FISH PARASITOLOGY - SHORT COMMUNICATION



You are how you eat: differences in trophic position of two parasite species infecting a single host according to stable isotopes

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Abstract

Parasitism is commonly recognised as a consumer strategy, although, the interaction of parasites in communities and ecosystems are generally poorly understood. As parasites are integral parts of food webs, analysis of the trophic interactions between parasites and hosts was assessed through comparison of stable isotope ratios of carbon $({}^{13}C/{}^{12}C)$ and nitrogen $({}^{15}N/{}^{14}N)$. Largemouth yellowfish (*Labeobarbus kimberleyensis*) infected with the Asian tapeworm (*Schyzocotyle acheilognathi*) were collected from the Vaal Dam. Signatures of $\delta^{13}C$ and $\delta^{15}N$ were assessed in host muscle and liver tissue, and cestodes using an elemental analyser coupled with an isotope ratio-mass spectrometer (EA-IRMS). Hosts were enriched by 4.1% in the heavy nitrogen isotope with respect to the *S. acheilognathi* and therefore occupy a higher trophic position than the parasite. Comparison of $\delta^{13}C$ indicates that dietary sources of carbon in cestodes are derived from the host liver. Comparison of stable isotope signatures between *Paradiplozoon ichthyoxanthon* (another common parasite of the Largemouth yellowfish in the Vaal River) and *S. acheilognathi* showed that the monogenean was enriched by 5.3% in ${}^{15}N$ which accounts for a difference of almost two trophic positions. Isotope differences in the host–parasite system considered indicate that differences can be related to the mode of nutrient acquisition employed by host and parasites. Cestodes, being depleted in both ${}^{13}C$ and ${}^{15}N$ relative to the host and monogenean (*P. ichthyoxanthon*), indicate that *S. acheilognathi* assimilates nutrients derived from the host metabolism which are released from the liver.

Keywords Monogeneans · Cestodes · Stable isotopes · Trophic relationships · Host-parasite system

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Introduction

Analysis of stable isotope ratios of carbon $({}^{13}C/{}^{12}C)$ and nitrogen (15N/14N) as a method for interpreting food web architecture and energy flows based on differences in isotope signatures (Doucett et al. 1999; Thompson et al. 2005; Wada 2009; Sures et al. 2019) has become increasingly popular in ecological studies. Differences in these isotopic signatures serve as a timeintegrated representation of the diet of organisms based on nutrient assimilation rather than ingestion of food materials (Fry and Sherr 1984). Generally, studies have shown that consumers are enriched in ¹⁵N by 3.4% per trophic level, and thus differences in nitrogen enrichment serve as a representation of the position of organisms in the trophic hierarchy (Minagawa and Wada 1984; Vander Zanden et al. 1997). Food sources incorporated into the diet of organisms can be determined through analysis of carbon (¹³C) signatures (Post et al. 2000). More recently, stable isotope analysis (SIA) has been applied as a means of better understanding the intricate relationship between

hosts and parasites (Sabadel et al. 2016; Nachev et al. 2017; Yohannes et al. 2017; Sures et al. 2019).

Parasites are important components of ecosystems and are known to shape and modify the structure and stability of food webs (Marcogliese 2003; Poulin 2010; Nachev et al. 2017). Despite their close link with the food web, they have been somewhat neglected in studies on trophic relationships in ecosystems (Thompson et al. 2005). This has been related to the fact that inclusion of parasitism in food web studies would ultimately result in invalidation of the trophic cascade model, as inclusion of parasitism in food webs leads to looping, increased omnivory and increased chain length (see Marcogliese and Cone 1997). Despite these effects, understanding how parasites interact with host organisms is crucial to understanding the ecology of parasites (Nachev et al. 2017). The relationship between parasites and hosts has mostly been synonymised with that of a predator-prey relationship (Poulin 2010; Nachev et al. 2017). This therefore suggests that parasites should be enriched in the heavier nitrogen isotope similarly to a predator (Poulin 2010). However, unlike predatory organisms, most parasites only derive nutrients from a single host at any one time in the life history, and as such only attain a fraction of the total biomass of their hosts (Lafferty and Kuris 2002).

Studies comparing isotope signatures between hosts and parasites have indicated that enrichment of the heavier nitrogen isotope in parasites is largely linked to the mode these organisms have developed to acquire nutrients (Nachev et al. 2017). For instance, endoparasites such as cestodes and acanthocephalans which acquire nutrients through assimilation of molecules derived from the hosts' metabolism are depleted in ¹⁵N isotope. Whereas, ectoparasites such as fleas (Boag et al. 1998), nymphs of Pteronarcys biloba (Doucett et al. 1999), ticks (Schmidt et al. 2011) and some monogeneans (Sures et al. 2019), which actively feed on host tissues are ¹⁵Nenriched compared to their hosts (Pinnegar et al. 2001; Deudero et al. 2002; Persson et al. 2007). In a study of 10 different species of fish hosts Deudero et al. (2002) found that cestodes and nematodes were depleted in ¹⁵N compared to their hosts, while copepod parasites were ¹⁵N-enriched. Similarly, Pinnegar et al. (2001) observed that Schistocephalus solidus (Cestoda) and Hysterothylacium aduncum (Nematoda) were ¹⁵N-depleted compared to their hosts. As for ectoparasites, in the same study, the authors however, observed that Lernaeocera branchialis (Copepoda) and Anilocra physodes (Isopoda) were not enriched in the heavier nitrogen isotope compared to their hosts. Persson et al. (2007) found that the cestode Eubothrium crassum was depleted in ¹⁵N. Sures et al. (2019) showed that, compared to two fish hosts infected with the monogenean Paradiplozoon ichthvoxanthon, parasites were ¹⁵N-enriched, which corroborated observations for other ectoparasites (Boag et al. 1998; Doucett et al. 1999; Deudero et al. 2002; Voigt and Kelm 2006; Schmidt et al. 2011; Welicky et al. 2017).

In South Africa, to date, a single study was performed to analyse the trophic relationship between an ectoparasite and its host fish inhabiting the Vaal Dam (Sures et al. 2019). Sures et al. (2019) demonstrated that the ectoparasitic monogenean. Paradiplozoon ichthvoxanthon, was enriched in ¹⁵N compared to two closely related fish species; Largemouth yellowfish (Labeobarbus kimberleyensis) and Smallmouth vellowfish (Labeobarbus aeneus). In addition to infection by P. ichthyoxanthon, Largemouth yellowfish is also host to the cestode Schyzocotyle acheilognathi (Bertasso and Avenant-Oldewage 2005; Retief et al. 2007). The aims of this study were therefore to compare stable isotope signatures of carbon and nitrogen within the L. kimberleyensis-S. acheilognathi system from the Vaal Dam in an attempt to better understand the trophic interactions between host and cestode. In addition, as the Largemouth yellowfish is also host to the monogenean, P. ichthyoxanthon comparisons between the stable isotope signatures detailed for L. kimberlevensis-P. ichthyoxanthon in Sures et al. (2019) and the host-cestode model in the present study were done to better understand the interactions of parasites which occupy different microhabitats within the same host organism and have different feeding strategies.

Materials and methods

Host and parasite collection

Collections of *Labeobarbus kimberleyensis* (n = 10) were conducted in the Vaal Dam, South Africa, around the University of Johannesburg (UJ) Island (26° 52′ 33.62″ S; 28° 10′ 25.76″ E) using gill nets (mesh size 45–170 mm) in March 2017. Fish were removed from the nets and immediately placed into 160-L plastic containers filled with aerated water from the dam. The fish were transported back to a field laboratory on UJ Island where weights and morphometric parameters (standard, fork and total length) were determined before dissection, for calculation of Fulton's condition factor (K; Eq. 1) according to Heincke (1908).

$$K = 100 \times \frac{SW}{TL^3} \tag{1}$$

with SW: as weight of the fish; TL: as total length.

The fish were euthanised by severing the vertebrae of the spinal cord, posterior to the head. Largemouth yellowfish were then dissected to expose the visceral organs; the intestines were opened using Dumont forceps to expose and collect *S. acheilognathi*. All cestodes were removed, counted and placed into 2-mL microcentrifuge tubes. Following removal of the intestines, muscle and liver tissue of the hosts were collected using plastic forceps and a ceramic knife and placed

into 15-mL centrifuge tubes. All host and parasite samples were frozen at -20 °C until stable isotope analysis. All procedures for the collection of fish and parasites from the Vaal Dam followed approval from the Faculty of Science Ethics Committee of the University of Johannesburg (Ethics reference number, 2016-5-03) and application of appropriate permits for the collection of sensitive species from the Gauteng Department of Agriculture and Rural Development (permit number, 009658).

Frozen host and parasite samples were freeze-dried using an Alpha 1-2 LDplus freeze drier (Martin-Christ Gefriertrocknungsanlagen, GmbH; Germany) for 48 h and then homogenised directly in the 2-mL microcentrifuge tubes using a stainless steel pestle. Analysis of stable isotopes in muscle and liver of the host was performed as muscle tissue has been mostly used for trophic position comparisons between organisms. The inclusion of liver tissue was due to the fact that cestodes rely on bile excreted from the liver for nourishment.

Stable isotope analysis

For stable isotope ratio analysis of carbon $({}^{13}C/{}^{12}C)$ and nitrogen $({}^{15}N/{}^{14}N)$, triplicate samples of homogenised host tissues and parasites were weighed (0.2–0.7 mg) into tin capsules (4 × 6 mm) and folded. Folded capsules were placed into a PYRO Cube elemental analyser (EA; Elementar Analysensysteme, Langenselbold, Germany) coupled with an IsoPrime 100 isotope ratio mass spectrometer (IRMS; Elementar Analysensysteme) for the quantification of the isotope ratios of carbon and nitrogen in host and parasite tissues. Stable isotope ratios in tissue samples and internal standards were reported as δ -notation by Eq. (2) following Werner and Brand (2001) and Nachev et al. (2017).

$$\delta^h E_{s,ref} = \frac{R \left({^{h_E}/_{l_E}} \right)_s}{R \left({^{h_E}/_{l_E}} \right)_{ref}} - 1 \tag{2}$$

where $R(\frac{h_E}{l_E})_s$: ratio of heavy and light isotopes for carbon and nitrogen in host and cestode tissues; $R(\frac{h_E}{l_E})_{ref}$: ratio in reference materials.

Normalisation of the internal working standard acetanilide (AcAn) was done by two point calibration against the international standards, USGS40 and USGS 41 (International Atomic Energy Agency). The international standards were used as reference materials for the normalisation of isotope ratios determined in AcAn and host and cestode tissue samples.

Statistical analyses

Normality of the data was tested using the Shapiro–Wilk test and as data was not normally distributed, non-parametric tests were used to compare differences in isotope signatures. Differences in isotope signatures between host and parasite were compared using the Kruskal–Wallis test. Post hoc comparison was performed using a pairwise comparison with Bonferroni correction.

Host–parasite discrimination factors ($\Delta^{h}E$) were estimated by subtracting isotope signatures in parasites from the signatures in host muscle and liver tissue (Eq. 3). The same equation was applied to assess isotope discrimination between the monogenean, *P. ichthyoxanthon* and *S. acheilognathi* ($\Delta^{P}E$ (P.i-S.a)).

$$\Delta^h E = \delta^h E_{\text{parasite}} - \delta^h E_{\text{host}} \tag{3}$$

where $\delta^h E$ are levels of ¹³C and ¹⁵N in host muscle and liver tissue and in cestodes. For comparison between parasites, the isotope signatures in the monogenean were subtracted from the isotope signatures in the cestodes.

Differences in trophic position (Δ TP) between host and cestode were calculated according to the following Eq. (4):

$$\Delta TP = \frac{\left(\delta^{15} N_{\text{parasite}} - \delta^{15} N_{\text{host}}\right)}{\text{TEF}}$$
(4)

For estimation of the trophic position of cestodes relative to the fish host, the average trophic enrichment factor (TEF = 3.4%) was used (Minagawa and Wada 1984). This was applied for isotope signatures between the parasite and muscle tissue, and not for the liver tissue of the host.

Results and discussion

Stable isotopes in host and parasite

The cestode, S. acheilognathi, was found to be depleted in both ¹⁵N and ¹³C isotopes relative to the liver and muscle tissue of the host fish, L. kimberlevensis (Table 1 and Fig. 1). Differences in stable isotope composition between S. acheilognathi and the muscle and liver of L. kimberlevensis were significant for $\delta^{15}N$ (Kruskal–Wallis test; H = 14.24; df = 2; P = 0.001) but not for δ^{13} C (Kruskal-Wallis test; H = 2.08; df = 2; P = 0.354). This pattern is in line with other studies (Table 2) on adult and larval cestodes (Boag et al. 1998; Iken et al. 2001; Pinnegar et al. 2001; Deudero et al. 2002; Power and Klein 2004; Persson et al. 2007; Navarro et al. 2014; Behrmann-Godel and Yohannes 2015; McGrew et al. 2015) and acanthocephalans (Nachev et al. 2017) which have similarly been found to be depleted in ^{15}N isotope compared to their hosts. Differences in ¹³C isotopes between host muscle tissue and cestode in the present study accounted for a difference of $1.2 \pm 2.1\%$ and therefore this difference is within the 1-2% enrichment factor reported in literature (DeNiro and Epstein 1981; Fry and Sherr 1984; Power and Klein 2004). However, this difference was not

	TL (cm)	W (kg)	k		$\delta^{13}C$	$\delta^{15}N$
L. kimberleyensis	42.4 (±3.1)	0.7 (±0.2)	0.9 (±0.1)	Muscle	-19.8 (±0.4)	17.5 (±0.8)
				Liver	$-20.9(\pm 2.7)$	15.8 (± 5.2)
S. acheilognathi	_	_	_		-20.9 (±2.2)	13.4 (±1.8)
$\Delta^{h}E$ (parasite-host)	_	_	_	Muscle	-1.17 (±2.14)	-4.10 (±1.11)
				Liver	$-0.04 (\pm 2.04)$	-2.47 (±5.92)
L. kimberleyensis*	35.8 (± 6.7)	0.43 (±0.2)	1.0 (±0.2)	Muscle	$-20.61 (\pm 0.28)$	16.42 (±0.32)
P. ichthyoxanthon*	_	_	_		$-20.83 (\pm 0.35)$	18.73 (±0.18)
$\Delta^{h}E$ (parasite-host)*	_	_	_		-0.22	-2.31
$\Delta^{\mathrm{P}}\mathrm{E}$ (P.i-S.a)	_	_	_		0.07	5.33

Table 1Fish morphometry and signatures of stable isotopes of carbonand nitrogen in the muscle and liver tissue of Labeobarbuskimberleyensisand cestode, Schyzocotyle acheilognathicompared with

data presented by Sures et al. (2019) for isotope signatures in *L. kimberleyensis–Paradiplozoon ichthyoxanthon* from the Vaal Dam

*Data from Sures et al. 2019

significant and is consistent with other studies where host and cestode parasites are relatively similar in terms of ¹³C enrichment (Power and Klein 2004), indicating that both host and parasite assimilate nutrients from similar sources (Power and Klein 2004).

The host fish muscle tissue was enriched by $4.1 \pm 1.1\%$ in the heavier nitrogen isotope with respect to the parasite, indicating that the host occupies a higher trophic position than the parasite (based on equation for Δ TP (4)). Liver tissue was similarly enriched in the heavier nitrogen isotope by $2.5 \pm 5.9\%$. For carbon isotopes, comparison with liver tissue of Largemouth yellowfish hosts showed cestodes were depleted in ¹³C by $0.04 \pm 2.04\%$ relative to the host and therefore, host liver and *S. acheilognathi* shared similar δ^{13} C signatures. These results are in line with the mode of nutrient acquisition observed in other endohelminths such as cestodes and



Fig. 1 Mean isotope ratios and standard deviations for δ^{15} N and δ^{13} C in *Labeobarbus kimberleyensis* (*L.k*)–*Schyzocotyle acheilognathi* compared with data from Sures et al. (2019) (*) for *L. kimberleyensis*–*Paradiplozoon ichthyoxanthon* and *L. aeneus* (*L.a*)–*P. ichthyoxanthon* collected from the Vaal Dam. Isotope ratios for the muscle (m) and liver (l) from *L. kimberleyensis* are compared for the present study. Data from the present study are indicated as bold font

acanthocephalans (Nachev et al. 2017). Cestodes lack a digestive system and are unable to metabolise complex molecules (Smyth and McManus 1989). As a result, these parasites acquire pre-digested nutrients from the hosts' intestine but also reprocessed compounds derived from the hosts' metabolites in bile salts. These compounds, which are assimilated across their highly specialised tegument, are depleted in heavier isotopes (Doucett et al. 1999; Nachev et al. 2017). This ultimately leads to endohelminths, such as cestodes, being depleted in heavier isotopes compared to their hosts (Nachev et al. 2017). Similarities in δ^{13} C in host liver and cestodes confirm that S. acheilognathi derives nutrients from metabolites released from the liver of the host and not through direct feeding on host tissue. Thus, cestodes as well as acanthocephalans might be assumed to have a somewhat more commensalistic and scavenger-type feeding strategy as they do not compete directly with the host for the same food source and do not feed actively on host tissues.

Large standard deviations in stable isotope levels were observed in the liver tissue of the host and cestodes. This variability could be related to the metabolic activity of the host and parasite tissues as well as variability in the feeding biology of the host fish (Neilson et al. 2005; Behrmann-Godel and Yohannes 2015; Yohannes et al. 2017). Behrmann-Godel and Yohannes (2015) suggest that variability in isotope levels of intestinal helminths is likely the result of the nutrient component they are exposed to in the intestine. Thus the variability in stable isotopes observed for S. acheilognathi may relate to shifts in the host diet. Yohannes et al. (2017) further indicated that higher variability in stable isotope levels of different organs of the perch (Perca fluviatilis) was related to the metabolic activity of the different organs. In their study, pike liver had higher isotope turnover rates than blood and muscle tissue. Comparison of host fish muscle and liver showed higher standard deviations in liver tissue than in muscle. Yohannes et al.

different environments						
Author	Host	Macroenvironment	Parasite	Developmental stage	Microhabitat	$\Delta^{\rm h} {\rm E}$ (parasite- host)
						Δ^{15} N Δ^{13} C
Boag et al. 1998	Oryctolagus cuniculus	Terrestrial	Cittotaenia denticulata	Adult	Intestine	-2.10 -1.32
			Mosgovoyia pectinata	Adult	Intestine	-3.10 - 3.66
Pinnegar et al. 2001	Gasterosteus aculeatus	Limnetic	Schistocephalus solidus	Plerocercoid	Peritoneal cavity	-2.41 - 0.14
Deudero et al. 2002	Rutilus rutilus	Limnetic	Ligula intestinalis	Plerocercoid	Peritoneal cavity	-1.42 - 0.43
Power and Klein 2004	Gasterosteus aculeatus	Limnetic	Schistocephalus solidus	Plerocercoid	Peritoneal cavity	-1.35 0.13
	Salvelinus fontinalis	Estuarine	Eubothrium salvelini	Adult	Intestine	-2.08 0.01
	Gadus ogae	Marine	Pyramicocephalus	Plerocercoid	Muscle*	-4.43 0.52
Neilson et al 2005	Orvetolaous cuniculus	Terrestrial	phocarum Mosoovovia nectinata	Adult	Intestine	-150 - 550
	annon an Smith		mining in for Scott			
			Cittotaenia denticulata	Adult	Intestine	-2.70 - 1.90
Persson et al. 2007	Salmo salar	Marine	Eubothrium crassum	Adult	Intestine	-2.45 1.61
Sánchez et al. 2013	Artemia	Brackish	Flamingolepis liguloides	Cysticercoid	Haemocoel	0.24 2.82
Navarro et al. 2014	parthenogenetica Chimaera monstrosa	Marine	Gyrocotyle uma	Adult	Gastrointestinal tract (stomach, spiral valve)	-3.33 -1.32
Behrmann-Godel and Yohannes 2015	Perca fluviatilis	Limnetic	Triaenophorus nadulosus	Plerocercoid	Liver	-0.79 - 2.17
Eloranta et al. 2015	Gasterosteus aculeatus	Limnetic	Schistocephalus solidus	Plerocercoid	Peritoneal cavity	-2.20 - 0.40
	Pungitius pungitius		Schistocephalus pungitii	Plerocercoid	Peritoneal cavity	-2.00 0.30
McGrew et al. 2015	Canis lupus	Terrestrial	Taenia krabbei	Adult	Intestine	-2.20 - 2.00
			Taenia hydatigena	Adult	Intestine	

Summary of the isotopic differences ($\Delta^{h}E$ (parasite-host)) for stable isotopes of nitrogen ($\Delta^{15}N$) and carbon ($\Delta^{13}C$) between hosts and cestodes of varying ontogenetic development from

(2017) related this to the higher metabolic activity of the liver compared with the other tissues analysed in their study and thus the same may apply in the present study.

As L. kimberlevensis was also often parasitised by the monogenean parasite, P. ichthvoxanthon, the results from the present analysis were compared to those performed by Sures et al. (2019). Labeobarbus kimberlevensis collected in the current study were of a larger size and more enriched in ¹⁵N than specimens analysed by Sures et al. (2019), but the opposite was observed for ¹³C signatures which were higher in smaller hosts. This finding can be related to the larger size of the yellowfish collected in the present study compared to those collected by Sures et al. (2019). As L. kimberleyensis grow, they undergo a dietary shift from a predominantly omnivorous diet to a predatory one where they feed on smaller fish specimens, even including juvenile yellowfish (Skelton 2001; Sures et al. 2019). This change in food source generally occurs when L. kimberlevensis attain a fork length of approximately 300 mm (Skelton 2001). Fish hosts collected in the present study were larger than 300 mm and therefore had likely undergone a dietary shift which had led to the observed shift in the ¹⁵N signatures. Comparison between hosts from each study showed that the larger L. kimberlevensis in the present study (present study, 700 g; Sures et al. 2019, 430 g) were more enriched in δ^{15} N (present study, 17.5%); Sures et al. 2019, 16.42%) than Largemouth yellowfish analysed by Sures et al. (2019), but δ^{13} C signatures did not differ within the sample from each study.

Comparison of stable isotope signatures between *P. ichthyoxanthon* and *S. acheilognathi* (Table 1; $\Delta^{P}E$ (P.i-S.a)) showed that ¹⁵N enrichment in *P. ichthyoxanthon* was 5.3% greater than in S. acheilognathi which accounts for a difference equal to almost two trophic positions. Differences in δ^{13} C between the parasites accounted for 0.07%, indicating that both parasites infecting L. kimberlevensis derive nourishment from the host and thus share a common food source (Table 1). Differences in stable isotope enrichment in P. ichthyoxanthon and S. acheilognathi can be related to the method of acquiring nutrients by each parasite. Paradiplozoon ichthyoxanthon behaves similarly to a predator and derives nutrients by feeding directly on host blood from the gills. Schyzocotyle acheilognathi on the other hand does not feed on host intestinal epithelium but rather by assimilation of metabolites released as components of bile by the host, as mentioned above. Nachev et al. (2017) observed a similar difference in isotope signatures between the nematode, Eustrongylides sp., and the acanthocephalan Pomphorhynchus laevis, which both infect Barbus barbus. The nematode was more enriched in ¹⁵N relative to the acanthocephalan, and similarly to P. ichthyoxanthon, feeds on host tissue. In contrast, *P. laevis* was depleted in ¹⁵N, and similarly to S. acheilognathi, assimilates by-products from the host across the tegument.

Overall, only a handful of studies have analysed differences in stable isotopes between hosts and cestodes (Table 2). From these studies, it appears that adult and larval stages of cestodes are variably depleted in nitrogen and carbon isotopes relative to their hosts. This variability in the enrichment of cestodes with either isotope may be related to the microhabitat of the parasite and the feeding strategy adopted by this group of parasites. From the enrichment factors of stable carbon and nitrogen isotopes, it is clear that both adult and larval cestodes absorb metabolites from the host and not through active feeding on the host. Most developmental stages of cestodes are depleted in ¹⁵N relative to the host organism indicating that regardless of the developmental stage, cestodes feed through absorptive routes. Adult stages were isotopically depleted relative to the hosts and, similarly, in the case of S. acheilognathi from the present study, this can be related to the food source utilised by this developmental stage as being isotopically depleted. Plerocercoid larvae infecting the peritoneal cavity were mostly depleted in ¹⁵N compared to their hosts but fractionation patterns of $\delta^{13}C$ are mixed compared to their hosts (Pinnegar et al. 2001; Deudero et al. 2002; Power and Klein 2004; Eloranta et al. 2015). However, Flamingolepis liguloides cysticercoid larvae infecting the haemocoel of the host, Artemia parthenogenetica, were isotopically enriched in both ¹⁵N and ¹³C relative to their hosts (Sánchez et al. 2013). The enrichment of larval stages with particular isotopes may point to differences in transmission pathways between systematically different groups of cestodes such as cyclophyllids and bothriocephalids, which have different larval stages, i.e. cysticercoides and procercoids and plerocercoids, respectively.

Conclusion

Results of the present study indicate that the cestode, S. acheilognathi, was isotopically depleted compared to the host fish L. kimberleyensis. This pattern corroborated previous studies on cestodes and can be linked to the source of nutrition being depleted in the heavier isotopes. This results in cestodes occupying at least 0.1-1.3 trophic positions below host. However, this varies between adult and larval stages. Additionally, this indicates that unlike other intestinal parasites such as adult nematodes, cestodes have adopted a somewhat commensalistic and scavenger-type feeding strategy and therefore do not compete directly with the host for nutrients. Stable isotope analysis not only serves as a useful tool for delineating the relationship between different life stages of parasites and hosts in the food web but may also be implemented as a means of tracing transmission pathways of larval parasites between intermediate and definitive hosts. To attain this, additional and intensive comparisons between the nutritional sources of hosts and parasites are required in future studies.

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Additional information All information gathered during the analysis of host and parasite samples are presented within the text.

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