



Trophic Dynamics of Zooplankton Before and After Polar Night in the Kongsfjorden (Svalbard): Evidence of Trophic Position Estimated by $\delta^{15}\text{N}$ Analysis of Amino Acids

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In polar ecology, zooplankton diets and survival rates vary according to the seasonality of solar radiation and oceanographic conditions. Each zooplankton species has evolved feeding strategies to survive in the diet-limited conditions of the “polar night.” Many zooplankton studies have reported seasonal adaptations in feeding activity during polar night based on their trophic niches. Nitrogen isotope analysis of amino acids has provided improved accuracy in estimates of trophic position (TP) in various marine species. In this study, field work was conducted in Kongsfjorden before (October 2017) and after polar night (April 2018). As representative zooplankton, an amphipod (*Themisto abyssorum*), euphausiids (*Meganycitiphanes norvegica* and *Thysanoessa* sp.), a chaetognath (*Parasagitta elegans*), and copepods (*Calanus* spp. and *Oithona similis*) were collected. trophic position values of each taxon were estimated using the nitrogen isotope ratio of glutamic acid ($\delta^{15}\text{N}_{\text{Glu}}$) and phenylalanine ($\delta^{15}\text{N}_{\text{Phe}}$). Results showed that TP values of *P. elegans* were relatively constant, averaging 3.2 in both seasons, likely due to continuous feeding activity during polar night. Trophic position values were also constant for *Calanus* spp., ranging 2.5–3.0 in both seasons, due to their ability to utilize stored high-energy wax. In contrast, average TP values for *O. similis*, an omnivorous zooplankton, were 2.9 in October and 2.3 the following April. Trophic position values for *O. similis* before polar night can be attributed to the relatively high availability of algae during longer periods of daylight. We found that TP variation in zooplankton before and after polar night differed according to feeding activities in diet-restricted circumstances.

Keywords: trophic position, nitrogen isotope, amino acid, zooplankton, Kongsfjorden, Svalbard

INTRODUCTION

The Arctic Ocean is an extreme environment, with seasonal cycles of solar radiation. This seasonality creates an extended period of darkness known as the polar night, which typically begins in late October and lasts until March. During polar night, primary production is low due to insufficient irradiation, and the diet available to consumers is limited. In the polar

marine ecosystem, the relationship between diet and survival among zooplankton in diet-limited conditions has been the subject of continuous study (Søreide et al., 2008; Varpe, 2012; Kraft et al., 2013). Zooplankton species have evolved over-wintering strategies that vary by their trophic niche or metabolism. Herbivorous zooplankton, such as some calanoid copepods, reportedly survive periods of food shortages by consuming accumulated storage lipids (Conover, 1988). Chaetognaths, which are carnivorous zooplankton, usually feed on calanoid copepods, surviving on a reduced diet during polar night (Grigor et al., 2014). For omnivorous zooplankton, seasonal feeding strategies can lead to changes in dietary composition, which in turn can alter a species' trophic position (TP).

Feeding strategies of zooplankton in polar and subpolar regions have been reported using feeding experiments (Atkinson, 1996), gut-content analysis (Kruse et al., 2010), and stable isotope analysis of bulk tissue (Søreide et al., 2008; Grigor et al., 2015). Recently, compound-specific isotope analysis of amino acids (CSIA-AAAs) has been used to estimate the TP of consumers (Bowes and Thorp, 2015; McMahon and McCarthy, 2016). Nitrogen isotope ($\delta^{15}\text{N}$) analysis of individual amino acids (AAs) can provide useful information on the trophic structure of the food web by separating the trophic elevation and isotopic baseline of ecosystems. The trophic elevation can be explained as trophic AAs, which show large $\delta^{15}\text{N}$ enrichment (3–8‰), and the isotopic baseline is represented by source AAs that exhibit smaller $\delta^{15}\text{N}$ enrichment (by 0–1‰). The TP can be estimated by the $\delta^{15}\text{N}$ values of glutamic acid and phenylalanine as representative of trophic and source AAs, respectively (Chikaraishi et al., 2009). Because CSIA-AAAs is independent of the baseline of the $\delta^{15}\text{N}$ ratio of the food web and is based on internal tissue measurements of the consumer itself, no basal food source sampling is required to estimate TP (Bowes and Thorp, 2015). In addition, a TP determined by CSIA-AAAs could be accurately estimated even with variation of nitrogen sources at the base of the food web (Hannides et al., 2009).

The TP can supply important information about food webs. Temporal or spatial differentiation of the TP of consumers usually reflects changes in dietary utilization (Vander Zanden et al., 1999; Pauly and Watson, 2005; Branch et al., 2010). Because the polar night results in a food-limited environment, most zooplankton can be expected to either reduce the amount of food they consume (Grigor et al., 2014) or rest until better conditions return. On the other hand, opportunistic zooplankton can easily switch from algae to protists (Iversen and Seuthe, 2011). Their TP would therefore change seasonally (Seuthe et al., 2011).

Kongsfjorden, which is a fjord on the island of Svalbard, is one of the most studied coastal areas in the entire Arctic (Hop et al., 2002; Svendsen et al., 2002). A large number of zooplankton studies have been conducted through the international scientific institutes and laboratories in Ny-Ålesund for several decades (Bämstedt, 1976; Scott et al., 2000; Berge et al., 2015a). In this study, we hypothesized that the TPs of zooplankton species can serve as an indicator of feeding activity before and after polar night. Temporal and spatial variation of representative zooplankton species abundance and their relative contributions in the inner and outer regions of Kongsfjorden and their TP

values were precisely compared through CSIA-AAAs before and after polar night.

MATERIALS AND METHODS

Study Site

Kongsfjorden lies on western Spitsbergen, Svalbard, in the European Arctic. Different environmental conditions in Kongsfjorden accompany seasonal and spatial changes in the zooplankton community (Walkusz et al., 2009) because the oceanographic conditions in Kongsfjorden are strongly influenced by sea-ice dynamics and currents flowing around the Svalbard Archipelago (Svendsen et al., 2002). Warm and saline Atlantic water, relatively cold Arctic water, and freshwater runoff from the glaciers make seasonally variable contributions to marine conditions in Kongsfjorden. According to Walkusz et al. (2009), the inflow of Atlantic water to Kongsfjorden is inhibited and a distinct frontal zone is dominated by winter-cooled water that forms in the spring. In contrast, most of the fjord is filled with Atlantic water inflow during the summer and autumn. Because each water mass contains indigenous zooplankton communities, Kongsfjorden experiences large seasonal and spatial variability in the zooplankton community (Varpe, 2012).

Sample Collection and Processing

Zooplankton samples were collected in Kongsfjorden (Figure 1) during the autumn before polar night (October 18 and 19, 2017) and the spring after polar night (April, 12–14 2018). Nine stations, except for St. 03 and St. 06, were selected from the inner to the outer fjord. Locations and depths of sampling sites are provided in Table 1. Vertical profiles of temperature and salinity were measured with conductivity, temperature, and depth (CTD) profiler. We used an SD204 CTD (SAIV A/S, rented from Marine Laboratory in Ny-Ålesund) in October and Concerto 3 loggers (RBR) in April. Zooplankton samples were collected at each station with a plankton net (mesh size of 200 μm , diameter in 50 cm, and 2.5 m in length) at a vertical velocity of 60 m/min. Detailed depths for zooplankton netting are provided in Table 1. In the harbor, additional zooplankton sampling was conducted by hand using a plankton net with the same dimensions. After transportation to the laboratory, zooplankton samples were sorted by microscope (LEICA DFC495), and dominant species were identified. Their abundances were counted and biomasses weighed. The calanoid copepod *Calanus* was defined as *Calanus* spp., which includes *Calanus finmarchicus*, *Calanus glacialis*, and copepodites of those two species (stage CIV–CV). The chaetognath *Parasagitta elegans* was separated into those with body lengths >20 mm and ≤ 20 mm according to a size-cohort grouping method described by Grigor et al. (2014). The amphipod *Themisto abyssorum* and euphausiid *Meganycitiphanes norvegica* were sorted in October. The euphausiid *Thysanoessa* sp. was sorted in April. After subsampling of the zooplankton, samples were freeze-dried for 24 h, homogenized, and stored at -20°C before further processing and analysis.

Oithona similis samples were sorted after chemical preservation because the species' body size was too small

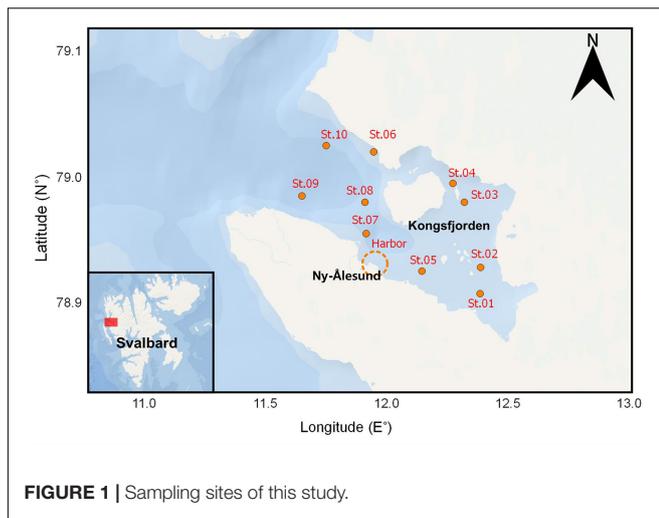


TABLE 1 | Detailed information of sampling stations.

Station	Latitude (°N)	Longitude (°E)	Water depth (m)	Sampling depth for zooplankton (m)
St.01	78.907	12.385	80	0–60
St.02	78.928	12.387	41	0–35
St.03	78.980	12.320	75	
St.04	78.995	12.273	61	0–50
St.05	78.925	12.145	110	0–50
St.06	79.020	11.946	148	
St.07	78.955	11.915	348	0–100
St.08	78.980	11.910	243	0–100
St.09	79.025	11.750	220	0–100
St.10	78.985	11.650	292	0–100
Harbor	78.929	11.936		

Note that St. 03 and St. 06 were not sampled.

(<1 mm) to isolate sufficient numbers of individuals in the available time. We preserved *O. similis* using formalin and Dess-Martin solution in October and April, respectively. Preserved *O. similis* samples were soaked for 24 h in distilled water, and oven-dried at 60°C for 24 h.

Preservation Effect on Samples

Formalin and ethanol fixation on samples reportedly does not affect $\delta^{15}\text{N}$ values of AAs in samples of fish and zooplankton (Bämstedt, 1976; Hannides et al., 2009; Ogawa et al., 2013; Berge et al., 2015a). As an alternative preservative for formalin or ethanol, Dess-Martin preservation was used for morphological and polymerase chain reaction analysis (Yoder et al., 2006). However, we could find no reports on the isotopic effects of AAs under Dess-Martin preservation. To identify isotopic effects on samples preserved in formalin and Dess-Martin solution, we compared the $\delta^{15}\text{N}$ of AAs in frozen (control) and preserved samples using lab-cultured shrimp (*Paratya* spp.). The Dess-Martin solution was prepared by mixing 20% dimethyl sulfoxide (DMSO) and 0.25 M disodium EDTA saturated with NaCl at pH 8.0 as reported by Seutin et al. (1991). Control samples were

frozen at -20°C in a refrigerator and freeze-dried for 24 h. We preserved each *Paratya* spp. specimen in formalin, 70% ethanol, and Dess-Martin solutions for periods of 1 week and 1 month. All preserved samples were soaked for 24 h in distilled water and dried at 60°C in an oven for 24 h.

Amino Acid Nitrogen Isotope Analysis

For nitrogen isotope analysis of AAs, we collected 5 or 6 individuals of *Calanus* spp., 2 or 3 *Parasagitta elegans* individuals, a single individual of the euphausiid, 2 or 3 amphipod individuals, and approximately 200 *O. similis* individuals. Nitrogen isotopes of AAs for all zooplankton samples were analyzed without any replication at each sampling site because of the need for collecting a large number of individuals. However, in this study, a total of 2 (*O. similis*) or 3 (*Calanus* spp. and *P. elegans*) replicates of representative zooplankton species were compared to discuss their trophic dynamics before and after polar night. Samples were hydrolyzed for 20 h with 12 M HCl at 110°C. After removing the hydrophobic contents with 3:2 n-hexane/dichloromethane (v/v), the residual HCl was dried with an N_2 purge at 70°C. In the case of the *O. similis* fixed in Dess-Martin solution, an additional purification procedure was needed due to the interference of residual ions dissolved in the solution. We therefore used cation-exchange chromatography for those samples before derivatization, following a method described by Takano et al. (2010). All samples were derivatized with 1:4 thionyl chloride/2-propanol (v/v) and 1:4 pivaloyl chloride/dichloromethane (v/v) in sequence. Amino acid derivatives were extracted with 3:2 n-hexane/dichloromethane (v/v) and stored at -20°C in a refrigerator before analysis. The nitrogen isotope composition of each AA was analyzed by gas chromatography (HP 6890N, Agilent) connected to a combustion furnace (GC5 interface, Elementar) and isotope ratio mass spectrometry (Isoprime, Elementar). A derivative of a standard mixture of 11 AAs obtained from SHOKO-Science and Indiana University (alanine, glycine, valine, leucine, norleucine, proline, aspartic acid, methionine, glutamic acid, phenylalanine, and hydroxyproline) was used to confirm $\delta^{15}\text{N}$ measurements. During the calibration procedure, $\delta^{15}\text{N}$ values of the AA standards displayed fine standard deviation of less than $\pm 1\text{‰}$. Trophic positions were estimated using an equation proposed by Chikaraishi et al. (2009):

$$TP = [(\delta^{15}\text{N}_{\text{Glu}} - \delta^{15}\text{N}_{\text{Phe}} - 3.4) / 7.6] + 1,$$

where $\delta^{15}\text{N}_{\text{Glu}}$ and $\delta^{15}\text{N}_{\text{Phe}}$ are the measured nitrogen isotope values of glutamic acid and phenylalanine of consumers, respectively. The value 3.4 indicates the isotopic difference between the glutamic acid and phenylalanine in primary producers (β value) and 7.6 is the trophic discrimination factor (TDF) for $\delta^{15}\text{N}$ enrichment of glutamic acid relative to phenylalanine with each trophic transfer.

Statistical Analysis

Hierarchical clustering using Bray–Curtis similarity (Bray and Curtis, 1957) was applied to zooplankton abundance and relative contributions to determine similarity among

sampling stations. Dominance (Simpson, 1949), diversity (Shannon and Weaver, 1949), evenness (Buzas and Gibson, 1969), and richness (Margalef, 1958) were calculated using the zooplankton abundance values. Species diversity values were analyzed by permutation tests to compare spatial differences among sampling stations. All statistical and ecological analysis related to zooplankton abundance and relative contributions were conducted by paleontological statistics (PAST) software (Hammer et al., 2001). We used independent *t*-tests for abundance and ecological parameters between sampling periods. Independent *t*-tests were also used for $\delta^{15}\text{N}_{\text{AAs}}$ and TP values between sampling periods and species body sizes. All independent *t*-tests were performed using IBM SPSS software v23.0.

RESULTS

Hydrography

Vertical profiles of temperature and salinity are displayed in **Figure 2**. In October 2017, a pycnocline was generated by freshwater run-off from glaciers and/or land, with low salinity of surface water (<33.5 psu) across the stations. By April, most of the water column was homogenized, with low water temperatures and a weak pycnocline. St. 01 and St. 02 showed slightly lower temperatures and less salinity than other stations, suggesting that strong winds and cold freshwater influenced the stations adjacent to land.

Zooplankton Community

Integrated abundance and relative contribution of the zooplankton community are shown in **Figure 3**. Zooplankton abundance and ecological parameters are provided in **Table 1** and **Supplementary Table S1**. Copepods were the most dominant taxon, representing more than 95% of the zooplankton community. Total abundance of zooplankton in October was greater than 10 times that of the following April. In the copepod group, the most abundant species was *O. similis*, which was also the most dominant cyclopoid, accounting for $85.21 \pm 6.78\%$ of copepods on average in October and $43.54 \pm 7.51\%$ in April. In the calanoid group, members of the genus *Pseudocalanus* were the most abundant. The *Calanus* copepodite was the most abundant within *Calanus*, but *Calanus* species represented less than 5% all of zooplankton. A Bray–Curtis similarity test of zooplankton composition showed a clear division between October and April (**Supplementary Figure S1**). Analysis of spatial distributions showed that zooplankton composition in October was less clearly separated between the inner (St. 01–05) and outer fjord (St. 07–10). In contrast, a significant separation was evident in zooplankton composition between the inner and middle-to-outer part of the fjord in April. *Calanus* species showed higher average abundance in the inner fjord (34.55 ± 8.19 individuals/m³ in October and 47.14 ± 24.37 individuals/m³ for April) compared with the outer fjord (14.31 ± 4.55 individuals/m³ in October and 11.27 ± 4.21 individuals/m³ for April). However, the relative contribution of the genus *Calanus* was lower in the inner fjord ($1.37 \pm 0.40\%$ for October and

5.33 ± 0.42 for April) than in the outer fjord ($2.34 \pm 1.14\%$ for October and $14.44 \pm 5.04\%$ for April). In both sampling periods, the number of taxon was similar. However, species diversity parameters showed significant differences in both October and April (**Table 2**). Average species diversity, evenness, and richness in April were higher than those in October. Species dominance in April was lower than that in October. More details of the spatial variation of species diversity as revealed by a permutation test was shown in **Supplementary Table S2**. In October, most sampling stations exhibited significantly different species diversity. In contrast, species diversity at each station in April was nearly homogeneous, except for St. 08.

Preservation Effect of Dess-Martin

The $\delta^{15}\text{N}_{\text{AAs}}$ values of *Paratya* spp. are displayed in **Supplementary Tables S3, S4**. As reported by a previous study, formalin fixation did not appear to influence $\delta^{15}\text{N}_{\text{AAs}}$ values, showing less than a $\pm 1\%$ difference from the control (Bämstedt, 1976; Ogawa et al., 2013; Berge et al., 2015a). In our results, a Dess-Martin solution also showed less than a $\pm 1\%$ difference in $\delta^{15}\text{N}_{\text{AAs}}$ from the control, confirming that Dess-Martin preservation had little isotopic effect on biota samples.

AA $\delta^{15}\text{N}$ Values and Trophic Position

$\delta^{15}\text{N}_{\text{AAs}}$ values and the TP of zooplankton are shown in **Table 2**. No significant spatial trend was observed in $\delta^{15}\text{N}_{\text{Phe}}$ values of zooplankton. For *O. similis*, a significant decrease was seen in $\delta^{15}\text{N}_{\text{Phe}}$ values. This suggests the $\delta^{15}\text{N}$ isotopic baseline of *O. similis* changed during polar night, as a result of changing diets. In *Calanus* species, a slight decrease in $\delta^{15}\text{N}_{\text{Phe}}$ values (1.2% on average) between the two seasons was apparent. For *P. elegans*, no significant difference in $\delta^{15}\text{N}$ between size cohorts in glutamic acid ($t_{0.05,4}$, $p = 0.68$) and phenylalanine ($t_{0.05,4}$, $p = 0.42$) in $\delta^{15}\text{N}_{\text{AAs}}$ trend was evident, nor between the two seasons ($\delta^{15}\text{N}_{\text{Glu}}$ $t_{0.05,10}$, $p = 0.15$; $\delta^{15}\text{N}_{\text{Phe}}$ $t_{0.05,10}$, $p = 0.34$).

Among all zooplankton, *T. abyssorum* exhibited the highest TP in our dataset. Trophic position values of three zooplankton collected in both sampling periods are displayed in **Figure 4**. *P. elegans* had a consistent TP temporally ($t_{0.05,10}$, $p = 0.87$) regardless of body size ($t_{0.05,10}$, $p = 0.71$). *Calanus* species showed an average TP increase of 0.2 from October to April but the difference was not statistically significant ($t_{0.05,4}$, $p = 0.14$). *O. similis* had the largest variation in TP, with a decrease of 0.5 from October to April.

DISCUSSION

Arctic zooplankton have evolved a variety of survival strategies, most of which are related to energy supply during polar night (Hagen and Auel, 2001; Berge et al., 2015b). Energy can be managed through storage (as well as diapause), and flexible feeding behavior (Hagen, 1999). The three zooplankton collected in this study, *O. similis*, *P. elegans*, and *Calanus* spp., exhibit species-specific over-wintering strategies. Such strategies allow researchers to track changing diet sources, and precise TP values can be applied to identify changes in feeding activity.

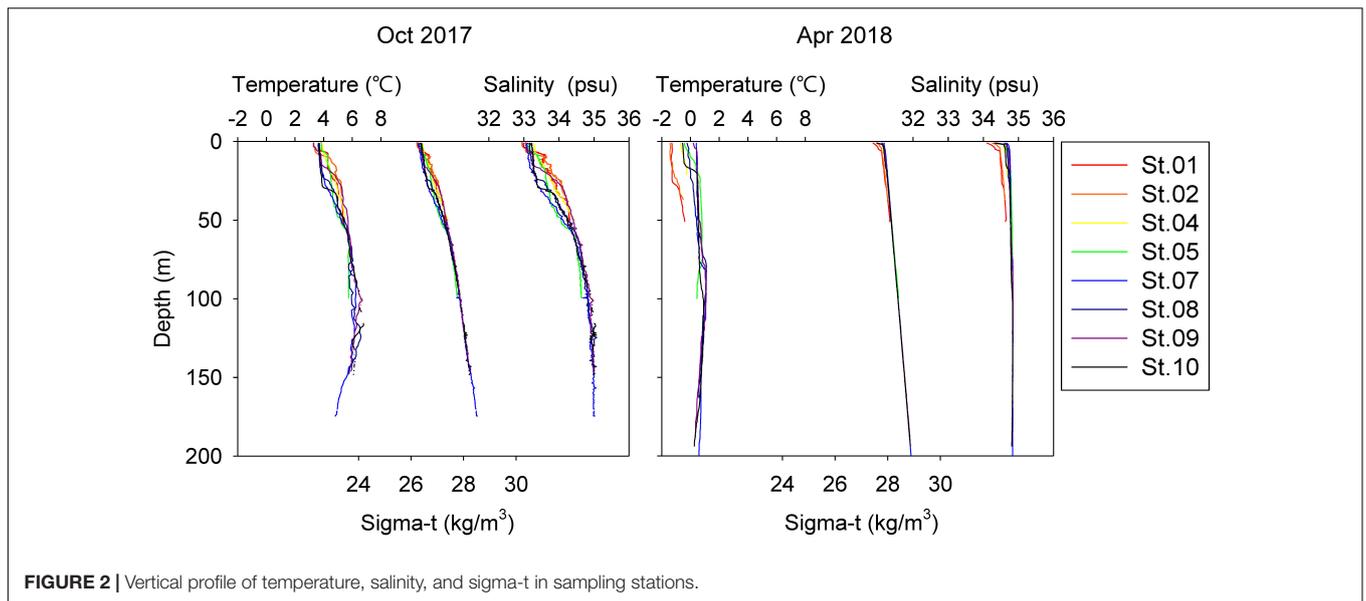


FIGURE 2 | Vertical profile of temperature, salinity, and sigma-t in sampling stations.

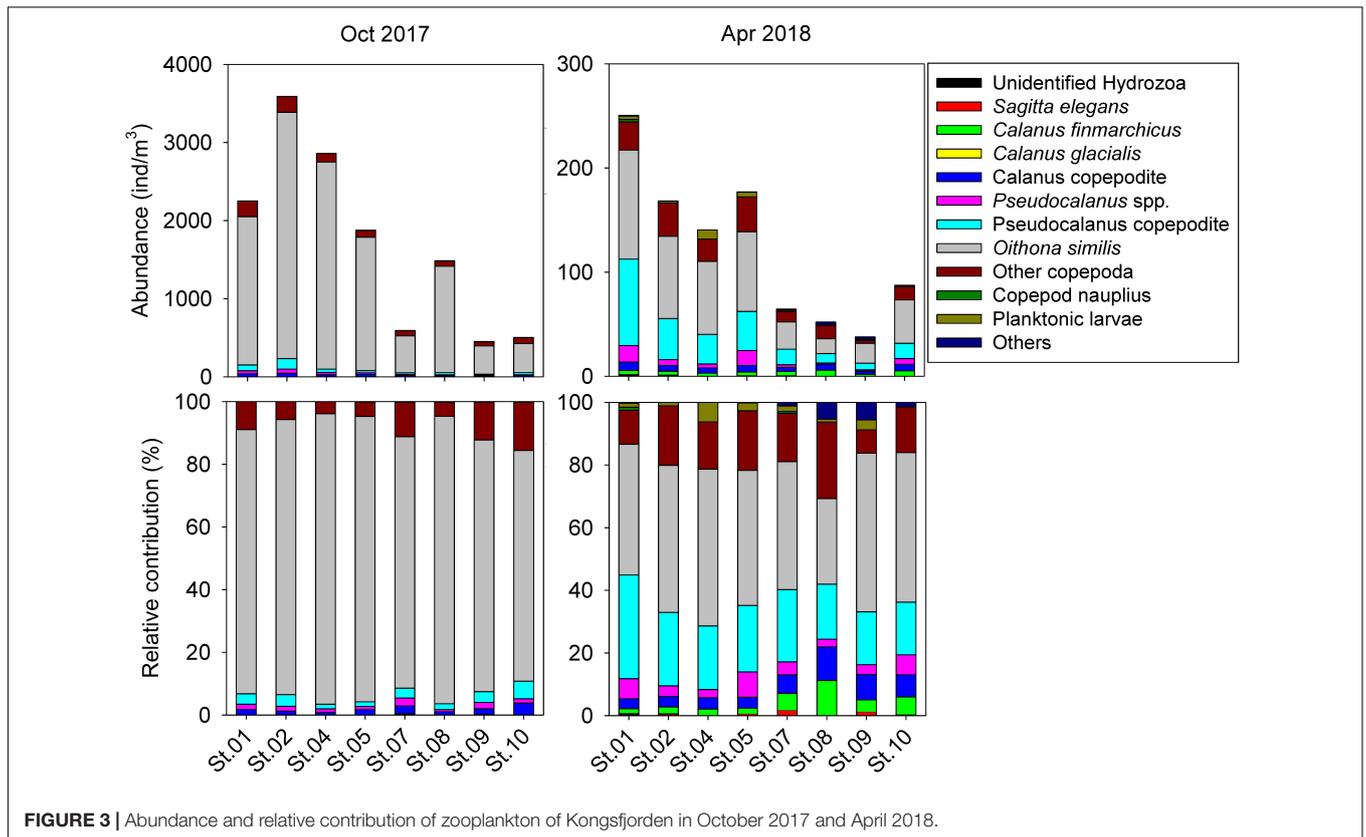


FIGURE 3 | Abundance and relative contribution of zooplankton of Kongsfjorden in October 2017 and April 2018.

Trophic dynamics in zooplankton species may indicate changes in integrated diets before and after polar night. Cases of temporal TP variation before and after polar night are displayed in **Figure 5** and inform the following discussion.

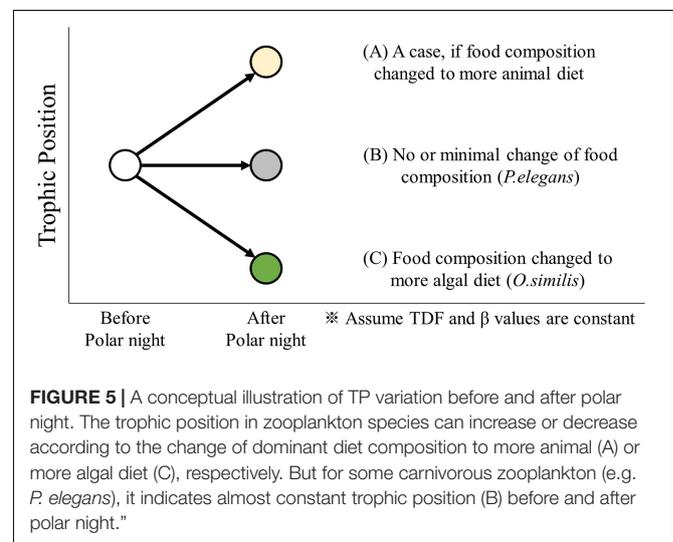
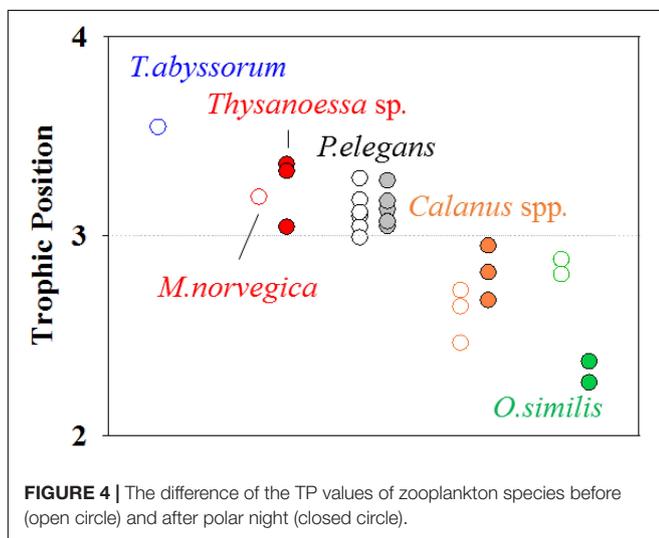
Hydrography between two sampling periods showed different patterns of water masses. In October, bottom water, probably

indicating the intrusion of the North Atlantic Ocean, displayed relatively high salinity and temperature compared with the surface water, which has a low salinity and temperature. In contrast, in April, the entire water column showed a homogeneous temperature and salinity. Our result showed those hydrographic properties were not clearly related to TP values

TABLE 2 | Mean abundance, biomass, and ecological parameters for zooplankton collected in October 2017 and April 2018.

	Oct 2017		Apr 2018		p-value
	Mean	Std	Mean	Std	
Unidentified hydrozoa	nd		0.10	0.28	
<i>Parasagitta elegans</i>	0.85	0.62	0.60	0.47	0.371
<i>Calanus finmarchicus</i>	0.18	0.27	3.67	1.33	<0.001*
<i>Calanus glacialis</i>	nd		nd		
<i>Calanus</i> copepodite	24.43	12.41	5.45	1.51	0.004*
<i>Pseudocalanus</i> spp.	22.75	16.76	6.28	5.73	0.027*
<i>Pseudocalanus</i> copepodite	46.42	40.48	29.20	25.10	0.324
<i>Oithona similis</i>	1497.20	1061.46	53.91	33.06	0.005*
Other copepods	107.55	60.02	19.01	11.19	0.005*
Copepod nauplius	0.40	0.75	0.35	0.84	0.899
Planktonic larvae	1.12	0.41	2.65	2.87	0.182
Other	0.35	0.56	1.01	1.01	
Total abundance (ind/m ³)	1701.26	1166.02	122.22	74.18	0.007*
Total biomass (wet mg/m ³)	49.60	26.43	41.58	18.34	0.492
Dominance (Simpson)	0.74	0.10	0.28	0.04	<0.001*
Diversity (Shannon)	0.58	0.19	1.55	0.13	<0.001*
Evenness (Buzas and Gibson)	0.22	0.04	0.55	0.10	<0.001*
Richness (Margalef)	1.01	0.21	1.73	0.37	<0.001*

Asterisk (*) indicates significant difference ($p < 0.05$) between October 2017 and April 2018.



of zooplankton species but were instead closely influenced to zooplankton community composition. Temporal variation of the zooplankton community in Kongsfjorden, which was caused by seasonal hydrography, was similar to results found in previous studies (Willis et al., 2006; Walkusz et al., 2009). In April, the total zooplankton abundance was much smaller than it was in the preceding October. As the April sampling period took place early in the month, which is considered the pre-bloom season, zooplankton was not yet abundant. With the benefit of two sampling periods, zooplankton communities were clearly separated. In both sampling periods, zooplankton abundance showed a continuous gradient from the inner to outer fjord. The

inner fjord usually has the highest abundance of zooplankton, as reported by Walkusz et al. (2009).

In both sampling periods, *O. similis* was the dominant zooplankton across all sampling stations, and our findings were similar to those of previous studies (Hop et al., 2002; Walkusz et al., 2009). Group *Oithona* was described as a cosmopolitan copepod group, suggesting a wide range of habitat throughout tropical to polar regions (Fransz and Gonzalez, 1995; Castellani et al., 2005). Thus, their population would be mainly dependent on seasonal reproduction rates, rather than hydrography. In October in particular, the relative contribution of *O. similis* was greater than 80% at all stations. A comparably

high abundance in the late autumn was previously reported in Kongsfjorden, as a result of reproduction from May to September (Lischka and Hagen, 2005). The contribution to the zooplankton community from *O. similis* was weaker in April, with increasing abundances of other copepod groups as a result of spring reproduction (Berge et al., 2015b). Although both sampling periods showed different spatial patterns of species diversity, which are estimated by permutation tests for the zooplankton community (Supplementary Table S2), spatial variation of TP values in zooplankton species (Table 3) were not related to species diversity.

Continuous energy supply based on flexible feeding behaviors may enable year-round reproduction of *O. similis* (Berge et al., 2015b). Cultivation experiments with *O. similis* have revealed a high clearance rate for ciliates, dinoflagellates (Atkinson, 1996; Zamora-Terol et al., 2013), and diatoms (Atienza et al., 2006; Pond and Ward, 2010). This flexible feeding by *O. similis* is likely connected to seasonally significant TP variation. In this study, as the TP value of *O. similis* decreased, their dominant diet appeared to shift after polar night from consumers to primary producers, with a case (C) in Figure 5. As a result, *O. similis* appeared to be connected with protozooplankton, making the dietary

composition primarily carnivorous in October and herbivorous in April. Iversen and Seuthe (2011) reported the integrated biomass of protozooplankton and phytoplankton were similar in pre-bloom seasons and before polar night in Kongsfjorden. When both diet sources are available to *O. similis*, it is likely that phytoplankton would be less preferred due to their low nutritional content compared with protozooplankton (Stoecker and Capuzzo, 1990; Atkinson, 1996). We found that the low solar elevation and short daylengths in October resulted in a diet dominated by protozooplankton. In contrast, when solar elevations were higher and days were longer in April, more phytoplankton was available.

Omnivorous zooplankton has different TP variations according to its habitat. For example, the copepod *Pleuromamma xiphias* is an opportunistic omnivore from the Sargasso Sea with a seasonal shift between carnivorous and herbivorous diets (Schnetzer and Steinberg, 2002). But in the north of Hawaii, *P. xiphias* showed similar TP values between summer and winter, as estimated by the CSIA-AAs method [2.8 ± 0.08 , Hannides et al. (2009)]. In addition, Hannides et al. (2009) reported that *Oithona* spp. in the north of Hawaii showed TP close to 2.0, with small seasonal variation. Their study suggested the diet of *Oithona* spp. would not consist solely of primary producers, in spite of a TP of 2.0, due to small ^{15}N discrimination in the rapid and tightly coupled microbial food web. Although our study produced results in opposition to those of Hannides et al. (2009), we were unable to directly compare TP values for *Oithona* between low and high latitudes. Because the environmental conditions differed between the two locations, the TDF could have changed, even by species, due to diet quality (Chikaraishi et al., 2015; McMahon et al., 2015). Opportunistic feeding activities of *Oithona* in various environments would therefore produce different TP values based on available diets and the magnitude of ^{15}N isotopic fractionation. Additional study would be required to determine the ecological relationship between *Oithona* and protozooplankton, and ciliates and dinoflagellates in particular.

Parasagitta elegans is a major carnivorous zooplankton in the Arctic, and its diet reportedly consists primarily of copepods (Samemoto, 1987; Terazaki, 2004). In this study, abundance of most copepods decreased with the onset of polar night, and the amount of available *P. elegans* consumption was therefore also lower. However, we found the abundance of *P. elegans* was constant between the two sampling periods, although temporal and spatial TP variations for the species were unclear. *P. elegans* requires additional lipids to achieve sexual maturity during spring algal blooms (Choe et al., 2003). Although growth rates and the magnitude of feeding activity of *P. elegans* decreased during winter compared with spring and summer, feeding activity continued (Grigor et al., 2014). Trophic position values of 2.9–3.0 for *P. elegans*, as estimated by bulk nitrogen isotope in January Grigor et al. (2015), were similar to our results. The constant TP before and after polar night can be explained by continuous feeding activity on copepods, as a case (B) in Figure 5. This also implies that the TP of copepods, which are available for consumption by *P. elegans*, would be constant before, during, and after polar night.

TABLE 3 | $\delta^{15}\text{N}$ of amino acids in zooplankton collected in the three stations of Kongsfjorden.

	October 2017			April 2018		
	$\delta^{15}\text{N}_{\text{Glu}}$	$\delta^{15}\text{N}_{\text{Phe}}$	TP	$\delta^{15}\text{N}_{\text{Glu}}$	$\delta^{15}\text{N}_{\text{Phe}}$	TP
<i>Parasagitta elegans</i> (>20 mm)						
St.01	22.4	3.1	3.1	23.6	3.5	3.2
St.07	23.8	3.0	3.3	24.0	5.9	2.9
St.10	23.6	4.6	3.0	25.1	6.0	3.1
<i>Parasagitta elegans</i> (<20 mm)						
St.01	22.8	4.3	3.0	24.0	3.3	3.3
St.07	23.2	3.3	3.2	23.3	3.4	3.2
St.10	24.5	5.0	3.1	24.4	5.3	3.1
<i>Calanus</i> spp.						
St.01	20.6	4.1	2.7	21.6	3.4	3.0
St.07	20.5	6.0	2.5	21.0	3.7	2.8
St.10	19.7	3.8	2.6	19.0	3.0	2.7
<i>Oithona similis</i>						
St.01	25.3	7.6	2.9	13.5	−0.3	2.4
St.07	26.0	8.9	2.8	14.0	1.0	2.3
St.10	22.0	4.4	2.8			
<i>Meganyctiphanes norvegica</i>						
St.01	23.4	3.3	3.2			
<i>Thysanoessa</i> spp.						
St.01				25.1	3.8	3.4
St.07				23.6	5.3	3.0
St.10				25.2	4.1	3.3
<i>T. abyssorum</i>						
Harbor	26.7	4.0	3.5			

All samples, except *O. similis*, were analyzed in triplicate and the average standard deviations of $\delta^{15}\text{N}_{\text{Glu}}$ and $\delta^{15}\text{N}_{\text{Phe}}$ were less than 0.5‰. *O. similis* samples were analyzed with a single injection due to the lack of a sufficient number of samples.

As a result of the TP estimates by CSIA-AAAs, *T. abyssorum* was identified as a carnivore in this study. *T. abyssorum* feeds primarily on *Calanus* (Kraft et al., 2013). But *T. abyssorum* was collected in the harbor only in October, making temporal comparisons of TP values impossible. The euphausiids *M. norvegica* and *Thysanoessa* sp. were collected during different sampling periods, and their TP values ranged from 3.0 to 3.4. The genus *Thysanoessa* has been reported as a herbivore or omnivore (Wold et al., 2011; Huenerlage et al., 2016). Similarly, *M. norvegica* is an omnivore, showing seasonally different diet preferences for diatoms and copepods in spring/summer and fall/winter, respectively (Schmidt, 2010). Trophic position values of euphausiids in this study suggest their diet was composed primarily of other zooplankton during our sampling periods.

In this study, copepodite was dominant in October and adult *C. finmarchicus* increased in April, showing similar results to a previous study (Willis et al., 2006). Continuous intrusion of North Atlantic waters into Kongsfjorden would result in a high contribution of *C. finmarchicus* to the group *Calanus* in October. In April, when winter-cooled water was dominant, the population of *Calanus* decreased and most individuals were resting in the deep layer (Conover, 1988). Previous studies have reported that the *Calanus* species store high-energy wax esters and utilize them during polar night (Scott et al., 2000). The wax material enhances survival over polar night, assuming minimal metabolic demands (Falk-Petersen et al., 2007). In our results, the TP values of *Calanus* spp. ranged from 2.5 to 3.0, suggesting a high contribution from primary consumers, and is in agreement with a reported gut-content analysis of *C. glacialis* (Cleary et al., 2017). *Calanus* spp. filter-feed mainly on phytoplankton, sea ice algae, detritus, protists, and small zooplankton (Frost, 1972; Levinsen et al., 2000; Falk-Petersen et al., 2007; Cleary et al., 2017). Their diets are therefore closely coupled with the composition of organic particles (both living and non-living) in the water mass, and all available diet components would be integrated and assimilated. According to previous studies, their TP values, as estimated by bulk isotope analysis, varied seasonally, ranging from the herbivorous to omnivorous position, and likely depend on diet available in the water column (Søreide et al., 2006, 2008). However, *Calanus* spp. appear to be omnivorous and carnivorous organisms, feeding on more heterotrophic species during the two non-productive seasons in Kongsfjorden, based on the TP values of 2.5 to 3.0 found in this study.

Compared with TP variation in *O. similis*, the average TP values of *Calanus* spp. showed little increase (0.2) after polar night, indicating no statistically significant difference. It is possible that different isotope turnover rates can be applied to *Calanus* spp. and *O. similis*. Both copepods showed high clearances on similar size classes (from 20 to 40 μm) of dinoflagellates and ciliates, suggesting available diet sizes for both copepods were not substantially different (Levinsen et al., 2000; Zamora-Terol et al., 2013). Instead, the slow growth rate of *Calanus* spp. may cause slow isotope turnover (Tamelander et al., 2006). The TP value of *O. similis* could shift earlier to a

phytoplankton diet compared to the TP of *Calanus* spp. during the cold-water conditions of the early spring.

CONCLUSION

The TPs of representative zooplankton species were measured to determine their feeding activity before and after polar night. We found that temporal TP variation of zooplankton reflected species-specific feeding strategies. In this study, nitrogen isotope analysis of AAs proved to be a useful approach to understanding changes in TPs of zooplankton species before and after polar night. For further study, additional application of CSIA-AAAs methods, including carbon isotope analysis of AAs, would help clarify the feeding strategy of zooplankton species under extreme environmental conditions, such as polar night.

DATA AVAILABILITY STATEMENT

All datasets generated for this study are included in the article/Supplementary Material.

AUTHOR CONTRIBUTIONS

HC, S-YH, SL, and J-HK participated in the field work. HC analyzed the nitrogen isotopes of amino acids in the samples and wrote the manuscript. SL and J-HK collected and sorted zooplankton samples for analysis. S-YH and K-HS designed the experiments and revised the manuscript. All authors contributed to the article and approved the submitted version.

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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.2020.00489/full#supplementary-material>

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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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