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# Helminth community structure of *Scinax nasicus* (Anura: Hylidae) from a South American subtropical area

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ABSTRACT: The main goal of this study was to evaluate the relative influence of season, year of study, host body size, and host sex on abundance of helminth species parasitic in the frog *Scinax nasicus*. A total of 273 frogs was collected between December 2004 and November 2006 over all seasons in Corrientes City, Province of Corrientes, Argentina. Helminth community included 21 taxa, and was dominated particularly by larval trematodes. Infected frogs harboured a maximum of 7 species. Host sex and season played no significant effect in determining infracommunity parasite species abundance. Similarly, species richness was similar for both host sexes, and across time (year and season). However, occurrence of the parasites *Opisthogonimus* sp. 2 and *Travtrema* aff. *stenocotyle* greatly varied over time. Host body size was the main factor for determining infrapopulation structure of *Centrorhynchus* sp. Species richness was significantly and positively correlated with host body size. Strong associations were observed mainly between metacercariae of some species. The transmission strategies of parasites suggest that this hylid acquires infections by ingestion of infective larvae and through direct contact with larval parasites from aquatic and terrestrial habitats.

KEY WORDS: Helminth community · Season · Study period · Host body size · Host sex · Scinax nasicus

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### **INTRODUCTION**

One of the most common ways of characterizing a community is to estimate its diversity, which reflects and combines 2 important aspects, species richness and evenness. Diversity can be measured at the level of either component community or infracommunity; in the latter, the species richness is a subset of the species occurring in the component community (Poulin 1997). In addition, local availability of parasite species and colonization abilities of parasites can affect parasite diversity (reviewed by Esch et. al. 1990), which in turn could affect the degree of interspecific interaction, and distribution of parasite species among infracommunities (see Holmes & Price 1986). One approach to examine this question would be to study potential species co-occurrence, i.e. testing for positive or negative associations between species present in infracommunities. However, according to Poulin (2005), the interpretation

of field study results through a matrix of correlation coefficients must be done cautiously, because when such correlations are found, they do not necessarily imply causation. However, this is one approach that can be taken to understand the functioning of complex natural systems.

Considerations of host sex have often listed as a factor that can influence the number and abundance of parasitic species in vertebrates (Poulin 1996). This approach would be supported by the physiological, behavioural and morphological differences between host sexes. For instance, high testosterone levels can cause immunosuppression in males (Folstad & Karter 1992), and could operate to create more infection levels than females (Poulin 1996). Females and males also differ with respect to behaviour, e.g. differing reproductive behaviours between sexes can influence exposure to parasite infective stages. The different behaviour of both sexes affects the course of the life cycle (Combes 1972), and can also be a strong determinant of differences in parasite occurrences (Gillilland & Muzzall 1999, Bolek & Coggins 2001). In addition, sexual dimorphism can also influence parasite intensity or species richness (McAlpine 1997); the larger sex might be more infected because greater host body size provides more surface area for colonization by parasites, i.e. greater probability of skin penetration by cercariae and larval nematodes. Several studies of parasite richness in relation to host body size support this concept (Dogiel 1964, Price 1980, Bell & Burt 1991, Gregory et al. 1996).

Studies of the influence of biotic factors and external environment on the structure of helminth parasite communities in amphibians have provided important information, showing that anthropogenic perturbations, such as watershed or landscape alterations, can have simultaneous negative and positive effects on distribution, and number of helminth parasite species in infracommunities (Hamann et al. 2006a, McKenzie 2007, King et al. 2008, Marcogliese et al. 2009). Similarly, habitat use (e.g. aquatic, arboreal, terrestrial), relative vagility, feeding preference in terms of foraging behaviour, range of prey species, and host size are key factors determining parasite species richness (Aho 1990, McAlpine 1997, Kehr et al. 2000, Muzzall et al. 2001, Bolek & Coggins 2003, Hamann et al. 2006b, Yoder & Coggins 2007).

Additionally, the effect of season on the parasite recruitment period, and the breeding season of the definitive host, have been identified as important factors regulating the structure of amphibian parasite communities (McAlpine 1997, Wetzel & Esch 1997, Bolek & Coggins 2001, 2003, Hamann 2004, 2006). The presence of helminths in the final host is also influenced by the effect of seasonal conditions on their intermediate hosts, i.e. changing availability of intermediate hosts and infective stages, as well as parasite transmission ability (Esch et al. 1997, 2002). The timing of larval release is also very important for helminth parasites, because it affects their survival during periods of unfavourable living conditions caused either by climatic factors or by the unavailability of a host (Moravec 1998).

Hamann et al. (2009) reported the helminth taxa of the frog *Scinax nasicus*, concluding that the adult parasite frequency was very low in the digestive tracts and that larval trematodes were the primary members of this helminth infracommunity; however, this study did not provide any information about the temporal dynamics of the helminth parasite infracommunity and the factors that explain them. In the present study, research was undertaken to test the following hypotheses: (1) helminth infracommunities vary significantly among sampling season and years; (2) large amphibians commonly have more individuals and species of helminths than small amphibians; and (3) helminth populations may serve as markers of diet and habitat use of the frog. These premises were tested by analyzing the composition and structure of parasite communities of S. nasicus. This frog is distributed in Paraguay, Northern Argentina, Uruguay, eastern Bolivia, and southwestern Brazil (Frost 2007). In the province of Corrientes, Argentina, S. nasicus is very abundant, and lives sympatrically with S. acuminatus; their habitat includes urban and rural areas, and is found mainly on arboreal environments, i.e. on small trees, shrubs and, especially, bromeliads. The S. nasicus diet is dominated by insects (e.g. dipterans, coleopterans), and these frogs principally employ a 'sit and wait' foraging strategy (Duré 1999).

Specifically, the main objectives in this study were to (1) evaluate the magnitude of the influence of season, year of study, host body size, and host sex on the abundance, and number of parasitic helminth species in the frog *Scinax nasicus*, (2) examine the species affinities (covariation and association) of helminth infracommunities, and (3) analyse the helminth cycles of transmission.

### MATERIALS AND METHODS

Study area. Our study area was located approximately 15 km east of the city of Corrientes Argentina (27° 30' S, 58° 45' W). The 30 ha site is characterized by its wide variety of habitats, containing numerous temporary and semipermanent ponds, and 1 permanent pond, 30 m long, 15 m wide and 0.80 meters deep. The vegetation cover is composed of Eichhornia azurea, Nymphaea sp., Hydrocleis nymphoides, Salvinia sp. and Ludwigia peploides; the pond is surrounded by grasslands (Andropogon lateralis). The forest vegetation consists of quebracho Schinopsis balansae, urunday Astronium balansae, and ñandubay Prosopis affinis, with herbaceous strata composed of areas that include gramineous *Elyonurus muticus*, and terrestrial bromeliads (Aechmea distichantha, Bromelia serra). The mean annual temperature is 23°C and the mean annual precipitation is 1500 mm, without a pronounced dry season; however, periods of rain shortages occur every 4 to 6 yr (Carnevali 1994). This area is surrounded by human activities, e.g. weekend houses, sports clubs and pig farming.

**Analytical procedure.** Adult specimens of *Scinax nasicus* were collected from terrestrial bromeliads in the study area from December 2004 to August 2005 (Year 1, n = 127), and January 2006 to November 2006 (Year 2, n = 146). The number of host population samples by year and season of capture and host sex is

shown in Table 1. Specimens were all hand-captured by 2 people between 18:00 and 21:00 h, using the 'visual encounters survey' sampling technique (Crump & Scott 1994). Frogs were transported live to the laboratory, killed in a chloroform (CHCl<sub>3</sub>) solution, and dissected following standard protocols (Goater & Goater 2001). Their snout-vent length (SVL) and body weight were recorded. At necropsy, hosts were sexed and the oesophagus, stomach, gut, lungs, liver, urinary bladder, kidneys, body cavity, musculature, skin, and brain examined for parasites. Helminths were observed in vivo, counted, killed in hot distilled water and preserved in 70% ethyl alcohol. 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 identified using Yamaguti (1961, 1963, 1975), Jones (1987), Anderson (2000), Gibson et al. (2002), and Jones et al. (2005). Specimens of various parasite species were deposited in the Helminthological Collection of the Centro de Ecología Aplicada del Litoral (CECOAL-CONICET), Corrientes City, Corrientes, Argentina (see Table 2).

**Statistical analysis.** 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 total number of helminth species (= richness), Shannon index (H') (Shannon & Weaver 1949), and evenness (J') as  $H'/H'_{max}$ . (Pielou 1966, Zar 1996). Mean helminth species richness is the sum of helminth species, per individual frog, divided by the total sample size. A distribution

Table 1. *Scinax nasicus*. Number of sampled populations by year of study, season of capture, and host sex. Season dates are summer (21 Dec-20 Mar), autumn (21 Mar-20 Jun), winter (21 Jun-20 Sep) and spring (21 Sep-20 Dec). Chi-square ( $\chi^2$ ) values in Table are for comparisons between numbers of each sex collected within seasons. For the comparison of total frogs collected in each year,  $\chi^2 = 1.33$ . All  $\chi^2$  values are non-significant (p > 0.05)

Season	Total frogs	No. of male	No. of female	$\chi^2$
Year 1 (2004–2005,	n = 127)			
Summer	53	26	27	0.04
Autumn	39	21	18	0.26
Winter	35	23	12	3.49
Year 2 (2006, n = 14	46)			
Summer	20	10	10	0.05
Autumn	37	14	23	2.22
Winter	53	23	30	0.94
Spring	36	16	20	0.47
Total	273	133	140	0.18

of infracommunity richness values was tested to see if they indicated some sort of structure between the number of helminth species vs. number of frogs. The diversity index was used with decimal logarithms ( $log_{10}$ ). Berger-Parker index of dominance (*d*) was used to determine the most abundant species (Magurran 2004).

Helminth communities were classified at the infracommunity (all helminth infrapopulations within a single Scinax nasicus) and component community (all helminth infracommunities within a population of S. *nasicus*) levels. Chi-square ( $\chi^2$ ) test was used for comparing the sex ratio of the frogs. Mann-Whitney's Utest was used to test for differences in helminth richness among the sex. Comparison of 2 proportions (Z-test) was used to test for differences in prevalence of helminth infection between sexes. Comparison of k proportion  $(\chi^2)$  was used to test for differences in prevalence between seasons. Kruskal-Wallis test (H)was used to test for differences in species richness between season and year. Pearson's coefficient correlation (r) was used to indicate the relationship between host body size, parasitic abundance, and infracommunity descriptors. Spearman's Rank test (r<sub>s</sub>) was used to calculate possible species co-variation. The species associations was analysed with a  $\chi^2$  test, with Yates correlation. The degree of aggregation of different species of parasites was calculated by the index of dispersion, which indicated the departure of the distribution from randomness (Elliot 1971, Southwood 1978):

$$\chi^2 = \frac{s^2(N-1)}{\overline{x}}$$

where  $s^2$  = variance, N = number of samples and  $\bar{x}$  = mean. This index of dispersion will often depart from unity, and the significance of these departures is assessed by reference to a table of  $\chi^2$  with n–1 degrees of freedom.

The abundance of infection of each parasite species and its relation with biotic and abiotic factors was analysed thought a multivariate analyses of variance (MANOVA), previously normalization of the data by  $log_{10}(x + 1)$  transformation. When MANOVA Wilks' Lambda was significant, a univariate test (ANOVA) was realized for each dependent variable; the probabilities were calculated according to the 'sequential Bonferroni' procedure (Holm 1979), because it provides great control over Type I errors. The spring data was not considered in MANOVA, because sampling in this season was only done in one year. For abundance vs. host body size, co-variation and associations of helminth species, and the effect of interacting factors, the species considered were those that had at least 10% occurrences in each of the amphibians' populations (6 species). The software used were Xlstat 7.5 (Addinsoft 2004) and SYSTAT 7.

#### RESULTS

#### **Community structure analysis**

The helminth component community for this frog population consisted of 21 species of helminths (Table 2). The predominant groups of parasites were the trematodes (71%) followed by the nematodes (19%); the other groups of parasites were represented by only 1 species (Cestoda: *Cylindrotaenia* sp.; Acanthocephala: *Centrorhynchus* sp.). Helminth diversity and evenness were 0.54 and 0.41, respectively. *Bursotrema tetracotyloides* was the dominant species (*d* = 0.59). The majority (76%) of the helminth parasite spe-

cies of *Scinax nasicus* showed a significant (p < 0.025) aggregated pattern of distribution. Parasites were found in kidneys, muscle, mesenteries, body cavity and pharyngeal zone at a high level of infection (Table 2). Of all the helminth species for which adults were examined, 4 are assumed to have indirect life cycles (*Glypthelmins* sp., *Catadiscus inopinatus, Gorgoderina rochalimai* and *Mesocoelium monas*), and 3 are assumed to have direct life cycles (*Cylindrotaenia* sp., *Cosmocerca podicipinus* and *Cosmocerca parva*) (Table 3).

At the infracommunity level, the mean helminth richness was  $2.07 \pm 1.26$  (maximum = 7) species per infected frog. The observed distribution of helminth in-

Table 2. Helminths infecting *Scinax nasicus*. Accession numbers of submitted helminth samples (Helminthological Collection, Centro de Ecología Aplicada del Litoral [CECOAL], Corrientes, Argentina), site of infection (Li: large intestine; Si: small intestine; UB: urinary bladder; K: Kidneys; Mu: muscle; Me: Mesenteries; Bc: Body cavity; Pz: Pharyngeal zone; L: liver; Ss: Serous of stomach), prevalence (%), mean ( $\pm$ SD) abundance, mean ( $\pm$ SD) intensity, and index of dispersion (*I*). *I*-values in **bold** are aggregated distribution (p < 0.025) and normal types are random distribution (0.025  $\leq$  p  $\leq$  0.975); degrees of freedom = 272 in all cases. NA: not available (specimen not submitted); –: mean intensity not calculated, because only 1 host was infected

Helminth species	Stage	CECOAL access. no.	Site of infection	Prevalence (%)	Mean abundance	Mean intensity	Ι
Trematoda							
<i>Catadiscus inopinatus</i> Freitas, 1841	Adult	05012412	Li	11.0	$0.15 \pm 047$	$1.33 \pm 0.65$	407.07
<i>Glypthelmins</i> sp.	Adult	05012416	Si	0.4	$0.01 \pm 0.18$	-	816.00
Gorgoderina rochalimai	Adult	06093013	Ub	0.4	$0.004 \pm 0.06$	-	272.00
Pereira and Cuocolo, 1940							
Mesocoelium monas	Adult	06083107	Si	0.4	$0.007 \pm 0.12$	-	571.20
(Rudolphi, 1819)							
<i>Bursotrema tetracotyloides</i> Szidat, 1960	Larva	05032809	К	42.1	17.20 ± 52.39	$40.83 \pm 74.50$	43411.20
Lophosyciadiplostomum aff. nephrocystis	Larva	04121324	K	0.4	$0.01 \pm 0.24$	-	1060.08
Travtrema aff. stenocotyle	Larva	06093005	Mu Me Bc Pz	28.9	$1.70 \pm 7.15$	$5.86 \pm 12.34$	8214.40
Cohn, 1902							
<i>Styphlodora</i> sp.	Larva	06083127	K Mu L	4.0	$0.15 \pm 1.17$	$3.73 \pm 4.51$	2475.20
<i>Opisthogonimus</i> sp. 1	Larva	06073113	Me Bc	2.9	$0.07 \pm 0.48$	$2.38 \pm 1.58$	897.60
<i>Opisthogonimus</i> sp. 2	Larva	06093008	Mu Me Bc Pz	62.3	$6.69 \pm 11.93$	$10.74 \pm 13.61$	5793.60
Nephrostomum sp.	Larva	05042119	L	1.1	$0.03 \pm 0.28$	$2.33 \pm 1.25$	788.80
Unknown echinostomatid species 1	Larva	NA	Pz	16.8	$2.34 \pm 10.89$	$13.89 \pm 23.30$	13763.20
Unknown echinostomatid species 2	Larva	NA	Κ	0.4	$0.01 \pm 0.24$	-	1060.80
Unknown plagiorchid species	Larva	05042118	Mu Bc	1.1	$0.08 \pm 0.87$	$7.33 \pm 4.03$	2556.80
Unknown strigeid species	Larva	05062909	Bc	2.2	$0.03 \pm 0.18$	$1.17 \pm 0.37$	326.40
Nematoda							
Cosmocerca podicipinus Bakor and Vauchor 1984	Adult	05072802	Li	1.5	$0.01\pm0.12$	$1.00\pm0.00$	252.96
Cosmocerca parva	Δdult	04121316	Τi	0.4	$0.004 \pm 0.06$	_	272.00
Travassos 1925	mun	04121010	11	0.4	0.004 ± 0.00		272.00
Physaloptera sp	Larva	06073104	Ss	0.4	$0.004 \pm 0.06$	_	272.00
Inknown rhabdochonid	Larva	05042115	Ss	1 1	$0.004 \pm 0.00$	$1.00 \pm 0.00$	272.00
species	Luivu	00042110	03	1.1	0.01 ± 0.10	1.00 ± 0.00	272.00
Acanthocephala							
<i>Centrorhynchus</i> sp.	Larva	06083118	Ss Me	27.1	$0.69 \pm 1.50$	$2.53 \pm 1.92$	897.60
Cestoda							
<i>Cylindrotaenia</i> sp.	Adult	06083118	Si	1.8	$0.07 \pm 0.64$	$4.00 \pm 2.53$	1496.00

Table 3. Summary of h	nelminth taxa,	life cycle, tra	nsmission of l	arval stages.	and definitiv	e host (DH)	of parasites in	Scinax	nasicus from
C	orrientes, Arge	entina. Source	s are for life cy	cles of cong	ener species o	species of t	he same family		

Helminth	Life cycle	Transmission	DH	Source
Catadiscus inopinatus	Indirect	Cercariae encysted on substrates (e.g. grass and aquatic vegetation) or on skin of tadpoles; metacercariae ingested with vegetation by definitive host, e.g. <i>Catadiscus uruguayensis</i> Freitas and Lent. 1939	Amphibian	Ostrowski de Núñez (1979a)
<i>Glypthelmins</i> sp.	Indirect	Skin penetration by cercariae in tadpoles and adult frogs. Oral ingestion by metacercariae encysted in sloughed skin of tadpoles by definitive best e.g. <i>C. guidt</i> Stafford 1900	Amphibian	Leigh (1946), Smyth & Smyth (1980)
Gorgoderina rochalimai	Indirect	Oral ingestion of cercariae by tadpoles or aquatic insect larvae; metacercariae encysted in body cavity ingested by definitive host e.g. <i>G. attenuat</i> Stafford, 1002	Amphibian	Bolek et al. (2009)
Mesocoelium monas	Indirect	Cercariae encysted in viscera of intermediate snail, or emerge and encyst on any object they contact, including vegetation, e.g. <i>M</i> monodi Dollfus, 1929	Amphibian	Thomas (1965)
Bursotrema tetracotyloides	Indirect	Presumably skin penetration by cercariae in amphibians, e.g. <i>Fibricola texensis</i> Chandler, 1947; metacercariae encapsulated in kidneys ingested by definitive bost	Mammal	Yamaguti (1975)
Lophosyciadiplo- stomum aff.	Indirect	Skin penetration by cercariae in tadpoles, e.g. <i>Neodiplostomum</i> <i>buteonis</i> Dubois and Rausch, 1948; metacercariae encapsulated in kidney ingested by definitive bost	Bird	Pearson (1960)
Travtrema aff. stenocotyle	Indirect	Skin penetration by cercariae in tadpoles; metacercariae encysted in muscle, mesentery, body cavity and pharyngeal zone ingested by definitive best of <i>a</i> , <i>T</i> , stangestyle Cohn, 1902	Snake	Ostrowski de Núñez (1979b)
<i>Styphlodora</i> sp.	Indirect	Skin penetration by cercariae in tadpoles, e.g. <i>Metaleptophallus</i> <i>gracillimus</i> (Lühe, 1909); metacercariae encysted in kidney, musele and liver ingested by definitive best	Snake	Grabda-Kazubska (1963)
<i>Opisthogonimus</i> sp. 1	Indirect	Skin penetration by cercariae in tadpoles, e.g. <i>Metaleptophallus</i> <i>gracillimus</i> (Lühe, 1909); metacercariae encysted in mesentery and bedy equity ingested by definitive heat	Snake	Grabda-Kazubska (1963)
<i>Opisthogonimus</i> sp. 2	Indirect	Skin penetration by cercariae in tadpoles, e.g. <i>Metaleptophallus gracillimus</i> (Lühe, 1909); metacercariae encysted in muscle, mesentery, body cavity and pharyngeal zone ingested by	Snake	Grabda-Kazubska (1963)
<i>Nephrostomum</i> sp.	Indirect	definitive nost Natural orifices, e.g. nostrils, mouth, penetrating by cercariae e.g. <i>Episthmium suspensum</i> (Braun, 1901); metacercariae	Bird	Ostrowski de Núñez (1974)
Unknown echino- stomatid species 1	Indirect	Natural orifices, e.g. nostrils, mouth, penetrating by cercariae in tadpoles, e.g. <i>Episthmium suspensum</i> (Braun, 1901); metacercariae encysted in pharyngeal zone ingested by definitive host	Reptile/Bird/ Mammal <sup>a</sup>	Ostrowski de Núñez (1974)
Unknown echino- stomatid species 2	Indirect	Natural orifices, e.g. nostrils, mouth, penetrating by cercariae in tadpoles, e.g. <i>Episthmium suspensum</i> (Braun, 1901); matacaraziae apprend in kidnov ingested by definitive best	Reptile/Bird/ Mammal <sup>a</sup>	Ostrowski de Núñez (1974)
Unknown plagiorchid species	Indirect S	Skin penetration of cercariae in tadpoles, e.g. <i>Metaleptophallus</i> <i>gracillimus</i> (Lühe, 1909); metacercariae encysted in muscle and body cavity ingrested by definitive best	Reptile/Bird/ Mammal <sup>a</sup>	Grabda-Kazubska (1963)
Unknown strigeid species	Indirect	Natural orifices, e.g. branchial, mouth, penetrating by cercariae in tadpoles, e.g. <i>Apharyngostrigea pipientis</i> (Oliver 1940); meta- cercariae encysted in body cavity ingested by definitive bost	Bird/ Mammal <sup>a</sup>	Yamaguti (1975)
Cosmocerca podicipipus	Direct	Skin penetration by infective larvae in amphibians, e.g. $C$ commutate (Diesing 1851)	Amphibian	Fotedar & Tikoo (1968)
Cosmocerca parva	Direct	Skin penetration by infective larvae in amphibians, e.g.	Amphibian	Fotedar & Tikoo (1968)
<i>Physaloptera</i> sp.	Indirect	Oral ingestion of eggs by insects. Larvae (L <sub>3</sub> ) encapsulated in intestine ingested by paratenic, e.g. amphibian, and definitive bosts $e \in P$ maxillaris Molin 1860	Reptile/Bird/ Mammal <sup>a</sup>	Cawthorn & Anderson (1976), Anderson (2000)
Unknown rhabdo- chonid species	Indirect	Oral ingestion of eggs by aquatic insects and amphipods. Larvae ( $L_3$ ) encapsulated in body cavity ingested by defini- tive hosts, e.g. <i>Rhabdochana phayini</i> Moravec, 1968	Fish/ Amphibian <sup>a</sup>	Moravec (1977), Moravec & Kaiser (1994)
<i>Centrorhynchus</i> sp.	Indirect	Oral ingestion of eggs by insects and probably terrestrial isopods. Larvae ( $L_3$ ) encapsulated in body cavity ingested by paratenic host	Bird	Schmidt (1985)
<i>Cylindrotaenia</i> sp.	Direct	e.g. amphibian, and demittive nost, e.g. <i>C. aluconis</i> Golvan, 1960 Probably, oral ingestion of embryonated eggs by definitive host, e.g. <i>C. americana</i> Jewell, 1916	Amphibian	Prudhoe & Bray (1982)
<sup>a</sup> Possible definitive	host			

fracommunity species richness showed a good fit to Poisson distribution ( $\chi^2 = 10.78$ , df = 6, p = 0.095) (Fig. 1). Mean values of helminth diversity and evenness were  $0.22 \pm 0.17$  and  $0.52 \pm 0.37$ , respectively. Two correlations between helminth species occurrence were significant: one was negative (*C. inopinatus* vs. *Bursotrema tetracotyloides*), and one was positive (unknown echinostomatid species 1 vs. *B. tetracotyloides*) (Table 4). Three associations were found among the 6 helminth species considered: *Opisthogonimus* sp. 2 + *B. tetracotyloides* ( $\chi^2 = 8.14$ , df = 1, p < 0.05), unknown echinostomatid species 1 + *B. tetracotyloides* ( $\chi^2 = 18.05$ , df = 1, p < 0.05) and *Opisthogonimus* sp. 2 + *Travtrema* aff. *stenocotyle* ( $\chi^2 = 6.30$ , df = 1, p < 0.05).

#### Infection in relation to host sex and body size

Of the 273 frogs examined, the infection prevalence was 89%; there was no significant difference in the prevalence of infected females (52%) vs. males (48%) (Z = 0.62, df = 1, p > 0.05). Parasite richness did not





show any relationship with host sex (Mann-Whitney U-test = 8212.50, p > 0.05,  $n_{female} = 126$ ,  $n_{male} = 118$ ).

Total length of the frogs ranged from 12.50 to 37.50 ( $24.47 \pm 5.18$ ) mm, and weight ranged from 0.15 to 3.78 ( $0.95 \pm 0.60$ ) g. Infracommunity descriptors and abundance of helminth species were primarily not significantly correlated with body size of the host (Table 5). Nevertheless, significant correlation were observed for richness vs. length (r = 0.149, n = 244, p < 0.05) and abundance of *Centrorhynchus* sp. vs. host body size (length: r = 0.226, n = 239, p < 0.05; weight: r = 0.231, n = 239).

Body sizes of males and female frogs did not show significant difference (length: Mann-Whitney *U*-test = 9918.50, p > 0.05,  $n_{female} = 140$ ,  $n_{male} = 133$ ; weight: Mann-Whitney *U*-test = 9873.00, p > 0.05,  $n_{female} = 140$ ,  $n_{male} = 133$ ). The length of female frogs ranged from 15.00 to 37.50 (25.09 ± 4.94) mm, and weight ranged from 0.10 to 3.78 (1.03 ± 0.65) g. These variables were not significantly correlated with most of the helminth parasites and infracommunity descriptors (Table 5). The length of male frogs ranged from 0.15 to 2.47 (0.92 ± 0.54) g. These variables were significantly correlated with infracommunity descriptors, and with the abundance of 2 larval stages of helminths (Table 5).

# Influence of year, season and sex on abundance and richness of parasites

MANOVA was performed using data from the year, season and sex as classification factors, while the dependent variable were the abundance of following 6 species: *Centrorhynchus* sp., *Catadiscus inopinatus, Bursotrema tetracotyloides*, echinostomatid species 1, *Opisthogonimus* sp. 2, and *Travtrema* aff. *stenocotyle*. These results showed that season (MANOVA Wilks' Lambda = 0.94,  $F_{12,378}$  = 0.96, p = 0.49), and sex (MANOVA Wilks' Lambda = 0.98,  $F_{6,189}$  = 0.67, p = 0.67) played no significant effect in determining infracommunity species abundance; nevertheless, sig-

Table 4. Covariation based on Spearman Rank correlations ( $r_s$ ), among 6 helminth species common in *Scinax nasicus* from Corrientes, Argentina. Significant values (p < 0.05) in **bold** 

Species	Table ID	(1)	(2)	(3)	(4)	(5)	(6)
Catadiscus inopinatus	(1)	1.000					
Unknown echinostomatid species 1	(2)	0.101	1.000				
<i>Opisthogonimus</i> sp. 2	(3)	0.001	0.098	1.000			
Travtrema aff. stenocotyle	(4)	-0.037	0.105	-0.127	1.000		
Bursotrema tetracotyloides	(5)	-0.159	0.381	-0.091	-0.029	1.000	
Centrorhynchus sp.	(6)	-0.087	-0.008	0.028	-0.041	-0.094	1.000

Helminth species	— Total frogs —		Ma	ale——	—— Female ——		
-	Length	Weight	Length	Weight	Length	Weight	
Parasitic abundance							
Catadiscus inopinatus	-0.037	-0.071	-0.025	-0.045	-0.057	-0.097	
Unknown echinostomatid species 1	-0.076	-0.077	0.039	0.042	-0.174	-0.170	
Opisthogonimus sp. 2	-0.060	-0.060	-0.042	-0.060	-0.044	-0.038	
Travtrema aff. stenocotyle	0.051	0.018	0.022	-0.034	0.094	0.073	
Bursotrema tetracotyloides	0.041	0.081	0.150	0.185	-0.034	0.006	
Centrorhynchus sp.	0.226	0.231	0.190	0.191	0.270	0.270	
Community descriptors							
Diversity	0.123	0.086	0.228	0.177	0.018	0.002	
Evenness	0.118	0.081	0.247	0.184	-0.028	-0.029	
Richness	0.149	0.122	0.264	0.220	0.038	0.039	

Table 5. Pearson correlation (r) between helminth parasites and sex size of Scinax nasicus. Significant values (p < 0.05) in **bold** 

nificant differences in the presence of helminth species were found between years (MANOVA Wilks' Lambda = 0.84,  $F_{6,189} = 5.82$ , p = 0.0001). In the last case, an *a posteriori* univariate test (ANOVA) was calculated (using probabilities according to 'sequential Bonferroni' criteria). These data showed significant differences between years in the parasite abundance

of *Opisthogonimus* sp. 2 (F = 23.86, df = 1, p = 0.001), and *T*. aff. *stenocotyle* (F = 6.70, df = 1, p = 0.010). Similar results were observed for mean abundance of both species (Fig. 2c,d). All factor interactions were non-significant (p > 0.05).

Results of k proportion comparison in helminth prevalence infections showed that season had a signif-



Fig. 2. Seasonal variation in mean (±SE) helminth parasite abundance in *Scinax nasicus*. (a) Unknown echinostomatid species 1,
(b) *Catadiscus inopinatus*, (c) *Opisthogonimus* sp. 2, (d) *Travtrema* aff. stenocotyle, (e) *Bursotrema tetracotyloides*, (f) *Centro-rhynchus* sp.

Table 6. Results of k proportions comparison in helminth prevalence infections among the seasons and years. Significant values (p < 0.05) in **bold** 

Helminth species	1	Year	1	γ	Year 2			
-	$\chi^2$	df	р	$\chi^2$	df	р		
Catadiscus inopinatus	5.49	2	0.064	3.79	3	0.285		
Unknown echinostomatid species 1	4.38	2	0.112	5.75	3	0.125		
<i>Opisthogonimus</i> sp. 2	4.48	2	0.106	2.45	3	0.484		
Travtrema aff. stenocotyle	1.14	2	0.566	9.10	3	0.028		
Bursotrema tetracotyloides	2.58	2	0.275	7.38	3	0.061		
Centrorhynchus sp.	1.35	2	0.510	12.30	3	0.006		

icant effect only on *Travtrema* aff. *stenocotyle* and *Centrorhynchus* sp. in the second year (Table 6, Fig. 3d,f).

Infracommunity richness was not related to years (Mann-Whitney *U*-test = 6713.00, p > 0.05,  $n_{Year 1} = 110$ ,  $n_{Year 2} = 134$ ), or seasons (Kruskal-Wallis *H*-test = 6.59, p > 0.05, df = 3).

## DISCUSSION

The analysis presented here shows that *Scinax nasicus* hosts a high richness (21 species) of larval and adult helminths, with a maximum of 7 species per frog, with *Bursotrema tetracotyloides* as the dominant species. Only 1 species (*Opisthogonimus* sp. 2) shows a high prevalence (>60%) of infection, for which *S. nasicus* represents a specific intermediate host; this

metacercarial infection occurs comparatively rarely in other sympatric amphibians (Hamann et al. 2006a,b). The potential definitive hosts are snakes (*Opisthogonimus* sp. 2) and mammals (*B. tetracotyloides*).

As part of the definition of a parasite, Crofton (1971) emphasized that the infection process produces an overdispersed distribution of parasites within the host



Fig. 3. Seasonal variation in helminth infection prevalence in *Scinax nasicus*. (a) Unknown echinostomatid species 1, (b) *Catadiscus inopinatus*, (c) *Opisthogonimus* sp. 2, (d) *Travtrema* aff. *stenocotyle*, (e) *Bursotrema tetracotyloides*, (f) *Centrorhynchus* sp.

population, i.e. only a few host individuals are heavily infected, and that this distribution can have at least 2 main origins: (1) when infective stages are released in discrete waves, the parasites become highly aggregated in their hosts, and (2) through the exposure to intermediate hosts in which infective stages are aggregated. For digenean species, aggregation in *Scinax nasicus* is probably generated by the host movement, which tends to increase exposure to free-living contagious cercariae distributed within the host's habitat.

Also, the results show that in most cases there was no temporal variation in terms of species richness in the parasite communities; similarly, the abundance and prevalence of infections only showed large variation in a few helminth parasitic species, e.g. Travtrema aff. stenocotyle and Centrorhynchus sp. varied in prevalence among seasons, and Opisthogonimus sp. 2 and Travtrema aff. stenocotyle in abundance over time. These variations could be related to temporal constraints on the activity and foraging patterns of the snakes and birds which act as definitive hosts (see Table 3), thus affecting colonization and transmission of the parasite. Moreover, the low temporal variations observed in the other helminth infracommunities may be accounted for by the recruitment of immature parasites throughout different seasons of the year, coinciding with the prolonged reproductive activity of Scinax nasicus, which breeds during most of the year (see Hamann et. al. 2009), and combined with the persistent infection by adult parasites throughout the year in the definitive host. In this regard, temperature is not a significant factor in determining strong seasonal patterns in helminth parasitism in Corrientes, Argentina (Hamann 1999, 2006). Furthermore, the availability of invertebrate species such as insects, which may act as intermediate hosts, may also be constant during the seasons, and therefore the helminths' parameters are more stable over time.

Regarding host size, large-bodied frogs harbour more helminth parasite species than small-bodied frogs (see Hamann et al. 2006a). In the present study, the size of amphibians was a determining factor for parasite species richness, which suggests that larger frogs have been exposed to parasites for a longer time and also offer a larger surface area for parasitic attack, especially for infective larval stages. We also found that larger males were more parasitized (in terms of species richness) than larger females, suggesting that reduced immune function and behavioural differences between males and females may also explain the increasing levels of parasites (Folstad & Karter 1992, Poulin 1996). Additionally, infrapopulations of Cen*trorhynchus* sp. increase in abundance in larger hosts; this can occur as the result of accumulation of cystacanth larvae, or an increase in the numbers of a single

prey type (e.g. coleopteran intermediate hosts) consumed. The host *Scinax nasicus* may be a good paratenic host for *Centrorhynchus* sp. (cystacanths) because it is eaten by birds, such as falconiformes (Smyth & Smyth 1980).

Most authors agree that the helminth infracommunity structure in amphibians is depauperate and isolationist (Aho 1990, Muzzall 1991, Bolek & Coggins 2003, Yoder & Coggins 2007, Ibrahim 2008), while others found no fixed pattern, i.e. communities could be characterized variously along this continuum rather than as isolationist or interactive (Luque et al. 2005, Hamann et al. 2006a,b). In the present study, the helminth infracommunities of Scinax nasicus exhibited few abundant species, and poor species richness. In fact, we noted associations for those larval species whose individuals were located in different frog organs (Bursotrema tetracotyloides + unknown echinostomatid species 1), and in the same organs of the frog (Opisthogonimus sp. 2 + Travtrema aff. stenocotyle). Likewise, we found a negative correlation between 2 species at different infection sites (e.g. Catadiscus inopinatus vs. B. tetracotyloides). There was no discernible pattern or structure to the infracommunities of *S. nasicus*, although these results provide evidence of their proximity to the isolationist extreme of the continuum, and they are depauperate when only the adult helminth species are considered. This might also indicate that there are numerous vacant niches in the frog digestive tract. On the other hand, the co-occurrence of metacercariae observed in this study could affect the structure of gastrointestinal helminths in the definitive hosts, such as snakes (see Poulin 2001).

Aho (1990) demonstrated that the habitat use and diet type (generalist vs. specialist) of the host can lead to associated differences in their parasite communities, and are important factors in the development of the latter. The present study also showed that the habitat preference plays an important role; in this sense, adult Scinax nasicus frogs live and feed almost exclusively in arboreal habitats, i.e. bromeliads, whereas they reproduce and develop into tadpoles in aquatic environments. Thus, the parasite fauna is dominated in both number of species and number of individuals by larval trematodes, particularly allogenic species whose definitive hosts are snakes, birds, and mammals, which can leave the immediate aquatic habitat thereby introducing the infective stage (Esch & Fernandez 1994, Esch et al. 1997). By contrast, adult trematodes are poorly represented, which could be related to the frog's diet type that includes dipterans as the dominant item (Duré 1999). The digeneans found are characterized by having heteroxenous life cycles that take place mainly in aquatic (e.g. Catadiscus inopinatus, Glypthelmins sp., and Gorgoderina rochalimai) or terrestrial (e.g. Mesocoelium monas; see Prudhoe & Bray 1982) environments. Similarly, the less diverse nematological fauna and lower abundance of parasitic infections in S. nasicus reflects the limited contact with the terrestrial environment where the infective forms of the nematodes occur, e.g. Cosmocerca parva, and Cosmocerca podicipinus, which infect through skin penetration (Anderson 2000). This result contrasts with those found in leptodactylid frogs captured in the same area but characterized by different habitat use (semiaquatic and terrestrial); the latter show higher values for the ecological descriptors and indices of the nematode community (Hamann et al. 2006a,b). Additionally, the low infections of these nematodes found in S. nasicus agree with the previous reports of Bursey et al. (2001), and Goldberg et al. (2007) from other American Scinax species.

In conclusion, the community of helminths in Scinax nasicus suggests that the 'sit and wait' feeding strategy and the diet in particular of this frog were determining factors in the less diverse infections of adult helminths with complex life cycles. At the same time, the low vagility and the arboreal habitat preference of S. nasicus contributes significantly to the low occurrence of infection by skin-penetrating nematodes, while its trematode fauna shows high larval infections, resulting from acquisition of parasites following penetration by infective larvae (see Table 3), when the frogs visit the water to breed. Moreover, the helminth communities of S. nasicus are dominated by generalist metacercariae species that infect diverse amphibian hosts in the same area (Hamann et al. 2006a,b); frogs are susceptible to infection with cercariae in the tadpole stage (Hamann & González 2009) and also after metamorphosis, which suggests that they are suitable second intermediate hosts for these metacercariae. In fact, these results may be alternatives to ensure transmission because S. nasicus occupies an intermediate position in the food web, being easy prey for potential definitive hosts, such as snakes, birds and mammals.

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