

Nematode infection patterns in a Neotropical lizard species from an insular mountain habitat in Brazil

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Abstract

Neotropical lizards are known to harbour rich nematode parasite faunas; however, knowledge of the diversity and patterns of infection are still lacking for many species. This is true for the genus *Tropidurus*, in which data on patterns of parasitism are known for only approximately 11 of its 30 species. We show that the nematode fauna associated with a population of *Tropidurus montanus* is composed of three species of host-generalist parasites with high overall prevalence. Male and female lizards did not differ in infection pattern and there was no relationship between host body size and intensity of infection for the most prevalent parasite species. Nevertheless, overall prevalence changed seasonally, with a higher proportion of parasitized individuals being found in the dry period than in the rainy period. We discuss our findings in the context of diet patterns of *T. montanus*, which we suggest may explain the similarities in prevalence and intensity of infection between the sexes. In addition, seasonal changes in diet are considered to be related to the observed differences in prevalence between dry and rainy periods.

Introduction

The major climatic factor influencing ecological systems in the Neotropics is seasonal variation between dry and rainy periods. Changes in the amount of rain and in duration of the rainy period can affect patterns of infection spread and persistence (see Altizer *et al.*, 2006). Therefore, parasitic interactions involving Neotropical species are expected to change over time in response to periodic

variation in rainfall, which can, in turn, result in seasonal patterns of infection. Lizards are ectotherms, thus adjusting their activity patterns and behaviours according to environmental conditions (e.g. Hatano *et al.*, 2001; Filogonio *et al.*, 2010), and so patterns of helminth infection can be expected to vary seasonally as well, reflecting the behavioural adjustments of these hosts (Ribas *et al.*, 1995; Vrcibradic *et al.*, 1999; Pereira *et al.*, 2012; Brito *et al.*, 2014).

Lizards are important components of the life cycle of many species of helminths, serving as definitive or intermediate hosts (e.g. Ávila & Silva, 2010). The helminth fauna that parasitizes lizards is diverse and with varied patterns of infection. In this sense, helminth richness can vary geographically among populations (Brito *et al.*,

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2014; Galdino *et al.*, 2014), and changes in the composition of component communities can be unrelated to geographical distance among populations (Bezerra *et al.*, 2016). Additionally, some studies have shown divergent results when considering whether infection rates and parasite loads of helminths vary between sexes, or whether parasite load is related to host body size. Hence, information is still needed for a more complete understanding of the epidemiological and ecological patterns of helminth infection in lizards.

The genus *Tropidurus* is speciose, with 30 species (Uetz & Hošek, 2016). The genus harbours a rich helminth fauna with nearly 30 different parasite species and with component communities hosting from 3 to 21 species (Ávila & Silva, 2010; Anjos *et al.*, 2012). Recently, there have been efforts to uncover the diversity and patterns of helminth infection of species of *Tropidurus* (e.g. *Tropidurus itambere*, Ávila *et al.*, 2011; *T. hygomi* and *T. psammonastes*, Lambertz *et al.*, 2012; *T. spinulosus*, Lunaschi *et al.*, 2012; *T. torquatus*, Pereira *et al.*, 2012; *T. oreadicus*, Santos *et al.*, 2013; *T. hispidus* and *T. semitaeniatus*, Brito *et al.*, 2014); however, parasitological data are missing for approximately 60% of the species of the genus.

The occurrence of the lizard *T. montanus* is restricted to the rocky outcrop formations of the mountaintop habitats of the Serra do Espinhaço (Espinhaço mountain range) in South America (Rodrigues, 1987). To the best of our knowledge, information on parasitism of this species by helminths is lacking. The aim of the present study was to describe the helminth species composition and infection levels in the Neotropical lizard *T. montanus*, relative to seasonality, host sex and size.

Materials and methods

We conducted the study in the rocky outcrop formation at c.1460 m above mean sea level in the Serra da Piedade (Piedade mountain range; 19°49'S; 43°40'W), a branch of the Espinhaço Range, in the state of Minas Gerais, Brazil. High-elevation open habitats of the mountains from eastern Brazil (which include the Espinhaço Range and related branches) are considered as isolated insular environments that play a role in biogeographical processes of plant and animal species (Chaves *et al.*, 2015; Neves *et al.*, 2016). The mean annual temperature is 21°C and the mean annual rainfall averages 1650 mm. The region experiences a marked distinction between rainy (from October to March) and dry (from April to September) periods (Alvares *et al.*, 2014).

Sampling of lizards took place during the rainy (December 2012) and dry months (June to August 2013). Lizards were sampled by performing visual encounter surveys throughout the period of activity of the species, and captured by noose or by shooting rubber bands. Captured lizards were killed by lethal injection of sodium thiopental, measured for body size (snout-to-vent length) with a Vernier caliper (nearest 0.1 mm), fixed and preserved following standard procedures, and deposited in the herpetological collection of the Museu de Ciências Naturais of Pontifícia Universidade Católica de Minas Gerais (MCNR 4893–4934; 5114–5145; 5159–5174), Belo Horizonte, Brazil.

Post-mortem examinations of the body cavity, digestive tract, liver and lungs were undertaken using a stereomicroscope. Nematodes were cleared with Aman's lactophenol (Andrade, 2000) and analysed using a computerized system for image analysis LAS 5.0 (Leica Application Suite), adapted to a DM 2500-Leica microscope with a differential interference contrast system (Leica, Wetzlar, Germany). All the parasitological terms used followed Bush *et al.* (1997). Worms were deposited in the helminthological collection of the Instituto de Biociências de Botucatu, Universidade Estadual Paulista (species batches: CHIBB 7861; CHIBB 7862; CHIBB 7863), Botucatu, Brazil.

Differences between the sexes in overall helminth prevalence were tested by using the binomial test for each of the periods (rainy and dry). The binomial test was also used to test for differences in prevalence between the two periods, considering all individuals in a period regardless of sex. To evaluate the effects of period (dry and rainy) and sex on the overall intensity of helminth infection we used a two-way analysis of variance (ANOVA). Effects of body size on intensity of infection were evaluated for the helminth species with higher prevalence by conducting a Generalized Linear Mixed Models analysis considering each individual as a random effect variable. In addition, we used a one-factor permutation test to evaluate the effect of period on infection intensity of the more prevalent helminth species (Hothorn *et al.*, 2008).

Results

The helminth component community of *T. montanus* ($n = 80$) was found to be composed of three species: *Parapharyngodon alvarengai* Freitas, 1957 (Pharyngodontidae) (prevalence = 59%; mean intensity of infection = 5.6 ± 5.1), found in both large and small intestines; *Physaloptera* sp. Rudolphi, 1819 (Physalopteridae) (prevalence = 12%; mean intensity of infection = 4.9 ± 8.2), found in the larval stage in the stomach and large intestine, and *Strongyluris oscar* Travassos, 1923 (Heterakidae) (prevalence = 44%; mean intensity of infection = 6.0 ± 7.2) found in lungs, large and small intestines. Therefore, *P. alvarengai* and *S. oscar* were the most prevalent nematodes found parasitizing individual lizards. Of the total of 80 lizards sampled, 64 were parasitized by at least one species, resulting in an overall prevalence of 80%. Regarding infracommunities, 64% of lizards were parasitized by a single parasite species, while 30% of the hosts were parasitized by two and 6% by three species. Nematode abundance averaged 7.4 ± 6.3 . The intensities of infection by *P. alvarengai* and by *S. oscar* were not related to host body size ($P = 0.07$, $n = 52$ and $P = 0.60$, $n = 50$, respectively).

We sampled 17 males and 20 females of *T. montanus* in the rainy period and 21 males and 22 females in the dry period. In the dry period, 95% of males and 100% of females were found to harbour helminths, with no difference in overall prevalence between the sexes ($P = 0.98$). During the rainy period, 70% of males and 50% of females were found to be parasitized and there was no difference in prevalence between males and females for the rainy period ($P = 0.34$). The population of *T. montanus* had higher overall prevalence during the dry period (97%) than

during the rainy period (59%) ($P < 0.001$). The overall intensity of helminth infection did not differ between seasons or between sexes (two-way ANOVA: $F = 0.03$, $P = 0.85$). The intensity of infection by *P. alvarengai* and *S. oscari* did not vary between dry and rainy periods (one-factor permutation test, $Z = 1.52$, $P = 0.14$ and $Z = -1.18$, $P = 0.26$, respectively).

Discussion

The nematode community parasitizing *T. montanus* was composed of three species. Of these, the nematode *P. alvarengai* (Pharyngodonidae) is known to infect other lizard species, including *Ameiva ameiva* (Padilha & Duarte, 1979), *Trachylepis atlantica* (Ávila & Silva, 2010), *Hemidactylus agrius* (Anjos *et al.*, 2011), *Urosaurus auriculatus* (Goldberg & Bursey, 2012), *Tropidurus hispidus* (Brito *et al.*, 2014; Galdino *et al.*, 2014) and *Tropidurus semitaeniatus* (Bezerra *et al.*, 2015). The nematode *S. oscari*, is considered to be a host generalist (i.e. parasitizing many host species), with a heteroxenic life cycle with arthropods as intermediate hosts (Barreto-Lima & Anjos, 2014), and is a common parasite of lizards of the genus *Tropidurus* (Ávila & Silva, 2010), having been found in *T. guarani*, *T. spinulosus*, *T. torquatus*, *T. melanopleurus* and *T. hispidus* (Bursey & Goldberg, 2004; Ávila & Silva, 2010; Brito *et al.*, 2014). Moreover, *S. oscari* is known to infect lizards of other taxa, such as *Ameiva ameiva*, *Anolis fuscoauratus*, *A. punctatus*, *A. transversalis*, *Enyalius bilineatus*, *Eurolophosaurus nanuzae*, *Brasiliscincus agilis*, *Plica plica* and *P. umbra* (Fontes *et al.*, 2003; Ávila & Silva, 2010; Barreto-Lima & Anjos, 2014). The third parasite species found, *Physaloptera* sp., is known to parasitize *A. ameiva*, *Glaucomastix littoralis*, *Ameivula ocellifera*, *H. mabouia*, *B. agilis*, *Psychosaura macrorhyncha*, *Polychrus acutirostris*, *Tropidurus etheridgei*, *T. torquatus*, *T. hispidus*, *Salvator meriane* and *Tupinambis teguixin* (Ávila & Silva, 2010; Ávila *et al.*, 2012; Pereira *et al.*, 2012). Therefore, the component community of helminths parasitizing the lizard *T. montanus* can be considered to be comprised of host generalist parasites, and our record extends hosts species for *P. alvarengai* and *S. oscari*.

The component community of parasites of *T. montanus* is species poor when compared to those found for other species of the genus (e.g. *T. torquatus* (Pereira *et al.*, 2012), *T. hispidus* (Brito *et al.*, 2014; Galdino *et al.*, 2014) and *T. semitaeniatus* (Brito *et al.*, 2014; Bezerra *et al.*, 2015)). The high-elevation formations of the mountains from eastern Brazil, such as the rocky outcrops from the highlands of the Espinhaço Range, are considered to be an archipelago of isolated insular mountaintop habitats (e.g. Chaves *et al.*, 2015). Therefore, the insularity of the mountaintop rocky outcrops of the Serra da Piedade is remarkable as this mountain is apart from the main Espinhaço Mountain Range where *T. montanus* is distributed. The helminth fauna of hosts inhabiting insular environments can be expected to be poorer than those parasitizing hosts on the mainland (e.g. Dobson *et al.*, 1992). For example, the pattern of helminths infecting rodent species differed between insular and continental populations, with insular populations having lower helminth richness (Kuhnen *et al.*, 2012). The same pattern

seems to hold true for lizards, with species-poor helminth infracommunities being found on hosts from insular environments (Dobson *et al.*, 1992; Martin & Roca, 2005). Therefore, the insular nature of the sampled area might constrain the available pool of helminth species that can infect individuals of *T. montanus*.

We found no intersexual difference in overall parasite prevalence. Intersexual differences in parasite prevalence are generally related to distinct social roles of male and female lizards, which in turn are associated with unequal blood hormone levels (Fuxjager *et al.*, 2011; see also Anjos *et al.*, 2012). Males of *T. montanus* use larger home ranges than females (Galdino *et al.*, in prep.), which is a proxy for different social roles between the sexes. Thus, one might expect that the use of larger areas by males might expose them to greater chances of infection, at least by *P. alvarengai*, a monoxenous parasite species. Therefore, our results cannot be explained by the different social role and/or space use of males and females of *T. montanus*. Concerning heteroxenous parasites, we regarded the similarities in prevalence between the sexes as being due to a broad diet overlap between males and females, as shown by Kiefer (1998). Therefore, the similarity in overall prevalence of *S. oscari* between the sexes might be related to males and females having the same dietary patterns.

We did not find any relationship between *T. montanus* body size and intensity of infection of either *P. alvarengai* or *S. oscari*. It is argued that hosts with a larger body size would provide more space for helminths to colonize, thereby leading to a positive relationship between the two variables (Poulin, 1997; Ribas *et al.*, 1998). It is also suggested that changes in infection intensity with size will be related to ontogenetic changes in diet (Fontes *et al.*, 2003). In the case of *T. montanus*, young and adult lizards have similar diets, and so the non-relationship between intensity of infection and lizard body size might be associated with ontogenetic conservatism in diet, which in turn might expose both young and adults to the same chances of infection. A relationship between lizard body size and intensity of infection was not found for other species of lizards (e.g. *T. torquatus* and *Brasiliscincus* (former *Mabuya*) *agilis* (Van Sluys *et al.*, 1997), *T. hispidus* (Galdino *et al.*, 2014) and *T. semitaeniatus* (Bezerra *et al.*, 2015)), suggesting that, in general, there is no trend between body size and intensity of infection for lizards.

Our results indicated that some patterns of helminth infection in lizards may vary seasonally. We found that the proportion of infected hosts was higher during the dry period than during the rainy period. Seasonal variation in helminth infection patterns has also been found for the Neotropical *T. torquatus*, with larger overall intensity of infection during the dry period (Pereira *et al.*, 2012). In the case of *T. torquatus*, Pereira *et al.* (2012) suggested that interseasonal differences in overall intensity of infection might be related to the consumption of a plant species with anthelmintic compounds. Although *T. montanus* increases its consumption of plant material during the dry period (Kiefer, 1998), we are not able to relate the observed seasonal variation in overall helminth intensity directly to plant ingestion. Alternatively, as the diet composition of *T. montanus* varies seasonally (Kiefer, 1998), the observed change in overall prevalence in our

study might be related to seasonal changes in the types of ingested food items.

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Conflict of interest

None.

Ethical standards

We followed the *Guidelines for the Euthanasia of Animals* (2013) of the American Veterinary Medical Association for euthanizing the lizards. The Instituto Chico Mendes de Conservação da Biodiversidade issued the permits (SisBio 35541 and 37266-3) to collect the animals.

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