 6.3 \left(\frac{\text{C:N}_{\text{bulk}} - 4.2}{\text{C:N}_{\text{bulk}}} \right)$$

TABLE 1 The dominant species, relative abundance and feeding habits of dominant neustonic copepods in the study areas (low salinity, A and high salinity, B) by months (June, August, and September).

Month	Area	Dominant species	Relative abundance	Feeding habit	Reference
June	A	<i>Calanus sinicus</i>	84.0	PF	Ohman and Runge (1994)
		<i>Labidocera rotunda</i>	7.6	C	Anraku and Omori (1963)
		<i>Pontellopsis yamadae</i>	0.2	C	Ohtsuka and Onbé (1991)
	B	<i>Paracalanus parvus</i> s. l.	46.7	PF	Al-Hanoun and Mayya (2020)
		<i>L. rotunda</i>	34.0	C	—
August	A	<i>Acartia pacifica</i>	33.6	PF	Anraku and Omori (1963)
		<i>Canthocalanus pauper</i>	16.7	PF	Hu et al. (2014)
		<i>L. rotunda</i>	13.9	C	—
		<i>Centropages furcatus</i>	8.3	PF	Anraku and Omori (1963)
		<i>Temora discaudata</i>	7.9	PF	Acros and Fleminger (1986)
		<i>T. turbinata</i>	8.7	PF	—
	B	<i>A. pacifica</i>	42.3	PF	—
		<i>L. acuta</i>	27.2	C	—
		<i>C. furcatus</i>	12.6	PF	—
		<i>T. discaudata</i>	6.5	PF	—
		<i>C. pauper</i>	4.4	PF	—
September	A	<i>C. pauper</i>	47.4	PF	—
		<i>C. furcatus</i>	11.8	PF	—
		<i>L. acuta</i>	8.7	C	—
		<i>Subeucalanus crassus</i>	8.2	PF	Price and Paffenhöfer (1986)
		<i>T. turbinata</i>	5.0	PF	—
		<i>T. discaudata</i>	4.2	PF	—
	B	<i>Undimula vulgaris</i>	28.2	PF	Turner (1986)
		<i>L. acuta</i>	27.3	C	—
		<i>C. pauper</i>	27.1	PF	—
		<i>T. discaudata</i>	2.2	PF	—
		<i>S. crassus</i>	1.0	PF	—
		<i>C. furcatus</i>	0.8	PF	—

It was assumed that the feeding habits of species without reference is the same as that of Genus.

The minor species were selected by considering their feeding relationship: area A (*Pontellopsis yamadae*) in June and area B (*Temora discaudata*, *Subeucalanus crassus* and *Centropages furcatus*) in August. PF, particle feeder; C, carnivorous. References are to the feeding habits of dominant species.

where $\delta^{13}C_{ex}$ is the predicted $\delta^{13}C$ value of the defatted copepod sample, and $\delta^{13}C_{bulk}$ and $C:N_{bulk}$ are the $\delta^{13}C$ value and the C:N ratio of the experimental samples, respectively.

2.4 Data analysis

Any outliers identified among the calculated stable isotope analytical values were excluded from further analysis. Before the statistical analyses, the normality and homogeneity of variance for all isotopic data were confirmed using the Shapiro–Wilk test and Levene’s test, respectively. The significance of the differences in the

$\delta^{13}C$ and $\delta^{15}N$ values of neustonic copepod communities observed between the different months and areas was assessed using a permutational multivariate analysis of variance (PERMANOVA) based on the Bray–Curtis similarity method (Anderson, 2017), as well as a multilevel pairwise PERMANOVA using adonis2 and pairwise adonis2 from the “vegan” package. The significance level for all statistical tests was set at 5%. Analyses were conducted using the R software (version 4.2.3) (R Core Team, 2013).

A hierarchical cluster analysis based on the $\delta^{13}C$ and $\delta^{15}N$ values of neustonic copepod species was performed using PRIMER, version 6.1.6, to identify trophic groups according to the Bray–Curtis similarity method. Because the isotope values within

individual trophic groups met the assumptions of normality and equality of variance, multivariate and univariate analyses of variance (MANOVA and ANOVA, respectively) were performed using the Bray–Curtis similarity method, to test the significance of the differences between trophic groups. Subsequently, ANOVA of $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ was followed by multiple comparisons using Tukey's honestly significant different (HSD) test, to detect differences among the trophic groups. These statistical tests were performed at $P < 0.05$ using IBM SPSS statistics (version 27.0 IBM Corp., Armonk, NY; George and Mallery, 2019).

The trophic niche of the dominant neustonic copepod community in the two areas according to the month of the year was described using stable isotope Bayesian ellipses in R (SIBER) (Jackson et al., 2011). The Layman metric of the total area of the convex hull (TA) can be directly converted to a measure of population niche area; however, it is very sensitive to sample size and, consequently, its value increases as the sample size increases (Layman et al., 2007). In contrast, the standard ellipse area (SEA) asymptote with a sample size of 30 and the TA increase as the number of added samples increases, with a standard ellipse containing 40% of the data, regardless of sample size (Syväranta et al., 2013). The corrected standard ellipse area (SEAc) provides a highly satisfactory estimate for all sample sizes (Jackson et al., 2011; Syväranta et al., 2013); therefore, we used SEAc as a measure of the mean core population isotopic niche. Other Layman metrics included NR ($\delta^{15}\text{N}$ range), to provide information on the trophic length of the community, and CR ($\delta^{13}\text{C}$ range), to estimate the diversity of the basal resources. TA and the mean distance to the centroid (CD) were used to indicate the niche width and diversity of the diet or trophic diversity of the population, respectively.

We estimated the relative contributions of the different sources [Yangtze River Water (YRW), TWC, and riverine freshwater discharge from the Korean peninsula] to the trophic groups of neustonic copepods (based on cluster groups) using a Bayesian isotope mixing model. We employed the Stable Isotope Mixing Model in R (SIMMR)—an upgraded version of the Stable Isotope Analysis in R (SIAR) (Parnell et al., 2013)—for this purpose. The $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ end-member values of POM originating from the YRW were obtained from the values reported from upstream to downstream (Supplementary Table 2). The $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ end-member values of suspended POM originating from the TWC in the eastern Jeju Island in summer were obtained from the values reported from subsurface and bottom layers. The $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ end-member values of suspended POM from the riverine freshwater discharge from the Korean peninsula were the values reported for the Seomjin River and Tamjin River, which are prominent rivers in the South Sea of Korea. Similar to SIAR, the SIMMR integrates stable isotope values from terminal end-members and consumers, together with the uncertainty of trophic enrichment factors, to generate credible intervals for potential dietary solutions (Parnell et al., 2010, 2013). Default settings for the number of iterations, burn-in, and Markov chain Monte Carlo chains were employed by SIMMR to explore possible solutions, excluding those that were not probabilistically consistent with the

data. The accuracy of the Gellman–Rubin statistic for curve analysis was as high as 1.00. In this study, trophic enrichment factors of 0.3‰ ($\pm 0.1\%$) for $\delta^{13}\text{C}$ and 2.2‰ ($\pm 2.1\%$) for $\delta^{15}\text{N}$ were used for the SIMMR calculation (McCutchan et al., 2003). The SIMMR results are presented as the median (50% quartile) and the 5%–95% Bayesian credible intervals of diet proportions.

3 Results

3.1 Isotope values of the primary sources of organic matter

The $\delta^{13}\text{C}$ values of pico- to microsized POM in the surface layer fell within narrow ranges of -23.7% to -25.4% , -21.9% to -24.8% , and -23.2% to -24.1% in both areas A and B in June, August, and September, respectively (Table 2). Their $\delta^{15}\text{N}$ values fell within nearly identical ranges of 4.9‰ to 7.3‰, 5.1‰ to 7.2‰, and 4.9‰ to 8.0‰ in both areas A and B at the three time points, respectively. The radiolarians were highly abundant in area B in June and had high $\delta^{13}\text{C}$ values, with a mean of -20.0% .

3.2 Isotope values of neustonic copepods

The $\delta^{13}\text{C}$ values of the dominant neustonic copepods were distinct from those of all size fractions of POM (Table 2). In area A, the mean $\delta^{13}\text{C}$ values of copepod were consistent on each sampling occasion: -21.3% to -20.8% in June, -17.9% to -16.5% in August, and -20.1% to -19.0% in September. The values were highly elevated in August compared with the values recorded in June and September. The mean $\delta^{13}\text{C}$ values of copepod were highly uniform in area B during the sampling period: -21.3% to -20.0% in June, -20.5% to -19.6% in August, and -21.0% to -19.4% in September.

The mean $\delta^{15}\text{N}$ values of the dominant neustonic copepods in area A varied from 5.1‰ to 7.9‰ in June, 8.2‰ to 9.4‰ in August, and 6.0‰ to 7.7‰ in September. The values were highly elevated in August compared with the remaining two time points. The mean $\delta^{15}\text{N}$ values of copepod in area B were also highly variable on each occasion: 2.7‰ to 7.9‰ in June, 5.8‰ to 8.0‰ in August, and 6.4‰ to 8.2‰ in September.

A multivariate two-way PERMANOVA revealed a strong interaction between area and month ($F_{2, 64} = 17.36$, $P < 0.001$) in terms of the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of the neustonic copepod community (Table 3). In turn, a univariate two-way PERMANOVA also denoted a strong interaction of area \times month for both $\delta^{13}\text{C}$ ($F_{2, 64} = 46.95$, $P < 0.001$) and $\delta^{15}\text{N}$ ($F_{2, 64} = 7.460$, $P < 0.001$) values. Pairwise comparisons within area showed significant differences in the $\delta^{13}\text{C}$ ($P < 0.001$) and $\delta^{15}\text{N}$ ($P < 0.001$, except for June vs. September) values between months in area A, but not in area B ($P > 0.054$ for all cases). Further pairwise comparisons within month revealed significant differences in the $\delta^{13}\text{C}$ ($P < 0.004$) and $\delta^{15}\text{N}$ ($P < 0.005$) values between areas in August and September, but not in June ($P = 0.202$ and 0.317 for $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$, respectively).

TABLE 2 Values of $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ of copepods and particulate organic matter according to area (A, low salinity; and B, high salinity) of the northeastern East China Sea.

Month	Area	Producers and consumers	$\delta^{15}\text{N}$		$\delta^{13}\text{C}$		Code
			Mean	SD	Mean	SD	
June	A	mPOM	5.4	0.5	-24.7	0.1	m
		nPOM	5.1	0.3	-24.5	0.1	n
		pPOM	4.9	0.4	-24.2	0.1	p
		<i>Calanus sinicus</i>	7.9	0.2	-20.8	0.1	CS
		<i>Labidocera rotunda</i>	5.1	0.1	-20.8	0.2	LR
		<i>Pontellopsis yamadae</i>	7.4	0.1	-21.3	0.3	PY
	B	mPOM	4.7	0.4	-25.1	0.3	m
		nPOM	5.7	0.4	-25.4	0.2	n
		pPOM	7.3	0.4	-23.7	0.1	p
		Radiolarian	6.3	0.2	-20.0	0.3	r
		<i>Paracalanus parvus</i> s. l.	2.7	0.2	-21.3	0.2	PP
		<i>L. rotunda</i>	7.9	0.1	-20.0	0.1	LR
August	A	mPOM	7.2	0.2	-23.4	0.4	m
		nPOM	6.0	0.2	-23.5	0.0	n
		pPOM	5.9	0.3	-23.8	0.5	p
		<i>Acartia pacifica</i>	9.4	0.0	-17.7	0.0	AP
		<i>Canthocalanus pauper</i>	9.2	0.2	-17.6	0.2	CP
		<i>L. rotunda</i>	8.7	0.2	-17.9	0.1	LR
		<i>Centropages furcatus</i>	9.3	0.1	-17.6	0.2	CF
		<i>Temora discaudata</i>	8.2	0.0	-17.1	0.1	TD
		<i>T. turbinata</i>	8.5	0.2	-16.5	0.6	TT
	B	mPOM	6.0	0.4	-21.9	0.4	m
		nPOM	5.1	0.1	-24.8	0.5	n
		pPOM	6.4	0.1	-22.3	0.3	p
		<i>A. pacifica</i>	8.0	0.1	-19.6	0.2	AP
		<i>L. acuta</i>	7.4	0.3	-20.3	0.1	LA
		<i>C. furcatus</i>	7.4	0.2	-20.0	0.2	CF
		<i>T. discaudata</i>	5.8	0.0	-19.8	0.2	TD
		<i>C. pauper</i>	7.7	0.1	-20.5	0.1	CP
		September	A	mPOM	5.1	0.1	-23.5
nPOM	8.0			0.2	-24.0	0.0	n
pPOM	7.1			0.2	-23.8	0.2	p
<i>C. pauper</i>	6.6			0.2	-20.1	0.1	CP
<i>C. furcatus</i>	6.5			0.1	-19.9	0.2	CF
<i>L. acuta</i>	7.7			0.2	-20.1	0.1	LA
<i>Subeucalanus crassus</i>	7.2			0.2	-19.0	0.1	SC
<i>T. turbinata</i>	6.0			0.2	-19.8	0.1	TT

(Continued)

TABLE 2 Continued

Month	Area	Producers and consumers	$\delta^{15}\text{N}$		$\delta^{13}\text{C}$		Code
			Mean	SD	Mean	SD	
		<i>T. discaudata</i>	6.3	0.0	-19.4	0.2	TD
	B	mPOM	7.2	0.6	-24.1	0.1	m
		nPOM	5.6	0.2	-23.7	0.1	n
		pPOM	4.9	0.1	-23.2	0.0	p
		<i>Undinula vulgaris</i>	7.7	0.0	-20.7	0.3	UV
		<i>L. acuta</i>	8.2	0.2	-20.6	0.2	LA
		<i>C. pauper</i>	7.6	0.0	-21.0	0.1	CP
		<i>T. discaudata</i>	6.4	0.0	-19.8	0.1	TD
		<i>S. crassus</i>	7.4	0.1	-19.4	0.2	SC
		<i>C. furcatus</i>	6.8	0.1	-20.3	0.1	CF

Data represent mean \pm 1 SD.

3.3 Identification of trophic groups

A cluster analysis based on the average $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values classified neustonic copepods into five distinct groups (designated as groups I, II, III, IV, and V) throughout the sampling occasions (Supplementary Figure 2). The copepod assemblages were separated into two groups in each area, except for area A in August (Figure 2). In June, group I consisted of carnivorous species (*L. rotunda* and *P. yamadae*) and a particle feeder (*C. sinicus*), whereas groups IV and V included a carnivorous species (*L. rotunda*) and a particle feeder (*P. parvus* s. l.) in areas A and B, respectively. In August, group I contained a carnivorous species (*L. acuta*) and particle feeders (*C. pauper*, *A. pacifica*, and *C. furcatus*), and group II consisted of a particle feeder (*T. discaudata*) in area B. In turn, group III comprised species of groups I and II and another carnivorous species (*L. rotunda*) and particle feeder (*T. turbinata*) in area A, with no segregation. In September, group I included a carnivorous species (*L. acuta*) and particle feeders (*C. pauper* and *S. crassus*) in both areas A and B, and particle feeders (*C. furcatus* and *U. vulgaris*) in area B. Group II consisted of a particle feeder (*T. discaudata*) in both areas A and B, and other particle feeders (*C. furcatus* and *T. turbinata*) in area A, with no carnivorous species identified in this group.

Significant differences were observed between trophic groups (one-way MANOVA, $F_{4, 65} = 56.77$, $P < 0.001$) in terms of the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values (Table 4). A subsequent univariate ANOVA revealed significant differences for both $\delta^{13}\text{C}$ ($F_{4, 65} = 77.29$, $P < 0.001$) and $\delta^{15}\text{N}$ values ($F_{4, 65} = 160.4$, $P < 0.001$). Tukey's HSD test of the $\delta^{13}\text{C}$ values between trophic groups revealed significant differences ($P < 0.001$), except for groups I vs. IV and IV vs. V). In fact, group III displayed considerably higher $\delta^{13}\text{C}$ values than the remaining groups ($P < 0.05$ for all cases). In contrast, the $\delta^{13}\text{C}$ values of groups II, IV, and V fell within the range of group I (-21.3‰ to -19.0‰). The $\delta^{15}\text{N}$ values of copepod assemblages were clearly differentiated between trophic

groups ($P < 0.05$ for all cases), except for group IV vs. V. Conversely, Tukey's HSD test of each trophic group revealed slight differences in both $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values, with the differences between areas and months being very small, within 1‰.

3.4 Community-wide trophic niche

The NR, CR, and CD values were lower in June and September (0.71‰ and 0.72‰, 0.43‰ and 0.70‰, and 0.42‰ and 0.51‰, respectively) than in August (1.60‰, 2.58‰, and 1.52‰, respectively) (Table 5). The Layman metrics further showed that the NR, CR, and CD values between two spatially different communities (i.e., low- vs. high-salinity areas in each month) were much larger in August than they were in June and September. The SEAc was much wider in area B in June than it was in the remaining months and in area A in all three months (Figure 3). The copepod communities in June and September exhibited a considerable overlap (17.0% and 13.6%, respectively) of trophic niches between areas A and B. In contrast, no niche overlap was observed between areas A and B in August.

3.5 Mixing-model quantification of source contribution

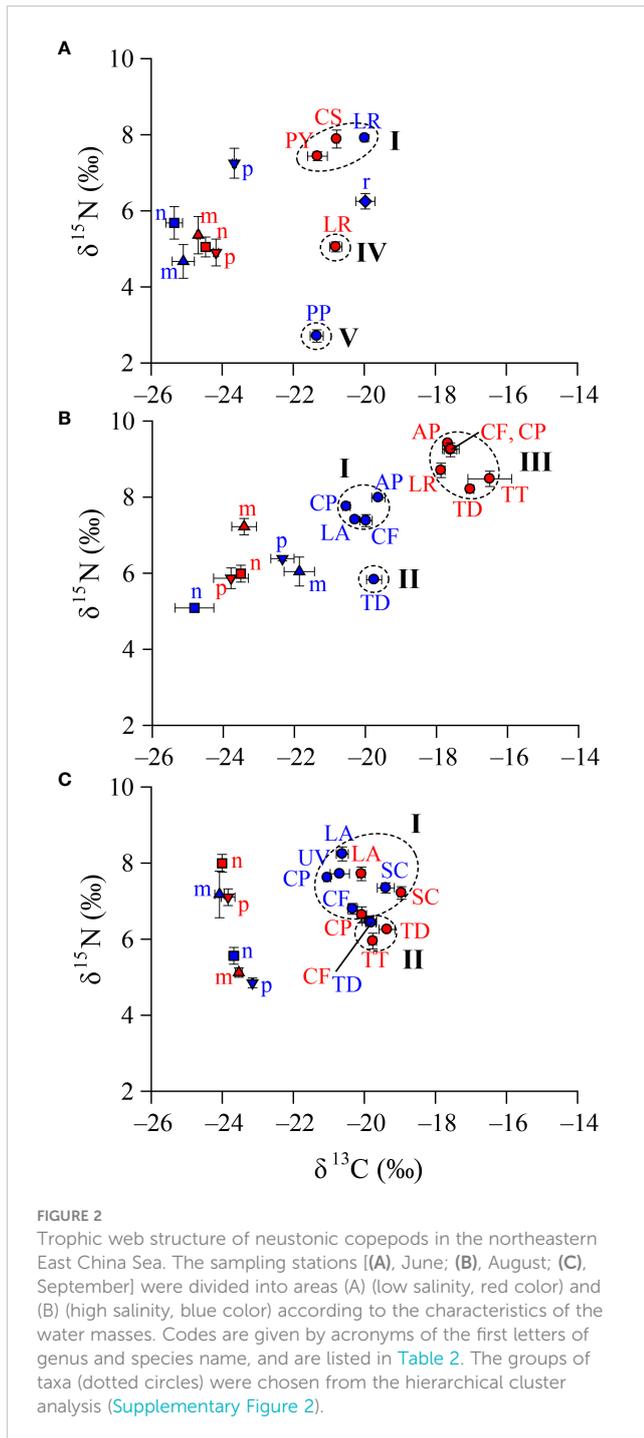
The SIMMR model calculation showed that almost all of the dietary nutrition of the neustonic communities in June, August, and September originated from the subsurface and bottom POM of the TWC rather than from the riverine discharge (Yangtze River and Korean rivers; Figures 4A, C). Furthermore, the nutritional source of the copepod community constituting the trophic group III (i.e., low-salinity area in August) was also calculated to be exclusively dependent upon the subsurface and bottom POM originating from the TWC (mean, 98.3%) than from the YRW (mean, 1.7%) (Figure 4B).

TABLE 3 Results of the two-way PERMANOVA (multivariate and univariate) and the pairwise comparison of the differences in the stable isotope ($\delta^{13}\text{C}$ and $\delta^{15}\text{N}$) values of the copepod community.

Multivariate							
$\delta^{13}\text{C}$ and $\delta^{15}\text{N}$							
Factors	df	SS	F	P			
Area	1	0.015	21.68	< 0.001			
Month	2	0.061	44.05	< 0.001			
Interaction	2	0.024	17.36	< 0.001			
Residual	64	0.045	—	—			
Univariate							
$\delta^{13}\text{C}$				$\delta^{15}\text{N}$			
Factors	df	SS	F	P	SS	F	P
Area	1	0.016	95.98	< 0.001	0.009	1.259	0.265
Month	2	0.034	100.8	< 0.001	0.208	14.34	< 0.001
Interaction	2	0.016	46.95	< 0.001	0.108	7.460	< 0.001
Residual	64	0.011	—	—	0.465	—	—
Pairwise "Area × Month" for pairs of levels of factor "Month" Within "Area"							
$\delta^{13}\text{C}$				$\delta^{15}\text{N}$			
Factors	df	SS	F	P	SS	F	P
Area A							
June vs. August	1, 19	0.003	229.3	< 0.001	0.082	27.85	< 0.001
June vs. September	1, 23	0.003	46.92	< 0.001	0.109	0.578	0.445
August vs. September	1, 30	0.005	175.2	< 0.001	0.045	108.4	< 0.001
Area B							
June vs. August	1, 15	0.003	3.734	0.074	0.323	4.402	0.088
June vs. September	1, 17	0.004	0.233	0.646	0.313	5.762	0.054
August vs. September	1, 24	0.004	3.487	0.063	0.058	0.126	0.739
Pairwise "Area × Month" for pairs of levels of factor "Area" Within "Month"							
$\delta^{13}\text{C}$				$\delta^{15}\text{N}$			
Factors	df	SS	F	P	SS	F	P
June							
Area A vs. Area B	1, 10	0.002	1.729	0.202	0.362	0.997	0.317
August							
Area A vs. Area B	1, 24	0.004	185.3	< 0.001	0.043	35.79	< 0.001
September							
Area A vs. Area B	1, 30	0.005	14.66	< 0.004	0.06	10.36	< 0.005

Bolds indicate significance at $P < 0.05$.

The tested factors included "area" (low salinity, A vs. high salinity, B) and "month" (June, August, and September), and their interaction.



4 Discussion

Despite the expected increase in the contribution of the POM originating from the YRDW during the summer, there were no observable conspicuous changes in the trophic niche (including trophic sources and positions) of neustonic copepods in the nECS. Our findings suggest that the primary sources of organic matter sustaining the neustonic copepod community were derived from POM produced within the TWC itself, rather than from river discharge. Although we observed slight temporal variations in trophic niche and Layman metrics, these variations were

attributed to an increased contribution of phytoplankton locally produced in situ, rather than POM originating from rivers. Consequently, we sought to assess the relative importance of two factors: (1) the exclusive contribution of local POM and (2) the minimal contribution of riverine POM to support of the production of neustonic copepods in the nECS, as evidenced by our SIMMR mixing-model estimations.

4.1 Importance of local production for the community trophic niche

The $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of POM in the surface water varied according to sampling month and area and depended on oceanographic characteristics and biogeochemical processes (Michener and Kaufman, 2007). The $\delta^{13}\text{C}$ values of marine POM typically vary from -18‰ to -23‰ , whereas terrestrial POM exhibits a range of -25‰ to -28‰ (Meyers, 1997). In this study, an average range of -21.9‰ to -25.4‰ of POM of various sizes indicated a mixture of POM from terrestrial and marine origins (Table 2). The $\delta^{13}\text{C}$ values of the POM derived from the Yangtze River estuary and riverine freshwater discharge from the Korean peninsula exhibited ranges of -23.7‰ to -26.4‰ and -24.1‰ to -26.9‰ , respectively (Tan et al., 1991; Gal et al., 2012; Kang et al., 2020). Notably, it has been observed that the $\delta^{13}\text{C}$ values tend to increase gradually with the increase in salinity from coastal regions to the open ocean (Tan et al., 1991; Wu et al., 2003; Gao et al., 2014). Consequently, the $\delta^{13}\text{C}$ values observed across all POM sizes suggest a discernible influence of the mixing of riverine and marine sources. The $\delta^{15}\text{N}$ values of POM across all size categories ranged from 4.7‰ to 8.0‰ , falling within a range that was comparable to that of the copepod community (2.7‰ – 9.4‰). Notably, the $\delta^{15}\text{N}$ values recorded in the vicinity of Jeju Island were also elevated, varying from 5.4‰ to 10.3‰ (Min et al., 2020). In coastal areas (low salinity), the $\delta^{15}\text{N}$ values of POM increase due to the influence of wastewater inflowing in through riverine water (McClelland et al., 1997). Conversely, the $\delta^{15}\text{N}$ values of POM in the open ocean (high salinity) are depleted due to the N_2 fixation prevails by diazotrophs and *Trichodesmium* (Montoya et al., 2002; Wu et al., 2018). Nevertheless, POM of area affected by the TWC showed high $\delta^{15}\text{N}$ values. Along the coast of Jeju Island, a large number of fish farms discharge wastewater with high nutrient concentrations, which contain high-protein fish tissue and feed, into the sea surrounding the island (Samanta et al., 2019; Lee et al., 2020). In addition, due to the recent increase in pig livestock, nitrate discharged from the farms flows into the under-ground water, and groundwater flows into the coastal seawater (Kim et al., 2021). For this reason, the high-concentration nutrients discharged from the Jeju Island may result in elevated $\delta^{15}\text{N}$ values of POM as they are transported and diluted through the surrounding TWC. The surface layer, although potentially rich in food resources, is expected to host a higher concentration of predators, both visual and nonvisual (Leising et al., 2005). Based on the feeding activities of predators, such as chaetognaths and planktivorous fish, including those in the larval stages, we inferred that POM exhibited elevated $\delta^{15}\text{N}$ values because of its higher organic matter content, including zooplankton detritus and proteins, compared with phytoplankton.

TABLE 4 Results of the one-way MANOVA, ANOVA, and Tukey's HSD test results, for testing the differences in the stable isotope ($\delta^{13}\text{C}$ and $\delta^{15}\text{N}$) values among the trophic groups of the northeastern East China Sea.

MANOVA							
$\delta^{13}\text{C}$ and $\delta^{15}\text{N}$							
Factors	df	Pillai's Trace	F	P			
Trophic group	4	1.555	56.77	< 0.001			
Residual	65	0.998	—	—			
ANOVA							
		$\delta^{13}\text{C}$			$\delta^{15}\text{N}$		
Factors	df	SS	F	P	SS	F	P
Trophic group	4	91.85	77.29	< 0.001	112.9	160.4	< 0.001
Residual	65	9827	—	—	918.9	—	—
Tukey's HSD test							
		$\delta^{13}\text{C}$			$\delta^{15}\text{N}$		
(I) Factors	(J) Factors	Mean Difference (I-J)	Std. Error	P	Mean Difference (I-J)	Std. Error	P
Group I	Group II	0.540*	0.175	< 0.024	1.349*	0.135	< 0.001
	Group III	2.796*	0.170	< 0.001	-1.393*	0.131	< 0.001
	Group IV	-0.583	0.327	0.392	2.491*	0.252	< 0.001
	Group V	-1.188*	0.395	< 0.030	4.835*	0.304	< 0.001
Group II	Group III	2.256*	0.210	< 0.001	-2.742*	0.162	< 0.001
	Group IV	-1.123*	0.349	< 0.017	1.142*	0.269	< 0.001
	Group V	-1.728*	0.414	< 0.001	3.486*	0.319	< 0.001
Group III	Group IV	-3.379*	0.347	< 0.001	3.884*	0.267	< 0.001
	Group V	-3.984*	0.412	< 0.001	6.228*	0.317	< 0.001
Group IV	Group V	-0.605	0.498	0.742	2.344*	0.383	< 0.001

The mean differences significant at the 0.05 level are marked with an asterisk. Bolds represent significance at $P < 0.05$.

The $\delta^{13}\text{C}$ values measured in neustonic copepods and the surface POM displayed a distinct discrepancy (Figure 2), indicating the complexity of the relationships among neustonic copepod communities. Notably, there was a clear distinction in neustonic copepods between areas A (low-salinity level) and B

(high-salinity level) (Table 3). The influx of monsoon-induced river discharge has been identified as a factor that reduces the density of marine copepods (Chen et al., 2023). In area A (low-salinity level), which is affected by river discharge, the neustonic copepod communities refrained from feeding on the surface POM because

TABLE 5 Comparisons of the isotopic niche areas of each area (A, low salinity; and B, high salinity) according to month (June, August, and September).

Month	NR	CR	CD	Area	TA	SEAc (% \cdot^2)	Nestedness (%)
June	0.711	0.427	0.415	A	1.659	1.757	17.0
				B	1.336	1.636	
August	1.596	2.581	1.517	A	1.329	0.698	0.0
				B	1.572	1.007	
September	0.723	0.704	0.505	A	1.890	0.907	13.6
				B	1.956	1.066	

Results of the metrics of the $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values pertaining to the month provided information on trophic diversity ($\delta^{15}\text{N}$ range, NR; $\delta^{13}\text{C}$ range, CR; mean distance to the centroid, CD). Each area estimated the total area (TA), the corrected standard ellipse area (SEAc), and the overlap in SEAc (nestedness).

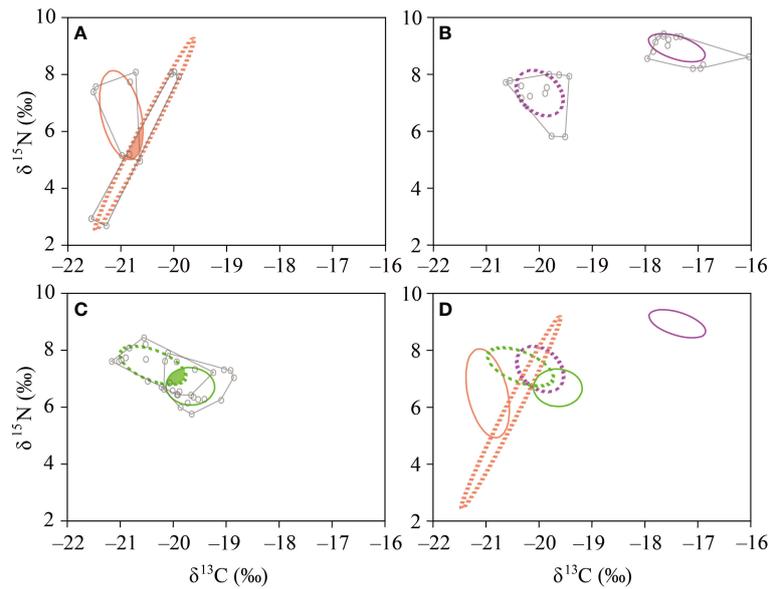


FIGURE 3

Isotopic niches of the neustonic copepod community. The lines and colors are as follows: areas A (low salinity, solid line), B (high salinity, dashed line), and total area (TA) of convex hull (gray line) of each area. (A) June (orange), (B) August (purple), (C) September (green), and (D) the entire study areas in the northeastern East China Sea.

of avoidance behaviors triggered by an osmotic and salinity acclimation stress (Saiz et al., 2022; Weissenberg et al., 2022). Conversely, in area B (high-salinity level), where the impact of river discharge is minimal, the neustonic copepod communities descended to avoid visual predators, such as chaetognaths and planktivorous fish, including those in the larval stages. Experimental evidence from a predator–prey system involving *Acartia hudsonica* revealed that, in the absence of visual predators, there was no vertical migration; however, such behavior was observed after the introduction of predators (Bollens and Frost, 1991). This pattern coincided with the arrival of krill and the subsequent response to predation by *Calanus finmarchicus*, which descended from the surface (Tarling et al., 2002). Considering the influence of predators and freshwater discharge, neustonic copepods are presumed to undergo vertical migration, selectively consuming POM in the subsurface

chlorophyll maximum (SCM) or bottom layers. This selective strategy is interpreted as a means to enhance feeding success by increasing individual survival rates while concurrently reducing mortality rates.

The $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values differed between areas in August and September, but not in June (Table 3). Approximately 60,000 tons of YRDW were discharged from the Yangtze River over the month of May (see “site description”). This discharge formed a tongue-shaped streamer that was propelled by persistent southeast monsoon winds, leading to the intermingling of areas A and B over time (Supplementary Figure 3). Subsequently, YRDW discharges decreased to 45,000 tons in August, whereas the TWC experienced a relative increase. In September, the intrusion of the TWC intensified, creating a broad frontal zone with the coastal waters of the SKCW. This water mass arrangement was also reflected in the trophic niche (Figure 3). The trophic-niche

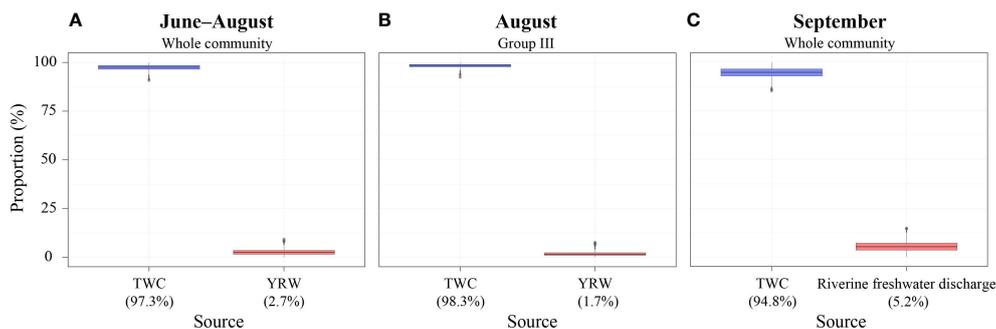


FIGURE 4

Comparison of the dietary compositions of the neustonic copepod community in (A) June–August, (B) August, and (C) September using a dietary mixing model. TWC, Tsushima Warm Current; YRW, Yangtze River Water.

variation in neustonic copepods revealed the presence of distinct feeding strategies in response to the contrasting water masses. Neustonic copepods exhibit trophic flexibility, with a complex trophic structure that is affected by the environment (López-Ibarra et al., 2018; Albuquerque et al., 2021). In June and September, when the contrasting water masses became mixed, the trophic niche of the neustonic copepod community overlapped between the two areas; in contrast, this was not observed in August (Table 5). The trophic-niche overlap resulting from the mixing of contrasting water masses likely signifies an increased prey diversity and potential competition among organisms. Conversely, the clear separation of the water masses observed in August suggests a trophic shift in the neustonic copepod community, indicating a high prey diversity in each area. This underscores the high adaptability of neustonic copepods as flexible feeders, even in areas with diverse water masses, where specific feeding types are better suited to variations in prey availability under local conditions.

4.2 Characterization of trophic groups

The cluster analysis of neustonic copepods based on $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values resulted in the broad categorization of the communities into two groups based on the month of collection (Figure 2). In June, *P. yamadae*, which is a neustonic copepod that is prevalent near the surface during the day and night, fell into group I, whereas *L. rotunda* was present in both groups I and IV. *Pontellopsis yamadae*, which is known for its raptorial feeding habit involving the consumption of copepodids or small copepods (such as *Acartia*, *Euterpina*, and *Paracalanus*) (Ohtsuka and Onbé, 1991), exhibited high $\delta^{15}\text{N}$ values, presumably indicating the *in situ* consumption of large copepods. *Labidocera rotunda*, possessing a mouth structure and feeding habits similar to *P. yamadae*, was considered carnivorous in this study (Ohtsuka and Onbé, 1991). Despite its robust and sharp teeth (Anraku and Omori, 1963), the genus *Labidocera* can feed on phytoplankton. *L. rotunda* in group I was found to consume the copepod nauplii, small copepods (*P. parvus* s. l.), and dominant radiolarians (Ohtsuka and Onbé, 1991), resulting in elevated $\delta^{15}\text{N}$ values (Supplementary Figure 1).

Conversely, group IV was linked to copepods that probably ingested phytoplankton particles with low $\delta^{15}\text{N}$ values, possibly as a response to a copepod avoidance behavior triggered by river discharge. *Calanus sinicus*, which was identified as an indicator species of the Yellow Sea Bottom Cold Water, displayed the highest $\delta^{15}\text{N}$ values (Wang et al., 2003). In coastal areas with nutrient input from river discharge, *C. sinicus* prefers nano- to micro-sized POM (Lee et al., 2012; Chen et al., 2018). Even in the Ulleung Basin, where river discharge is minimal, *C. sinicus* leans toward autotrophic and heterotrophic protozoa in the nano- to microsize range, rather than toward picosized prey (Yang et al., 2009). With the capability to consume substantial quantities of nonphytoplankton prey (Yang et al., 2009; Jang et al., 2010), *C. sinicus* exhibits higher $\delta^{15}\text{N}$ values than carnivorous copepods because of the ingestion of a diverse range of prey with elevated $\delta^{15}\text{N}$ values in the study area. In group V, *P. parvus* s. l., which has been characterized as a particle feeder, is believed to consume small-sized phytoplankton particles using

short, sharp-edged teeth (Al-Hanoun and Mayya, 2020). During feeding experiments, *Paracalanus* species exhibited an increased rate of feeding ciliates and dinoflagellates within the size range of 15–20 μm , whereas the feeding rate on particles exceeding 40 μm was decreased (Suzuki et al., 1999). The observed low trophic position of *P. parvus* s. l. likely reflects its feeding preference for picoplankton, which is characterized by low $\delta^{15}\text{N}$ values.

In August, the influence of contrasting water masses led to the clear differentiation of groups I, II, and III. In group I, *L. acuta* exhibited a feeding habit similar to that observed for *L. rotunda* in June. Among the particle feeders, it was reported that *C. furcatus* can feed on microzooplankton, including ciliates (Jang et al., 2010). The assessment of the food-source contribution of *Centropages* species confirmed their association with micro-sized POM (Chen et al., 2018). *Acartia pacifica*, which is a member of group I, selectively feeds on nanosized plankton (>20 μm) (Lee et al., 2012). *Acartia pacifica* can rapidly consume nanoflagellates more efficiently than large prey and shows a preference for tintinnid ciliates with shells among ciliates (Vargas and González, 2004; Gifford et al., 2007). Also in this group, *C. pauper* is known to feed on a variety of particles, ranging from phytoplankton to metazoan individuals (Hu et al., 2014), although the specific prey-size preferences have not been established well. Nevertheless, similar to that observed for *C. sinicus*, it is believed that they exhibit a selective feeding behavior on plankton larger than 20 μm (Jang et al., 2010; Lee et al., 2012). In group II, *Temora discaudata* displayed the ability to consume diatoms and dinoflagellate particles in the range of 3–106 μm , with a particular preference for food with a size between the pico- and nanorange (Turner, 1984a). The presence of groups I and II seemed to indicate the mitigation of interspecific competition by consuming prey with the sizes preferred by each taxon.

Surprisingly, group III, which was expected to exhibit decreased $\delta^{13}\text{C}$ values because of the influence of river discharge, displayed enriched $\delta^{13}\text{C}$ values compared with groups I and II. The copepods in group III exhibited high $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values, potentially indicating the consumption of ^{13}C -enriched phytoplankton, such as diatoms (Fry and Wainright, 1991), which can have relatively high $\delta^{15}\text{N}$ values compared with other phytoplankton (Montoya and McCarthy, 1995). The significant increase in the $\delta^{15}\text{N}$ values of *T. discaudata* in group II suggested a substantial consumption of diatoms, even though they are herbivorous prey. In particular, the $\delta^{15}\text{N}$ values of *T. discaudata* in group II were significantly increased. A fatty acid and gut-content analysis revealed that species of the genus *Temora* are capable of feeding on small flagellates and copepod nauplii (Gentsch et al., 2009). This suggests that they act as particle feeders, consuming not only picosized phytoplankton but also both autotrophic and heterotrophic prey.

In September, the neustonic copepod species became intertwined because of the mixing of contrasting water masses. *Undinula vulgaris*, which was part of group I, exhibited an indiscriminate particle-feeding habit, as it consumed phytoplankton of various sizes and shapes (Turner, 1986). Specifically, *U. vulgaris* fed predominantly on nanosized plankton, dinoflagellates, and diatoms (Kleppel et al., 1996). *Subeucalanus crassus* displayed the ability to consume prey of diverse sizes,

encompassing living and dead phytoplankton (diatoms and dinoflagellates), sediment, fecal pellets, and crustaceans (Turner, 1984b). Notably, this species exhibited a preference for nanosized plankton over other phytoplankton (Kleppel et al., 1996). Unlike most species that capture live prey, Eucalanidae species, including *S. crassus*, can also detect and consume dead prey (Turner, 1984b; Paffenhöfer and Van Sant, 1985), which is a feeding behavior that is considered a survival strategy to minimize the potential competition between species within the water column. *Canthocalanus pauper* and *C. furcatus* were allocated to groups I and II, respectively. Although the constituent species of groups I and II were intermingled, areas A and B appeared to be distinct. As mentioned previously, *C. pauper* and *C. furcatus*, which belong to group I in area B (with high salinity), primarily consumed nano- and micro-sized plankton, exhibiting high $\delta^{15}\text{N}$ values. These species were presumed to have consumed plankton with lowered $\delta^{15}\text{N}$ values, possibly because of the adverse effects of river discharge in group II located in area A (with a lower salinity).

Our SIMMR mixing-model calculation supports that neustonic copepod community rejects to feeding surface POM and reflecting vertical migration to subsurface and bottom layers caused by the possible influences of factors such as river discharge or the presence of visual and nonvisual predators. The SCM layer within the range of vertical migration contains phytoplankton that can be consumed by the neustonic copepods, but also contains toxic diatoms, dinoflagellates, or other harmful algae (Miralto et al., 1999; Pierson et al., 2005). So they can migrate to deep layer to find nutrient-replete and non-toxic preys (Im and Suh, 2019). Therefore, we suggest that further research is needed on feeding in the subsurface and bottom layers of the neustonic copepods. In the future, it is necessary to approach the prey and feeding ecology of neustonic copepod species through research such as culture experiments using video cameras to determine the sinking prey feeding behavior of copepods, DNA sequence analysis of gut-contents, and fatty acid analysis (Cleary et al., 2017; Van der Jagt et al., 2020; Cui et al., 2021).

5 Conclusion

We observed variations in the food-source composition of neustonic copepods according to the trophic group, rather than the salinity levels of the water masses. Although a distinct trophic-niche shift was observed between high- and low-salinity environments in August, the contribution of riverine POM to the diets of neustonic copepods was minimal, even in the low-salinity area. The neustonic copepod community exhibited no association with surface POM. This suggests the presence of intricate feeding relationships among trophic groups, reflecting vertical migration caused by the possible influences of factors such as river discharge or the presence of visual and nonvisual predators. Furthermore, our examination of trophic groups within the neustonic copepod community revealed differences in individual preferred prey sizes and types. These copepods, which were primarily characterized as opportunistic particle feeders, exhibited a capacity for navigating both inter- and intraspecific competition within water masses through selective feeding. Our SIMMR

estimation, aimed at identifying the source water mass that affected the food consumed by neustonic copepods, highlighted a predominant reliance on dietary items derived from the TWC. This aligns with the existing literature, supporting the notion of a minimal influence from river-derived material on ocean phytoplankton productivity (Mutia et al., 2021). Consequently, it can be inferred that the marine neustonic copepod community is not significantly dependent on freshwater-originated food. In future studies, a comprehensive examination incorporating other zooplankton taxonomic groups, together with neustonic copepods, would provide a more detailed understanding of the entire planktonic food web.

Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding authors.

Ethics statement

The manuscript presents research on animals that do not require ethical approval for their study.

Author contributions

JHC: Writing – review & editing, Writing – original draft, Visualization, Validation, Methodology, Investigation, Formal analysis, Data curation, Conceptualization. YK: Writing – review & editing, Methodology, Conceptualization. C-KK: Writing – review & editing, Validation, Conceptualization. HYS: Writing – review & editing, Funding acquisition.

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Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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Supplementary material

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fmars.2024.1378025/full#supplementary-material>

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SUPPLEMENTARY FIGURE 1

The dominant radiolarians in the area B in June. The radiolarians were observed in gut of *Labidocera rotunda*.

SUPPLEMENTARY FIGURE 2

Hierarchical clustering of the average $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values of neustonic copepods in June, August, and September for study areas A and B using the Bray–Curtis similarity (average grouping method).

SUPPLEMENTARY FIGURE 3

(A) Ocean prediction data based on marine numerical models provided by the ocean and fisheries research institute in the Jeju special self-governed province showing salinity distribution. (B) Water temperature–salinity diagram to determine the point of introduction of the Yangtze River Diluted Water into the areas A and B in northeastern East China Sea. The dotted line indicates the criteria for water with abnormally low salinity ($>22\text{ }^{\circ}\text{C}$, 25–30) (Hyun and Pang, 1998). During the study period, approximately 60,000 tons of the YRDW were discharged continuously from the Yangtze River a month before the survey (end of May) was conducted (see 'site description'). Furthermore, the southeast monsoon winds blow to form a tongue-shaped 'streamer'. The 'streamer' phenomenon was observed in area A in June (Figure 1). Accordingly, water temperature–salinity diagram was drawn to identified how much YRDW had flowed into the study area in June, and abnormally low-salinity water inflow was observed according to water temperature ($>22\text{ }^{\circ}\text{C}$) and salinity (25–30). Because of the continuous inflow of YRDW and active surface mixing caused by the persistent southeast monsoon winds, it was judged that there was little difference between the $\delta^{13}\text{C}$ values of POM in areas A and B in June.

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