

Predatory role of the commander squid *Berryteuthis magister* in the eastern Bering Sea: insights from stable isotopes and food habits

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ABSTRACT: Squid are an important component of many marine food webs, and they can impact other species through predation and competition. However, quantifying their influence on other food web components requires knowledge of their trophic position and trophic ontogeny, which are unknown in many ecosystems. The eastern Bering Sea (EBS) is a highly productive region that supports large commercial fisheries, and a modicum of knowledge exists on the ecological role of squid in this region. We combined stomach content and stable isotope analyses of muscle tissue ($\delta^{15}\text{N}$ and $\delta^{13}\text{C}$) to identify the feeding ecology of the commander squid *Berryteuthis magister* in the EBS continental slope ecosystem. We also use a novel methodology to elucidate potential finer-scale variation in squid trophic ecology by reconstructing feeding chronologies of individual *B. magister* from concentric eye lens layers. Our analyses indicate that the position of *B. magister* in the EBS food web increases by approximately 1 trophic level between juvenile and adult stages. Also, in contrast to many squid species, we found that predation by *B. magister* is not constrained by prey body size and that *B. magister* are more likely to share prey resources with commercially valuable fishes, particularly walleye pollock *Theragra chalcogramma*, than to prey upon their juvenile stages. Further, the reconstructed feeding chronologies indicate substantial variability in squid feeding patterns that are not captured on the time scales of the conventional analyses. Together, the findings of this study contribute to a better understanding of the ecological role of *B. magister* and the trophic linkages and energy flow within the EBS food web.

KEY WORDS: Squid · Trophic interactions · Stable isotopes · Prey size spectra · Feeding chronology · *Berryteuthis magister* · Eastern Bering Sea

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INTRODUCTION

Squid occupy a central position in marine food webs. They are a valuable prey resource for fishes, seabirds, and marine mammals and are also voracious predators of crustaceans, squid, and fishes (Clarke 1996, Croxall & Prince 1996, Klages 1996, Smale 1996). Because of their high feeding rates and generalist feeding strategy, squid can potentially exert trophodynamic control on the recruitment of the early life stages of fishes (Dawe 1988, Rodhouse & Nigmatullin 1996, Hunsicker

& Essington 2008). Moreover, their importance in food webs may be changing in response to fisheries-induced alterations to trophic structures and climate change (e.g. Humboldt squid *Dosidicus gigas*, Field et al. 2007, Zeidberg & Robinson 2007). An understanding of the potential impacts of these changes on marine systems is presently hampered by gaps in our understanding of the ecological role of squid.

As in many ecosystems, we have a limited understanding of the trophic roles of squid in the eastern Bering Sea (EBS). The EBS is a highly productive

ecosystem that supports some of the world's largest commercial fisheries as well as the production of species of conservation interest (National Research Council, NRC 1996). Over the past few decades, the structure of this ecosystem has shifted in response to human activities and changing environmental conditions (NRC 1996, Alexander 1999, Springer 1999). To understand how anthropogenic and natural forcing may impact the trophic linkages and energy flow within this system, much effort has focused on identifying the trophic relationships of commercial and conservation species. However, the ecological roles of non-target species, such as mesopelagic squid, have been understudied.

The goal of our work is to provide a better understanding of the trophic ecology of the commander squid *Berryteuthis magister* (Cephalopoda, Gonatidae), by examining its predatory role in the EBS continental slope ecosystem. *B. magister* is the most abundant squid species in the EBS (Radchenko 1992, Sinclair & Stabeno 2002), where it inhabits the outer edge of the continental shelf and slope, reaching depths of 1500 m (Nesis 1998). Juveniles mostly occur in midwater at mesopelagic depths, while adults mainly inhabit the bottom layers in the upper bathyal zone (Nesis 1997, 1998, Arkhipkin et al. 1998): both life stages undertake diel vertical migrations with night ascent (Gorbatenko et al. 1995, Arkhipkin et al. 1998). Previous work indicates that *B. magister* is a key component in the food webs of the western and central Bering Sea (Frost & Lowry 1981, Lowry et al. 1982, Radchenko 1992, Nesis 1997, 1998, Aydin et al. 2007). Food web models of EBS also indicate that squid are an important link in the transfer of energy from lower to higher trophic levels (Aydin et al. 2007). Further, squid can be important predators or competitors of juvenile stages of gadoids and other high trophic level fishes (Hunsicker & Essington 2006, 2008, Field et al. 2007), and commercially valuable gadoid fishes, particularly walleye pollock *Theragra chalcogramma*, dominate the fish biomass in the EBS.

Knowledge of the trophic ontogeny of *Berryteuthis magister* and the size-structure of their feeding relationships is paramount for determining their ecological role. Body size dictates predator–prey relationships in many marine systems (Lundvall et al. 1999, Jennings et al. 2001, Shin & Cury 2004), and squid exhibit ontogenetic, size-dependent patterns in predation (Collins & Pierce 1996, Lordan et al. 1998, Hunsicker & Essington 2006). By identifying the prey size spectra of *B. magister*, we can determine the prey species and sizes that are vulnerable to squid predation. This information is valuable for assessing the potential consequences of *B. magister* feeding habits and for exploring how the strength of their trophic interactions may change if

predator and (or) prey body sizes are altered by ecosystem perturbations (Woodward et al. 2005).

Identifying the feeding habits of squid is challenging owing to their feeding behavior and rapid digestion of prey (Nixon 1987, Dawe et al. 1997). Squid masticate their food and therefore may only consume a portion of their prey at feeding. Here, we attempted to overcome those challenges by using multiple pathways to characterize the trophic role of *Berryteuthis magister*. We used traditional stomach content analyses to identify prey taxa, estimate the prey size spectra of *B. magister*, and determine how prey composition changes with squid ontogeny. Stomach content analysis provides high detail on prey contents; however, due to squid feeding attributes, it does not provide a good assessment of their trophic positions (i.e. the average levels, relative to primary producers, at which an organism feeds). Also, this analysis only provides information on an animal's last feeding event and reveals no information on the feeding habits of sampled predators with empty or unidentifiable stomach contents. In contrast, stable isotope ratios ($^{15}\text{N}:$ ^{14}N and $^{13}\text{C}:$ ^{12}C) of animal tissues can provide more precise estimates of long-term average trophic position. These ratios provide less detail on digested prey items, but they provide a measure of feeding that is integrated over several weeks or months (Ruiz-Cooley et al. 2006). Stable isotope analyses have been a valuable tool for investigating the trophic ecology of squid (Takai et al. 2000, Cheral & Hobson 2005, Ruiz-Cooley et al. 2006, 2010, Parry 2008, Cheral et al. 2009). We used stable isotopes as a complementary method to identify ontogenetic shifts in *B. magister* trophic position and to evaluate their trophic position with respect to marine fishes inhabiting the EBS.

In addition to these 2 conventional approaches, we further explored whether reconstructed feeding chronologies from concentric eye lens layers can be used as an alternative method to evaluate the ecological role of *Berryteuthis magister*. Feeding chronologies from squid hard parts, such as eye lenses (Parry 2003) and gladii (Ruiz-Cooley et al. 2010) are a valuable tool because they offer several measures of individual feeding events that are integrated over time scales intermediate to stomach content and stable isotope analysis of muscle tissue. For instance, fish lens cells are nearly ametabolic because most of the cells (all but those in the youngest, differentiating layer) lack the cellular functions of protein synthesis and catabolism (Wistow & Piatigorsky 1988, Nicol 1989, Horwitz 1992, Dove 1999). The same is likely to be true for squid eye lenses (see West et al. 1995); thus, with little to no protein turnover in the lens cells, isotope signatures laid down in the lens layers may essentially be locked in place during their formation (Parry 2003). From a sam-

pling standpoint, reconstructed feeding chronologies could improve our ability to evaluate the feeding ecology of squid life stages or species that are difficult to capture with traditional sampling gear. Similarly, feeding chronologies would allow us to explore the trophic ontogeny of squid with far fewer samples than would be needed for stomach content analysis or muscle tissue stable isotope analysis.

In this study, we addressed the following questions to better understand the role of *Berryteuthis magister* in the EBS continental slope ecosystem. Do *B. magister* exhibit ontogenetic shifts in prey composition and trophic position? Are the predation patterns of *B. magister* dictated by size constraints? Is there evidence of predation by *B. magister* on higher trophic level fishes, including species of commercial value? Are there variations in *B. magister* trophic patterns revealed by feeding chronologies that are not captured by the conventional analyses of predator food habits? Our findings provide a strong foundation for identifying the potential influence of *B. magister* on other nekton through direct and indirect food web pathways.

MATERIALS AND METHODS

Sample collection. *Berryteuthis magister* were collected near the edge of the continental shelf and slope in the EBS (Fig. 1). Squid were sampled during summer and winter seasons with the intention of collecting a wide size range to account for seasonal and ontogenetic shifts in diet composition. Summer samples were collected through research surveys conducted by NOAA Fisheries, Alaska Fisheries Science Center (NOAA - AFSC; June to July 2004) and the University of Washington (August 2007). The summer samples were collected mainly from daytime bottom trawls (cod end mesh size 13 mm) at discrete depths ranging from 330 to 755 m

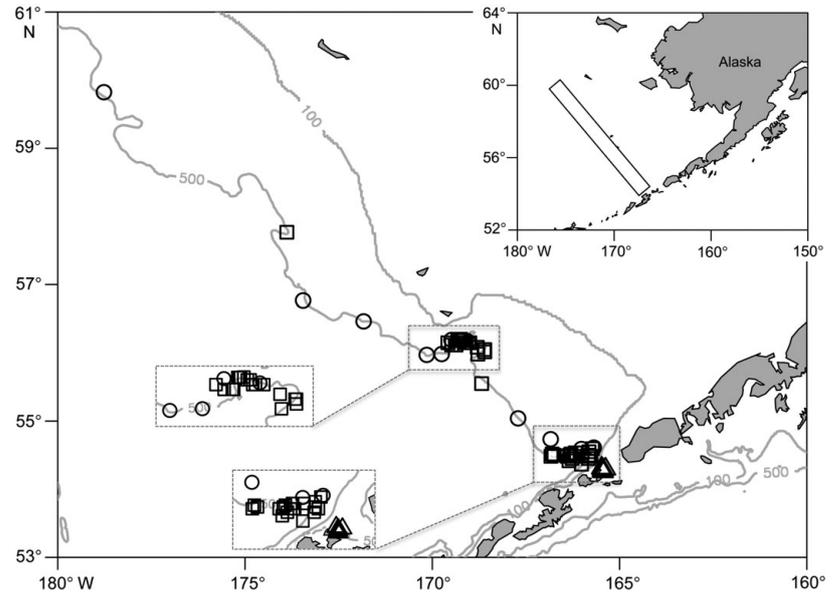


Fig. 1. Capture locations of *Berryteuthis magister* in the eastern Bering Sea. (O) Summer 2004, (□) winter 2007, and (Δ) summer 2007. The 100 and 500 m depth contours are shown. The insets (dotted lines) provide a closer view of the multiple capture locations that are in proximity to each other. The rectangle in the inset map of Alaska indicates the region of the capture locations within the eastern Bering Sea

Table 1. *Berryteuthis magister*. Mean capture depths and body sizes analyzed for stomach contents (n = sample size). The percent frequency of occurrence of *B. magister* prey items is shown by season and year, and all years combined. The scientific names of prey identified from the stomach contents of *B. magister* and the numbers of otoliths and beaks identified to each prey are also shown by season and year

	All years n = 697	Summer 2004 n = 107	Winter 2007 n = 336	Summer 2007 n = 254
Capture depth (m)	394 ± 103	393 ± 92	466 ± 90	321 ± 61
Dorsal mantle length (mm)	219 ± 97	172 ± 74	227 ± 127	230 ± 47
Body mass (g)	360 ± 216	243 ± 249	338 ± 210	389 ± 192
Frequency of occurrence (%):				
Fish	40.75	20.56	42.56	46.85
Squid	19.94	17.76	16.37	25.59
Crustacean	7.32	14.95	2.68	10.24
Empty	30.70	34.58	34.52	24.02
Unidentifiable prey only	13.20	19.63	8.63	16.54
Number of identified hard parts:				
Fishes (otoliths)				
<i>Leuroglossus schmidti</i>	309	3	214	92
<i>Stenobranchius leucopsarus</i>	7	2	0	5
<i>Lampanyctus jordani</i>	2	0	2	0
<i>Benthabella dentata</i>	2	2	0	0
Post-larval liparid	1	1	0	0
Cyclopteridae sp.	1	0	0	1
<i>Theragra chalcogramma</i>	1	1	0	0
Squid (beaks)				
<i>Berryteuthis magister</i>	17	2	12	3
<i>Gonatopsis borealis</i>	4	0	4	0
<i>B. magister</i> / <i>G. borealis</i>	36	0	17	19
<i>Gonatus</i> sp.	46	30	11	5
<i>Gonatus onyx</i>	9	0	0	9
<i>Gonatus madokai</i> / <i>Gonatus middendorffi</i>	6	1	4	1

in summer 2004 and 200 to 400 m in summer 2007 (Table 1). The winter samples were collected through the North Pacific Groundfish Observer Monitoring Program (January to March 2007). These squid were randomly sampled by observers from pelagic trawls primarily targeting pollock (cod end mesh size ~75 to 115 mm) and fishing at discrete depths ranging from 350 to 985 m (Table 1). All samples were frozen and stored at -20°C after collection. Once defrosted, squid dorsal mantle lengths were recorded to the nearest 1 mm, and total wet weights were measured to the nearest 0.1 g.

General diet analysis. Approximately 700 *Berryteuthis magister* collected during summer 2004, summer 2007, and winter 2007 were dissected for stomach content analysis (Table 1). Prey items of *B. magister* were identified from undigested hard parts found in squid stomachs. Digested fish were identified from otoliths, scales, and bones, and squid prey were identified from beaks, gladii, hooks, and statoliths (Vinogradov & Noskov 1979, Santos & Haimovici 1998, Hunsicker & Essington 2006). Crustaceans were identified by eyes, appendages, mandibles, and exoskeletons (Karpov & Cailliet 1978, Santos & Haimovici 1998). Prey items were identified using a dissection microscope and were initially grouped by visual identification of hard parts and soft tissues as fish, squid, crustaceans, or unknown. We recorded the wet weights of squid stomach masses and individual prey items. We also noted freshly digested prey items and excluded them from our analyses to omit bias due to predation inside trawl nets. Prey hard parts were stored dry or in 80% ethanol for subsequent taxonomic classification. The diet composition of *B. magister* was quantified by calculating the percent frequency of occurrence of each prey item in the squid stomach contents. Squid masticate their food; thus it is nearly impossible to quantify their prey composition by percent number or mass.

We used logistic regression analysis to detect ontogenetic shifts in the diet of *Berryteuthis magister*. Separate analyses were conducted to gauge the influence of body size on the frequency of occurrence of prey categorized as fish, squid, or crustacean. This analysis relates the probability of occurrence of a specific prey item (fish, squid or crustaceans) to the mantle lengths of *B. magister* collected in the summer and winter seasons. We evaluated the ontogenetic shifts in diets during both seasons by calculating the 95% confidence intervals of the slope. We then repeated this analysis to examine whether *B. magister* might also exhibit dietary shifts related to capture depth or location. We simply replaced the explanatory variable with capture depth and latitude.

Prey size spectrum. Fish and squid consumed by *Berryteuthis magister* during summers of 2004 and 2007 and winter 2007 were identified to lowest possi-

ble taxon by comparing the morphological features and size ratios of digested fish otoliths and squid beaks to those in the reference collection of the National Marine Mammal Laboratory (NMML - NOAA - AFSC). The sizes of fish and squid prey were then estimated using species-specific otolith length to body length relationships and lower rostral beak length (LRL) to mantle length relationships, respectively (Zeppelin et al. 2004, Gudmundson et al. 2006, W. Walker unpubl. data). Note that the beaks of *B. magister* and *Gonatopsis borealis* with LRL of <1 mm are nearly indistinguishable. Similarly, the small beaks (LRL <1 mm) of *Gonatus* species as well as *Gonatus middendorffi* and *G. madokai* are difficult to discern from each other. Thus, beaks with LRLs <1 mm were identified to coarser taxonomic groups, i.e. *B. magister*/*G. borealis*, *Gonatus* sp., or *G. middendorffi*/*G. madoakai*.

We used quantile regression to analyze the relationship between *Berryteuthis magister* mantle lengths and their prey sizes. The 10th and 90th quantiles were used as the upper and lower bound regression quantiles in this analysis (Rogers 1992, Scharf et al. 1998). This technique has been used previously to estimate the relationship between squid body size and the minimum or maximum sizes of their prey (Hunsicker & Essington 2006). Further, we used the Student's *t*-test to determine if there was a significant difference in upper bound regression lines of fish and squid prey. A difference in slope values may indicate that maximum prey size is dependent upon the prey type (Scharf et al. 2000, Juanes 2003).

Stable isotope analyses. To determine the trophic position of *Berryteuthis magister* in the EBS, we analyzed $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures of squid mantle muscle tissue. The muscle tissue of squid reflects predator feeding habits (Stowasser et al. 2006), and mantle tissue has been used to quantify the relative trophic position of squid in past studies (Takai et al. 2000, Parry 2008). A small portion ($\sim 13\text{ mm}^3$) of tissue was excised from the mantles of 50 squid caught in summer 2004 and 50 squid collected in winter 2007. We did not analyze the isotope signatures of squid collected during summer 2007. The tissue samples were dried at 64°C , ground with mortar and pestle into a homogeneous powder, and weighed according to the guidelines of the Colorado Plateau Stable Isotope Laboratory (CPISL; www.mpcer.nau.edu/isotopelab/). Lipids were not removed from squid tissues prior to the stable isotope analysis. Lipids do not contain nitrogen and thus do not affect nitrogen isotope signatures, though variations in the lipid content within tissues can significantly influence $\delta^{13}\text{C}$ values and hence bias the interpretation of the carbon isotope results (Kiljunen et al. 2006, Sweeting et al. 2006). However, the C:N ratios of the squid tissues analyzed in this study ranged

between 3 and 3.5, indicating that the lipid content in these tissues was quite low (Kiljunen et al. 2006). For that reason, we expect that the $\delta^{13}\text{C}$ values of *B. magister* muscle tissue would not be confounded by individual variation in lipid densities. The isotope analyses were conducted at the CPISL and were carried out in continuous-flow mode using a Thermo-Finnigan Delta^{plus} Advantage gas isotope-ratio mass spectrometer interfaced with a Costech Analytical ECS4010 elemental analyzer. Isotopic ratios are expressed using standard \times notation, where $\delta^{15}\text{N}$ or $\delta^{13}\text{C} = [(R_{\text{sample}} / R_{\text{standard}}) - 1] \times 1000$ and R is the ratio of $^{15}\text{N}:^{14}\text{N}$ or $^{13}\text{C}:^{12}\text{C}$ in the sample and standards, respectively. Atmospheric nitrogen and Pee Dee Belemnite were used as nitrogen and carbon standards, respectively. The data were normalized using 4 internationally accepted isotope reference standards (IAEA CH6, CH7, N1, and N2). External precision on laboratory standards was $\pm 0.20\%$ or better for $\delta^{15}\text{N}$ and $\pm 0.10\%$ or better for $\delta^{13}\text{C}$. We used the Spearman rank correlation test to determine whether the $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures of *B. magister* muscle tissue differed by capture depth or location. Regression analysis was also done to evaluate the relationship between squid mantle lengths and the isotopic signatures of muscle tissues.

To evaluate the trophic level of *Berryteuthis magister* within a broader ecosystem context, we compared $\delta^{15}\text{N}$ signatures of juvenile (<120 mm dorsal mantle length) and adult squid (≥ 120 mm, i.e. size at maturation; Nesis 1997) collected in summer 2004 to the $\delta^{15}\text{N}$ signatures of the dorsal white muscle tissue of commercially important fishes inhabiting the eastern Bering Sea (C. Kurlle, E. Sinclair, A. Edwards, C. Gudmundson unpubl. data). The fishes were collected in summer 2005 through the North Pacific Groundfish Observer Monitoring Program. The fishes were captured at different times and locations than the *B. magister* samples analyzed in this study. We cannot show the specific capture locations of the fishes because the observer data could not be aggregated in a manner that protects the confidentiality of private commercial fishing interests (see www.afsc.noaa.gov/FMA/fma_database.htm). We compared the nitrogen isotopes of *B. magister* to the following fish species: juvenile and adult walleye pollock *Theragra chalcogramma*, adult yellowfin sole *Limanda aspera*, adult northern rock sole *Lepidopsetta polyxystra*, adult flathead sole *Hippoglossoides elassodon*, adult arrowtooth flounder *Atheresthes stomias*, and adult Pacific cod *Gadus macrocephalus*. Again, lipids were not removed from fish tissues prior to stable isotope analysis; thus comparisons between squid and fish $\delta^{13}\text{C}$ signatures could be biased due to differences in lipid content. For that reason, we limited our comparisons between squid and fishes to $\delta^{15}\text{N}$ signatures, which are not affected by

lipids (Kiljunen et al. 2006). Our comparisons were also restricted to the summer season because at the time of our study the only available estimates of fish $\delta^{15}\text{N}$ signatures were from fishes captured during this period. With the exception of pollock, we were not able to stratify the $\delta^{15}\text{N}$ signatures of the fishes by size or life stage because the fishes were of similar sizes. Box plot graphs of the summer $\delta^{15}\text{N}$ values of *B. magister* and fishes were created to compare their trophic positions.

Feeding chronology through eye lens analysis. Following the methods of Parry (2003), we evaluated the squid feeding chronology by examining changes in isotopic signatures ($\delta^{15}\text{N}$ and $\delta^{13}\text{C}$) as a function of eye lens radius. The squid eye is a spherical structure comprised of distal and proximal lenses and has a recognizable layering of thin tissue. Similar to many fishes, squid exhibit chronological deposition of concentric layers to the outer surface of their eye lens as they grow (Nicol 1989, West et al. 1995, Dove 1999). The center of the lens contains the oldest layer, whereas the youngest layer lies on the outer rim (Fig. 2). The proximal lens is the larger of the 2 lenses, and we analyzed the isotopic signatures of this tissue to reconstruct the feeding chronologies of 20 summer squid (year 2004 only) and 21 winter squid (year 2007). Preparations for isotope analysis proceeded in 3 steps. First, we cleaved the eyeball of an adult squid while it was thawing to isolate the proximal lens and measured the longest width of this lens using an ocular micrometer on a dissecting microscope at 8 \times power. Mantle lengths of the squid ranged from 220 to 310 mm, and total widths of their proximal lens ranged from 6.5 to 10.5 mm. Second, a segment of tissue layers was

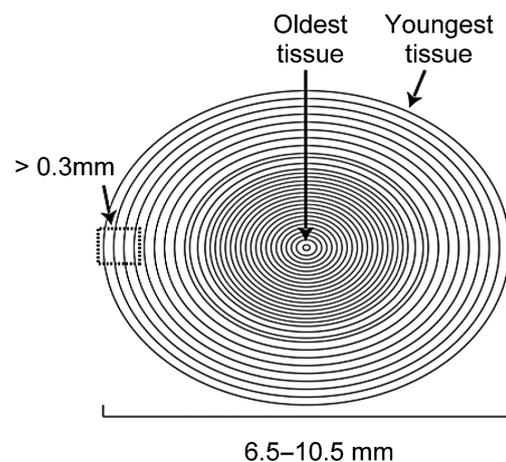


Fig. 2. *Berryteuthis magister*. Squid proximal eye lens. The youngest tissue lies on the outer edge of the lens, and the oldest tissue is in the center of the lens. The width of the eye lenses examined in our study ranged between 6.5 and 10.5 mm. Six to 11 lens segments, each >0.3 mm in width, were extracted from the eye lenses for stable isotope analysis.

Each segment contained multiple individual lens layers

excised from the outermost edge of the lens, rinsed with distilled water, and set aside for subsequent preparation. As it was nearly impossible to remove single layers of lens tissue, we removed relatively precise fixed groups of lens layers to get an approximation of intermediate feeding chronology (Fig. 2). To ensure that there was a sufficient amount of tissue for the isotope analyses, each segment was >0.3 mm in width. The number of segments extracted from an individual eye lens ranged between 6 and 11 among squid samples due to size differences of the squid eyes. Third, upon removal of each segment, the width of the proximal lens was re-measured. To avoid contamination, the scalpels and forceps used to separate and measure the individual lens layers were cleaned with ethanol following each excision. Also, if the proximal lens began to dry, making it more difficult to excise lens segments, we rinsed the eye lens with distilled water. The second and third steps were repeated until the final lens segment at the center of the squid eye was removed. All lens segments were dried, ground, weighed (0.6 to 1.2 mg), and analyzed for $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signature in the same manner as the *Berryteuthis magister* mantle muscle tissue.

To evaluate the trophic record of individual *Berryteuthis magister*, we related the isotopic signatures of the lens segments to particular squid body sizes. First, a regression for squid eye lens width to squid mantle length was developed from the proximal lenses of 50 summer squid (year 2004 only) and 50 winter squid (year 2007) of various body sizes. Squid mantle lengths ranged from approximately 65 to 135 mm, and total widths of the proximal lenses ranged from 2.5 to 10.5 mm. For each lens segment that was analyzed for isotopic signatures, we used these regressions to predict the approximate length range of the squid at the time a lens segment was laid down. Next, we generated coarse estimates of the time scales over which the lens segments were laid down. Length at age was estimated for individual *B. magister* captured during summer 2004 (Drobny 2008); however, it was not possible to fit a growth model to the data because the estimates were highly variable. This precluded us from using growth curves to back-calculate the age of squid to the time when the tissues were formed. Thus, we instead calculated the time over which the prey isotope signatures were integrated within the lens segment by dividing the estimated age of an individual squid (Drobny 2008; ~120 to 180 d) by the number of lens segments examined. Age estimates of *B. magister* were only available for the summer 2004 samples; therefore, we used the average integration time estimated from these samples to back-calculate the approximate time at which the first lens segments of winter- and summer-caught squid were laid down.

We used multiple approaches to evaluate the observed patterns of squid feeding chronologies. Our intent was to first determine if the $\delta^{15}\text{N}$ signatures revealed by the individual chronologies exhibited the monotonic increase with mantle length that is often observed from isotopic signatures of squid muscle tissue (e.g. Parry 2008, Cherel et al. 2009). To this end, we regressed the $\delta^{15}\text{N}$ values of the lens segments on the predicted mantle lengths and then quantified the frequency of individual records that demonstrated positive slope values that were at least marginally statistically significant ($p < 0.10$) and those that did not. Next, we estimated the average slopes of individual eye lens $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values among and within seasons by fitting linear mixed effects models to the isotope data. The $\delta^{15}\text{N}$ or $\delta^{13}\text{C}$ value was the response variable and squid mantle length and individual squid ID were the fixed and random effects, respectively. We also determined if the average relationship of lens $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values and estimated squid mantle lengths differed between seasons by adding season and season \times mantle length as fixed effects terms to the linear mixed effects models. The interaction term is useful for identifying whether the average slopes differed between winter and summer seasons. We fit the models using the nlme package in R version 2.10.2 (R Development Core Team 2010). Lastly, we examined whether the average slopes of eye lens $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values among the individuals were similar to the slopes estimated from muscle tissue analysis.

RESULTS

General diet analysis

The mantle lengths of the summer 2004 and summer 2007 squid samples ranged from 62 to 334 mm and 90 to 335 mm, respectively, and the winter 2007 samples ranged from 100 to 335 mm (Table 1). Of all of the stomachs analyzed, 56% contained contents that were identifiable to coarse taxonomic groupings (i.e. fish, squid, or crustacean), 13% contained only unidentifiable prey items, and 31% were empty (Table 1). The percent occurrence of fish in summer and winter *Berryteuthis magister* stomach contents was the highest of all prey items, followed by squid and crustaceans. The percent occurrence of fish in squid diets was as high as 47%. The frequencies of empty *B. magister* stomachs or those with only unidentifiable prey items were relatively similar across summer and winter seasons (Table 1).

Results of the logistic regression analysis indicated that *Berryteuthis magister* underwent an ontogenetic shift in diet composition. The consumption of fish and

squid by *B. magister* increased with mantle length, while the consumption of crustaceans decreased. This ontogenetic shift was evident during the summer (years 2004 and 2007 combined) and winter (year 2007) seasons (Fig. 3). The regression analysis highlighted the substantial frequency (>0.10) of piscivory by *B. magister* over all body sizes. In the summer, the median estimates of the probability of piscivory increased from 10 to 65% from the smallest to largest observed squid body sizes. For winter, the median estimates ranged from 31% for squid at mantle lengths of 130 mm to 51% for the largest size squid. The regression analysis further revealed a seasonal difference in the pattern of piscivory by *B. magister* with capture

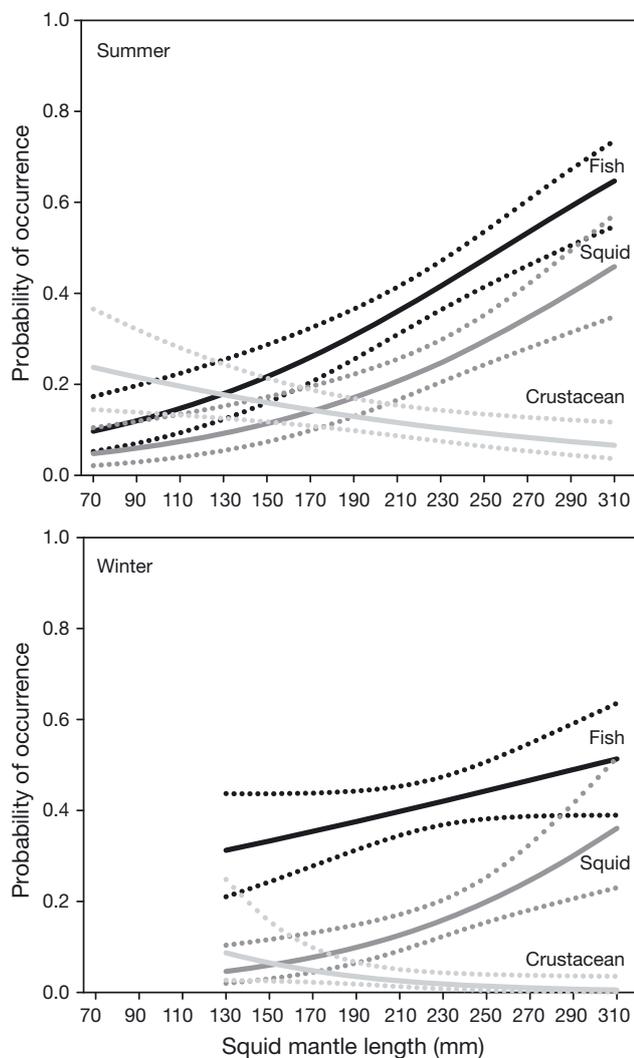


Fig. 3. *Berryteuthis magister*. Predicted probability of consumption of fish (black lines), squid (dark gray lines), and crustaceans (light gray lines) by *B. magister* at different mantle lengths. Logistic regression analyses were conducted for squid collected in summer (2004 and 2007 combined) and winter (2007). The 5th and 95th confidence intervals (dotted lines) are shown

depth (data not shown). The median estimates of the probability of piscivory for summer *B. magister* decreased from 36 to 4% over the primary capture depths (i.e. mean \pm 1 SD; 250–450 m). In contrast, the frequency of piscivory by winter *B. magister* was higher at deeper capture depths and gradually declined with increasing depth: the median estimates of the probability of piscivory ranged from 54 to 40% over the primary capture depths (i.e. mean \pm 1 SD; 350 to 550 m). The consumption of squid by *B. magister* in relation to capture depth followed the same trends as those described for fish. The median estimate of the probability of squid consumption by *B. magister* declined with capture depth from 42 to 8% in the summer and from 23 to 11% in the winter. The frequency of consumption of crustaceans increased slightly from 10 to 16% with summer capture depths and was $<4\%$ across all winter capture depths. We did not observe any meaningful shifts in the diets of *B. magister* with respect to latitude at capture.

Identification of digested hard parts revealed that *Berryteuthis magister* specimens preyed upon deep-water slope fishes and a variety of squid species, including conspecifics. The percent occurrence of otoliths and statoliths in non-empty *B. magister* stomachs was 16 and 9%, respectively. We identified a total of 323 otoliths and 118 statoliths, but the northern smooth-tongue *Leuroglossus schmidti* was by far the most common prey identified from the stomach contents (Table 1), representing 95% of the total number of identified otoliths. The remaining otoliths belonged to northern lampfish *Stenobranchius leucopsaurus*, brokenline lanternfish *Lampanyctus jordani*, northern pearleye *Benthabella dentata*, lumpfishes (Cyclopteridae sp.), snailfishes (liparids), and pollock *Theragra chalcogramma*. In contrast to the fish otoliths, the squid beaks consumed by *B. magister* were not dominated by a single taxonomic group (Table 1). Approximately 14% of the lower beaks were identified as *B. magister*, 7% were *Gonatus onyx*, and 3% were *Gonatopsis borealis*. The coarser taxonomic grouping of *Gonatus* sp. represented 40% of the digested beaks whereas the *B. magister*/*G. borealis* and *G. madokiai*/*G. middendorffi* groups represented 31 and 5% of the beaks, respectively.

Prey size spectrum

Quantile regression analysis of all prey sizes consumed by *Berryteuthis magister* during summers 2004 and 2007 and winter 2007 revealed a non-significant slope of the upper bound (90th quantile; $p = 1.000$) and lower bound regression lines (10th quantile, $p = 0.339$; Fig. 4). However, the regression analysis of *B. magister*

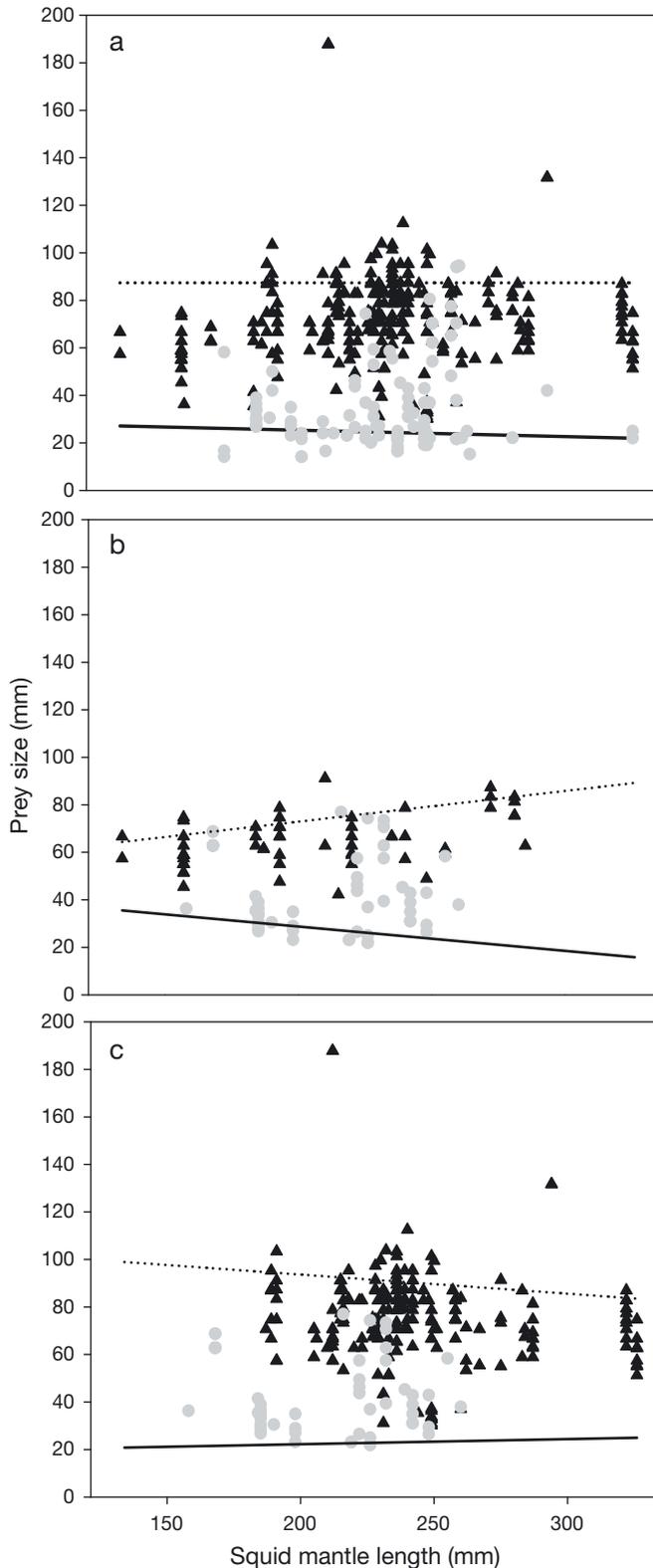


Fig. 4. *Berryteuthis magister*. Relationship between mantle length and estimated body sizes of consumed fish (\blacktriangle) and squid (\bullet) in (a) summer and winter, (b) winter only, and (c) summer only. Estimated quantile regression lines $t = 0.10$ (solid lines) and 0.90 (dotted lines) are superimposed on the scatterplots

prey sizes differed among seasons. We found non-significant upper (90th quantile, $p = 0.610$) and lower bound regression lines (10th quantile, $p = 0.071$) of summer prey sizes, whereas the winter prey sizes showed a positive, significant slope of the upper bound regression line (90th quantile, $p < 0.001$) and a non-significant lower bound (10th quantile, $p = 0.228$; Fig. 4). Also, non-significant trends were observed when we analyzed the prey size spectra of fish prey alone across both seasons (90th quantile, $p = 1.000$; 10th quantile, $p = 0.578$), whereas the size spectra of squid prey had positive, significant slopes for the upper (90th quantile, $p < 0.001$) and lower (10th quantile, $p < 0.001$) bound regression lines. Statistical comparisons of the regression lines for different prey types indicated that the slope of the upper bound of the squid prey size–predator size scatter plot was significantly greater than the slope for fish prey ($t = 25.830$, $df = 415$, $p < 0.001$). The minimum, maximum, and average sizes of squid consumed by *B. magister* were 15, 95, and 42 ± 19 mm, and the fish prey sizes typically ranged between 30 and 112 mm with an average size of 72 ± 18 mm.

Stable isotope analyses

The $\delta^{15}\text{N}$ signatures of squid muscle tissue indicated that the food web position of *Berryteuthis magister* spanned roughly 1 trophic level over the sampled size range (winter 2007: 105 to 310 mm, summer 2004: 72 to 334 mm). The $\delta^{15}\text{N}$ signatures of winter 2007 squid samples increased from approximately 11 to 15‰ over increasing squid mantle length (Fig. 5a). This range in $\delta^{15}\text{N}$ falls within the range of trophic fractionation (3 to 4‰; Minagawa & Wada 1984). The $\delta^{15}\text{N}$ signatures of summer 2004 samples followed the same trend as the winter samples, ranging from approximately 10.50‰ for small squid to 13.45‰ for large squid. In contrast to the $\delta^{15}\text{N}$ signatures, the $\delta^{13}\text{C}$ of winter 2007 and summer 2004 squid samples did not exhibit any distinguishable trends with body size (Fig. 5b). The $\delta^{13}\text{C}$ values of winter samples ranged from -19.50 to -21.20 ‰ and the summer samples ranged from -18.90 to -21.25 ‰.

Overall, we found a significant, negative correlation between the $\delta^{13}\text{C}$ signatures of *Berryteuthis magister* muscle tissue and latitude and depth at which they were captured (Table 2). This was found for both summer 2004 and winter 2007 samples. We did not find a significant correlation between the $\delta^{15}\text{N}$ of *B. magister* muscle tissue and the latitude at which squid were captured (Table 2). However, there was an overall significant, positive correlation between the $\delta^{15}\text{N}$ signatures of *B. magister* and capture depths (Table 2). The

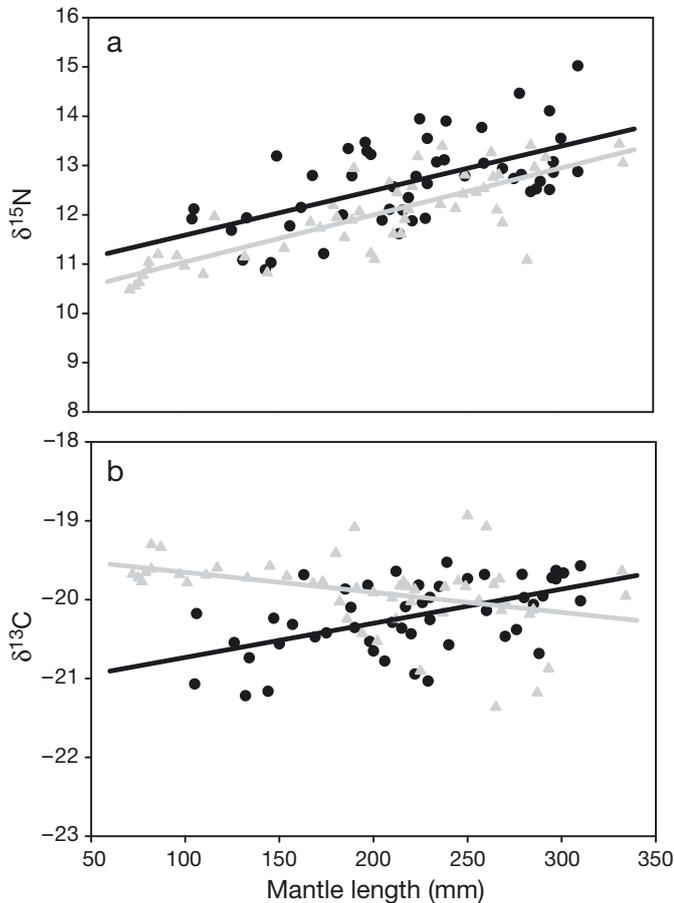


Fig. 5. *Berryteuthis magister*. (a) $\delta^{15}\text{N}$ and (b) $\delta^{13}\text{C}$ signatures of mantle muscle tissue. The isotopic signatures were quantified from squid collected in summer 2004 (\blacktriangle) and winter 2007 (\bullet). Equations of regression lines are: $y = 10.071 + 0.010x$ ($r = 0.79$, $F = 89.26$, $p < 0.001$) and $y = 10.69 + 0.009x$ ($r = 0.55$, $F = 25.18$, $p < 0.001$) for $\delta^{15}\text{N}$ values of summer and winter muscle tissue versus mantle length, respectively, and $y = -19.40 - 0.003x$ ($r = 0.50$, $F = 8.53$, $p = 0.005$) and $y = -21.16 + 0.004x$ ($r = 0.55$, $F = 20.38$, $p < 0.001$) for $\delta^{13}\text{C}$ values of summer and winter muscle tissue versus mantle length, respectively

average $\delta^{13}\text{C}$ value of winter samples was significantly higher than the average summer $\delta^{13}\text{C}$ value. The opposite trend was found for the $\delta^{15}\text{N}$ signatures of *B. magister* muscle tissue. The average C:N mass ratios for summer 2004 and winter 2007 samples did not differ significantly between seasons (Table 2).

Comparisons between the isotopic signatures of squid and fish muscle tissues indicated that the $\delta^{15}\text{N}$ values of *Berryteuthis magister* and walleye pollock were similar. The median $\delta^{15}\text{N}$ value of summer 2004 *B. magister* samples was 11.96‰ ($n = 50$). When grouped by size, the median $\delta^{15}\text{N}$ values of juvenile (<120 mm, $n = 11$) and adult (≥ 120 mm, $n = 39$) squid were 10.90 and 12.19‰, respectively. The median $\delta^{15}\text{N}$ values of juvenile and adult pollock captured in summer 2005 were 11.66‰ ($n = 10$) and 12.03‰ ($n = 10$),

respectively (Fig. 6). The $\delta^{15}\text{N}$ values of the remaining fishes collected in summer 2005 were substantially greater than *B. magister*, ranging between 14.16 and 16.09‰ (Fig. 6).

Isotope analysis of squid eye lenses

The number of individual lens segments analyzed per squid eye (proximal lens) varied between 6 and 11, and the average width of these lens segments was 0.79 ± 0.41 mm. To approximate the mantle lengths of *Berryteuthis magister* at the time the segments were laid down, we developed squid proximal eye lens width to squid mantle length regressions. We applied linear regression to summer 2004 *B. magister* data and found a strong, positive relationship ($R^2 = 0.94$, $a = 0.026$, $b = 1.075$, $df = 46$, $F = 757.7$, $p < 0.001$) between squid eye lens width and body size. The winter 2007 samples did not exhibit a linear relationship, thus we fit the data using a non-linear Gompertz function ($\text{Asym} = 8.380$, $p < 0.001$; $b_2 = 11.342$, $p = 0.117$; $b_3 = 0.977$, $p < 0.001$). The Gompertz function was not chosen based on any presumed mechanistic basis to relate eye lens to body size, but rather because it is a flexible function that could fit the observed relationship. The relationship between squid eye lens width and body size differed between seasons. The widths of the summer lenses continued to increase over increasing body sizes, while the widths of the winter eye lenses appeared to have leveled off at squid mantle lengths of >200 mm.

The eye lens segments revealed variations in the trophic relations of squid at fine temporal scales. We estimated that, on average, the individual eye lens segments represented approximately a 16 ± 2.0 d period. Back calculations based on this estimate indicated that the $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures of winter 2007 squid lenses reflected prey consumed by squid over lifetimes that spanned early to mid-autumn through early to mid-winter (Fig. 7a,b). In a similar manner, we found that the $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures of summer 2004 squid eye lenses reflected prey consumed by *Berryteuthis magister* from early winter through early to mid-summer (Fig. 7a,b). From the reconstructed trophic records we were able to detect increases and declines in $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ of individual *B. magister* at short time intervals over their entire life span.

The $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures of the squid eye lenses also revealed that the trophic record of *Berryteuthis magister* varied in notable ways among individuals (Fig. 7a,b). We classified the squid $\delta^{15}\text{N}$ trophic records as 2 trends: (1) a 'conventional' monotonic increase in $\delta^{15}\text{N}$ with increasing body size or (2) a non-monotonic trend with body size (i.e. an initial increase in $\delta^{15}\text{N}$ fol-

Table 2. *Berryteuthis magister*. Capture depths and body sizes analyzed for seasonal $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures of mantle muscle tissue (n = sample size). Student's t -tests were performed to identify potential seasonal differences between capture depths, squid body sizes, $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures, and C:N ratios. The Spearman correlation test was used to identify significant correlations between *B. magister* isotope signatures and latitude and capture depth

	All seasons n = 100	Summer 2004 n = 50	Winter 2007 n = 50	Statistics	
				t	p
Capture depth (m)	438 ± 113	390 ± 62	488 ± 132	-4.68	<0.001
Dorsal mantle length (mm)	210 ± 66	198 ± 72	222 ± 58	-1.80	0.075
Body mass (g)	371 ± 304	336 ± 304	409 ± 302	1.18	0.242
Mantle tissue					
$\delta^{15}\text{N}$ (‰)	12.32 ± 0.93	12.67 ± 0.88	11.96 ± 0.86	4.07	<0.001
$\delta^{13}\text{C}$ (‰)	-20.05 ± 0.48	20.20 ± 0.45	-19.89 ± 0.50	3.31	0.001
C:N mass ratio	3.50 ± 0.10	3.51 ± 0.12	3.50 ± 0.07	-0.22	0.820
Spearman correlation test					
	ρ (p)	ρ (p)	ρ (p)		
Latitude, $\delta^{15}\text{N}$ (‰)	-0.073 (>0.05)	0.255 (>0.05)	-0.278 (>0.05)		
Latitude, $\delta^{13}\text{C}$ (‰)	-0.315 (<0.01)	-0.323 (<0.05)	-0.419 (<0.01)		
Depth, $\delta^{15}\text{N}$ (‰)	0.366 (<0.001)	0.588 (<0.001)	-0.150 (>0.05)		
Depth, $\delta^{13}\text{C}$ (‰)	-0.471 (0.001)	-0.464 (<0.001)	-0.362 (<0.05)		

lowed by a decrease or vice versa). Based on estimated slope values, we found that 45% of the total squid samples demonstrated a significant, positive slope ($p < 0.10$) or a 'conventional' pattern. When we quantified the occurrence of significant, positive slope for each sampling season, we found that the variability in feeding chronologies may be season dependent: 70% of the winter samples exhibited the 'conventional' pattern, whereas only 20% of the summer samples demonstrated a significant linear increase in $\delta^{15}\text{N}$. We also found seasonal differences in the variation of $\delta^{15}\text{N}$ as a function of mantle length (e.g. ~3‰ for Novem-

ber 2006 and ~5 to 6‰ for July 2004; Fig. 7a) as well as within season sub-groupings (e.g. early July 2004; Fig. 7a).

Overall, the slope relating eye lens $\delta^{15}\text{N}$ to body size was positive ($t = 4.12$, $df = 305$, $p < 0.001$). The degree to which lens $\delta^{15}\text{N}$ values increased with mantle length differed between summer 2004 and winter 2007 samples ($t = 3.14$, $df = 304$, $p = 0.002$). The average lens $\delta^{15}\text{N}$ value did not significantly differ between the 2 periods ($t = 0.58$, $df = 37$, $p = 0.560$). The slope of lens $\delta^{13}\text{C}$ values across all individuals was not significantly different from 0 ($t = 0.57$, $df = 305$, $p = 0.573$); however, the slope and intercept of lens $\delta^{13}\text{C}$ values differed between summer 2004 and winter 2007 samples ($t = 4.60$, $df = 304$, $p = 0.000$ and $t = 7.01$, $df = 37$, $p = 0.000$). Lastly, the average slope of eye lens $\delta^{15}\text{N}$ values was lower than the slope of $\delta^{15}\text{N}$ values estimated from muscle tissue analysis (0.004 and 0.009, respectively). The slopes of winter eye lens $\delta^{15}\text{N}$ values were similar to winter muscle tissue signatures (0.007 and 0.009), but the average slope of summer eye lens $\delta^{15}\text{N}$ values was lower than the slope of summer muscle tissue signatures (0.001 and 0.010). The slopes of eye lens and muscle tissue $\delta^{13}\text{C}$ values showed opposite trends in both winter (-0.004 and 0.004) and summer (0.004 and -0.003) squid samples.

DISCUSSION

Our study on the trophic role of *Berryteuthis magister* in the EBS continental slope ecosystem produced 3 main findings. First, stomach content and stable isotope analyses revealed that *B.*

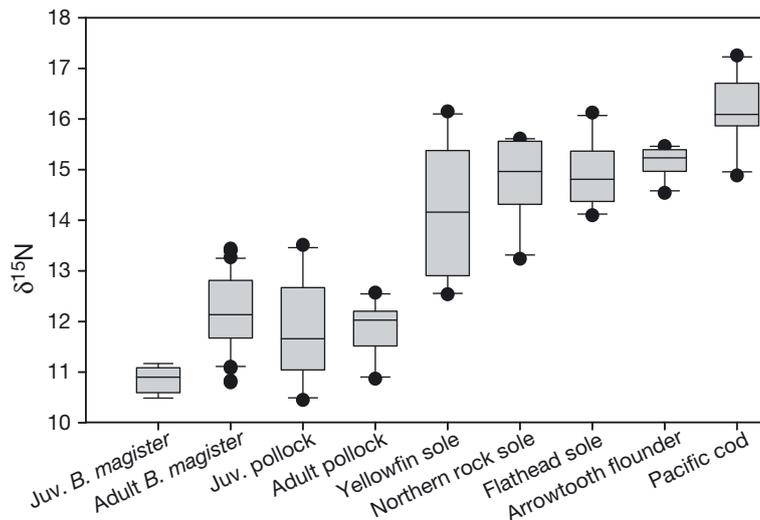


Fig. 6. $\delta^{15}\text{N}$ signatures of *Berryteuthis magister* and commercial fishes (C. Kurle, E. Sinclair, A. Edwards, C. Gudmundson unpubl. data) captured in the eastern Bering Sea during summer 2004 and summer 2005, respectively. The isotopic signatures were quantified from squid and fish muscle tissue. The box boundaries indicate the 25th and 75th percentiles. Whiskers below and above the box indicate the 10th and 90th percentiles

magister exhibits an overall increase in trophic position with ontogeny. The incidence of fish and squid prey in *B. magister* stomach contents gradually increased with increasing squid mantle length, and the muscle $\delta^{15}\text{N}$ signatures revealed that the food web position of *B. magister* increased by approximately 1 trophic level between juvenile and adult life stages. Second, *B. magister* did not exhibit size-dependent patterns of piscivory. The body sizes of fish prey consumed by *B. magister* remained constant with increasing predator mantle length. Third, in our attempt to reconstruct the

feeding chronologies of individual *B. magister*, we found substantial variability in squid trophic records. The feeding chronologies did not consistently reflect a monotonic increase in squid trophic position; non-monotonic patterns were relatively common. Our results also indicate that the variability in feeding chronologies may be season dependent. These findings suggest that there may be finer-scale spatiotemporal variations in squid trophic records, possibly related to seasonal prey availability, that are not captured by traditional isotope analyses of squid muscle tissue.

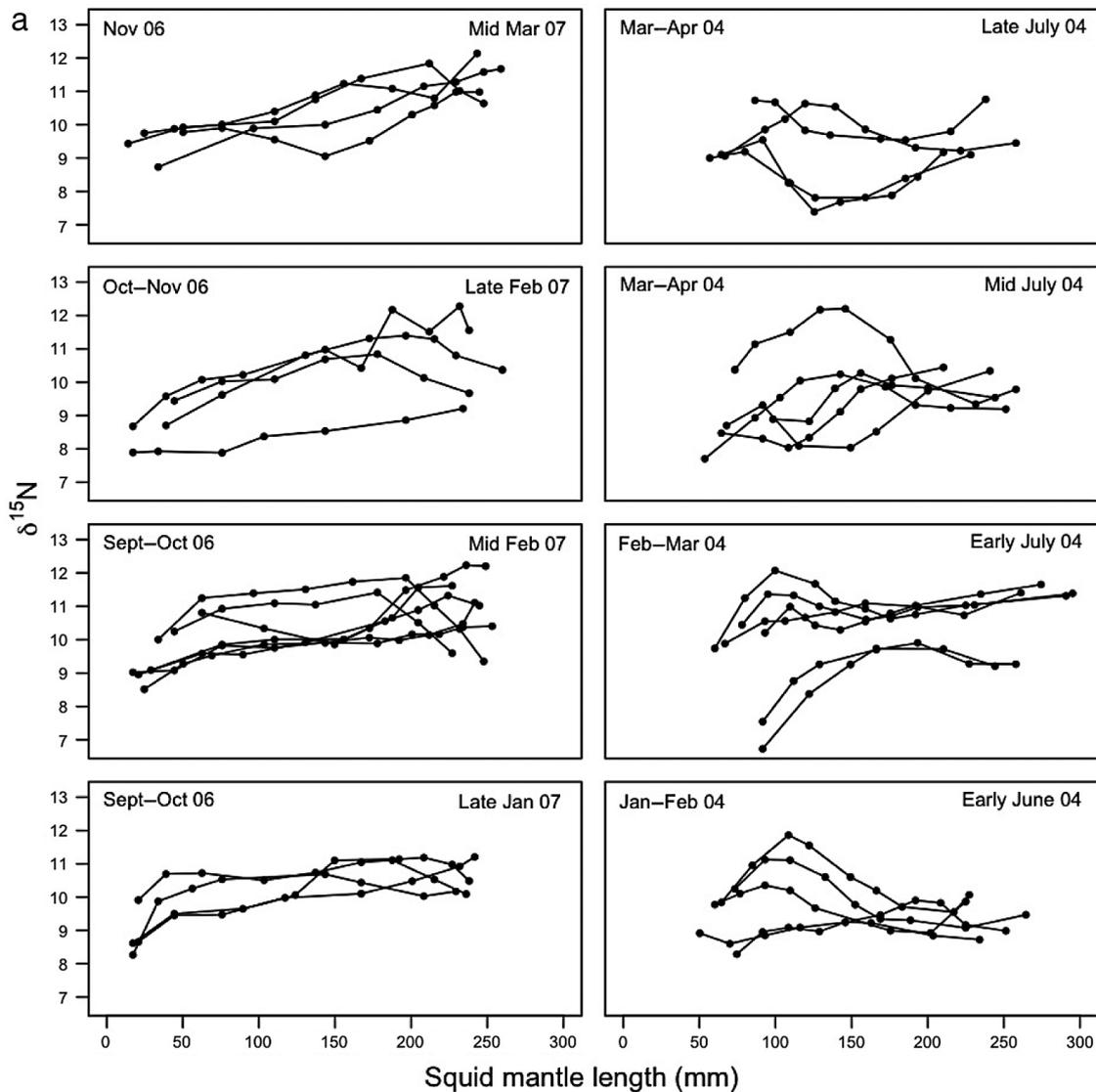


Fig. 7. *Berryteuthis magister*. Reconstructed feeding chronologies of individual squid captured in summer 2004 (right-hand columns) and winter 2007 (left-hand columns). The lines reflect the relationship between the (a) $\delta^{15}\text{N}$ signatures and (b, next page) $\delta^{13}\text{C}$ signatures of *B. magister* eye lens segments and the estimated mantle lengths of squid when the segments were deposited. Each line represents an individual squid. The squid are grouped by capture date (month and year; upper right-hand corner of box) and approximate time at which the first lens segment was laid down (month and year; upper left-hand corner of box). The regression equations are $y = 9.50 + 0.001x$, $y = 9.28 + 0.007x$, and $y = 9.38 + 0.004x$ for summer, winter, and all squid eye lens $\delta^{15}\text{N}$ values versus mantle length, respectively. The regression equations are $y = -20.36 - 0.004x$, $y = -18.31 - 0.005x$, and $y = -19.31 - 0.001x$ for summer, winter, and all squid eye lens $\delta^{13}\text{C}$ values versus mantle length, respectively

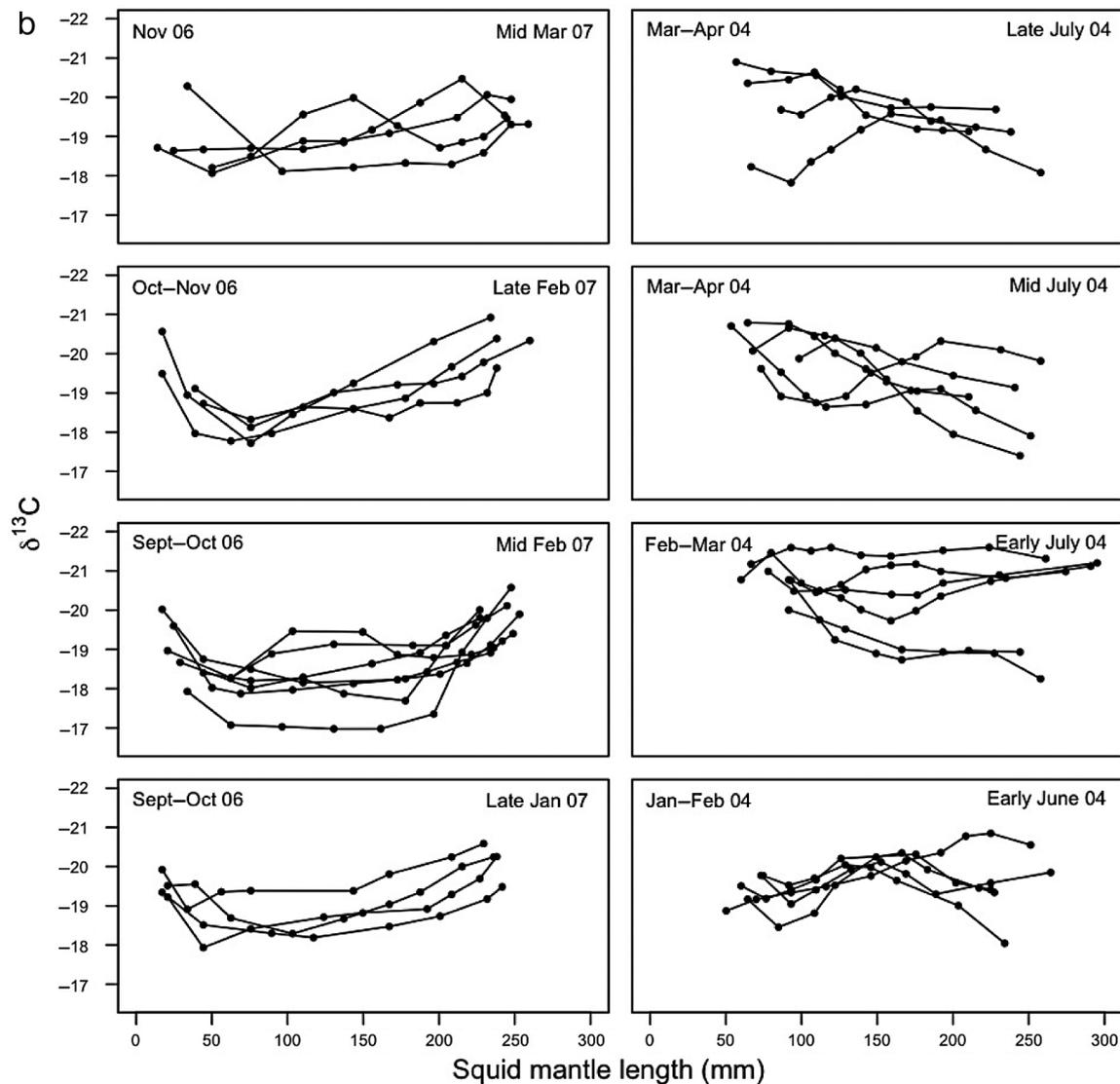


Fig. 7 (continued)

General diet analysis

The stomach content analyses indicate that *Beryteuthis magister* does not demonstrate feeding habits typified by squid species that are potentially important predators of higher trophic level fishes. For example, in the northwest Atlantic Ocean, the longfin inshore squid *Loligo pealeii* exhibits a rapid onset of piscivory, preys on the juvenile stages of its predators, including commercially valuable species, and demonstrates size-related patterns of piscivory (Hunsicker & Essington 2006). *L. pealeii* and other squid species exhibiting similar feeding traits have been described as voracious predators that may constrain the recruitment success of commercially targeted fishes (Rodhouse & Nigmatullin 1996, Dawe et al. 1997, Hunsicker & Essington 2006, 2008, Field et al. 2007). However, the diet con-

tents of *B. magister* revealed a slower ontogenetic shift to piscivory, and this species preyed mostly upon deep-water forage fishes and showed little evidence of size-related feeding constraints. These findings suggest that *B. magister* may interact with higher trophic level fishes through indirect food web pathways (i.e. sharing or competing for prey resources) rather than directly as a predator. The discrepancy in the types of fish prey consumed by *B. magister* and other well-studied squid may be explained by the differences in their migratory behaviors. Many squid species, including *L. pealeii*, undergo seasonal onshore and offshore migrations (Serchuk & Rathjen 1974, Ogawa et al. 1983, Hatfield & Rodhouse 1991, 1994, Augustyn et al. 1993, Roberts & Sauer 1994). As a result, they likely experience high spatiotemporal overlap with the early life stages of high trophic level fishes at spawning and/or nursery

grounds. In contrast, *B. magister* inhabit the edge of the continental shelf and slope over most of their lifetime and do not undergo shelf migrations (Kubodera et al. 1983, Nesis 1985, Okutani et al. 1988). Thus, *B. magister* probably encounter a higher abundance of deepwater slope fishes (i.e. myctophids) than the juvenile stages of many commercially harvested fish species that inhabit shelf waters.

Prey size spectrum

The lack of evidence for size-related constraints on piscivory by *Berryteuthis magister* could be due to prey availability. Previous studies of squid feeding ecology have shown that squid predation is dictated by predator and prey body sizes (Rocha et al. 1994, Collins & Pierce 1996, Phillips et al. 2003, Hunsicker & Essington 2006). Prey body sizes typically increase with increasing squid mantle length because squid feeding becomes less constrained by morphological limitations as squid grow. However, our results show that body size constraints may not be an important agent dictating the frequency of piscivory by *B. magister*. We observed an increase in the occurrence of fish in *B. magister* stomach contents over ontogeny, but the sizes of fish consumed by *B. magister* were independent of squid mantle length. A potential explanation is that *B. magister* encountered a limited size range of fish prey in their habitat, and thus our analysis reflects prey availability rather than size constraints. We expect quantile regression to be an effective technique for identifying size constraints when squid are feeding opportunistically on prey items of any size. However, in this study a large majority of the fishes consumed by *B. magister* were northern smooth-tongue *Leuroglossus schmidti*, suggesting that *B. magister* mainly encountered schools of this fish species. Size-related constraints may have been more evident if we had prey size estimates for a wider size range of *B. magister*. Identifiable otoliths were only found in the stomachs of squid with large body sizes.

Stable isotope analyses

Stable isotope analysis of squid muscle tissue has been used successfully as a tool for investigating the role of squid as predator and prey (Hobson et al. 1994, Takai et al. 2000, Cherel & Hobson 2005, Ruiz-Cooley et al. 2006, Stowasser et al. 2006, Parry 2008, Cherel et al. 2009). Our results are consistent with many of these studies. For example, the ontogenetic increase in $\delta^{15}\text{N}$ observed from *Berryteuthis magister* muscle tissue has been found for many squid species (Ruiz-Cooley et al.

2004, Parry 2008, Cherel et al. 2009). Similar to *B. magister*, the food web position of 3 oceanic squids increases by at least 1 trophic level with ontogeny: the $\delta^{15}\text{N}$ values of *Todarodes filippovae*, *Dosidicus gigas*, and *Ommastrephes bartramii* muscle tissues were shown to increase by ~ 3 , ~ 4 , and $>5\%$, respectively (Parry 2008, Cherel et al. 2009, see Ruiz-Cooley et al. 2010). Also, consistent with a recent study (Cherel et al. 2009), we found little variation in the $\delta^{13}\text{C}$ signatures of squid muscle tissue with increasing mantle length. This was expected because the enrichment of carbon isotopes per trophic level is low (i.e. $\sim 1\%$; DeNiro & Epstein 1978, Tieszen et al. 1983) compared to nitrogen isotopes (i.e. ~ 3 to 4% ; Minagawa & Wada 1984). Further, similar to Takai et al. (2000), we found a significant negative correlation between the $\delta^{13}\text{C}$ signatures of *B. magister* muscle tissue and intra-specific capture location, but did not find a significant correlation between $\delta^{15}\text{N}$ and latitude. The $\delta^{13}\text{C}$ signatures likely reflect the latitudinal characteristics of phytoplankton carbon isotopes, whereas the large enrichment in $\delta^{15}\text{N}$ per trophic level may have weakened any correlation between squid $\delta^{15}\text{N}$ values and latitudinal variation in phytoplankton $\delta^{15}\text{N}$ signatures (Takai et al. 2000). We note that our findings on squid carbon isotopes need to be interpreted with caution, particularly when making comparisons between species or taxa, because lipids (^{13}C -depleted) were not extracted from *B. magister* muscle tissues prior to analysis.

One unexpected finding in our study was the correlation between squid isotope signatures and capture depth. It is possible that the capture location (i.e. latitudinal effect) confounds the negative relationship between $\delta^{13}\text{C}$ and depth while the positive relationship between $\delta^{15}\text{N}$ and capture depth is related to squid body size. For instance, the adult stages of *Berryteuthis magister* are known to inhabit deeper depths than the juveniles (Nesis 1997, 1998, Arkhipkin et al. 1998), and thus we may expect to observe higher $\delta^{15}\text{N}$ signatures with increasing depths. The positive correlation between $\delta^{15}\text{N}$ and capture depth could also be attributed to differences in the $\delta^{15}\text{N}$ values of deep- and surface-water prey items. Several authors have found that the $\delta^{15}\text{N}$ values of marine predators feeding on mesopelagic prey resources are higher than the $\delta^{15}\text{N}$ of predators feeding on epipelagic prey (Graham et al. 2007, Ménard et al. 2007, Lorrain et al. 2009). Graham et al. (2007) hypothesized that this trend reflects the higher $\delta^{15}\text{N}$ values of deep-water prey species, which is an effect of nutrient cycling. The nitrogen isotopes and stomach contents of *B. magister* indicate that squid consumed mostly deep-water prey in the winter period, whereas epipelagic and mesopelagic prey were of similar importance in squid diets during the summer. The summer samples (year 2004) exhibited a

positive correlation between $\delta^{15}\text{N}$ values and capture depth, and no correlation was found for winter samples. Also, fish dominated the diets of winter squid over all capture depths, while the occurrence of fish and squid in summer squid diets was similar across depths. Together, these findings suggest that the summer squid fed across a strong $\delta^{15}\text{N}$ depth gradient, whereas the winter squid fed more consistently on mesopelagic fishes (e.g. northern smooth-tongue) with similar $\delta^{15}\text{N}$ signatures. Standardized sampling of *B. magister* and prey isotope signatures is needed to better examine the influences of body size, capture depth, and latitude on squid $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values. Analyses of squid and prey isotope signatures from multiple years and seasons are also needed to determine if our findings are indicative of a season or year effect. Interpretations of our findings are limited by the opportunistic manner in which *B. magister* samples were collected (i.e. different years, depths, and latitudes).

Comparisons of $\delta^{15}\text{N}$ values from squid and fish muscle tissues revealed that *Berryteuthis magister* might occupy a trophic position similar to or lower than several commercial fishes inhabiting the shelf and slope regions. Specifically, the isotope signatures indicate that adult *B. magister* could occupy a similar trophic level as adult and juvenile walleye pollock (>100 mm and ≤ 100 mm, respectively). The $\delta^{15}\text{N}$ values of these 2 species are not directly comparable because the squid and fish samples were collected at different locations in the EBS and during consecutive years (2004 and 2005, respectively). Our interpretation of these results is limited because we cannot show fish capture locations due to the NOAA Fisheries Observer Program confidentiality statutes. However, diet analyses also show that *B. magister* and pollock consume similar prey items, such as northern smooth-tongue, lampfish, and lanternfish (Lang et al. 2005). Further, *B. magister* and walleye pollock are known to overlap spatially and temporally in the EBS (www.afsc.noaa.gov/FMA). These findings suggest that these species share prey resources and that there could be potential for competition if resources became limited. Quantifying the extent of the dietary overlap of *B. magister* and walleye pollock should be a focus of future work.

Isotope analysis of squid eye lenses

Stable isotope analyses of the *Berryteuthis magister* eye lenses provide a richer picture of squid trophic ontogeny. Our results indicate that squid diets may not simply increase from lower to higher trophic level prey items with increasing body size, but that there can be substantial and sustained deviations over time or body size. The slopes of $\delta^{15}\text{N}$ in squid muscle tissue and

eye lenses in relation to squid mantle length suggest higher variation among the $\delta^{15}\text{N}$ signatures estimated from squid eye lens layers. Also, the individual feeding chronologies revealed finer temporal variations in $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures that were not evident from analysis of muscle tissues. Short-term changes in squid $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ ratios could be due to changes in individuals' feeding habits, prey availability, or localized differences in prey isotope signatures. Temporal variation in prey $\delta^{15}\text{N}$ levels can be quite pronounced among short-lived, high turnover species (e.g. zooplankton), but would generally not be revealed in the $\delta^{15}\text{N}$ tissues of predators if taken from tissues with slower turnover times (Jennings et al. 2008). Indeed, our analysis of muscle tissue suggested a monotonic increase in $\delta^{15}\text{N}$ values with increasing squid mantle length: the turnover time of squid muscle tissue varies from a few weeks to a few months (Ruiz-Cooley et al. 2006). While we cannot discern whether feeding habits or the underlying prey $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures are ultimately responsible for the observed trends in feeding chronologies, our results do point to the presence of intermediate-scale variability in the food webs that support squid.

The variability in *Berryteuthis magister* trophic records revealed by squid eye lenses is supported by earlier studies. Parry (2003) developed the eye lens method to reconstruct the feeding chronologies of 2 squid species in the north Pacific sub-tropical gyre, *Ommastrephes bartramii* and *Sthenoteuthis oualaniensis*. These squid also exhibited individual variability and non-monotonic trends of lens $\delta^{15}\text{N}$ values as a function of mantle length. However, these species also demonstrated greater shifts in lens $\delta^{15}\text{N}$ values over their lifetimes than *B. magister*. For instance, the $\delta^{15}\text{N}$ values for individual *S. oualaniensis* and *O. bartramii* varied by as much 6.67‰ (mean 4.25 ± 1.91 ‰) and 13‰ (mean 8.85 ± 3.27 ‰) across mantle lengths, respectively. The maximum range of $\delta^{15}\text{N}$ across *B. magister* across body size was 3.6‰ (mean 2.08 ± 0.63 ‰). These differences among species could reflect differences in their trophic ecology and/or migration patterns. For example, *O. bartramii* exhibit large, seasonal migrations between spawning and feeding grounds (Murata & Nakamura 1998), whereas *S. oualaniensis* and *B. magister* do not undergo any known seasonal migrations. As a result, *O. bartramii* are likely to encounter greater latitudinal shifts in isotope signatures over their lifetime and assimilate a wider range of $\delta^{15}\text{N}$ in their tissues (Parry 2003).

The work of Ruiz-Cooley et al. (2010) further corroborates our findings of *Berryteuthis magister* trophic records. These authors reconstructed the feeding chronologies of individual Humboldt squid *Dosidicus gigas* using the $\delta^{15}\text{N}$ signatures of proostracum sec-

tions sub-sampled from squid gladii. Consistent with the *B. magister* eye lenses, the $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ signatures of *D. gigas* gladii were highly variable over squid body size, and there was an overall increase in the $\delta^{15}\text{N}$ values of sampled sections as squid became larger. The squid gladii exhibited greater differences in the mean isotope values of young and old segments than the *B. magister* eye lenses. For example, based on estimated regression models, the mean $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ values of squid gladii differed by 1.6 and 0.6‰, respectively, between new (6 cm) and old sections (40 cm), whereas the mean isotope differences of *B. magister* lens layers laid down at juvenile (6 cm) and adult (30 cm) stages differed by 0.96‰ for $\delta^{15}\text{N}$ and 0.24‰ for $\delta^{13}\text{C}$. Again, this finding could be an effect of differences in trophic ecology and/or lifetime migration patterns. In contrast to *B. magister*, *D. gigas* are large, voracious predators and are highly mobile (Nigmatullin et al. 2001). This finding could also reflect the differences in the molecular structure and isotope fractionation rates of squid eye lenses and gladii. Our results and those of Parry (2003) show that $\delta^{15}\text{N}$ signatures of squid eye lenses are often lower than the $\delta^{15}\text{N}$ of squid muscle tissue. The ontogenetic shift in isotope values measured from squid eye lenses was also lower than those measured from squid muscle tissue (Parry 2003, present study). Similarly, the findings of Chérel et al. (2009) and Ruiz-Cooley et al. (2006, 2010) indicate that the isotopic signatures of squid hard parts (beaks and gladii) had lower $\delta^{15}\text{N}$ signatures and demonstrated smaller ontogenetic shifts in isotope values than soft tissue. These authors suggest that the differences are due to the presence of chitin in squid hard parts, which is a ^{15}N -depleted molecule. A comprehensive study that examines how squid isotope signatures vary among eye lenses, beaks, gladii, and muscle tissue is needed to make better comparisons between squid isotope studies.

The reconstructed feeding chronologies of *Berryteuthis magister* might reflect a shift in seasonal foraging strategies. The variation among individual trophic records was higher among squid caught in summer than those captured in winter. Also, the winter squid more often exhibited conventional trophic patterns; squid trophic level increased with increasing body size. These differences might be attributed to the seasonal availability of prey to *B. magister* over their short lifetime. For instance, the maximum age (post-embryonic growth) of squid captured in the EBS was estimated to be ~180 d (Drobny 2008). Thus, it is possible that *B. magister* collected in the EBS during summer months were exposed to highly productive ocean conditions (spring and summer) over much of their life span. The high abundance of different prey types available to the summer-caught squid would have allowed them to feed

opportunistically, potentially resulting in more variation among the trophic history of individual squid. In contrast, *B. magister* caught during the winter months were likely to encounter less productive conditions (autumn and winter) and would have been more limited by prey choice. Under such conditions, squid may need to search for additional food items in deeper waters where prey $\delta^{15}\text{N}$ values may be inherently higher. They might also shift toward more specialized feeding strategies when prey abundance is low to reduce intra-specific competition and maximize energetic return. Previous authors have documented similar shifts in the seasonal foraging patterns of demersal fishes (Reum & Essington 2008), and others have suggested that spatial or temporal variation in prey availability and the phenotypic differences among individuals that influence foraging success could promote the divergence of a generalist population into alternate foraging specialists (Partridge & Green 1985, Magurran 1986, Estes et al. 2003). It is plausible that squid also shift their foraging strategy in response to changes in seasonal and/or annual prey availability. This notion is supported by the findings Ruiz-Cooley et al. (2010), which suggest that the variable but increasing $\delta^{15}\text{N}$ and $\delta^{13}\text{C}$ along the proostacum of *Doscoedius gigas* gladii is an effect of prey availability and optimal foraging strategy. Feeding chronologies of squid captured in multiple seasons and years are needed to evaluate if the *B. magister* trophic records reflect seasonal environmental conditions or are instead an effect of sampling year (e.g. 2004 versus 2007).

Our study demonstrates that squid eye lenses can provide a chronology of trophic position over time. We chose to analyze squid eye lenses because the tissue layers are readily distinguishable over the entire lens structure, and the numerous segments of the lens tissue laid down over a squid's lifetime can be sampled with ease. Feeding chronologies reconstructed from squid eye lenses and gladii are a valuable tool, as they could improve our ability to evaluate the trophic ecology of squid life stages or species that are difficult to capture with traditional sampling gear. For example, adult *Berryteuthis magister* are frequently captured in bottom trawl surveys or as by-catch in commercial fisheries, but the juvenile stages of *B. magister* are more widely dispersed in pelagic habitats (Arkhipkin et al. 1998, Nesis 1998) and are not often captured by traditional sampling methods. To be able to apply the eye lens method in a more rigorous fashion, direct experimental work is needed to validate the assumptions we made here. Also, future work should attempt to verify how often layers of eye lenses are laid down and to relate the age of squid to the time when the tissues were formed. This information can be used to develop more precise estimates of the time scales over which

prey isotope signatures are measured and to determine whether there is any synchrony among the age of individuals and their trophic signatures. We could not use squid growth models to back-calculate the age of squid to the time when the tissues were formed due to the high variability surrounding the estimated ages of *B. magister* inhabiting the EBS (Drobny 2008). However, we did use these estimates to generate coarse approximations of the time at which the first eye lens segments were laid down. We emphasize that these results need to be interpreted with caution because there is some discrepancy between the estimates used in our study and those from an earlier study. Arkhipkin et al. (1996) aged *B. magister* from the western Bering Sea and found that they live longer than 1 yr.

In summary, our study used 3 complementary methods to provide valuable information on the trophic position of *Berryteuthis magister* in the EBS and to reveal the individual variation in trophic ontogeny that is masked by traditional stomach content and stable isotope analyses. Future work should aim to concurrently sample and identify the isotope signatures of *B. magister* and their prey over multiple seasons and years. This would allow one to better evaluate the seemingly important effects of body size, depth, latitude and season on squid trophic ecology. Such work would build on the present study and further improve our knowledge of the role of *B. magister* in the EBS food web and their potential to impact juvenile fishes through indirect food web pathways. Better knowledge of *B. magister* trophic interactions is important for broadening our understanding of how human and environmental forcing can affect the EBS ecosystem.

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