

Helminth community structure in the Argentinean bufonid *Melanophryniscus klappenbachi*: importance of habitat use and season

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Abstract The goal of this study was to evaluate the relative influence of terrestrial habits, season, and host body size on the species richness and abundance of helminth parasites in the toad *Melanophryniscus klappenbachi*, for which a greater abundance of nematode parasites was expected. A total of 90 toads were collected in the Chaco Province, Argentina. The helminth community found in infected toads included 17 taxa and was dominated particularly by larval parasites. Contrary to our expectations, nematode species showed lower values of infection parameters. Infected toads harbored a maximum of seven species, and the mean helminth richness was 3.16 ± 1.66 species per infected toads. Season played a significant effect on determining the species richness and abundance of the parasite infracommunity. Similarly, the prevalence of infection of several helminth species (8/47 %) varied greatly over time. Host body size was the main factor in determining the infrapopulation structure of helminth parasites. Species richness was significantly and negatively correlated with host body size. Strong associations were observed mainly between larvae of some species. The transmission strategies of parasites suggest that this bufonid acquires infections through direct contact with larval parasites from aquatic and terrestrial habitats and by ingestion of infective larvae. The characteristic of the host tegument, such as the presence of alkaloids, could significantly contribute to the low occurrence of infection by skin-penetrating nematodes. Results also showed that diets and mobility of the host and the local microhabitat conditions play an important role in parasitic infections of toads.

Keywords Helminth community · Season · Habitat use · *Melanophryniscus klappenbachi* · Chaco Province

Introduction

In the study of the helminth community, no single factor is generally responsible for the level of parasite in a community. One of the most important considerations is the range of life cycles involved: some parasites lack intermediate hosts, others move sequentially among different host species to complete their life cycles, and several larval stages develop in different aquatic and terrestrial microhabitats. Thus, the helminth community depends a lot more on transmission, i.e., parasites vary in how they are acquired by the host, penetrate, or get eaten. This would suggest that the life history of the host would determine where and how it picks up its helminths through its diet or spending time in the water or on land (Esch 1971). A small change in the parasite transmission level causes great changes in the number of adults as well as changes in the parasite community structure (Marcogliese and Cone 1997; Esch et al. 2002). Kennedy (1975) emphasized the importance of the host diet in the control of infection level, but its seasonal changes, which are indirectly influenced by temperature, and the seasonal maturation cycle among parasites induce seasonal changes in the levels of infection. Chubb (1979) also indicated that the development of helminth parasites can show a marked seasonality in mid-latitude climates, i.e., the cold season causes changes in acquisition of the parasite by the host. However, these results may also vary among regions with subtropical climates, where the temperature is not a significant factor to determine strong seasonal patterns of helminth development. For instance, ecological studies regarding helminth parasites in Argentinean vertebrate hosts from subtropical environments have shown

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that there is more time for amphibian hosts to become infected during the year because there is no or little freezing temperature unlike the temperate region of the world (Hamann 1999, 2006).

Other important factors in the helminth community structure are species richness and their relative abundance, which are specifically related to the interactions between helminth parasites at the infracommunity level. In this sense, Holmes and Price (1986) considered that interactive communities are species-rich, with species regularly cooccurring at densities such that there is often evidence of interactions between the species, and that isolationist communities are species-poor, with species being independent of each other. Thus, communities may be located anywhere along this continuum, and those that parasitize amphibians tend to be found towards the isolationist, i.e., show little evidence of parasitic interactions (Aho 1990; Muzzall 1991; Bolek and Coggins 2003; Luque et al. 2005; Yoder and Coggins 2007; Ibrahim 2008; Hamann et al. 2006a, b, 2010, 2012, 2013a). Notably, the relationships between the richness species and host body size and between host mobility and ability to be in different microhabitats, together with the ability to feed on different intermediate host species (which leads to a greater likelihood of acquiring food-borne parasites), provide a particular structure to parasite species communities.

In recent decades, the study of communities of toad amphibian parasites and host-parasite relationships has gained interest in diverse regions of South America (Burse et al. 2001; Goldberg and Bursey 2003; Iannacone 2003; Luque et al. 2005; Lux Hoppe et al. 2008; Pinhão et al. 2009; Santos et al. 2013). Hamann et al. (2013a) have previously reported the first study of the ecology of helminth parasite communities in the Argentinean toad amphibian *Rhinella fernandezae* (Gallardo, 1957). The present study was also performed in bufonids but with different habitat use and different host behavior (e.g., adult toads have diurnal habits and live on dry land with a predominance of terrestrial bromeliads in the Chaco region), which consequently provide the parasite communities and indicate the diversity of intermediate and definitive hosts in this local environment (Poulin and Morand 2004). No taxonomic and ecological studies have been carried out on helminth infections in this bufonid toad.

Based on certain assumptions, the approach used in this current study predicts (1) that the terrestrial habit of toads determines greater abundance of nematode fauna, (2) that the short winters allow the development of helminths most of the year, and (3) that the helminth community is influenced by the diet and size of the host. These premises were tested by analyzing the definitive host *Melanophryniscus klappenbachi* Prigioni & Langone (2000), a member of the family Bufonidae. *M. klappenbachi* is a typical amphibian species of the Chaco region, and its distribution in Argentina covers Chaco, Formosa, Santa Fe, and Santiago del Estero provinces

(Frost 2014). This species has typically diurnal habits and breeds in temporary aquatic environments. Several breeding periods occur during the rainy season. After heavy rainfall, there is an explosive breeding, and a large number of individuals congregate in these sites. The couples perform multiple ovipositions, depositing eggs in small groups that are attached to submerged vegetation by the male. These toads are small to medium in size (22 to 34 mm) and have aposematic coloration type, with patches of yellow, red, and orange on dark backgrounds. The behavior associated with this type of color is called “unken-reflex.” When facing potential predators or capture, the specimens have members upwards, bending the body and exposing the ventral spots (see Manzano et al. 2004). A large number of lipophilic alkaloids (termed “dendrobatid alkaloids”) with pharmacological potential have been isolated from the skin glands of these amphibians. These substances, which are either synthesized de novo or obtained from their diet (see Daly et al. 2008), are considered to provide a defense against predation during the mainly diurnal breeding episodes of toads (Santos and Grant 2011). Feeding studies indicate that adults feed mainly on ant and scarce acari (unidentified mites) (Daly et al. 2008).

The objectives of this study were (1) to determine the richness and diversity of helminth parasites at the component and infracommunity levels, (2) to analyze the helminth cycles of transmission, (3) to examine the species affinities (covariation and association) of helminth infracommunities, and (4) to evaluate the influence of season on the abundance and number of parasitic helminth species in the toad *M. klappenbachi*.

Materials and methods

Study area

The study area (50 ha) is located within the Chaco biogeographic province, Chaco Province, Argentina (27° 23' S, 58° 58' W). The habitat is characterized by deciduous dry woodland with herbaceous grasses and numerous cacti and terrestrial bromeliads. The river bank is dominated by alder forests (palo bobo) or creole sauce or gallery forest logging. The forest vegetation consists of quebracho (*Schinopsis balansae*), urunday (*Astronium balansae*), and ñandubay (*Prosopis affinis*), with herbaceous strata composed of areas that include gramineous (*Elionurus muticus*) and terrestrial bromeliads (*Aechmea distichantha*, *Bromelia serra*). The mean annual temperature is between 20 and 23 °C and the mean annual precipitation between 500 and 1,200 mm (Cabrera and Willink 1980). Table 1 shows the values of temperature and precipitation by seasons during sampling period, in the study area.

Table 1 Summary of seasonal data for temperature minimum/maximum (mean±standard deviation), total precipitation, and *Melanophryniscus klappenbachii* body size from Chaco, Argentina

Variables	Autumn	Winter	Spring	Summer
Abiotic				
Temperature (°C)	1.7/34.1 (20.5±4.0)	-0.4/39.5 (18.2±5.7)	2.5/40.0 (24.4±4.1)	12.5/40.0 (25.4±3.3)
Precipitation (mm)	252	141	385	599
Biotic				
Toad length (mm)	22/34 (28.7±3.1)	28/34 (30.6±1.9)	23/35 (29.2±2.4)	25/31 (28.6±1.9)
Toad weight (g)	1.1/4.8 (2.2±31.0)	1.0/4.8 (2.6±1.0)	1.0/4.8 (1.9±0.8)	1.4/2.3 (1.9±0.2)

Collection and examination of host and parasites

Adult specimens of *M. klappenbachii* were collected in autumn 2011 ($n=21$) and in winter ($n=22$), spring ($n=26$), and summer ($n=21$) 2012–2013. Specimens were hand-captured always by two people between 1400 and 1700 hours. Toads were transported live to the laboratory and killed in a chloroform (CHCl_3) solution. Their snout-vent length (SVL) and body weight were recorded. At necropsy, hosts were sexed and the esophagus, stomach, gut, lungs, liver, urinary bladder, kidneys, body cavity, musculature, skin, and brain examined for parasites. Helminths were observed *in vivo*, counted, and killed in hot distilled water and preserved in 70 % ethyl alcohol. For identification, the digeneans, cestodes, and acanthocephalans were stained with hydrochloric carmine, cleared in creosote, and mounted in Canada balsam. Nematodes were cleared in glycerine or lactophenol and examined as temporary mounts. Helminths were deposited in the Helminthological Collection of Centro de Ecología Aplicada del Litoral (CECOAL) Consejo Nacional de Investigaciones Científicas y Técnicas (CONICET), Corrientes city, Corrientes, Argentina [accession numbers CECOAL, 12082218, *Glythelmins palmipedis* (Lutz, 1928); 12082217, *Catadiscus longicoecalis* (Caubisens Poumarau, 1965); 12082208, *Styphlodora* sp.; 13031117, *Bursotrema tetracotyloides* (Szidat, 1960); 12082201, *Travtrema* aff. *stenocotyle* (Cohn, 1902); 12082215, *Opisthogonimus* sp.; 13031107, *Cylindrotaenia* sp.; 12082207, *Mesocercoides* sp.; 11051215 and 12082204, *Cosmocerca* sp. 1; 12082202, *Cosmocerca* sp. 2; 11051202, *Physaloptera* sp.; 13031118, Seuratoidea gen sp.; 13031103, *Spiroxys* sp.; 13031116 Rhabdochoniidae gen. sp.; 12113009, *Centrorhynchus* sp.]. Toads were deposited in the Herpetology Collection of CECOAL-CONICET, with accession number LHC, 5120. The approval for using animals in this research was given by the Secretary of Natural Resources, Department of Fauna and Protected Areas from Production Ministry, Chaco Province, Argentina.

Statistical analysis

The infection prevalence, intensity, and abundance were calculated for helminths according to Bush et al. (1997). The measures of community richness and diversity employed included the following: the total number of helminth species (=richness), Shannon index (H') (Shannon and Weaver 1949), and evenness (J') as H'/H' maximum (Zar 2010). The Brillouin index (BH) and evenness (E) were used at the infracommunity level (Pielou 1966). The diversity index was used with decimal logarithms (\log_{10}). Species richness is the number of helminth species, and mean helminth species richness is the sum of helminth species, per individual toads, divided by the total sample size. A distribution of infracommunity richness values was tested to see if they indicate some sort of structure between the numbers of helminth species vs. number of toads. Berger-Parker index of dominance (d) was used to determine the most abundant species (Magurran 2004). The Kruskal–Wallis test, and posteriori Dunn tests (Zar 2010), was used to compare richness of helminths between seasons. The probabilities were calculated according to the Bonferroni procedure (Hochberg 1988), because it provides great control over type I error.

Helminth communities have been classified at the infracommunity (all helminth infrapopulations within a single *M. klappenbachii*) and component community (all helminth infracommunities within a population of the *M. klappenbachii*) levels (Bush et al. 1997). Comparison of k proportion (χ^2) was used to test for differences in prevalence of helminths between seasons. Kruskal–Wallis test (H) was used to test for differences in size of toads between seasons. Spearman's rank test (r_s) was used to indicate the relationship between host body size, parasitic abundance, and infracommunity descriptors. Spearman's rank test (r_s) was used to calculate possible species covariation. The species associations was analyzed with a chi-square test (χ^2), with Yates correlation. The degree of aggregation of different species of parasites was calculated by the relation variance (s^2) and mean (\bar{x}). The helminth community structure was examined according to the methodology outlined by Thul et al. (1985), in which helminth species are classified into four groups (dominant, codominant,

subordinate, and unsuccessful) by taking into account their prevalence, intensity, and maturity factor (equal to 1.0 if at least one mature specimen of species j is found and equal to 0 if otherwise), which is related to the degree of host specificity.

The non-metric multi-dimensional scaling (NMDS), an ordination technique, was used to study the pattern in helminth community structure based on abundance of parasite species composition. For that, a similarity matrix was constructed based on the Bray–Curtis measure. The abundance of infection of each parasite species in each host and its relation with season was analyzed through a one-way analysis by means of a non-parametrical permutational ANOVA (PERMANOVA) test (Anderson et al. 2008).

For abundance vs. host body size and covariation and associations of parasites, the species considered were those that had at least 8 % occurrences in the amphibian population (seven species). The softwares used were Xlstat 7.5 (Addinsoft 2004) and Primer v6 (Anderson et al. 2008).

Results

Helminth community characteristics

The helminth component community for this toad population consisted of 17 species of helminths (Table 2). The predominant groups of parasites were trematodes (8/47 %) followed by nematodes (6/35 %); the other groups of parasites were represented by very few species (two Cestoda and one Acanthocephala). Helminth diversity (H') and evenness (J') were 0.59 and 0.48, respectively. *Centrorhynchus* (L.) was the most abundant species ($d=0.57$). Most (94 %) of the helminth parasite species of *M. klappenbachi* showed an aggregated pattern of distribution. Five species (29 %) were reported at adult stage and use *M. klappenbachi* as definitive host, and 12 species (71 %) were reported at larval stage and infect this bufonid toad as intermediate or paratenic hosts. Parasite transmission occurs to the toad host by skin penetration (53 %) and oral ingestion (47 %) (Table 2). Parasites were found in different sites of infection in the host (Fig. 1). The most parasitized organs in this host were the serous membrane of the stomach with *Centrorhynchus* sp. (12 %), followed by the kidneys with *Styphlodora* sp. (7 %), and the small intestine with *G. palmipedis* (7 %) (Fig. 1).

At the infracommunity level, the mean helminth richness was 3.16 ± 1.66 (maximum=7) species per infected toad. Multiple infections were observed in 79 % of toads, with 2, 3, 4, 5, 6, and 7 species occurring in 15, 9, 24, 19, 3, and 1 toads, respectively. The distribution of helminth infracommunity species richness observed did not show good fit to Poisson distribution ($\chi^2=21.21$, $df=6$, $P=0.002$). Mean

helminth value of diversity (HB) and evenness (E) were 0.25 ± 0.16 and 0.52 ± 0.32 , respectively.

Interspecific relationships in the infracommunity

Five correlations between helminth species occurrence were significant (three were negative and two were positive) (Table 3). Three associations were found among the seven helminth species considered: *Styphlodora* sp./*C. longicoecalis* ($\chi^2=4.70$, $df=1$, $P<0.05$), *Mesocestoides* sp./*Styphlodora* sp. ($\chi^2=3.85$, $df=1$, $P<0.05$), and *Centrorhynchus* sp./*Styphlodora* sp. ($\chi^2=21.26$, $df=1$, $P<0.05$).

Parasitic infection related to size of host

The infection prevalence of the 90 toads examined was 97 %. Total length of the toads ranged from 22.00 to 34.50 mm (29.30 ± 2.37), and weight ranged from 1.00 to 4.80 g (2.12 ± 0.83). The abundances of helminth species and ecological descriptors were primarily not significantly correlated ($P>0.05$) with host body size. Nevertheless, significant negative correlation was observed for *Styphlodora* sp. vs. length of the host ($r_s=-0.26$, $n=86$, $P<0.05$).

There were no significant differences in the weight of frogs between season (Kruskal–Wallis $H=6.83$, $df=3$, $P=0.08$), but the length of frogs were different between seasons (Kruskal–Wallis $H=9.04$, $df=3$, $P=0.03$). In the last case, posteriori multiple pairwise comparisons using Dunn's procedure were calculated. These data showed that the length of frogs was only different between winter and summer ($P<0.05$) (Table 1). Infection related to host body size per season showed predominantly a negative correlation with host body size. Significant negative correlation was observed for species richness vs. host length (winter $r_s=-0.46$, $n=22$, $P<0.05$; spring $r_s=-0.49$, $n=26$, $P<0.05$), and a significant negative correlation was also observed for abundance of helminths vs. host length (*Cosmocerca* sp. 1/autumn $r_s=-0.54$, $n=17$, $P<0.05$; *C. longicoecalis*/ winter $r_s=-0.47$, $n=22$, $P<0.05$; *Styphlodora* sp./spring $r_s=-0.46$, $n=26$, $P<0.05$). Nevertheless, significant positive correlation was observed for *G. palmipedis* vs. length of the host (summer $r_s=0.53$, $n=21$, $P<0.05$).

Importance of species within the community

Helminth species were classified according to community importance values (Table 2); two species were strongly characteristic of the community (e.g., *C. longicoecalis*), and two species contributed significantly to the community, although to a lesser extent (e.g., *Cosmocerca* sp. 1). One species occurred infrequently and did not contribute significantly to the community (e.g., *Cylindrotaenia* sp.), and 12 that are able to enter the host but not to attain maturity therein contributed

Table 2 Helminths recorded in *Melanophryniscus klappenbachi*

Helminth species	<i>t</i>	<i>I</i>	<i>d</i>	%	Abundance		$S^2\bar{x}$	Seasons							
					Mean, SD	Min–max		Autumn		Winter		Spring		Summer	
								%	MI, SD	%	MI, SD	%	MI, SD	%	MI, SD
Trematoda															
<i>Catadiscus longicoecalis</i> ^a	I	2.212*	0.03	43	0.89, 1.64	0–12	3.06	38	1.75, 0.83	64	2.21, 2.80	19	2.80, 2.10	57	2.00, 1.00
<i>Glythelmins palmipedis</i> ^a	P	1.718*	0.02	47	0.63, 0.90	0–5	1.29	24	1.40, 0.80	45	1.50, 0.80	50	1.30, 0.70	67	1.00, 1.00
<i>Styphlodora</i> sp. ^b	P	0.000†	0.08	42	2.24, 3.97	0–16	7.03	–	–	50	5.80, 5.20	58	5.60, 4.70	57	5.00, 4.00
<i>Bursotrema tetracotyloides</i> ^b	P	0.000†	0.04	11	1.28, 4.77	0–29	17.7	–	–	5	–	8	15.0, 14.0	33	12.0, 8.00
<i>Travtrema</i> aff. <i>stenocotyle</i> ^b	P	0.000†	0.01	4	0.19, 1.13	0–9	6.79	5	–	5	–	8	3.50, 2.50	–	–
<i>Opisthgonimus</i> sp. ^b	P	0.000†	0.01	7	0.29, 1.34	0–10	6.14	10	5.50, 4.50	5	–	8	4.50, 0.50	5	–
Strigeidae gen. sp. 1 ^b	P	0.000†	0.00	2	0.07, 0.53	0–5	4.00	–	–	–	–	–	–	10	3.00, 2.00
Strigeidae gen. sp. 2 ^b	P	0.000†	0.00	3	0.04, 0.25	0–2	1.48	–	–	–	–	–	–	14	1.00, 0.00
Cestoda															
<i>Cylindrotaenia</i> sp. ^a	I	0.001‡	0.00	1	0.01, 0.10	0–1	0.00	–	–	–	–	–	–	5	–
<i>Mesocestoides</i> sp. ^b	I	0.000†	0.21	58	6.22, 9.12	0–41	13.4	43	10.4, 12.8	41	17.4, 11.2	58	6.67, 6.10	91	11.0, 8.00
Nematoda															
<i>Cosmocerca</i> sp. 1 ^a	P	0.025	0.01	8	0.16, 0.61	0–4	2.41	10	2.00, 1.00	9	2.50, 1.50	8	1.50, 0.50	5	–
<i>Cosmocerca</i> sp. 2 ^a	P	0.070	0.00	6	0.08, 0.34	0–2	1.49	–	–	14	1.00, 0.50	8	1.50, 0.50	–	–
Rhabdochonidae gen. sp. ^b	I	0.000†	0.00	4	0.04, 0.21	0–1	0.95	–	–	–	–	–	–	19	1.00, 0.00
<i>Physaloptera</i> sp. ^b	I	0.000†	0.00	2	0.03, 0.23	0–2	1.64	10	1.50, 0.50	–	–	–	–	–	–
Seuratoidea gen. sp. ^b	I	0.000†	0.00	4	0.11, 0.57	0–4	2.89	–	–	9	2.50, 1.50	8	2.50, 0.50	–	–
<i>Spiroxyis</i> sp. ^b	I	0.000†	0.00	2	0.03, 0.23	0–2	1.64	–	–	–	–	4	–	5	–
Acanthocephala															
<i>Centrorhynchus</i> sp. ^b	I	0.000†	0.57	70	16.8, 9.17	0–101	21.6	–	–	91	15.70, 10.90	89	21.80, 15.30	95	35.0, 23.0

Transmission to toad (*t*), classification and importance values (*I*), dominance (*d*), prevalence (%), mean abundance±1 standard deviation (SD), dispersion ($S^2\bar{x}$, and seasonal variation in prevalence (%) and mean intensity±SD are shown. Transmission, *I* oral ingestion; *P* skin penetration

Classification of helminth * $I \geq 1.0$, dominant species; † $I = 0$, unsuccessful pioneer species; ‡ $0 < I < 0.01$, subordinate species; $0.01 \leq I < 1.0$, codominant species

^a Adult helminth species

^b Larval helminth species

little to the community and are characteristic of a different host (e.g., *Mesocestoides* sp.).

Influence of season in the helminth community

A non-parametric PERMANOVA test was performed using data from the season as classification factors and the abundance of all helminth parasites (17 species) as dependent variables. Results showed that season (ANOVA pseudo $F = 13.44, P = 0.0001$) indicated significant differences among the four groups of seasons in terms of their infracommunity species abundance. The factor interaction was not significant only between winter and spring ($P < 0.05$) (Table 4). The NMSD ordination of helminth communities, represented in two dimensions (stress=0.18), indicated potential differences

among the abundance of parasitic species composition among the four groups of seasons (Fig. 2).

The seasonal variations of helminth species in relation to parasitological descriptors are shown in Table 2. Results of comparison of *k* proportion in helminth prevalence infections showed that seasons had a significant effect on 47 % of the species (Table 5).

Species richness in component community showed no significant differences between seasons (autumn vs. winter $\chi^2 = 0.94, P > 0.05$; autumn vs. summer $\chi^2 = 1.37, P > 0.05$; autumn vs. spring $\chi^2 = 1.37, P > 0.05$) (Table 6); however, infracommunity species richness was related to seasons (Kruskal–Wallis H test=38.4, $df = 3, P < 0.05$). In the last case, posteriori multiple pairwise comparisons using Dunn’s procedure were calculated. These data showed that the infracommunity species richness was significantly lower in

Fig. 1 Relative frequency in percentage per infection site (*Gm*, gastric mucosa; *Ss*, serous of stomach; *Bc*, body cavity; *Me*, mesenteries; *Ub*, urinary bladder wall; *Mu*, muscles; *Pz*, pharyngeal zone; *Liv*, liver; *Fb*, body fat; *Ki*, kidney; *Si*, small intestine; *Li*, large intestine) of helminth species (total worm frequency, $N=538$). Y-axis scales are different for each helminth species

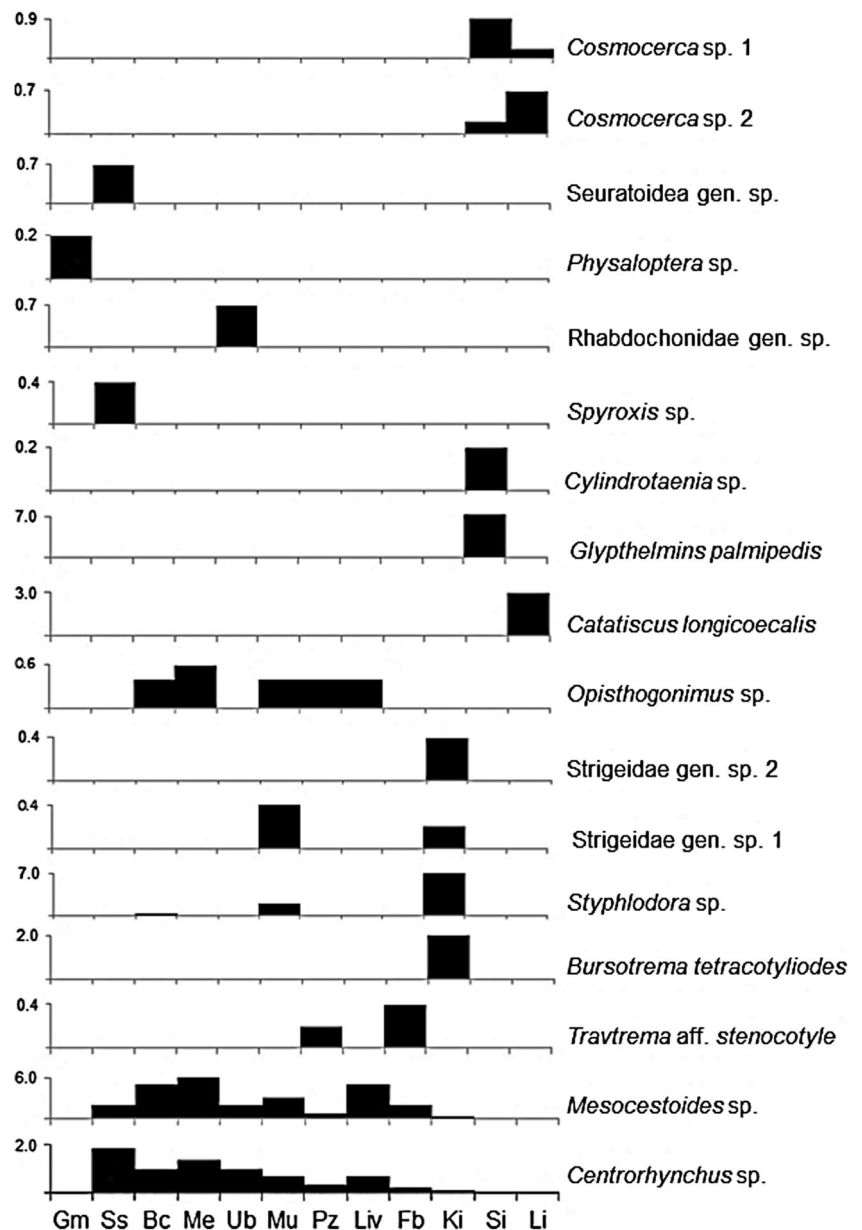


Table 3 Covariation base on Spearman coefficient correlations (r_s) among seven common helminth species in *Melanophryniscus klappenbachi* from Chaco, Argentina

Helminth species	TableID	(1)	(2)	(3)	(4)	(5)	(6)	(7)
<i>Centrorhynchus</i> sp.	(1)	1						
<i>Catadiscus longicoecalis</i>	(2)	0.164	1					
<i>Glypthelmins palmipedis</i>	(3)	0.138	0.091	1				
<i>Bursotrema tetracotyloides</i>	(4)	0.224*	-0.087	-0.028	1			
<i>Styphlodora</i> sp.	(5)	0.421*	0.197	0.094	-0.046	1		
<i>Mesocestoides</i> sp.	(6)	0.212	0.204	-0.063	0.060	0.188	1	
<i>Cosmocerca</i> sp. 1	(7)	-0.252*	-0.130	-0.052	0.037	-0.251*	-0.252*	1

* $P < 0.05$

Table 4 Summary of main results of the non-parametrical permutational ANOVA (PERMANOVA) relating abundances of parasite species of *Melanophryniscus klappenbachi* and seasons

Source	df	SS	MS	Pseudo <i>F</i>	<i>P</i> (perm)
Seasons	3	51048	17016	13.44	0.0001*
Res	86	1.0888E ⁵	1266		
Total	89				
Groups				<i>t</i>	
Autumn×winter				4.3069	0.0001*
Autumn×spring				4.7167	0.0001*
Autumn×summer				5.3784	0.0001*
Summer×spring				2.0638	0.0006*
Summer×winter				1.8165	0.0071*
Winter×spring				1.0792	0.3224

**P*<0.01

autumn than in the other seasons and that the species richness in summer was higher than that in spring (*P*<0.05).

Discussion

The results of this study indicate that *M. klappenbachi* hosts a high richness of helminth species (larvae+adults), with a maximum of seven species per toad. However, the richness of the adult parasite community was lower than that of other South American bufonid toads (Table 7). Particularly, the importance of the species in the community shows that *C. longicoecalis* and *G. palmipedis* would be favored by the transmission, because both trematodes release their infective stage in the water (Smyth and Smyth 1980), where the warm temperature and high rainfall have an important effect on the

Fig. 2 Non-metric multi-dimensional scaling (NMDS) plot results of all seasons (1, autumn; 2, winter; 3, spring; and 4, summer) in terms of their parasitic abundance

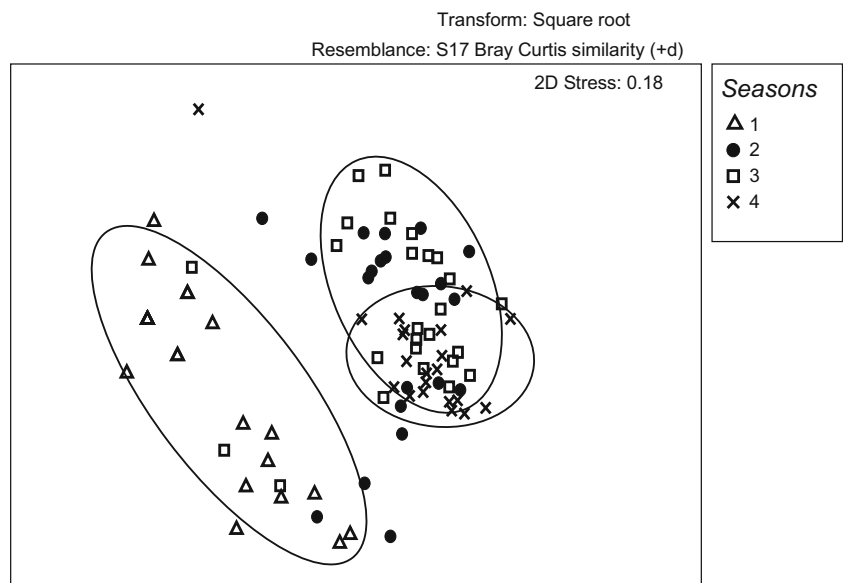


Table 5 Results of *k* proportions comparison of helminth prevalence infections of *Melanophryniscus klappenbachi* among the seasons

Helminth species	Prevalence infections between seasons		
	χ^2	df	<i>P</i>
<i>Catadiscus longicoecalis</i>	11.71	3	0.01*
<i>Glythelmins palmipedis</i>	7.91	3	0.05*
<i>Styphlodora</i> sp.	20.36	3	0.001*
<i>Bursotrema tetracotyloides</i>	14.39	3	0.01*
<i>Travtrema</i> aff. <i>stenocotyle</i>	1.63	3	0.65
<i>Opisthogonimus</i> sp.	0.60	3	0.90
Strigeidae gen. sp. 1	6.72	3	0.08
Strigeidae gen. sp. 2	10.20	3	0.05*
<i>Cylindrotaenia</i> sp.	3.32	3	0.35
<i>Mesocestoides</i> sp.	13.69	3	0.01*
<i>Cosmocerca</i> sp. 1	0.41	3	0.94
<i>Cosmocerca</i> sp. 2	5.44	3	0.14
Rhabdochonidae gen. sp.	13.75	3	0.01*
<i>Physaloptera</i> sp.	3.38	3	0.35
Seuratoidea gen. sp.	3.72	3	0.29
<i>Spiroxys</i> sp.	3.32	3	0.35
<i>Centrorhynchus</i> sp.	64.17	3	0.001*

**P*<0.05

development of early stages (e.g., cercariae) as well as on intermediate hosts as snails and tadpoles (Kennedy 1975; Chubb 1979; Esch and Fernandez 1994; Hamann 2006; Hamann et al. 2010). Therefore, *M. klappenbachi* becomes infected either by ingestion of metacercariae with aquatic vegetation (e.g., *C. longicoecalis*) or by skin penetration of cercariae (e.g., *G. palmipedis*) either in tadpoles or adult frogs. In addition, three larval species (*Styphlodora* sp.,

Table 6 Ecological indices of helminth communities of *Melanophryniscus klappenbachi* from Chaco, Argentina

Variables	Autumn	Winter	Spring	Summer
Component community				
Richness	7	11	12	13
Diversity (<i>H'</i>)	0.47	0.60	0.54	0.52
Evenness (<i>J</i>)	0.55	0.58	0.50	0.47
Infracommunity				
Richness	1.35±1.17	3.35±1.27	3.00±1.50	4.70±1.16
Diversity (<i>HB</i>)	0.08±0.13	0.26±0.16	0.23±0.17	0.32±0.13
Evenness (<i>E</i>)	0.26±0.41	0.54±0.30	0.50±0.35	0.53±0.16

Mesocestoides sp., and *Centrorhynchus* sp.) have a high prevalence of infection (>40 %) for which *M. klappenbachi* represents an intermediate host (e.g., *Styphlodora* sp.) and/or a paratenic host (*Mesocestoides* sp. and *Centrorhynchus* sp.) in the parasite life cycles. The potential definitive hosts for these larval stages are snakes, birds, and mammals, respectively (Grabda-Kazubska 1963; Smyth and Smyth 1980; Prudhoe and Bray 1982). The helminth infections in *M. klappenbachi* showed a typical aggregated pattern of distribution for many of the parasites (94 %). The aggregated distributions among these toads and heterogeneity in susceptibility to infection or in exposure to parasites are the main factors of aggregation and explained also by the clumped distributions of helminth species (see Poulin 2013).

Most authors consider that parasite communities in toad amphibians are depauperate and isolationist (Bolek and

Coggins 2003; Yoder and Coggins 2007; Ibrahim 2008), while others consider no fixed pattern, i.e., communities could be arranged along an axis from isolationist communities to interactive communities (Luque et al. 2005; Hamann et al. 2013a). In this study, the helminth infracommunities of *M. klappenbachi* exhibited few abundant species and poor species richness. An individual toad generally harbors only three helminth species, and in most cases, the prevalence is low, so that opportunities for interspecific interactions are limited. For instance, the scarce species affinity was evident. On the one hand, we noted associations for those larval and adult species whose individuals were located in different toad organs (*Styphlodora* sp.+*C. longicoecalis*), and on the other hand, we found a negative correlation between two species at different infection sites (e.g., *Mesocestoides* sp./*Cosmocerca* sp. 1). Thus, *M. klappenbachi* infracommunities showed a proximity to the isolationist extreme of the continuum, as well as the presence of depauperate infracommunities when only the adult helminth species were considered. Finally, the cooccurrence of metacercariae observed in this amphibian host could affect the structure of gastrointestinal helminths in the definitive hosts and can also serve as indicators of the biodiversity surrounding the habitat, such as the abundance of several major animal groups as turtles, snakes, birds, and mammals (Marcogliese and Cone 1997).

With regard to amphibian body size, several authors suggest that this variable positively influences species richness and abundance of the helminth parasites (Muzzall 1991; McAlpine 1997; Bolek and Coggins 2000; Yoder and Coggins 2007; Ibrahim 2008; Hamann et al. 2012, 2013a).

Table 7 Helminth species richness at the component community level of bufonid toads from South America

Amphibians	Number of samples	Total richness	Helminth parasites								Locality	References
			Adult				Larval					
			N	T	C	A	n	t	c	a		
<i>M. klappenbachi</i>	90	17	2	2	1	–	4	6	1	1	Argentina	Present study
<i>R. fernandezae</i>	65	22	4	4	–	–	4	8	1	1	Argentina	Hamann et al. (2013b)
<i>R. icterica</i>	60	12	7	2	1	–	–	–	1	1	Brazil	Santos et al. (2013)
<i>R. fernandezae</i>	90	13	5	3	1	1	–	1	1	1	Brazil	Santos and Amato (2010)
<i>R. icterica</i>	15	5	2	–	–	1	1	–	–	–	Brazil	Pinhão et al. (2009)
<i>R. icterica</i>	58	12	5	4	2	1	–	–	–	–	Brazil	Lux Hoppe et al. (2008)
<i>R. schneideri</i>	42	6	5	–	1	1	–	–	–	–	Brazil	Lux Hoppe et al. (2008)
<i>R. icterica</i>	32	15	13	2	–	–	–	–	–	–	Brazil	Luque et al. (2005)
<i>R. typhonius</i>	27	7	5	–	–	–	1	–	–	1	Peru	Burseley et al. (2001)
<i>R. glaberrimus</i>	1	2	2	–	–	–	–	–	–	–	Peru	Burseley et al. (2001)
<i>R. marina</i>	5	5	3	–	1	–	1	–	–	–	Peru	Burseley et al. (2001)
<i>A. bomolochos</i>	33	7	3	1	1	–	1	–	–	1	Peru	Iannacone (2003)
<i>A. spurrelli</i>	35	3	1	–	–	–	2	–	–	–	Colombia	Goldberg and Bursey (2003)

Adult/larval helminths N/n nematodes, T/t trematodes, C/c cestodes, A/a acanthocephalans

Indeed, it is important to point out that in the present study, the infection of helminth species decreased in larger amphibians. This result suggests that larger toads have less helminth parasites because they probably offer a large number of lipophilic alkaloids (termed “dendrobatid alkaloids”) with pharmacological potential in a body surface area that could avoid the parasitic attack, especially of infective larval stages that actively penetrate into the host through the skin (e.g., cosmocerids). Another possible reason of the limited penetration of parasites could be related to that adults have a thicker integument compared with juvenile toads and by the immune response of the host. On the other hand, the infrapopulations of *Mesocestoides* sp. and *Centrorhynchus* sp. that infect the toads by the foot web via predator–prey interactions also decrease in abundance in larger hosts. This can occur as the result of an increase in the amount of a single prey type consumed, e.g. mainly Formicidae. Additionally, the preference of this food item was reflected in the production of alkaloid profiles in the skin (Daly et al. 2008). Lastly, these infections are highly dependent on the diet of toads and hence on the nature of the microhabitat.

As demonstrated in a previous research (Hamann et al. 2013b) in amphibian hosts from a variety of habitats in Corrientes, Argentina, the diets and mobility of the host and the local microhabitat conditions may have considerable influence on the trematode parasite communities and are important factors in the development of the latter. The results of the present study also showed that the microhabitat preference plays an important role in parasitic infections. Specifically, adult *M. klappenbachi* toads live and feed almost exclusively on terrestrial habitats and reproduce and develop into tadpoles in temporary marshes in remnants of deciduous xerophilous forest with a predominance of grasses and terrestrial bromeliads. The parasite fauna is dominated by high richness of larval trematodes, particularly allogenic species whose definitive hosts (e.g., snakes, mammals, and birds) are temporary visitors to the aquatic microhabitat thereby introducing infective stage. Additionally, these larvae provide a discontinuous but seasonally predictable source of infective parasites to the habitat and therefore for the trematode community (Esch and Fernandez 1994). By contrast, adult trematodes are poorly represented regarding the species richness, which could be related to the toad diet type, which includes mainly ants and, to a lesser extent, oribatid mites and isopods (M.I. Duré per obs.). The digeneans found are characterized by having heteroxenous life cycles that take place in aquatic environments, and their cercariae could infect the toad host in all life stage (Hamann and González 2009). On the other hand, the poorer nematological fauna and lower abundance of parasitic infections reflect the limited penetration of the infective forms of nematodes, e.g., the genus *Cosmocerca*, which can infect through skin penetration (direct life cycle) (Anderson 2000). This result is different from those found in bufonid

amphibians in temperate region (Bolek and Coggins 2003; Yoder and Coggins 2007) and tropical region (Burseley et al. 2001; Goldberg and Burseley 2003; Luque et al. 2005; Ibrahim 2008; Lux Hoppe et al. 2008; Pinhão et al. 2009; Santos and Amato 2010; Hamann et al. 2013a), where nematode species dominate the community. Specifically, when compared to the helminth community of bufonids (e.g., *R. fernandezae*) from Corrientes, Argentina, the latter showed an equal richness of nematodes and trematodes but was better represented by skin-penetrating nematodes (e.g., *Cosmocerca podicipinus* Baker & Vaucher, 1984) as dominant species (Hamann et al. 2013a).

The temperature is likely to provide the vital synchronization of the life cycle of helminth parasites in their definitive and intermediate hosts and to affect the temporal and spatial distribution of helminth parasites (Chubb 1979). The results of the present study show that, in most cases, the prevalence of plathyhelminths and acantocephalans showed a temporal variation. Similarly, species richness and abundance in the parasite infracommunities showed variation among seasons. These results could be related to colonization, i.e., temporal constraints on the activity and foraging patterns of the definitive hosts, as well as to the parasite transmission cycles. The availability of invertebrates may also be irregular during the seasons, and therefore, the helminth parameters are not similar (Esch et al. 2002). Specifically, the lowest infection parameters for many species in autumn compared with summer may be related to the low level of rainfalls. This dry season induces changes in the behavior of these amphibians, i.e., less frequency of the toads in the breeding site and consequently a decrease in the exposure time to helminth larvae. So, the temporal variations observed in the helminth infracommunities may be accounted for by the recruitment of immature parasites throughout the different seasons of the year, which coincides with the explosive breeding of *M. klappenbachi* after heavy rainfall and by the timing of the larval release by adult parasites throughout the year. In this regard, climatic factors such as warm temperature and high rainfall in summer are significant in determining the high level of parasitic infection of *M. klappenbachi* from Chaco, Argentina.

In conclusion, the features of the community of helminths in *M. klappenbachi* suggest that the characteristic of the tegument (e.g., the presence of alkaloids in the skin glands) could contribute significantly to the low occurrence of infection by skin-penetrating nematodes. In this case, more studies under laboratory conditions are needed to confirm this hypothesis. In contrast, the trematode fauna shows high infections resulting from acquisition of species by penetration of infective larvae in smaller amphibians since it spends more time in water and therefore have a higher probability of contact with these parasites. At the same time, the local host factors and their biology, i.e., terrestrial microhabitat preference, explosive breeding in ephemeral bodies of water, and the diet type intermediate between that of a generalist and a

specialist play an important role in parasitic infections of these toads.

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