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Predatory impact of the myctophid fish community on zooplankton in the Scotia Sea (Southern Ocean)

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ABSTRACT: Myctophids are the most abundant mesopelagic fishes in the Southern Ocean, although their trophic role within the predominantly krill-based food web in regions south of the Antarctic Polar Front (APF) is poorly resolved. This study therefore examined the diets of 10 species of myctophid fishes: Electrona antarctica, E. carlsbergi, Gymnoscopelus braueri, G. fraseri, G. nicholsi, Krefftichthys anderssoni, Protomyctophum bolini, P. tenisoni, P. choriodon and Nannobrachium achirus, in the Scotia Sea, together with their predatory impact on the underlying zooplankton community. Myctophids and their prey were sampled in different seasons by scientific nets deployed across the Scotia Sea from the sea-ice zone to the APF. Based on the percentage index of relative importance, myctophids had high overlap in their diets, although the data indicate dietary specialisation in some species. There was also a distinct switch in diet, from copepods to euphausiids and amphipods, with increasing myctophid size. Myctophid predation impacted daily copepod production by between 0.01 and 5%, with Calanus simillimus being most impacted. Total annual consumption of copepods was around 1.5 million t (Mt) per year. All myctophids preyed upon the euphausiid Thysanoessa spp., consuming ~12% of its daily productivity and around 4 Mt per year. However, only larger myctophid species preyed upon Antarctic krill Euphausia superba, consuming 2% of its daily productivity, which could amount to as much as 17 Mt per year. Themisto gaudichaudii was also an important dietary component, with 4 % of its daily productivity being consumed, amounting to around 2 Mt per year. This study demonstrates that myctophids link secondary productivity to higher predators both through krill-dependent and krill-independent trophic pathways.

KEY WORDS: Myctophidae · Predation rates · Feeding ecology · Scotia Sea · Southern Ocean

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INTRODUCTION

The estimated global biomass of mesopelagic fish is in excess of 11000 million tons (Mt), making them a major contributor to the function of oceanic ecosystems and global biogeochemical cycles (Irigoien et al. 2014). Mesopelagic fish transfer energy through pelagic food webs, linking primary consumers and omnivorous macro-zooplankton to higher marine predators. They also contribute to the export of carbon from the sea surface to mesopelagic depths through their extensive vertical migrations (Pakhomov et al. 1996, Smith 2011, Irigoien et al. 2014). Nevertheless, despite their ecological importance, this group of fish remains one of the least investigated components of the oceanic ecosystem, with major uncertainties in their abundance, biology and ecology. Of the mesopelagic fishes, myctophids (family Myctophidae) are considered one of the most diverse and numerically abundant families (Gjøsaeter & Kawaguchi 1980). Determining the ecology of myctophids therefore constitutes an important step towards understanding the operation of oceanic ecosystems at both regional and global scales.

Our understanding of myctophids is confounded primarily due to difficulties in sampling them appropriately at the necessary spatial and temporal scales, particularly in remote, high-latitude regions such as the Southern Ocean. One example of a high-latitude region where myctophids are considerably understudied is the Scotia Sea in the Atlantic sector of the Southern Ocean; one of the most productive regions of the Southern Ocean (Holm-Hansen et al. 2004). This region is also subject to broad-scale, long-term environmental change, with marked increases in sea-surface temperatures and substantial reductions in both winter sea ice extent and Antarctic krill stocks (de la Mare 1997, Curran et al. 2003, Atkinson et al. 2004, Murphy et al. 2007a, Whitehouse et al. 2008). There is therefore an imminent need for more information on all components of the Scotia Sea pelagic ecosystem, particularly myctophids, in order to understand and predict the manifestations of this change, both in the Scotia Sea and throughout the Southern Ocean.

There are 33 species of myctophid fish in the Scotia Sea comprising an estimated biomass of 4.5 million t (Mt) (Collins et al. 2012). Although the food web of the Scotia Sea is predominantly centred on Antarctic krill Euphausia superba (Murphy et al. 2007b), it is clear that other trophic pathways are both regionally and seasonally important, with myctophids providing a key alternative (Murphy et al. 2007b, Stowasser et al. 2012). Myctophids in the Scotia Sea are the primary prey of king penguins Aptenodytes patagonicus, elephant seals Mirounga leonina and squid Martialia hyadesi, and are important dietary components for many other predators, including fur seals Arctocephalus gazella, Cape petrels Daption capense and toothfish Dissostichus eleginoides (Olsson & North 1997, Casaux et al. 1998, Brown et al. 1999, Dickson et al. 2004, Reid et al. 2006, Collins et al. 2007). In turn, they are predators of copepods, amphipods and euphausiids, including Antarctic krill (Pusch et al. 2004, Shreeve et al. 2009, Saunders et al. 2014, 2015a). Under a scenario of regional ocean-warming and declines in krill stocks, the role of myctophids in food webs may become increasingly important. However, the extent to which myctophids can potentially support the ecosystem against such change is unknown, primarily due to uncertainties in their distribution of abundance and trophodynamics.

Determining diet is essential to understanding food web dynamics and resource partitioning (Ross 1986), but studies of Southern Ocean myctophid diets have been predominantly restricted to the most abundant species on limited spatial and temporal scales, often with very small sample sizes (Rowedder 1979, Naumov et al. 1981, Kozlov & Tarverdiyeva 1989, Gerasimova 1990, Pakhomov et al. 1996, Gaskett et al. 2001, Pusch et al. 2004, Shreeve et al. 2009). Recent studies have cast new light on the diet and feeding ecology of myctophids in the Scotia Sea at more appropriate spatial and temporal scales (Saunders et al. 2014, 2015a,b), but parameters that are important for determining their trophic role, such as daily rations, have rarely been estimated (Gerasimova 1990, Pakhomov et al. 1996, Pusch et al. 2004, Shreeve et al. 2009). Furthermore, only a few studies considered predation impact of Southern Ocean myctophids on their prey species, focussing on a small range of prey species at limited spatial and temporal scales (Williams 1985, Pakhomov et al. 1996, Pusch et al. 2004, Shreeve et al. 2009).

In this study, we examine and compare the diets of the most abundant myctophid species across the entire latitudinal extent of the Scotia Sea (63°S to 50°S), spanning the sea-ice zone (SIZ) to the Antarctic Polar Front (APF). Furthermore, we integrate over the austral spring, summer and autumn to gain a seasonally averaged perspective. Vertical distributions of myctophids are compared with those of their prey species to investigate the spatial overlap between predators and prey and to assess the extent of prey selectivity. The predation impact of myctophids on prey assemblages was also estimated and sensitivity analyses used to determine confidence intervals around these estimates. These data are the most comprehensive for any region of the Southern Ocean to date and provide important parameterisations for new food web and ecosystem studies in the region. They also contribute to resolving the composition and dynamics of the global mesopelagic fish community that is a prerequisite for understanding global ecosystem and biogeochemical processes.

MATERIALS AND METHODS

Oceanographic, acoustic and biological data were collected in the Scotia Sea during 3 research cruises on board RRS James Clark Ross in October–December 2006 (JR161, austral spring), January–February 2008 (JR177, austral summer) and March–April 2009 (JR200, austral autumn). The study area covered regions from the SIZ to the APF, with sampling stations distributed across several prevailing water masses and frontal zones (Fig. 1). Six nominal sta-

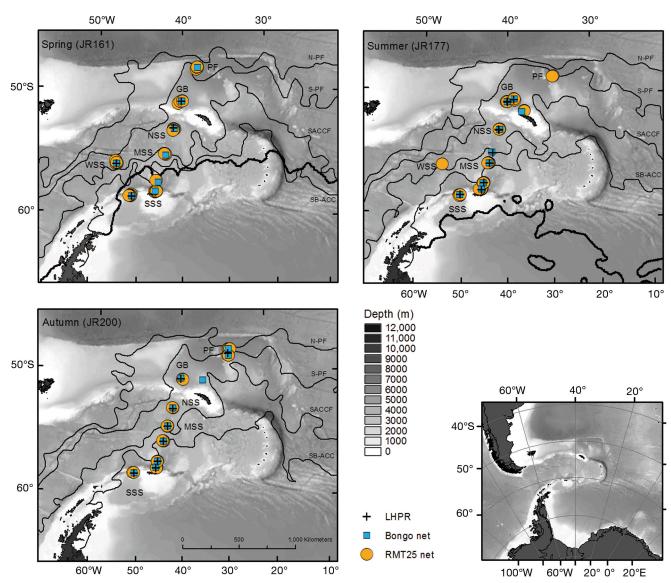


Fig. 1. Locations of 25 m² rectangular midwater trawls (RMT25), Longhurst-Hardy Plankton Recorder (LHPR) trawls and Bongo net hauls during 3 surveys. Sampling stations included Southern Scotia Sea (SSS), Western Scotia Sea (WSS), Mid-Scotia Sea (MSS), North Scotia Sea (NSS), Georgia Basin (GB) and Polar Front (PF). Mean frontal positions (thin black lines) determined during the cruises from dynamic height data (Venables et al. 2012) include the northern Antarctic Polar Front (N-PF), southern Antarctic Polar Front (S-PF), South Antarctic Circumpolar Current Front (SACCF) and Southern Boundary of the Antarctic Circumpolar Current (SB-ACC). The heavy black line shows the position of the 15 % ice-edge cover for 24 October 2006 (JR161) and for 15 January 2008 (JR177). The ice-edge occurred south of the transect during autumn 2009 (JR200). Bathymetry data from GEBCO_08 grid (version 20091120, www.gebco.net)

tions were sampled repeatedly across the study site during the surveys: Southern Scotia Sea (SSS), Mid Scotia Sea (MSS), Western Scotia Sea (WSS), Northern Scotia Sea (NSS), Georgia Basin (GB) and the Polar Front (PF).

Net sampling

Mesopelagic fish were collected with a 25 m^2 rectangular midwater trawl net (RMT25) (Piatkowski et al. 1994). Depth stratified hauls were undertaken at each station covering depth intervals between 0–200, 200–400, 400–700 and 700–1000 m. The hauls were repeated day and night in spring and summer, but only during hours of darkness in the autumn. The abundance and vertical distribution of the zooplankton prey were characterised by oblique Longhurst-Hardy Plankton Recorder (LHPR) tows to 1000 m depth during both day and night. The LHPR was equipped with a 0.38 m diameter nose cone and a 200 µm mesh net and filtering gauzes. The gauze advance mechanism was set to 90 s during the spring and 120 s during summer and autumn, which resulted in a depth resolution of around 20-25 m per patch. The prey field was further characterised using a paired Bongo net (180 mm diameter mouth) fitted with 53 µm mesh. Bongo nets were deployed to 400 m depth and hauled vertically to the surface during hours of daylight. Further details of the net samplers, haul deployments and analyses are described in Collins et al. (2012) and Ward et al. (2012).

Sample processing

RMT25 net haul catches were sorted onboard to the lowest possible taxonomic level (Hulley 1990). Total catch weights per fish species were recorded using a motion-compensated balance and all fish were measured to the nearest mm using standard length (SL). Stomachs were dissected from a random sub-sample of 25 fish per net haul, or from each specimen where catches were small. All stomachs were frozen for subsequent microscopic analysis. LHPR samples were frozen at -20°C and transported back to the laboratory where species were identified and enumerated under a stereomicroscope. Counts were averaged into the same depth horizons as used for the RMT25 net hauls to enable direct comparisons of vertical distributions. Bongo net samples were preserved in 4 % formalin and seawater solution and subsequently aliquots were analysed under a stereomicroscope back at the laboratory.

Stomach contents analysis

Following Shreeve et al. (2009), fish stomach contents were thawed and sorted to the lowest taxonomic level that the state of digestion would allow. Individual prey items were enumerated and weighed. If the prey was highly disaggregated, the weights of component species were estimated as a proportion of the weight of the total contents.

Diet was expressed using 4 measures: (1) percentage frequency of occurrence (%F), (2) percentage mass (%M), (3) percentage number (%N) and (4) percentage index of relative importance (%IRI) (Cortes 1997). The %IRI was calculated for prey species and %IRI_{DC} was calculated for prey categories (Main et al. 2009, Shreeve et al. 2009). The initial prey categories used in the analysis were defined according to order (Amphipods, Copepods, Euphausiids, Ostracods, Molluscs, Urochordata and other taxa), but a more detailed analysis was performed subsequently for the most numerically dominant prey categories: the copepods *Metridia* spp., *Pleuromamma robusta*, *Rhincalanus gigas*, *Calanoides acutus*, *Calanus simillimus*, *Paraeuchaeta* spp., 'other copepods', the euphausiids *Euphausia superba*, *Thysanoessa* spp., 'other euphausiids', the amphipod *Themisto gaudichaudii* and 'other taxa' (mostly unidentified crustaceans, Mollusca, Ostracoda, Urochordata). The %IRI was calculated as:

$$\text{\%IRI}_{i} = \frac{(\%N_{i} + \%M_{i}) \times \%F_{i}}{\sum_{i=1}^{n} (\%N_{i} + \%M_{i}) \times \%F_{i}} \times 100 \tag{1}$$

where i is prey item.

95% confidence limits for the mean %IRI of each prey category were calculated using a bootstrapping technique, whereby each species dataset (individual stomachs) was re-sampled (with replacement) 1000 times (Main et al. 2009).

Diet comparison between myctophid species

Similarities in the diets of the myctophid species were examined using the PRIMER (version 6) software package (Clarke & Warwick 2001). The %IRI values for each diet component for each myctophid species were first square root transformed and a Bray-Curtis similarity index was then calculated for each pair of species. Hierarchical agglomerative cluster analysis was performed on this data set using the group average linking method and a SIMPER routine was used to determine which prey species contributed most to the resulting cluster groupings.

Predation impact of myctophids

Following Shreeve et al. (2009), we used the following function to determine the proportion of prey productivity consumed by each myctophid species:

$$I_{i,j} = \frac{N_{i,j} C_i P_j \left(\frac{24}{G}\right)}{Z_i F_i}$$
(2)

where $I_{i,j}$ is the proportion of production of prey species *i* consumed by myctophid species *j* per day, $N_{i,j}$ is the number of individuals of prey species *i* in the stomachs of myctophid species *j*, C_i is the carbon mass of prey species *i*, P_j is the depth-integrated concentration of predator species *j* (ind. m⁻²), *G* is the gut passage time (h), Z_i is the depth-integrated concentration of prey species *i* (ind. m⁻²), and F_i is the growth rate of prey species i (µg C d⁻¹). We extended this calculation to estimate total consumption of each prey taxon by myctophids using the equation:

$$Q_{i} = A D 365 R \left(\frac{\sum_{j} N_{i,j} C_{i} P_{i}}{\sum_{i,j} N_{i,j} C_{i} P_{i}} \right)$$
(3)

where Q_i is the total annual consumption of prey taxon *i*, *A* is the approximate area of the Scotia Sea (2 million km²), *D* is the mean density of myctophids (2.23 t km² ± 0.79 SD), and *R* is the daily food intake of myctophids as a percentage of body mass (1.5%) All values were taken from Collins et al. (2012). *R* is a mean daily ration (% dry body weight) calculated from data presented in Pakhomov et al. (1996) for Antarctic and high sub-Antarctic myctophids. CI (95%) were calculated around our annual consumption estimates to represent the variation in mean myctophid density observed in the Scotia Sea.

We used the approach of Shreeve et al. (2009) to derive the most plausible estimates of $I_{i,i}$ and their upper and lower bounds. The upper bound is based on the upper estimate of the number of prey items *i* eaten by myctophid *j*, the upper estimated concentration of myctophid *j*, the lower estimated concentration of prey *i*, and the fastest gut passage time. Conversely, the lower bound is derived from the lower estimate of the number of prey species *i* in the stomachs of myctophid species *j*, the lower estimated concentration of myctophid j_i , the upper estimated concentration of prey species *i*, and the slowest gut passage time. The most plausible estimate uses the median values for each of the above parameters. Each of these parameter values were calculated as detailed below.

Numbers of individuals of prey species *i* in the stomachs of myctophid $j(N_{i,j})$

Ten myctophid species were considered in our analysis: *Electrona antarctica, E. carlsbergi, Gymnoscopelus braueri, G. fraseri, G. nicholsi, Protomyctophum bolini, P. tenisoni, P. choriodon, Krefftichthys anderssoni* and *Nannobrachium achirus*. The dataset was restricted to the most common prey taxa found in the myctophid stomachs: the amphipod *Themisto gaudichaudii*, the euphausiids *E. superba, Euphausia frigida* and *Thysanoessa* spp., the copepods *Metridia* spp., *R. gigas, Calanoides acutus, Calanus simillimus, Pleuromamma robusta, Paraeuchaeta* spp., and *Oncaea* spp., ostracods, salps and pteropods.

The following non-parametric bootstrapping technique was used to generate the upper and lower bounds; for each myctophid species, 30 individuals were extracted at random and the mean number of items of each prey species in this subset was calculated and the process was repeated 100 times. The median of the series was used as the best estimate value, with the 25th and 75th percentiles comprising the lower and upper bounds, respectively.

Depth-integrated myctophid concentrations (P_i)

Myctophid concentrations were determined from the RMT25 net catches that were aggregated for all surveys and regions across the Scotia Sea. Only night-time hauls were used in the analysis to avoid potential bias due to daylight net avoidance in the upper regions of the water column (Collins et al. 2012). A total of 86 stratified net hauls were deployed during this time. At each station, the entire water column between 0-1000 m was sampled in depth-discrete intervals. Net catch concentrations (ind. m⁻³) were therefore multiplied by the respective depth interval (m) and combined to give a depth-integrated concentration per net (ind. m⁻²) between 0 and 1000 m. Our best estimate value for P_i was the median of the pooled net concentrations, with the 25th percentile representing the lower bound and the 75th percentile comprising the upper bound.

Depth-integrated prey species concentrations (Z_i)

A total of 24 LHPR deployments were undertaken during the study, each sampling the whole water column between 0–1000 m at a depth resolution of approx. 20–25 m. Net catch concentrations of prey species (ind. m^{-3}) were multiplied by the respective depth interval and summed to give depth-integrated concentrations (ind. m^{-2}) per haul between 0– 1000 m. All LHPR hauls were pooled for all surveys and the median of this series was used as the best estimate value, the 25th percentile value as the lower bound and the 75th percentile value as the upper bound.

Prey species abundance estimates (standardised to ind. m^{-2}) were also calculated from 65 Bongo net hauls deployed between 0–400 m. These data were pooled for all surveys and the median, 25^{th} and 75^{th} percentile values were selected to represent the best estimate values and their associated upper and lower bounds. We assumed that all zooplankton sampling devices would most likely underestimate the actual concentrations of prey species present in the water column. Therefore, the median LHPR and Bongo net values were examined and the highest estimates for each species were selected for use in our calculations. This approach, which applied mostly to copepods, was adopted to provide the most conservative estimates of myctophid predation rates on the prey field. Some prey species exhibited a high degree of patchiness during the surveys and were absent in several of the net hauls. On occasion, this resulted in 25^{th} percentile values of zero for these species (see Table 1) and in such instances, it was not possible to calculate an upper bound for $I_{i,i}$.

Growth rate of prey species (F_i)

Following Shreeve et al. (2009), species-specific growth rates (μ g C d⁻¹) were estimated from direct measurements of carbon weight, multiplied by the weight-specific growth rate of each species using the functions provided by Hirst et al. (2003). Mean carbon weight measurements were calculated from around 10 to 60 individuals of each species during the surveys. For the copepod species, we used a weight-specific growth rate function appropriate for adult broadcast spawning copepods at 5°C. A growth rate function covering all crustaceans (excluding copepods) at 5°C was selected for the euphausiids, amphipods and ostracods, whilst a function suitable for Thaliaceans at 15°C was used for salps. Although these functions were derived at temperatures greater than those of our study region, particularly for Thaliaceans, they are the most appropriate functions available in the scientific literature to date. We consider estimates derived from these functions to represent an upper limit to zooplankton production, which means that our calculations represent a minimum of the predatory impact of myctophids on zooplankton. We assumed that the majority of pteropod species collected during the surveys were most probably Limacina spp., therefore the growth rate function provided by Bednaršek et al. (2012) was used for this prey group.

Gut passage time (G)

The temperature-specific gut passage time function detailed in Shreeve et al. (2009) was used in our analysis:

$$y = 4.50 + 24.92^{(-0.265x)} \tag{4}$$

where *y* is gut passage time (h) and *x* is temperature.

This model was derived from data on the gut passage time of a number of different planktivorous fish from various locations with different ambient water temperatures (Pakhomov et al. 1996). In our calculations, temperature data collected at each station during the surveys (Venables et al. 2012) were collated and averaged to provide an estimate of the overall ambient temperature between 0-1000 m across the Scotia Sea. The mean temperature in the region was 0.67°C, giving an estimated gut passage time of 25.4 h, which was used as our best estimate value. Mean temperature values varied between -0.30 to 2.0°C, giving an estimated slowest gut passage time of 31.2 h and a fastest gut passage time of 19.1 h. This level of variance simulates to a degree the variance in gut passage time between prey species found in other studies (Andersen 1999, Andersen & Beyer 2008), although further investigations are required to provide more robust species-specific gut passage times for Southern Ocean zooplankton.

RESULTS

Myctophid distribution

Detailed descriptions of the horizontal and vertical distributions of the myctophids are given in Collins et al. (2012) and Saunders et al. (2014, 2015a,b), therefore only an overview is given here. These studies also provide information on their seasonal and regional biomass. Electrona antarctica and Gymnoscopelus braueri were the most abundant species encountered on the surveys (Fig. 2). These 2 species occurred throughout the Scotia Sea, including the sea ice sectors, where E. antarctica was most abundant. G. nicholsi had a similar distribution pattern, but occurred only in small numbers. Krefftichthys anderssoni, Protomyctophum bolini and E. carlsbergi were the most abundant species in the northern Scotia Sea, but they seldom occurred at the southernmost stations. P. tenisoni, Nannobrachium achirus, G. fraseri and P. choriodon were also distributed predominantly in the northern regions, with the abundance of P. tenisoni and N. achirus being highest in regions associated with the APF, and G. fraseri and P. choriodon highest around the Georgia Basin.

Only nighttime data were used here to illustrate the vertical distribution of the myctophid species because of possible daytime net avoidance in the upper water column (Fig. 3). Six species were distributed predominantly in the upper 400 m of the water column, with *E. carlsbergi*, *P. bolini*, and *P.*

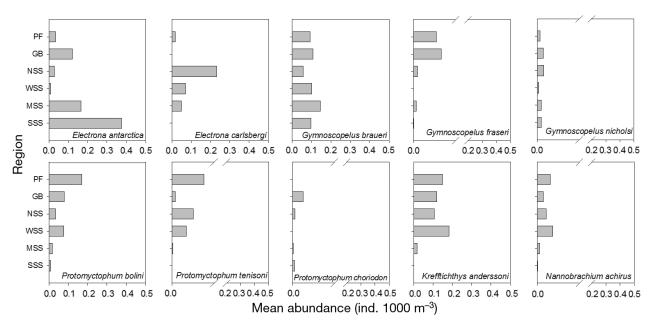


Fig. 2. Mean abundance of myctophid fish at each station located in the Polar Front (PF), Georgia Basin (GB), North Scotia Sea (NSS), West Scotia Sea (WSS), Mid Scotia Sea (MSS), and South Scotia Sea (SSS) regions during 3 surveys. The breaks in the abundance axis start at 0.05 ind. 1000 m⁻³. Comprehensive descriptions of these species distribution patterns are given in Collins et al. (2012) and Saunders et al. (2014, 2015a,b)

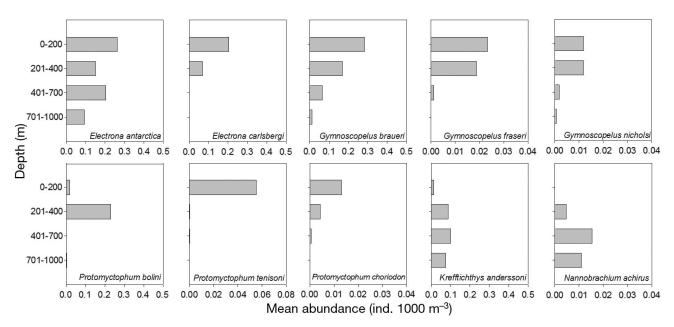


Fig. 3. Nighttime depth distributions of myctophid fish caught in the RMT25 net hauls during the 3 surveys. Data modified from Saunders et al. (2014, 2015a,b)

tenisoni restricted exclusively to this zone, and *P. choriodon*, *G. fraseri*, and *G. nicholsi* occurring only in low abundance in regions deeper than 400 m. *E. antarctica*, *G. braueri* and *K. anderssoni* were caught throughout the sampled depth range, whilst *N. achirus* was distributed predominantly below 400 m.

Abundance and vertical distribution of zooplankton prey species

Best estimates (median values) of depth-integrated macrozooplankton abundance varied between 37 ind. m^{-2} for *Euphausia frigida* to 636 ind. m^{-2} for *E. superba* (Table 1). All euphausiid species occurred

Table 1. Depth-integrated net catch concentrations of the most abundant myctophid fish and zooplankton taxa in the Scotia Sea during the 3 surveys. Concentration estimates are the 25th percentile (lower), **median** and 75th percentile (upper)

Taxon	Sam	pling	Cond	centration (ind. m ⁻²)
	Device	Depth (m)	Lower	Median	Upper
		,			
Myctophidae					
Electrona carlsbergi	RMT25	0-1000	0.002	0.015	0.207
Electrona antarctica	RMT25	0-1000	0.003	0.155	0.586
Gymnoscopelus fraseri	RMT25	0-1000	0.002	0.007	0.048
Gymnoscopelus nicholsi	RMT25	0-1000	0.002	0.004	0.015
Gymnoscopelus braueri	RMT25	0-1000	0.002	0.078	0.431
Krefftichthys anderssoni	RMT25	0-1000	0.002	0.067	0.346
Nannobrachium achirus	RMT25	0-1000	0.003	0.006	0.033
Protomyctophum tenisoni	RMT25	0-1000	0.002	0.006	0.084
Protomyctophum bolini	RMT25	0-1000	0.002	0.032	0.143
Protomyctophum choriodon	RMT25	0-1000	0.002	0.003	0.030
Amphipoda					
Themisto gaudichaudii	Bongo	0-400	0.000	235.740	628.672
Copepoda					
Calanoides acutus	LHPR	0-1000	569.040	1018.730	2187.315
Calanus simillimus	Bongo	0-400	0.000	117.900	7858.400
<i>Metridia</i> spp.	Bongo	0-400	3143.360	11237.512	21570.210
Oncaea spp.	Bongo	0-400	196.460	6522.472	71664.960
Pleuromamma robusta	Bongo	0-400	78.580	12180.520	46207.392
Paraeuchaeta spp.	Bongo	0 - 400	117.876	275.044	471.504
Rhincalanus gigas	Bongo	0-400	157.168	1178.760	5343.440
Euphausiacea	-				
Euphausia frigida	LHPR	0-1000	1.218	37.340	482.553
Euphausia superba	LHPR	0-1000	0.000	636.693	13021.204
Thysanoessa spp.	LHPR	0-1000	0.000	134.571	1150.767
Ostracoda					
Ostracods	Bongo	0-400	628.640	943.008	1729.200
Mollusca	Dongo	0-400	020.040	310.000	1720.200
	Domero	0 400	600.000	2020.024	14459.456
Pteropods	Bongo	0-400	628.800	2829.024	14459.456
Urochordata					
Salps	LHPR	0-1000	0.000	46.957	766.109

predominantly in the upper 200 m of the water column along with the amphipod *Themisto gaudichaudii* (Fig. 4), which had a depth-integrated abundance of 236 ind. m⁻². Salps were found mainly above 400 m and had a depth-integrated abundance of 47 ind. m⁻². Pteropod counts were only available from the Bongo net hauls, so it was not possible to examine their vertical distribution. These organisms had a depth-integrated concentration of 2829 ind. m⁻². Ostracods comprised a depth-integrated abundance of 943 ind. m⁻² and were spread throughout the water column, with the greatest concentrations above 400 m.

Copepods generally occurred in greater concentrations than macrozooplankton, with best estimates of depth-integrated abundance ranging between 118 and 12181 ind. m^{-2} . The most abundant copepod species were *Pleuromamma robusta*, *Metridia* spp. and Oncaea spp. (Table 1). These 3 species were found throughout the water column, but the highest concentrations occurred mostly above 400 m (Fig 4). Calanoides acutus, Calanus simillimus, and Paraeuchaeta spp. were found at all depths, but maximal concentrations were found in the upper 200 m. Rhincalanus gigas occurred predominantly above 700 m, with the greatest concentrations spread between the surface and 400 m.

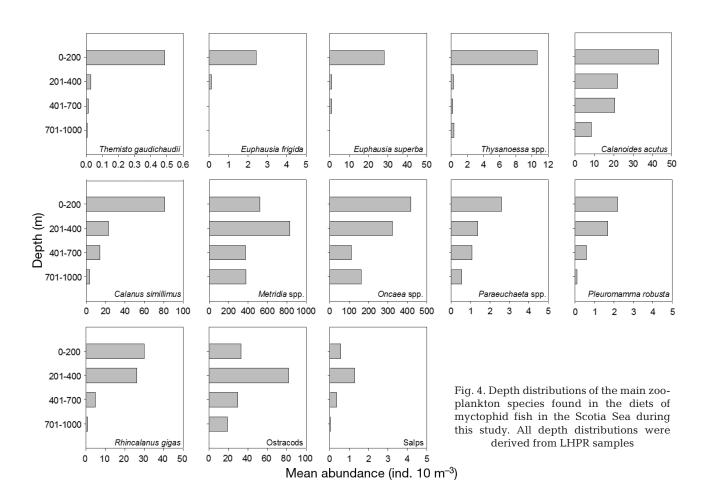
Diet compositions

A total of 1804 myctophid stomachs contained prey items and were used in the analysis (Table 2). Empty stomachs were excluded from the analysis. For each myctophid species, the size ranges, depths and locations of the sampled fish were representative of those found previously in the Scotia Sea region (Hulley 1981, McGinnis 1982, Pusch et al. 2004, Collins et al. 2008).

Planktonic crustaceans dominated the diets of all myctophid species (Fig. 5, Tables S1–S4 in the Supplement at www.int-res.com/articles/ suppl/m541p045_supp.pdf). The diet of *Electrona antarctica* (24–115 mm SL) was dominated by *Euphausia*

superba and *T. gaudichaudii* (Fig. 5, Table S1). These species were distributed predominantly in the upper 200 m, a region that *E. antarctica* appeared to occupy only at night. By contrast, *Electrona carlsbergi* was found in greatest abundance above 200 m at night and had a smaller size range (68–90 mm SL). *E. carlsbergi* was predominantly a copepod feeder (93% IRI) with *R. gigas, Metridia* spp., and *Oncaea* spp. the most preyed upon species (Fig. 5, Table S1).

The 3 *Gymnoscopelus* species had diets that were dominated by copepods and euphausiids, although there were some differences in their respective diets (Fig 5, Table S2). *G. braueri* (mean = 82 mm SL) reached its maximum abundance in the upper 200 m at night and had a diet dominated by the copepod *Metridia* spp. and the euphausiid *Thysanoessa* spp. (Table S2). *T. gaudichaudii* and *Euphausia superba* also formed an important part of this species' diet



(~5% IRI). Similarly, the abundance of *G. fraseri* (mean = 67 mm SL) was highest between 0–200 m at night and the species preyed mostly upon *Metridia* spp., although *R. gigas* formed a substantial part of the diet (10% IRI) and *E. superba* was absent. By contrast, *G. nicholsi* (mean = 126 mm SL), which was spread between the surface and 400 m at night, had

a diet dominated by *Metridia* spp., *R. gigas*, and *E. superba* (Table S2). This species also took substantial proportions of *P. robusta* (10% IRI).

Protomyctophum bolini (mean = 49 mm SL) was mainly caught between 200–400 m at night and fed mostly on copepods (Table S3, Fig. 5). The principle prey species were *Metridia* spp., *R. gigas* and *Thysa*-

Table 2. Numbers of myctophid stomachs containing prey items from each station during the 3 surveys in the regions: South Scotia Sea (SSS), Mid Scotia Sea (MSS) West Scotia Sea (WSS), North Scotia Sea (NSS), Georgia Basin (GB) and Polar Front (PF). The mean size (standard length; SL) and SL ranges of the fish specimens from which the stomachs were extracted are also given

Species	SSS	MSS	WSS	NSS	GB	PF	Total	Mean SL (mm)	Range SL (mm)
Electrona antarctica	228	83	3	8	133	30	485	71	24-115
Electrona carlsbergi	0	51	0	102	2	30	185	77	68-90
Gymnoscopelus braueri	96	81	9	36	64	86	372	82	34-162
Gymnoscopelus fraseri	0	0	0	2	58	43	103	67	39-115
Gymnoscopelus nicholsi	10	10	1	8	5	6	40	126	34-165
Protomyctophum bolini	20	17	28	28	76	62	231	49	23-66
Protomyctophum tenisoni	0	0	9	15	0	22	46	42	32-55
Protomyctophum choriodon	0	0	0	0	30	7	37	70	55-85
Krefftichthys anderssoni	2	24	18	79	108	50	281	51	15-74
Nannobrachium achirus	1	1	3	4	9	6	24	132	65-167

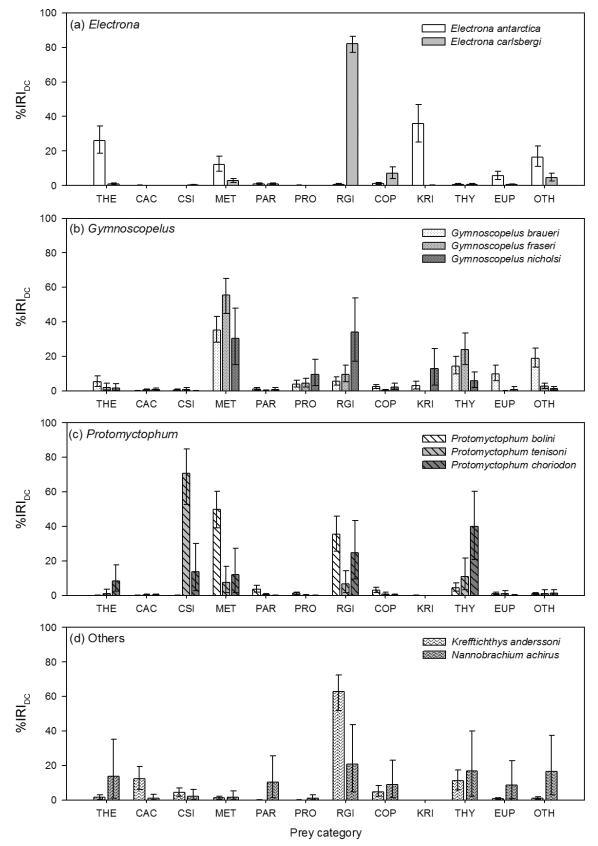


Fig. 5. Diet composition of 10 myctophid species: (a) *Electrona* spp., (b) *Gymnoscopelus* spp., (c) *Protomyctophum* spp. and (d) others, in the Scotia Sea, expressed as the percentage index of relative importance for prey categories (%IRI_{DC}) — *Themisto gaudichaudii* (THE), *Calanoides acutus* (CAC), *Calanus simillimus* (CSI), *Metridia* spp. (MET), *Paraeuchaeta* spp. (PAR), *Pleuromamma robusta* (PRO), *Rhincalanus gigas* (RGI), other copepods (COP), *Euphausia superba* (KRI), *Thysanoessa* spp. (THY), other euphausiids (EUP), and other taxa (OTH) comprising predominantly unidentified crustaceans, ostracods and pteropods. Error bars are the bootstrapped 95% CI

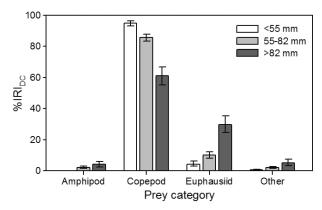


Fig. 6. Diet composition, expressed as percentage index of relative importance by prey category (% IRI_{DC}) of all myctophid species grouped by size class (standard length; mm). The Other category was dominated by unidentified crustaceans, ostracods, pteropds and salps. Size classes were derived from the 25th and 75th percentiles of the pooled length-frequency data

noessa spp. P. tenisoni (mean = 42 mm SL) occurred in the top 200 m at night and also preyed upon copepods, particularly Calanus simillimus (75%IRI), together with substantial proportions of the euphausiid Thysanoessa spp. (10% IRI). By contrast, the main copepod prey species of P. choriodon (mean = 70 mm SL) was R. gigas and this myctophid species preyed upon much greater proportions of Thysanoessa spp. (42%IRI) than P. bolini and P. tenisoni (Table S3). P. choriodon abundance was greatest above 200 m at night and Themisto gaudichaudii also comprised an important component of its diet (5% IRI).

K. anderssoni (mean = 51 mm SL), which was most abundant between 200 and 700 m, fed mostly on copepods, particularly *R. gigas* (59% IRI). This myctophid also took relatively high proportions of Calanoides acutus (16% IRI) and the euphausiid Thysanoessa spp. (14% IRI) (Fig. 5, Table S4). N. achirus (mean = 132 mm SL) was the largest myctophid species studied and it occurred in highest abundance below 400 m. The sample size was relatively small for this species, but the available data indicate that it fed upon copepods, euphausiids and amphipods, with R. gigas (25% IRI), Thysanoessa spp. (25%IRI), and unidentified non-hyperiid amphipods (6% IRI) comprising the main dietary components within these groups (Fig. 5, Table S4). N. achirus also took relatively high proportions of the copepod Paraeuchaeta spp. (15% IRI) and was the only species to prey upon fish (9% IRI).

Copepods were the dominant prey items in all myctophid size classes, although there was a distinct change in diet with size (Fig. 6). The smallest sized fish (<55 mm SL) consumed significantly more copepods than the larger size classes, with the older copepodite stages usually predominant (CV and CVI stages of Metridia spp., Calanoides acutus, and Calanus simillimus). A greater range in developmental stages was only apparent for Paraeuchaeta spp., with stages from CII upwards being present and the CIII stage being the most abundant in myctophid diets. Euphausiids and amphipods increased proportionally in the diet with increasing fish size. Euphausiids $(\sim 30\% \text{ IRI}_{\text{DC}})$ and amphipods $(\sim 5\% \text{ IRI}_{\text{DC}})$, including the species E. superba and Themisto gaudichaudii, were most abundant in the largest sized fish (>82 mm SL) (Fig. 6). There was a further increase in diet breadth with increasing size, as other taxa became more prevalent in larger sized fish. The 'other taxa' category was dominated by unidentified crustaceans, ostracods, pteropods and salps.

Impact on prey productivity

The majority of stomachs examined contained more than 1 species of prey, with some myctophids containing more than 5 prey species. For most myctophid species, each copepod prey species was consumed in numbers of 10 or more, whilst the main macrozooplankton taxa preyed upon were commonly found in numbers of 5 or more. However, when averaged out for a particular myctophid species, the number of prey items was mostly <1 because of the large numbers of stomachs from which a prey species was absent (Table 3). The exception were some of the copepod species, particularly Metridia spp. and R. gigas, which were found in relatively high numbers in the stomachs of the predominant copepod feeders, such as Electrona carlsbergi, G. nicholsi and G. fraseri. In these instances, the average prey numbers per stomach were >1. *Thysanoessa* spp. was the only macrozooplankton prey item to be taken in sufficient quantities such that the average prey numbers per stomach was greater than 1 (Table 3). This prey item was most abundant in the stomachs of Protomyctophum choriodon and G. fraseri.

Best estimates of average depth-integrated concentration across all 10 myctophid species in the upper 1000 m ranged between 0.003 and 0.155 ind. m^{-2} (Table 1). In the best estimate, myctophids consumed up to ~5% of the daily productivity (C m⁻² d⁻¹) of key copepod taxa in the Scotia Sea, with *K. anderssoni* having the greatest overall impact, consuming ~2% of the *C. simillimus* production (Table 4). The impact of myctophid predation on macrozooplankton

Table 3. Estimates of the number of individuals of key prey taxa within the stomachs of different myctophids in the Scotia Sea. Estimates represent the 25 th percentile (upper) of the dataset
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species	Esti- mate	Themisto Euphau- gaudi- sia chaudii frigida	Euphau- sia frigida	Euphau- sia superba	<i>Thysa-</i> <i>noessa</i> spp.	Cala- noides acutus	Calanus simil- limus	<i>Metri-</i> dia spp.	<i>On-</i> <i>caea</i> spp.	Pleuro- mamma robusta	<i>Para-</i> <i>euchaeta</i> spp.	Rhin- calanus gigas	Ostra- cods	Ptero- pods	Salps
Electrona	Lower	0.10	0.00	0.00	0.17	0.03	0.30	1.47	1.06	0.10	0.33	11.99	0.03	0.00	0.13
carlsbergi	Median	0.23	0.00	0.00	0.27	0.10	0.43	1.87	2.42	0.13	0.50	13.78	0.03	0.10	0.60
)	Upper	0.51	0.03	0.03	0.50	0.23	0.67	2.50	4.02	0.23	0.67	15.04	0.07	0.27	0.97
Electrona	Lower	0.27	00.00	0.20	0.03	0.00	0.00	0.46	0.00	0.00	0.03	0.03	0.03	0.03	0.00
antarctica	Median	0.38	0.00	0.43	0.07	0.03	0.02	0.63	0.00	0.03	0.10	0.10	0.10	0.10	0.00
	Upper	0.54	0.03	1.84	0.14	0.10	0.03	0.87	00.00	0.07	0.20	0.20	0.13	0.30	0.00
Gymnoscopelus	Lower	0.06	00.00	0.00	1.35	0.14	0.23	8.24	0.00	0.77	0.06	1.29	0.33	0.00	0.00
fraseri	Median	0.11	0.00	0.00	1.73	0.24	0.53	10.08	0.00	1.04	0.10	1.75	0.46	0.00	0.00
	Upper	0.21	00.00	0.00	2.27	0.37	0.70	11.93	00.00	1.34	0.16	2.48	0.57	0.00	0.00
Gymnoscopelus	Lower	0.17	00.00	0.27	0.83	0.30	0.10	6.91	0.03	2.96	0.36	5.58	0.17	0.03	0.00
nicholsi	Median	0.27	0.23	0.35	1.00	0.43	0.17	0.00	0.03	4.07	0.43	10.13	0.23	0.03	0.10
	Upper	0.30	0.23	0.44	1.17	0.60	0.20	11.75	0.07	5.01	0.54	13.05	0.31	0.07	0.10
Gymnoscopelus	Lower	0.03	00.00	0.03	0.23	0.00	0.03	0.79	0.00	0.17	0.03	0.13	0.13	0.00	0.00
braueri	Median	0.07	0.00	0.07	0.30	0.03	0.07	1.13	0.00	0.23	0.07	0.23	0.17	0.03	0.00
	Upper	0.10	0.03	0.07	0.40	0.03	0.13	1.47	0.00	0.33	0.10	0.38	0.23	0.07	0.03
Krefftichthys	Lower	0.07	00.00	0.00	0.77	2.82	0.63	0.37	00.00	0.00	0.00	4.85	0.00	0.00	0.00
anderssoni	Median	0.17	0.00	0.00	1.13	4.62	1.12	0.58	0.00	0.00	0.02	6.02	0.00	0.00	0.00
	Upper	0.83	00.00	0.00	1.67	6.97	1.74	0.80	0.00	0.04	0.07	7.57	0.00	00'0	0.00
Nannobrachium	Lower	0.00	00.00	0.00	0.30	0.03	0.10	0.07	0.00	0.03	0.20	0.37	0.10	0.00	0.00
achirus	Median	0.03	0.00	0.00	0.37	0.10	0.12	0.10	0.00	0.07	0.30	0.50	0.13	0.03	0.00
	Upper	0.07	00.00	0.00	0.47	0.10	0.17	0.13	0.00	0.10	0.33	0.63	0.17	0.07	0.00
Protomyctophum	Lower	0.13	00.00	0.00	0.40	0.07	7.48	1.00	0.00	0.00	0.03	0.53	0.00	0.00	0.00
tenisoni	Median	0.25	0.00	0.00	0.57	0.13	9.03	1.53	0.13	0.00	0.10	0.70	0.03	0.00	0.00
	Upper	0.40	00.00	0.00	0.70	0.21	10.50	1.90	0.13	0.07	0.17	0.83	0.10	00'0	0.00
Protomyctophum	Lower	0.00	00.00	0.00	0.17	0.00	0.00	5.23	0.00	0.17	0.27	2.15	0.00	0.00	0.00
bolini	Median	0.00	0.00	0.00	0.23	0.00	0.03	6.67	0.00	0.30	0.37	2.97	0.07	0.00	0.00
	Upper	0.03	00.00	0.00	0.30	0.03	0.10	8.59	0.00	0.50	0.53	3.43	0.13	00'0	0.00
Protomyctophum	Lower	0.73	0.00	0.00	3.47	0.17	5.76	2.06	0.00	0.00	0.03	4.38	0.13	0.00	0.00
choriodon	Median	0.93	0.00	0.00	4.28	0.30	7.53	6.12	0.00	0.07	0.07	6.07	0.23	0.00	0.00
	Ilnnar	1 78	000	000	11	0 11	10 50			~ ~ ~ ~		100		000	

:t of myctophid predation on the production of the key zooplankton taxa expressed as a percentage of daily production consumed ($\mu g \ C \ m^{-2} \ d^{-1}$) by	becies caught in the Scotia Sea during the study. Estimates represent the 25 th percentile (lower), median, and 75 th percentile (upper) values of	the data set. Dash denotes insufficient data (i.e. where 25 th percentile estimates were zero) to generate a confident estimate
Table 4. The impact of myctophid pr	each myctophid species caught in the Scotia Sea d	the data set. De

Myctophid species	Esti- mate	Themisto Euphau- gaudi- sia chaudii frigida	Euphau- sia frigida	Euphau- sia superba	Thysa- noessa spp.	Cala- noides acutus	Calanus simil- limus	Metri- dia spp.	<i>On-</i> <i>caea</i> spp.	Pleuro- Para- mamma euchaeta robusta spp.	<i>Para-</i> euchaeta spp.	Rhin- calanus gigas	Ostra- cods	Ptero- pods	Salps
Electrona carlsbergi	Lower Median Upper	0.00 0.16	0.00 - 0	0.00 -	0.00 0.32 -	0.00 0.00 0.38	0.00 0.18 -	0.00 0.01 0.71	0.00 0.01 6.93	0.00 0.00 2.93	0.01 0.13 7.61	0.02 0.80 -	0.00 0.00 0.04	0.00 0.00 0.03	0.00 0.10 -
Electrona antarctica	Lower Median Upper	0.01 2.75 -	0.00 - 0	0.00 2.26 -	0.00 0.82 -	0.00 0.02 0.45	0.00 0.07	0.00 0.03 0.69	0.00 0.00 0.00	0.00 0.00 2.36	0.00 0.27 6.46	0.00 - 06	0.00 0.02 0.22	0.00 0.00 0.09	0.00
<i>Gymnoscopelus</i> fraseri	Lower Median Upper	0.00 -	0.00 0	0.00 -	0.02 0.90 -	0.00 0.01 0.14	00.0	0.00 0.02 0.78	0.00 0.00 0.00	0.00 0.00 3.86	0.00 0.01 0.42	0.00 0.04 -	0.00 0.00 0.08	0.00 0.00 0.00	0.00
<i>Gymnoscopelus</i> <i>nicholsi</i>	Lower Median Upper	0.01 0.05	0.00 0.23	0.00 0.05	0.01 0.31 -	0.00 0.01 0.07	0.00 0.02 -	0.00 0.01 0.24	0.00 0.00 0.01	0.01 0.00 4.60	0.01 0.03 0.45	0.01 0.15 -	0.00 0.00 0.01	0.00 0.00 0.00	0.00 - 00. 0
Gymnoscopelus braueri	Lower Median Upper	0.00 0.24 -	0.00 - 000	0.00 0.18	0.00 1.86 -	0.00 0.01 0.11	0.00 0.14 -	0.00 0.03 0.86	0.00 0.00 0.00	0.00 0.01 8.70	0.00 0.09 2.38	0.00 0.07	0.00 0.02 0.28	0.00 0.00 0.02	0.00 - 00.0
Krefttichthys anderssoni	Lower Median Upper	0.00 0.52	0.00 	0.00 - 000	0.01 6.06 -	0.01 1.01 18.71	0.00 2.02	0.00 0.01 0.38	0.00 0.00 0.00	0.00 0.00 0.87	0.00 0.02 1.27	0.01 1.54 -	0.00 0.00 0.00	0.00 0.00 0.00	0.00 - 00.0
Nannobrachium achirus	Lower Median Upper	0.00 0.01	0.00 - 000	0.00 -	0.01 0.18 -	0.00 0.00 0.03	0.00 0.02	0.00 0.00 0.01	0.00 0.00 0.00	0.00 0.00 0.20	0.00 0.03 0.61	0.00 0.01	0.00 0.00 0.02	0.00 0.00 0.00	0.00 - 00.0
Protomyctophum tenisoni	Lower Median Upper	0.00 0.08	0.00 - 0	0.00 -	0.01 0.29 -	0.00 0.00 0.14	0.00 1.57 -	0.00 0.00 0.22	0.00 0.00 0.09	0.00 0.00 0.34	0.00 0.01 0.77	0.00 0.02	0.00 0.00 0.02	0.00 0.00 0.00	0.00 - 00.0
Protomyctophum bolini	Lower Median Upper	0.00 -	0.00 - 0	0.00 - 0	0.00 0.59 -	0.00 0.00 0.04	0.00 - 0.03	0.00 0.06 1.67	0.00 0.00 0.00	0.00 0.00 4.32	0.00 0.21 4.20	0.00 0.36 -	0.00 0.00 0.05	0.00 0.00 0.00	0.00
Protomyctophum choriodon	Lower Median Upper	0.02 0.12 -	0.00 	0.00 -	0.05 0.96 -	0.00 0.00 0.13	0.00 0.57 -	0.00 0.00 0.30	0.00 0.00 0.00	0.00 0.00 0.20	0.00 0.00 0.12	0.01 0.07 -	0.00 0.00 0.03	0.00 0.00 0.00	0.00
Total	Lower Median Upper	0.05 3.97 -	0.00 0.23 -	0.00 2.49 -	0.12 12.29 -	0.01 1.06 20.19	0.01 4.70 -	0.01 0.17 5.86	0.00 0.01 7.04	0.02 0.02 28.38	0.02 0.82 24.29	0.04 3.12 -	0.00 0.05 0.74	0.00 0.00 0.14	0.00 0.11 -

Table 5. Estimated total annual consumption of zooplankton biomass (t yr⁻¹) for the whole Scotia Sea. The 95% CI around these estimates reflect the level of variation in myctophid density observed during the study

Estimate	Themisto	Euphau-	Euphausia	Thysano-	Cala-	Calanus	Metridia	Oncaea	Pleuro-	Paraeu-	Rhin-	Ostra-	Ptero-	Salps
	gaudi-	sia	superba	essa	noides	similli-	spp.	spp.	mamma	chaeta	calanus	cods	pods	
	chaudii	frigida		spp.	acutus	snur			robusta	spp.	gigas			
T 05 0/ 0/	GOG AEE	1016	6127600	1 1 47 4 40	010 00	1 4 4 6 0	62010	LC	0600	01000	090916	100	C V	67 011
PUMPI A3 70	000	4010	070101C	1 14 <i>1</i> 440	010000	14 4J9	01000	10	0000	01067	340 900	100	40	110 10
Mean	2245883	14120	16808493	3754095	110723	47305	176078	121	28136	95922	$1\ 135\ 180$	1083	140	220222
Upper 95 %	3805311	23924	28479466	6360750	187604	80152	298338	206	47672	162525	1923393	1835	237	373133

production was also relatively high (Table 4), with a best estimate of consumption of ~4% of Themisto gau*dichaudii* daily production and ~12% of Thysanoessa spp. daily production. Themisto gaudichaudii and Thysanoessa spp. were impacted most by Electrona antarctica and K. anderssoni, respectively. Myctophids also consumed around 2% of Euphausia superba daily production, where Electra antarctica had the highest impact on this prey species. The impact of myctophids on salps and ostracods accounted for up to 0.1% of these prey species' daily production, but their impact on pteropods was negligible.

Annual consumption of zooplankton

Estimates of the total annual consumption of zooplankton across the whole Scotia Sea were dominated by the diet of Electrona antarctica, the most common myctophid species. Our data suggest that the main taxa consumed by myctophids were Euphausia superba, Thysanoessa spp. and Themisto gaudichaudii, with 16.8, 3.8, and 2.2 Mt yr⁻¹ of these species being eaten, respectively (Table 5). The estimated annual consumption of all key copepods was around 1.5 Mt yr⁻¹, where *R. gigas* was the most preved upon copepod species (1.1 Mt yr^{-1}). The estimated consumption of the other main macrozooplankton taxa, such as salps and ostracods, was $< 0.5 \text{ Mt yr}^{-1}$ (Table 5).

Diet comparisons between species

Hierarchical cluster analysis produced 5 clusters at the 60% similarity level, although 2 of these clusters were comprised of single species (Cluster 1: *Electrona antarctica*, Cluster 2: *N. achirus*) (Fig. 7). *G. braueri*, *G. fraseri*, *G. nicholsi* and *P. bolini* were grouped in Cluster 3, where the copepod *Metridia* spp. was the predominant prey species of this group (36%) (Table 6). Cluster 4 contained E. carlsbergi and K. anderssoni in which diets were dominated by *R. gigas* (54%), and P. tenisoni and P. choriodon were grouped in Cluster 5 that was dominated by the consumption of C. simil*limus* (25%) and *Thysanoessa* spp. (22%). There was substantial overlap between the composite length-frequency distributions of fish within each cluster dominated by copepod consumption, indicating that this clustering reflected differences in feeding selectivity rather than size-related differences in feeding patterns (e.g. the median fish size for clusters 3, 4 and 5 was 72, 73 and 64 mm SL, respectively). However, there was also a high degree of overlap in the overall diets of Clusters 3, 4 and 5, as R. gigas, Metridia spp. and Thysanoessa spp. all occurred within the top 3 to 4 most consumed prey species in each cluster, contributing a total of ~57-69% to the groupings (Table 6). Themisto qaudichaudii and C. simillimus were also common to the 3 clusters, suggesting that other, less dominant species were important contributors to these clusters. Most notably, P. robusta, ostracods and E. superba were unique in the grouping of Cluster 3 (contributing 13%, collectively), as were unidentified euphausiids and unidentified crustaceans in the grouping of Cluster 4 (contributing ~8%, collectively).

DISCUSSION

The present study provides a comprehensive analysis of myctophid diets and their predatory impact on zooplankton communities in the Southern Ocean, and represents one of the most detailed studies undertaken to date on the trophic role of myctophids in any oceanic region. These results must be placed within a context of the associated sampling issues inherent with net-based surveys of mesopelagic fish

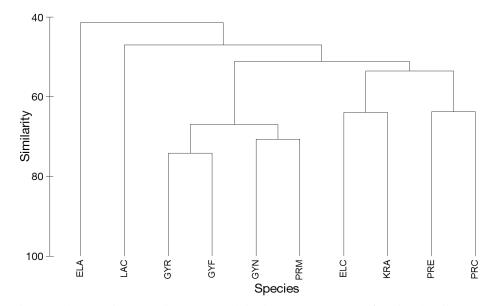


Fig. 7. Cluster diagram of a Bray-Curtis similarity matrix of the dietary composition (%IRI data for all prey items) of the 10 myctophid species caught in the Scotia Sea. Cluster 1 = *Electrona antarctica* (ELA), Cluster 2 = *Nannobrachium achirus* (LAC), Cluster 3 = *Gymnoscopelus braueri* (GYR), *G. fraseri* (GYF), *G. nicholsi* GYN), *Protomyctophum bolini* (PRM), Cluster 4 = *E. carlsbergi* (ELC), *Krefftichthys anderssoni* (KRA), Cluster 5 = *P. tenisoni* (PRE), *P. choriodon* (PRC). The average similarity in the diets of the grouped myctophid species is also shown

Table 6. Results of SIMPER analysis showing percentage contributions of prey species to the myctophid groupings identified by agglomerative hierarchical cluster analysis (see Fig. 7). The average similarity in the diets of the grouped myctophid species is also shown

Cluster group/ Myctophid species	Prey species	Average abundance	Percentage contribution	Cumulative percentage
3				
Gymnoscopelus braueri	Average similarity: 68.82			
Gymnoscopelus fraseri	Metridia spp.	6.89	35.59	35.59
Gymnoscopelus nicholsi	Rhincalanus gigas	4.28	18.10	53.68
Protomyctophum bolini	Thysanoessa spp.	3.46	14.44	68.12
1 1	Pleuromamma robusta	2.15	8.84	76.96
	Paraeuchaeta spp.	1.11	3.79	80.75
	Themisto gaudichaudii	1.24	3.70	84.45
	Ostracods	0.92	2.49	86.94
	Calanus simillimus	0.57	1.92	88.85
	Euphausia superba	1.52	1.91	90.77
4				
Electrona carlsbergi	Average similarity: 64.01			
Krefftichthys anderssoni	Rhincalanus gigas	8.59	54.02	54.02
-	Metridia spp.	1.79	7.95	61.97
	Thysanoessa spp.	2.31	7.24	69.21
	Themisto gaudichaudii	1.05	6.84	76.05
	Unidentified euphausiids	0.84	5.52	81.57
	Calanus simillimus	1.36	5.22	86.80
	Paraeuchaeta spp.	0.39	2.61	89.41
_	Unidentified crustaceans	0.46	2.27	91.68
5				
Protomyctophum tenisoni	Average similarity: 63.77			
Protomyctophum choriodon	Calanus simillimus	6.16	25.43	25.43
	Thysanoessa spp.	4.82	21.83	47.26
	Metridia spp.	3.02	17.91	65.16
	Rhincalanus gigas	3.76	17.02	82.18
	Themisto gaudichaudii	1.66	6.34	88.52
	Calanoides acutus	0.41	2.85	91.37

and zooplankton. Such issues include active net avoidance by myctophids and the patchy nature of both myctophid and zooplankton aggregations which may, for example, impact estimates of abundance averaged over relatively broad spatial and temporal scales. Indeed, recent acoustic studies have reported that the abundance of mesopelagic fishes may be at least an order of magnitude greater than previously assumed from net survey data, suggesting that the role of mesopelagic fish in oceanic ecosystems may be underestimated in net-based trophodynamics studies (Irigoien et al. 2014). A further consideration is that seasonal variations were not resolved in the study since the data were integrated over the 3 seasons. Although this approach does not provide a seasonal synopsis, it does provide a more accurate view of the average situation during the productive months because the data are more representative of myctophid diets over the longer-term.

Niche partitioning

The results of our study show that myctophids consume a range of mesozooplankton and macrozooplankton, particularly copepods, euphausiids and amphipods, which is consistent with studies carried out in other parts of the Southern Ocean (Naumov et al. 1981, Kozlov & Tarverdiyeva 1989, Gerasimova 1990, Pakhomov et al. 1996, Gaskett et al. 2001, Pusch et al. 2004, Shreeve et al. 2009) and on the myctophid community elsewhere (Hopkins & Gartner 1992, Williams et al. 2001, Suntsov & Brodeur 2008, Pepin 2013, Tanaka et al. 2013).

Resource partitioning is key to minimising interspecific competition and enabling the coexistence of species in a region (Schoener 1974), and such partitioning has been demonstrated in highly diverse low latitude myctophid communities (Clarke 1980, Hopkins & Gartner 1992) and at high and temperate latitudes (Watanabe et al. 2002, Sassa & Kawaguchi 2005, Shreeve et al. 2009, Cherel et al. 2010). However, species tend to exhibit a high degree of overlap in their diets in high latitude regions and it has been suggested that inter-species food competition is avoided because of high regional food availability (Pakhomov et al. 1996). In the present study, there was evidence of dietary segregation and specialisation for some myctophid species that is linked, in part, to horizontal and vertical distribution and individual size (see Shreeve et al. 2009 for an overview of the size ranges of myctophids and their prey species). Electrona antarctica, for example, occurred mostly in

the sea-ice sectors and, unlike the other myctophids, had a diet dominated by Euphausia superba and Themisto gaudichaudii. Also, Nannobrachium achirus was the largest species encountered and was caught predominantly below 400 m, and had a diet that included substantial amounts of deep-water amphipods and small fish. Thus these species appear to have different niches from the other myctophids. Furthermore, similarity analysis identified 3 clusters that were dominated by copepod consumers, but preferential selection of certain copepod species appeared to separate their niches. Of the predominantly smaller myctophid species, Electrona carlsbergi and Krefftichthys anderssoni, which had different depth distributions, targeted mostly Rhincalanus gigas, whilst Protomyctophum tenisoni and P. choriodon favoured Calanus simillimus. In contrast, the group comprising the 3 larger-sized Gymnoscopelus species and P. bolini preyed mostly upon Metridia spp. These results are broadly consistent with concurrent studies using trophic biomarkers, such as stable isotopes and fatty acids, which provide complimentary time-integrated synopses of predator diets and habitats (Stowasser et al. 2012, Tarling et al. 2012). Similar niche partitioning was also observed for most of the studied myctophid species at lower latitudes (Kerguelen Islands, southern Indian Ocean) using these techniques, where strong segregation between the genera Electrona, Gymnoscopelus and Protomyctophum was observed (Cherel et al. 2010). However, there was also a high degree of overlap in the overall diets of all myctophids in our study, with R. gigas, Metridia spp., and Thysanoessa spp. preyed upon substantially by all species. This suggests that inter-specific competition for these prey items may be reduced in the Scotia Sea because of their high availability in the water column (Pakhomov et al. 1996).

Prey selection

The overall distribution patterns of *K. anderssoni* and *E. carlsbergi* broadly matched that of its main prey, *R. gigas*, as did the distribution patterns of *P. tenisoni* and *P. choriodon* and their preferred prey species, *C. simillimus*. These myctophids and prey items occurred mostly in the northern regions of the Scotia Sea and were less abundant in regions south of the South Antarctic Circumpolar Current Front (Ward et al. 2012, Saunders et al. 2014). Similarly, *E. antarctica* occurred in highest abundance in the SIZ where its main prey species, *Euphausia superba*, was

also most abundant. The trend was less obvious for the *Gymnoscopelus* species and *P. bolini*, however, the abundance of these species was generally higher in the northern regions, which broadly matched the distribution pattern of *Metridia* spp. in the region.

The most abundant copepod species in the region, the small copepods Oithona spp. and Ctenocalanus spp., were seldom preyed upon by any of the myctophids. These prey species may either be too small to be retained by the gill rakers or too unprofitable to exploit (Shreeve et al. 2009). The exception to this was the consumption of Oncaea spp. by Electrona carlsbergi, which suggests that myctophids are capable of retaining small copepods, but there is a high degree of prey selectivity. Further evidence of prey selectivity within the copepod community was apparent, as all myctophids tended to select the older copepodite stages, particularly CVI females that are generally considered to be more lipid-rich than other stages (Hagen & Schnack-Schiel 1996, Shreeve et al. 2009). A relatively high degree of selectivity was also apparent in the macrozooplankton component of the prey field. Myctophids appeared to select the euphausiid Thysanoessa spp. in preference to Euphausia frigida, which is a similar-sized euphausiid and had a similar depth distribution and abundance in the Scotia Sea. Likewise, E. triacantha, a euphausiid similar in size to E. superba, was seldom consumed by any of the larger myctophid species even though its abundance was relatively high in the region (Saunders et al. 2014). These euphausiids have comparable energy content in terms of total lipids, although there are some differences in component lipid composition, which may be important in resource selectivity by myctophids (Reinhardt & Vanvleet 1986, Ruck et al. 2014). Differences in euphausiid aggregation and escape behaviour may also be an important factor in myctophid predation on these organisms (Daly & Macaulay 1988, Brierley et al. 1998).

Body size effects on diet

The results showed that myctophid size was an important determinant of diet, as larger-sized fish clearly preyed upon a broader range of prey taxa and took larger prey items, such as *E. superba* and *Themisto gaudichaudii*. Adult Antarctic krill are probably one of the largest prey species that can be preyed upon by myctophids, and as a consequence, were only consumed by the largest myctophids. An increase in trophic level with increasing myctophid

size was also detected during stable isotope analyses (Tarling et al. 2012). The ability to prey upon largersized organisms is most likely controlled by gape size and body size such that only the larger-sized myctophids are able to capture and consume these animals (Karpouzi & Stergiou 2003).

Food-web implications

The significance of krill in the diet of Southern Ocean myctophids has been the source of debate in the scientific literature (Williams 1985, Lancraft et al. 1989, Pakhomov et al. 1996, Pusch et al. 2004). Our results support the concept that the myctophids, particularly the small species, provide an important krillindependent link between secondary production and higher predators (Murphy et al. 2007b). Myctophid predation accounted for approximately 2% of the daily krill productivity in the Scotia Sea, with Electrona antarctica consuming the majority of this productivity. Whilst this level of predation impact is relatively low, it is still indicative of major quantities of krill biomass being consumed by myctophids in the Scotia Sea on an annual basis. Collins et al. (2012) estimated that zooplankton consumption by myctophids in the Scotia Sea was approx. 25 Mt y⁻¹. We used our diet data to partition this consumption estimate amongst prey taxa to estimate the cumulative impact of myctophid predation on their prey biomass throughout the year. The data suggests that myctophids in the Scotia Sea consume around 17 ± 6 (mean ± SD) Mt of Euphausia superba per year, supporting the notion that large myctophids are possibly the main consumers of this species in the region (Lancraft et al. 1989, Pusch et al. 2004, Hill et al. 2007).

Myctophids consume both larval and adult stages of krill. However, there are currently no independent estimates of krill biomass or production that encompass all the developmental stages of krill that myctophids consume. Our estimate of krill density (637 ind. m⁻²), which encompassed larval and post-larval stages, is higher than that reported for post-larval krill in the Scotia Sea (16–256 ind. m^{-2}), suggesting that krill biomass and production are also higher than that estimated in the region (Hewitt et al. 2004, Atkinson et al. 2009). Determination of biomass of the whole life-cycle of krill, together with the predatory impact of myctophids on the specific developmental stages, is a necessary further step towards understanding high latitude Southern Ocean food webs and ecosystem function.

Our result showed that myctophid predation on the daily productivity of Thysanoessa species was high. These smaller euphausiids comprised a substantial proportion of the diets of all myctophids, particularly *K. anderssoni*, indicating that they have a key role in the Southern Ocean ecosystem. Thysanoessa species, such as T. macrura and T. vicini, are the most consistently found euphausiid in Antarctic waters (Nordhausen 1994, Boltovskoy 1999, Haraldsson & Siegel 2014) and often exceed E. superba in abundance in some regions (Daly & Macaulay 1988). These smaller euphausiids are an important dietary component of penguins, seabirds and mackerel ice fish (Brown & Klages 1987, Kock et al. 1994, Main et al. 2009, Pichegru et al. 2011), but information on the trophic role of Southern Ocean Thysanoessa species within Antarctic ecosystems is limited. Given their importance in the diet of Southern Ocean myctophids, resolving the trophodynamics of Thysanoessa species in this region is an important part of predicting how myctophids will respond in this rapidly changing environment (Flores et al. 2012). Myctophids also preyed upon a substantial proportion of the daily productivity of Themisto gaudichaudii, and the ecological importance of this species was highlighted by Shreeve et al. (2009) and Bocher et al. (2001), for the northern Scotia Sea and sub-Antarctic latitudes, respectively.

Even though copepods were the main prey item of myctophids, myctophid predation had relatively little impact on the productivity of most copepod species in the Scotia Sea region. The exceptions were the larger copepods R. gigas and C. simillimus of which myctophids consumed between 3-5% of their daily productivity. The myctophid species that had the greatest impact on these copepods was K. anderssoni due to its relatively high abundance in the northern Scotia Sea. This predominant APF species was one of the smallest myctophids encountered on the surveys, but it also consumed the greatest proportions of Thysanoessa spp. productivity and was the secondhighest consumer of Themisto gaudichaudii productivity. K. anderssoni is the primary prey of king penguins (Olsson & North 1997, Bost et al. 2002, Cherel et al. 2002) and an important dietary component of other predators (Rodhouse et al. 1992, Casaux et al. 1998, Deagle et al. 2008, Cherel et al. 2010), indicating that it has an important role in the operation of the Scotia Sea ecosystem, despite it being a species that typically resides in waters of the APF. Given that K. and erssoni and the other sub-Antarctic species (e.g. Electrona carlsbergi, Gymnoscopelus fraseri and P. tenisoni) are possibly expatriates, or seasonal

migrants, in the Scotia Sea (Hulley 1981), it is clear that further studies are warranted in regions north of the APF in order to gain better insight into the trophodynamics and ecology of these myctophids, which are likely to have a direct bearing on ecosystem dynamics in regions at higher latitudes, such as the Scotia Sea.

In conclusion, the myctophid community in the Scotia Sea maintained a large dietary breadth, but there was some evidence of dietary segregation between species, related to their horizontal distribution, inter-specific variations in body size, variations in vertical migratory behaviour and depth selection. These differences potentially minimise the impact of seasonal changes in the prey field and minimise competition and the exhaustion of any one particular food resource. There is likely to be a considerable flux of biomass through the Scotia Sea myctophid community, which appears largely independent of Antarctic krill. This indicates that the myctophid community is a robust component of the Southern Ocean mesopelagic system that is able to exploit a wide range of food resources and provide a major link between lower and upper trophic levels in the Southern Ocean.

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