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RESEARCH NOTE

Infection of the lesser spotted dogfish with *Proleptus obtusus* Dujardin, 1845 (Nematoda: Spirurida) reflects ontogenetic feeding behaviour and seasonal differences in prey availability

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Abstract

Proleptus obtusus Dujardin, 1845 is the most common parasite infecting the gut of the lesser spotted dogfish (*Scyliorhinus canicula*, Linnaeus). This nematode is trophically transmitted from an intermediate crustacean host to the definitive elasmobranch host. Sexual and age-related differences in habitat occupancy and feeding behaviour of the lesser spotted dogfish make this parasite-host dyad ideal for testing which aspects of host biology influence parasite transmission. Here, the relationship between *P. obtusus* burden and host condition, sex and age were investigated in lesser spotted dogfish captured in the Northeast Atlantic. Prevalence of *P. obtusus* was of 94.8% with a mean abundance of 23.3 worms per host. Our results indicate that parasite burden is best explained by the interaction between ontogenetic differences in foraging behaviour of the lesser spotted dogfish and seasonal differences in prey availability.

Keywords

Scyliorhinus canicula, parasite trophic transmission, ontogenic feeding behaviour

Introduction

Nematodes are one of the most common pathogens infecting sharks (Garner, 2013). *Proleptus* Dujardin, 1845 (Spirurina: Physalopteridae) is one such nematode genus, with species commonly parasitising the gut of elasmobranchs (Moravec *et al.*, 2002). *Proleptus* spp. rely on predator-prey interactions to complete their life cycle, with early larval stages infecting decapod crustaceans as obligate intermediate hosts but sexual maturity is only attained in the definitive elasmobranch host (Moravec 2007). Clinical signs of infections with *Proleptus* sp. are usually mild and restricted to local inflammation of the stomach wall, although repeated attachment of the nematodes can seriously wound the host (Heupel and Bennett, 1998a).

To the best of our knowledge, six parasitological studies have been conducted on the lesser spotted dogfish, *Scyliorhinus canicula* (Linnaeus), and only three parasitic nematodes were reported, all infecting the digestive tract: adult *Proleptus obtusus* Dujardin, 1845, and third stage larvae of both *Anisakis simplex* (Rudolphi, 1809) and *Pseudoterranova decipiens* (Krabbe, 1878) (see Table I for a summary). *P. obtusus* was the only species reported in all studies and was by far the most common, with 67–100% prevalence and up to 177 parasites per host (Lloret *et al.*, 2012). The reported prevalence of *A. simplex* is low and *P. decipiens* is considered to be rare, so far only found in lesser spotted dogfish from Wales and southern England (see Table I). To date, only one study has tested the relationship between infection intensity and host condition (Lloret

Locality	Number of hosts	Nematode parasites species (Prevalence %)	Reference
Plymouth, UK (NE Atlantic)	37	Proleptus obtusus (100) Anisakis simplex (10.8) Pseudoterranova decipiens (5.4)	Moore 2001
Cardigan Bay, UK (NE Atlantic)	49	Proleptus obtusus (100) Anisakis simplex (2) Pseudoterranova decipiens (53.1)	Moore 2001
Eastern Solent, UK (NE Atlantic)	15	Proleptus obtusus (100) Anisakis simplex (13.3)	Moore 2001
Western coast of Portugal (NE Atlantic)	3	Proleptus obtusus (66,6)	de Oliveira Rodrigues et al. 1973
SW coast of Ireland (NE Atlantic)	28	Proleptus obtusus (100)	Casadevall et al. 2010
Blanes, Spain (NW Mediterranean)	130	<i>Proleptus obtusus</i> (96.9) Anisakidae larvae (0.8)	Casadevall et al. 2010
NW Mediterranean	128	Proleptus obtusus (97.7)	Lloret et al. 2012
NW Spain (NE Atlantic)	57	Proleptus obtusus (91.2) Anisakis simplex (3.5)	Sanmartin-Duran et al. 1989
Galway bay, UK (NE Atlantic)	144	Proleptus obtusus (88.2) Anisakis simplex (4.2)	Henderson and Dunne 1998
Póvoa do Varzim, Portugal (NE Atlantic)	58	Proleptus obtusus (94.8) Anisakis simplex (1.7)	Present study

Table I. Summary from the literature of nematode parasites recorded in the digestive tract of the lesser spotted dogfish

et al. 2012). In that study, involving 128 individuals caught in the northwestern Mediterranean, no relationship was found between host condition, length and parasite intensity. Instead, the authors suggested that parasite intensity was related to feeding frequency and availability of infected prey (intermediate host).

The lesser spotted dogfish occurs throughout the eastern North Atlantic, from Senegal up to Norway, and in the Mediterranean Sea. The species is very popular for human consumption in the Mediterranean Sea (Capapé *et al.* 2008), and it is a common fisheries by-catch in the eastern North Atlantic being the main shark species captured in northern Spain and northern France (Cavanagh 2005). Although populations have generally either remained stable or even increased over the last few decades (Jukic-Peladic *et al.* 2001; UNEP-MAP-RAC/SPA 2013; Coll *et al.* 2014; McHugh *et al.* 2011), declines in lesser spotted dogfish landings have been reported for the Aegean Sea (Damalas and Vasilopolou, 2011), the Adriatic Sea (Barrausse *et al.*, 2014) and the Tyrrhenian Sea (Ferretti *et al.* 2005).

The lesser spotted dogfish is considered a generalist predator feeding mostly on crustaceans and pelagic bony fishes, but there are recognized ontogenetic differences in feeding behaviour with adults consuming more teleosts than juveniles (Olaso *et al.* 2005; Valls *et al.* 2008; Martinho *et al.* 2012; Šantić *et al.* 2012). Martinho *et al.* (2012) also hypothezised that there could be habitat segregation between juveniles and adults during winter off Portugal, based on their different diets at this time of year, in sharp contrast to the other seasons. Other ontogenetic differences have been reported for this species, with juveniles occurring in shallower waters than adults in the Cantabrian Sea (Olaso *et al.* 2005). Sex-related differences in feeding patterns were also observed, with adult females feeding less during spring and summer, in contrast to adult males and juveniles of both sexes that increase their feeding rate during these seasons thus matching the higher abundance of prey (Martinho *et al.* 2012). A high degree of spatial segregation with depth has been described for this species, with males and females in southwest Ireland occupying different depth-related habitats (Sims *et al.* 2001). Importantly, there is a known effect of depth in the diet of the species: individuals captured on the continental shelf feed on reptantians, polichaetes and teleosts, whereas individuals from deeper habitats feed mostly euphausiids (Valls *et al.* 2008).

The present study aimed to assess the gastro-intestinal helminths of lesser spotted dogfish captured in Northern Portugal. Specifically, we hypothesized that sex and ontogenicrelated differences in feeding behaviour and habitat use influence parasite burden (e.g. Sims *et al.* 2001; Martinho *et al.* 2012). We also expected to find seasonal differences in parasite abundance since host feeding activity is reportedly higher in spring and summer due to the higher availability of prey, except in the case of adult females which are expected to decrease feeding activity during this period (Martinho *et al.* 2012).

Materials and Methods

A total of 58 lesser spotted dogfish caught by coastal artisanal fisheries were purchased from a fish auction on five separate

occasions between November 2013 and March 2014 at Póvoa do Varzim (Northern Portugal, see Table II for details). Total length (from tip of the snout to tip of caudal fin, TL) and total weight (TW) were recorded to the nearest millimeter and gram, respectively. Sex of each shark was determined visually by presence/absence of claspers, and maturity stage was determined using the criteria of ICES (2013). Upon dissection, weight of liver and gonads (testes and ovaries) was recorded. Hepatosomatic (HSI) and gonadosomatic indices (GSI) were calculated using the following formulas: HSI = $(LW/TW) \times 100$ and $GSI = (GW/TW) \times 100$. These two indices were used as proxies for overall body condition (e.g. Lambert and Dutil 1997). Stomach and gut contents were emptied into petri dishes and the contents carefully searched for any helminths, which were recovered and stored in 96% ethanol. Parasites were observed under a light microscope (Olympus CX4). Proleptus obtusus were identified based on the presence of a cephalic collar and the conical shape of the caudal region (Moravec et al. 2002). Furthermore, in females, the placement of the vulva was located on the posterior tip of the animal, very close to the anal opening (Moravec et al. 2002). Anisakis simplex were identified following the diagnosis of Hurst (1984). To validate morphological identification, we sequenced a portion of the 18S rRNA gene of 22 randomly selected P. obtusus collected from 5 different hosts (two adult females collected in winter and spring, two adult males collected in winter and one pre-adult female collected in the autumn), for direct comparison with a Proleptus sp. sequence available on GenBank (Accession Number JF934733). The hotSHOT DNA extraction protocol was used (Montero-Pau et al., 2008) as well as the polymerase chain reaction (PCR) procedure of Perera et al. (2013). Finally, since the taxonomy of Proleptus is controversial because of high intraspecific morphological plasticity (Moravec et al. 2002, 2007), and to assess levels of genetic variability, a portion of the 28S rRNA was sequenced from the same individuals. PCR primers were the 28S rD1.2a and 28Sb of Whiting (2002), and the PCR protocol was as follows: preliminary denaturation for 3 min at 94°C, followed by 35 cycles of 30 s at 94°C, 30 s at 55°C and 1 min at 72°C; and a final extension at 72°C for 10 min. Since the 28S rRNA sequences showed no variation (see Results and Discussion of details), we also sequenced the mitochondrial "barcoding" cytochrome oxidase subunit 1 (COI) gene for 15 of the individuals initially analysed for the 18S and 28S rRNA, plus another 39 individuals randomly selected from the remaining hosts. For COI amplification, the primers and PCR conditions were used according to Prosser et al. (2013). Unique haplotypes for 18S rRNA, 28S rRNA and COI were deposited in GenBank (Accession Numbers: KY411561-KY411576).

A generalized linear model (GLM) was used to test the effects of TW, TL, GSI, HSI, host sex, season and maturity stage on parasite abundance. Collinearity between variables was inspected using pairwise correlation coefficients and variance inflation factors (VIF). VIF values were obtained using

Table II. SdevelopmenHepatosom	ummary of d nt stage (subs atic Index; G	lata col adult ar SI- Go	Table II. Summary of data collected for lesser spotted dogfish and <i>Proleptus obtusus</i> sample sizes by sex and season, as well as mean host length, weight and somatic indexes per development stage (subadult and adult). N- total number of lesser spotted dogfish s sampled, with number of subadults given in brackets; TL-Total length; TW- Total weight; HSI-Hepatosomatic Index; GSI- Gonadosomatic Index	potted dogfish number of les x	1 and <i>Proleptus c</i> sser spotted dogf	<i>obtusus</i> samp ish s sampled	ish and <i>Proleptus obtusus</i> sample sizes by sex and season, as well as mean host length, weight and somatic indexes per lesser spotted dogfish s sampled, with number of subadults given in brackets; TL -Total length; TW- Total weight; HSI-	ınd season, as if subadults g	s well as mean h iven in brackets;	ost length, w. TL-Total ler	eight and somation ath; TW- Total	c indexes per weight; HSI-
Host sex Season	Season	Z	N Mean parasite abundance	abundance	Mean TL (mm)	(mm)	Mean TW (g)	W (g)	Mean HSI	ISH	Mean GSI	ISE
			Subadults	Adults	Subadults	Adults	Subadults	Adults	Subadults	Adults	Subadults	Adults

Host sex	Season	z	Host sex Season N Mean parasite abundance	e abundance	Mean TL (mm)	L (mm)	Mean TW (g)	(W (g)	Mean HSI	ISH	Mear	Mean GSI
			Subadults	Adults	Subadults	Adults	Subadults	Adults	Subadults	Adults	Subadults	Adults
	Autumn	2 (0)	1	12 (± 3)	I	549 (± 16)	I	555 (± 33)	I	8.5 (± 0.3)	I	2.7 (± 0.7)
Males	Winter 15 (1)	15(1)	Ι	$16 (\pm 13)$	I	566 (± 23)	I	614 (± 74)	I	7 (± 2.4)	I	$3 (\pm 0.8)$
	Autumn 10 (5)	10 (5)	9.4 (± 5)	7 (± 3)	484 (± 25)	527 (± 8)	462 (± 60)	514 (± 58)	514 (± 58) 13.2 (± 1.9)	19.6 (± 7)	$1.2 (\pm 0.5)$	3.8 (± 1.3)
	Winter	14 (6)	Winter 14 (6) 24.3 (± 22) 15 (± 17)	15 (± 17)	508 (± 15)	515 (± 20)	515 (± 20) 465 (± 61)	547 (± 76)	547 (\pm 76) 10.5 (\pm 3.4) 15.1 (\pm 9) 0.4 (\pm 0.5) 2.7 (\pm 1.5)	15.1 (± 9)	$0.4~(\pm 0.5)$	2.7 (± 1.5)
remaies	Spring	17 (7)	Spring 17 (7) 69.4 (± 45.6) 26 (± 19)	26 (± 19)	501 (± 17)	523 (± 10)	467 (± 62)	502 (± 41)	501 (\pm 17) 523 (\pm 10) 467 (\pm 62) 502 (\pm 41) 8.8 (\pm 1.1)	8.5 (± 2)	$0.9 (\pm 0.7)$	2.6 (± 1.3)

the 'vif' function implemented in R package 'car' (Fox *et al.* 2012). A Poisson regression was fitted to the data and since overdispersion was detected (overdispersion parameter >1) standard errors were corrected using a Quasi-Poisson model (Zuur *et al.* 2009). To find the optimal model the drop1 command was used (Zuur *et al.* 2009).

Results

Table II summarizes morphological and parasitological data collected on lesser spotted dogfish from Northern Portugal. Nematodes were found in 94.8% of sharks (n=58), most in the stomach rather than in the intestine. *Proleptus obtusus* numbers ranged from 1 to 169, with a mean abundance of 23.3. Additionally, two individuals of *Anisakis simplex* were found co-infecting a single host. Host total length (TL) was highly correlated with sex and total weight (TW) (correlation coefficients of 0.7 and 0.8, respectively). Likewise, the gonadossomatic index (GSI) was highly correlated with maturity stage (correlation coefficients of -0.7). All other variables had moderate correlation coefficients ($\leq \pm 0.6$). For this reason both GSI and TL were not included in the analysis. For the remaining variables, the VIF values were below 3 and as such

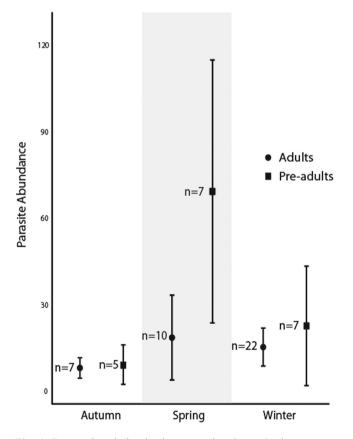


Fig. 1. Seasonal variation in the mean abundance (and corresponding 95% confidence interval) of *Proleptus obtusus* infecting preadult and adult lesser spotted dogfish Numbers of individuals included in each class are also depicted

there was no reason to exclude them a priori (Zuur *et al.* 2009). The drop1 command sequentially excluded variables HSI (p = 0.428), sex (p = 0.139) and TW (p = 0.07). For this reason, the final model retained only the effects of season and maturity stage. Post-hoc comparisons of season × maturity stage groups revealed a significantly higher abundance of parasites in pre-adults in spring (p<0.05, Fig. 1).

Some individuals of *P. obtusus* (61 from a total of 1278) were sequenced for at least one gene. The 18S rRNA sequences (837 bp) were identical to each other, and had 99% identity match with the only *Proleptus* sp. sequence available on GenBank (with 100% query coverage and differing by only 3 point mutations). Likewise, sequences from the 28S rRNA (1060 bp) showed no variation. In contrast, 14 different haplotypes were identified in 54 COI sequences (428 bp), with a total of 14 parsimony informative sites. Uncorrected p-distances between COI sequences varied between 0–3.5%, which is within the range of intraspecific variation reported for several parasitic nematodes (0–4.3%, see Hu and Gasser, 2006 for a review).

Discussion

In line with previous parasitological studies on the lesser spotted dogfish, the prevalence of P. obtusus was high (94.8%) and that of A. simplex was low (1.7%) in the current study. Usually, in vertebrates, larger hosts tend to harbour more parasites (e.g. Lo et al. 1998; Poulin and George-Nascimento 2007), and the higher parasite abundance in larger (likely older) hosts could result from increased feeding rates (i.e. larger sharks eat more intermediate and presumed paratenic hosts) and/or parasite accumulation over the host's lifetime (e.g. Heupel and Bennett 1998a). In previous studies, the abundance of Proleptus worms has been positively correlated with host size, as in the case of Proleptus australis Bayliss, 1933 infecting the epaulette shark, Hemiscyllium ocellatum (Bonaterre) (see Heupel and Bennett, 1998a), Proleptus acutus Dujardin, 1845 infecting the Chilean catshark, Schroederichthys chilensis (Guichenot) (see George-Nascimento and Vergara, 1982), and P. obtusus infecting the dark shyshark, Haploblepharus pictus (Muller and Henle) (see Yeld, 2009). However, no such correlation was found for P. obtusus infecting Haploblepharus edwardsii (Schinz) and Poroderma africanum (Gmelin) captured off South Africa (Yeld, 2009), or the lesser spotted dogfish in the Mediterranean region (Lloret et al., 2012). These disparities in results could be indicative that Proleptus burden is highly correlated to the feeding frequency on infected intermediate hosts, i.e. decapods (e.g. George-Nascimento et al. 1994). In fact, in the cases of the epaulette shark, the Chilean catshark and the dark shyshark, there is a recognized increase in the preference to feed on decapods with age (Dainty 2002; George-Nascimento and Vergara 1982; Heupel and Bennet 1998b), which supports the fact that larger sharks of these species are more heavily parasitized with *Proleptus* sp.. *P. africanum* seems to be an exception, as although dietary information indicates an increase in the consumption of decapods with age (Dainty 2002), Yeld (2009) showed that this preference is not always translated in higher parasitic burden in older/bigger sharks. On the other hand, South African populations of *H. edwarsiii* seem to prefer decapods throughout their entire life (Dainty, 2002), which is concordant with the fact that in these populations no relationship between host size and abundance of *Proleptus obtusus* was detected (Yeld, 2009).

The results from the current study showed that smaller/subadult sharks were more heavily infected than larger adults (see Figure 1), hence supporting our initial hypothesis that ontogenetic differences in foraging behaviour could translate in different parasite abundances. Crustaceans are the intermediate hosts of *Proleptus* (see Moravec, 2007) and the main prey of lesser spotted dogfish (Olaso *et al.* 2005; Valls *et al.* 2008; Martinho *et al.* 2012; Šantić *et al.* 2012). However, adult sharks consume a higher proportion of teleosts than juveniles (Olaso *et al.* 2012), and thus could be less prone to infection. Additionally, the impact of depth segregation with ontogeny on the feeding strategy of this shark, and thus on parasite abundance, cannot be discarded (Olaso *et al.* 2005; Valls *et al.* 2005; Valls *et al.* 2005; Valls *et al.* 2005; Valls

Habitat use and behavioural segregation between sexes occurs in lesser spotted dogfish with males resting during the day in deep waters and feeding during the night in shallow areas, whereas females rest during the day in shallow waters and feed at night in deeper habitats (Sims et al. 2001). Additionally, adult males and juveniles of both sexes are reported to feed more during spring and summer due to higher prey availability whereas mature female sharks decrease feeding activity during this period (Martinho et al. 2012). In contrast to what we expected, host sex did not affect parasite abundance, indicating that both males and females are feeding with the same intensity on intermediate hosts of P. obtusus despite potential differences in habitat use. Likewise, no significant effects on parasite burden were associated with adult female hosts. However, juveniles did show higher parasite abundances during spring, which is probably a consequence of a higher feeding activity perhaps triggered by higher abundance of prey (including intermediate hosts) in spring (Martinho et al. 2012). Finally, in the present study there was no effect of host somatic indices on parasite abundance, which indicates that host condition does not affect infection ability of parasites.

Overall, our data confirms some of the previous observations made by Lloret *et al.* (2012) on Mediterranean lesser spotted dogfish, as no relationship was found between the abundance of *P. obtusus* and host sex or somatic indexes, which serve as a proxy for host condition. Our findings suggest that parasite load is best explained by the interaction between age-related foraging behaviour and seasonal differences in prey availability. Acknowledgements. The authors would like to thank Dr Antigoni Kaliontzopoulou for all the help regarding the statistical analysis. The authors would also like to thank two anonymous reviewers for all the comments which helped improve this manuscript. This work was partially funded by the European Regional Development Fund (ERDF) through COMPETE program and by National Funds through FCT -Foundation for Science and Technology (projects EXPL/MAR-BIO/1034/2012, PTDC/MAR-BIO/4458/2012 and FCOMP-01-0124-FEDER- 029939); and partially funded by Norte Portugal Regional Operational Programme (NORTE 2020), under the PORTUGAL 2020 Partnership Agreement, through ERDF under project MarInfo (NORTE-01- 0145-FEDER- 000031). RX and AV are supported by FCT under the Programa Operacional Potencial Humano - Quadro de Referência Estratégico Nacional funds from the European Social Fund and Portuguese Ministério da Educação e Ciência (RX. IF-FCT contract IF/00359/2015; AV post-doctoral grant SFRH/BPD/77487/ 2011).

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