

Disentangling the influences of habitat structure and limnological predictors on stream fish communities of a coastal basin, southeastern Brazil

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In stream environments habitat structure and limnological factors interact regulating patterns of energy and material transfer and affecting fish communities. In the coastal basins of Southeastern Brazil, limnological and structural characteristics differ between clear and blackwater streams. The former have a diversity of substrate types, higher water velocities, and lower water conductivity, while the latter have sandy substrate, tea-colored and acidic waters, and low water velocities. In this study, we verified the relative importance of habitat structure and limnological variables in predicting patterns of variation in stream fish communities. Eight first to third order streams were sampled in the coastal plain of Itanhaém River basin. We captured 34 fish species and verified that community structure was influenced by physical habitat and limnology, being the former more important. A fraction of the variation could not be totally decomposed, and it was assigned to the joint influence of limnology and habitat structure. Some species that were restricted to blackwater streams, may have physiological and behavioral adaptations to deal with the lower pH levels. When we examined only the clearwater streams, all the explained variation in fish community composition was assigned to structural factors, which express specific preferences for different types of habitats.

Em ambientes de riacho, fatores relacionados à estrutura dos habitats e limnologia interagem regulando os padrões de transferência de energia e matéria, afetando a composição da comunidade de peixes. Em bacias costeiras do sudeste do Brasil as características limnológicas e estrutura dos habitats diferem entre riachos de águas claras e pretas. Os primeiros são compostos por uma variedade de tipos de substrato, possuem velocidades de corrente mais elevadas e baixa condutividade, enquanto os últimos apresentam substrato arenoso, baixas velocidades de corrente e águas escuras e ácidas. Neste trabalho analisamos a importância relativa da estrutura dos habitats e das variáveis limnológicas como preditores dos padrões de composição em comunidades de peixes de riachos. Oito riachos de primeira a terceira ordem foram amostrados na planície costeira da bacia do rio Itanhaém. Capturamos 34 espécies e verificamos que a composição das comunidades foi influenciada por fatores estruturais e limnológicos, sendo os primeiros mais importantes. Uma fração de variação que não pode ser totalmente decomposta, deve-se à influência conjunta da limnologia e estrutura dos habitats. Algumas das espécies restritas aos riachos de águas pretas provavelmente apresentam adaptações fisiológicas e comportamentais para lidar com os baixos níveis de pH. Quando foram examinados somente os riachos de águas claras, toda a variação explicada na composição da comunidade de peixes foi atribuída aos fatores estruturais, devido a preferências específicas por diferentes características de habitats.

Key words: Blackwater streams, Clearwater streams, East basin, Restinga forest, Variance partitioning.

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Introduction

The importance of limnology and physical habitat as predictors of the structure of fish communities is fully established in the current ecological literature (Schlosser, 1982; Taylor *et al.*, 1993; Jackson *et al.*, 2001; Gerhard *et al.*, 2004; Suárez & Petrer Jr., 2007). Among habitat predictors we may point out the stability of lateral banks, substrate heterogeneity, presence of wood debris and packs of leaf litter that provide the necessary conditions for fish survivor by offering camouflage against predators, protection from currents, source of food, and substrate for eggs and larval development. In addition, limnological variables including water temperature, pH, dissolved oxygen, turbidity, among others, influence fishes due to their physiological tolerances (Angermeier & Karr, 1984; Pulsey & Arthington, 2003; Allan & Castillo, 2007).

In streams ecosystems, these variables interact through complex pathways regulating energy and material transfer. Due to these interactions, Buisson *et al.* (2008) argue that identifying isolated responses of each group is a challenge in observational studies. A way to deal with this complexity is to consider a hierarchical view where the variables are linked to each other according to a network of potential causal effects (Pulsey & Arthington, 2003). For instance, Cruz *et al.* (2013) considered the riparian zone as the higher level predictor followed by predictors at intermediate levels as bank slope, shade availability, substrate composition, water velocity and water quality. Functional fish composition consisted of the lowest level in this network, which was directly and indirectly affected by the higher hierarchical levels. The authors concluded that bank slope and substrate composition were the only direct influences and that water limnological traits do not affect the functional fish composition.

Another way to deal with this complexity is to partialling out the variance explained by each set of predictors. Buisson *et al.* (2008) applied a partitioning analysis to a large fish data set in French rivers and verified that in 18 out of 28 fish species, the variance assigned the joint influence of thermal regimes and structural factors was higher than the variance assigned independently to each factor. In community ecology, variance partitioning was popularized by Borcard *et al.* (1992) as a method to extract the spatial and environmental components of community variation. In this context, most applications are motivated to put forward evidences for niche *versus* neutral theories of metacommunity regulation (Falke & Fauch, 2010; Landeiro *et al.*, 2012; Siqueira *et al.*, 2012; Grönroos *et al.*, 2013). In addition, few studies have also compared the influence of different sets of environmental predictors related for example to aquatic and riparian habitats, while others had included factors related to predation pressure, resource availability or biotic interactions (Peltonen *et al.*, 2007; Holmes *et al.*, 2011; Siqueira *et al.*, 2012).

In this paper, we studied the stream fish communities of the Itanhaém River basin in SE Brazil. Hydrology and limnology are the main features used to classify the rivers and streams in this region. Mountain streams drain steep areas and have clear waters with high levels of oxygen concentration, whereas some lowland streams draining the Restinga forest have black, poor oxygenated, and acidic waters (pH < 4) (Camargo *et al.*, 1996; Por, 2004). Given these differences, we tried to understand how much of variation in the stream fish communities can be assigned solely to limnological variables (pH, water temperature, turbidity, and conductivity), or to physical habitat traits (water velocity, substrate composition, stream width, and depth), and how much of the variation due to limnological conditions are also structured by the habitat variables (joint variability).

Material and Methods

Study area

The studied streams are located in the coastal plain of the Itanhaém River basin, State of São Paulo SE Brazil. This watershed is the second largest coastal basin of the state with an area of 954 km² within two metropolitan areas (Baixada Santista and Grande São Paulo). According to Köppen-Geiger classification, the southeastern Brazil has mostly a warm temperate climate (Type - C with average temperature in the coldest month between 0 and 18°C and average temperature in the hottest month above 10°C) (Peel *et al.*, 2007). Nonetheless, in Itanhaém River basin, the climate near the sea coast line is classified as Af (tropical humid and without dry season), while throughout higher elevations in the mountain ranges of Serra do Mar it is classified as Cfa (humid mesothermal without dry season and with hot summers). The rainfall levels are among the highest in the state, and although the rainfall is distributed throughout the year, it peaks between January and March (260 mm/month) with minimum values between June and August (90 mm/month) (São Paulo, 2012).

The soil is covered by the Atlantic rainforest and, while most of the headwaters are located inside preservation areas, the lower portions have a mixture of preserved and degraded areas due to agriculture (banana plantation) and urbanization. The main tributaries of Itanhaém River are Preto, Branco, Mambu, and Aguapeu rivers (Fig. 1). Alike other basins in the region these rivers and streams can be classified as clear or blackwaters due to their limnological and structural characteristics. Clearwater streams have a variety of bottom types from sandy to rocky substrates, highly oxygenated water and a diverse range of mesohabitats such as pools, riffles and runs. Blackwater streams draining the Restinga forest have tea-colored and acidic waters with low oxygen levels, low water velocities and fine-particulate substrate.

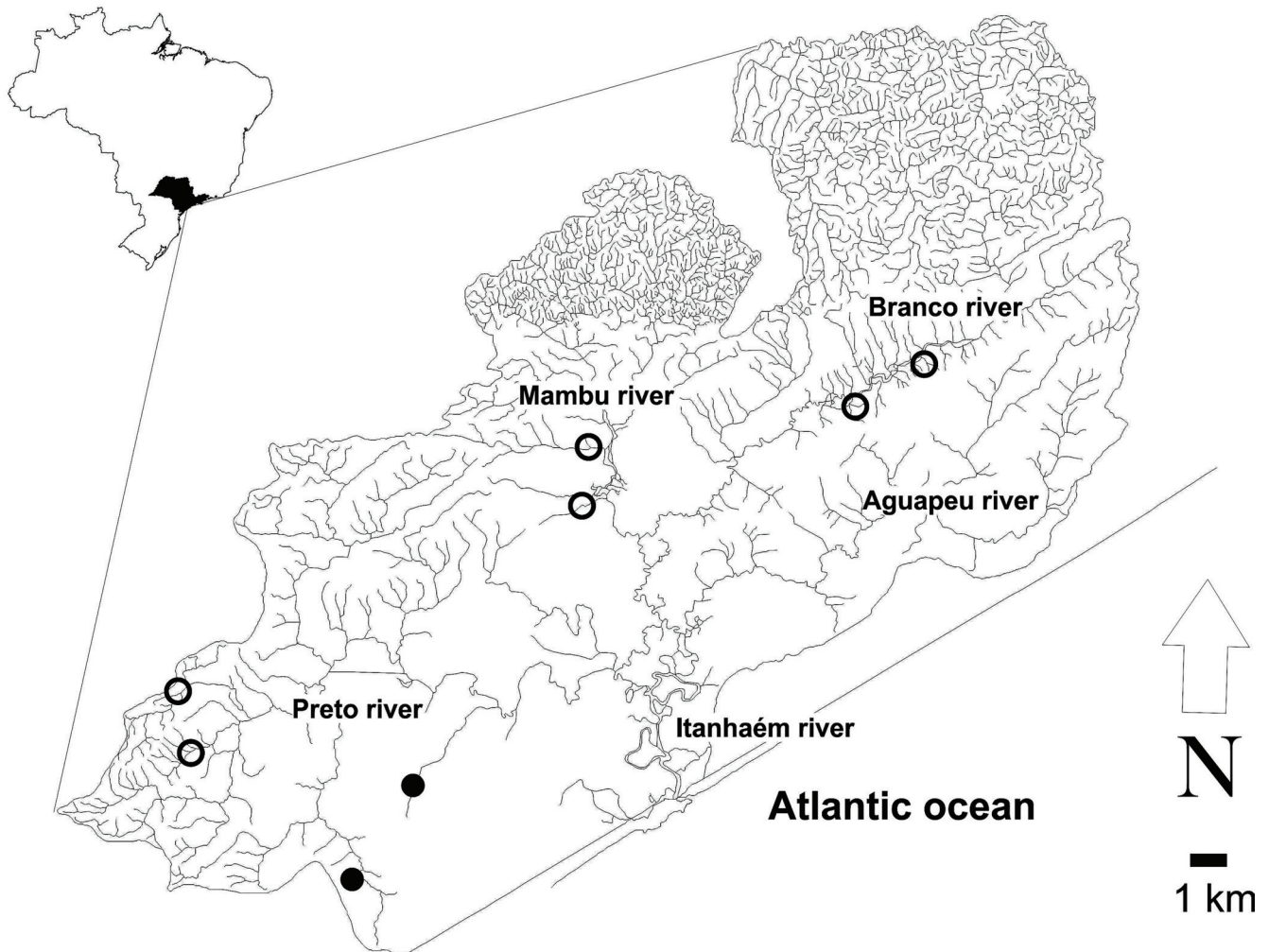


Fig. 1. Hydrography and location of the sampling sites (first to third-order streams) in the coastal plain of Itanhaém River basin, southeastern Brazil. Open circles: clearwater streams. Black circles: blackwater streams.

Fish and habitat sampling

Fishes were sampled in eight first to third-order streams between 11 and 69 m above the sea level where the Characidae family is the dominant group (Ferreira & Petrere Jr., 2009). The clearwater stream sites were near the confluences with the main channels of Branco, Mambu, and Preto rivers (fourth or higher orders) (Fig. 1), while the blackwater streams were inside the Restinga forest. Fishes were caught along 50-m long-stretches (without block nets) by electric fishing gear using a current rectifier (connected to two dip nets) powered by a generator (Yamaha, model EF2600, 2.3 kVA, 60 Hz, 600 volts, direct current). The samples were carried out by one passage of two people handling the dip nets in the opposite direction to the stream flow. Six clearwater and two blackwater streams were sampled quarterly in 2006 (February to November). However, the blackwater streams could not be sampled in February due to the high water levels, while four clearwater sites were excluded from the analyses in November

due to equipment failure measuring the limnological variables.

Fishes were immediately fixed in 10% formalin, and in the laboratory, the individuals were identified at species level, weighed (g), and preserved in alcohol 70%. Voucher specimens were deposited in the fish collection of the Departamento de Zoologia e Botânica, Universidade Estadual Paulista “Júlio de Mesquita Filho”, câmpus de São José do Rio Preto, São Paulo State (DZSJRP) (Table 2). Prior the fish sampling the limnological variables (water temperature, pH, turbidity, and conductivity) were measured in the middle stretch of the sampling sites with an electronic device (Horiba U-10 model). The habitat structure was visually assessed along the 50-m long-stretches. Average channel width was calculated from individual transects taken at each 5 meters. Along these transects and spaced every 1 m, we measured stream depth, water velocity and substrate type. Water velocity was visually classified in 5 categories (V1 - water movement almost imperceptible, practically stagnated; V2 - perceptible movements, but without alterations in the surface; V3 -

altered surface with small undulations, smaller than 0.5 cm; V4 - surface quite altered, with undulations larger than 0.5 cm; and V5 - turbulent water). The substrate was classified as clay, sandy, gravel, pebble, or boulder. We transformed the qualitative measures of water velocity and substrate in quantitative indexes after Gonçalves & Braga (2012). The indexes were calculated by the weighed mean according to the formula:

$$x = \sum (f_i x_i) / \sum f_i,$$

where f_i is the numerical frequency and x_i is the weight given to substrate (clay = 1, sand = 2, gravel = 3, pebble = 4, boulder = 5), or water-velocity categories (V1 = 1, V2 = 2, V3 = 3, V4 = 4, V5 = 5).

Data analyses

In order to provide an initial description of how the clear and blackwater streams differ in habitat structure and limnology, a Principal Component Analysis (PCA) was used to ordinate the sampling sites and to describe the pattern of correlation among environmental variables. Turbidity and conductivity were log-transformed to remove the influence of extreme values (Legendre & Legendre, 2012). Prior to PCA, the variables were centered to zero mean and standardized to unit variance. The mean values for each environmental variable were then compared between clear and blackwater streams by individual t tests.

Partial Redundancy Analysis (pRDA) was used to quantify the isolated effects of each set of predictors (limnological or structural) on community composition. The Hellinger transformation (

$$\sqrt{\frac{n_{ij}}{\sum_{j=1}^S n_{ij}}},$$

where n_{ij} is the abundance of the species i in the site j and S the species richness) was applied to the matrix of species abundance, which is indicated to make the matrices of community composition suitable for applying linear methods such as pRDA (Peres-Neto *et al.*, 2006; Legendre, 2008). The data structure used in pRDA was $Y \sim X|W$, where Y is the response variable (community composition), X the predictor matrix and W the conditioning matrix. The objective of the pRDA is measure the influence of X on Y after remove the effect of W . Thus, to measure the influence of pure limnological variables we used pH, water temperature, conductivity and turbidity as the predictor group, and the habitat variables as the conditioning matrix. Conversely, to measure the influence of pure structural variables we used the water velocity, substrate, depth, and width as the predictor group and the limnological factors as the conditioning variables. By using pRDA we partitioned the community's total variation in the

following fractions: pure limnological variation [a], shared variation (limnological and structural factors) [b], pure structural variation [c], and unexplained variation [d]. The results, presented as proportions of total variance [a+b+c+d], were based on adjusted coefficients of determination (R^2_{adj}). A randomization procedure (10.000 permutations) was used to test the hypothesis of statistical significance of fractions [a] and [c], and to measure the statistical significance of each environmental variable on pRDA (Peres-Neto *et al.*, 2006).

Variance partitioning was firstly proposed in the context of community ecology to partialling out the spatial from the environmental component of community variation (Borcard *et al.*, 1992). Thus, to decide whether space should be included in pRDA, we run a Mantel test (10.000 permutations) to compare the dissimilarity in species composition (using the Bray-Curtis index) with the watercourse distance (in kilometers) among streams.

As some sites were sampled repeatedly throughout 2005, the observations at the same site but in different months were not completely independent, and ignoring this temporal data structure in the statistical tests would increase the type I error rates. To deal with this in the Mantel test, we did not include the pairwise distances between the same sites sampled at different months. As previously verified, these pairwise comparisons had major influences on the final results as the spatial distances between these samples were actually zero. For the other statistical analysis (t tests and pRDA) the p-values were calculated under restricted randomization (Manly, 2007). Restricted randomization is often applied when simply shuffling the data at random is inappropriate, as when there is some temporal structure, or in block designs (Simpson, 2012). Under this scheme, permutation of the sampling units is restricted by some factor, in our case, the month at which each sample was obtained. Thus, when generating the null distributions, we did not allow for an observation to be shuffled among different months, but only among different sites within months. Considering our sampling design, this procedure provide us with a better control over the type I error rates for testing the spatial effect (*i.e.*, clear *versus* blackwater sites).

All analyses were performed with R software (R Development Core Team, 2012). Variance partitioning and pRDA was carried out using functions `varpart` and `rda` available in `vegan` package version 2.0-4 (Oksanen *et al.*, 2012). Restricted randomization were done using the options available in `permute` package, version 0.7-0 (Simpson, 2012).

Results

The first axis of the PCA isolated black from clearwater streams, accounting for 46.1% of the variation in the patterns of correlation among environmental variables. Blackwater streams were characterized by fine-particulate substrate, lower velocities, and were deeper and narrower than the clearwaters.

Table 1. Mean and standard deviation (in parenthesis) for limnological and structural variables in clear (n = 21) and blackwater (n = 6) streams, including the results of the t tests.

		Clearwaters	Blackwaters	t	p
Limnological variables	Temperature (°C)	20.82 (1.90)	20.19 (1.58)	0.74	0.503
	pH	6.91 (0.67)	4.56 (0.45)	7.92	0.001
	Turbidity (NTU)	14.49 (12.06)	10.33 (3.07)	0.07	0.939
	Conductivity (mScm ⁻¹)	0.031 (0.027)	0.075 (0.015)	5.16	0.001
Structural variables	Width (cm)	598.01 (268.18)	287.23 (188.19)	2.64	0.015
	Depth (cm)	13.33 (3.12)	20.91 (2.83)	5.02	0.001
	Substrate	3.94 (0.44)	2.10 (0.21)	9.67	0.001
	Water velocity	2.85 (0.27)	1.67 (0.68)	6.57	0.001

They also had higher conductivity and lower pH levels. The second axis accounted for 16.2% of variation, describing the gradients of temperature and turbidity, but did not distinguished black from clearwater streams (Fig. 2, Table 1).

We caught 34 fish species, 29 in clearwater and 19 in blackwater streams, with 14 common species. The four most abundant species were *Mimagoniates lateralis*, *Deuterodon iguape*, *Characidium* sp., and *Kronichthys heylandi*, being the last three exclusively from or predominantly captured in clearwaters. On the blackwater streams *M. lateralis*, *Scleromystax macropterus*, *Pseudotothyris obtusa*, *Hyphessobrycon reticulatus*, and *Acentronichthys leptos* were the most abundant. The first three were exclusively captured in blackwater streams, while *H. reticulatus* and *A. leptos* had occasional occurrence in clearwaters (Table 2).

With all sites included, the environmental predictors accounted for 46.2% of total variation in community composition due to the influence of pH, width, depth and substrate. Pure structural predictors explained 19.3% of total variation ($p = 0.024$) while pure limnological factors accounted for a small but statistically significant fraction of 3.1% ($p < 0.001$). The highest explained fraction (23.8%) was assigned to the shared variation. Removing the blackwater streams, pure structural factors were the unique influence on community composition, accounting for 34.7% ($p < 0.001$) of the explained variation, while pure limnological and shared fractions had small and negative values (-0.05, -0.004) (Table 3). The watercourse distances among sites were not correlated with community dissimilarity (Mantel test: $r = 0.01$, $p = 0.426$).

Considering the influence of pure limnological predictors with all sites included (Fig. 3a), pH was significantly correlated to community composition ($p = 0.0473$), while temperature ($p = 0.154$), turbidity ($p = 0.158$), and conductivity ($p = 0.113$) was not. Lower values of pH were correlated with the abundance of *M. lateralis*, *S. macropterus*, *H. reticulatus*, *A. leptos*, *P. obtusa*, and *Characidium* sp. while higher values were correlated with the abundance of *Phalloceros* cf. *reisi*, *Rineloricaria kronei*, *Awaous tajasica*, *Hollandichthys multifasciatus*, and *Bryconamericus microcephalus*.

Concerning the effects of pure structural predictors (Fig. 3b), community composition was significantly correlated with width ($p < 0.001$), depth ($p < 0.001$) and substrate ($p = 0.006$), but not with water velocity ($p = 0.109$). The abundance of the catfishes *K. heylandi* and *Rhamdioglanis transfasciatus* were correlated with shallow streams, *D. iguape* with coarser substrate, *Characidium* sp. and *M. microlepis* with wider streams, and *H. multifasciatus* and *Phalloceros* cf. *reisi* with narrow channels and fine-particulate substrate.

Discussion

We verified that environmental conditions in clear and blackwater streams differ in several aspects. The blackwater streams were on average deeper, narrower, predominantly sandy with lower water velocities. They also had higher water conductivities and lower pH levels. These results evidenced that part of the variation in limnology is structured

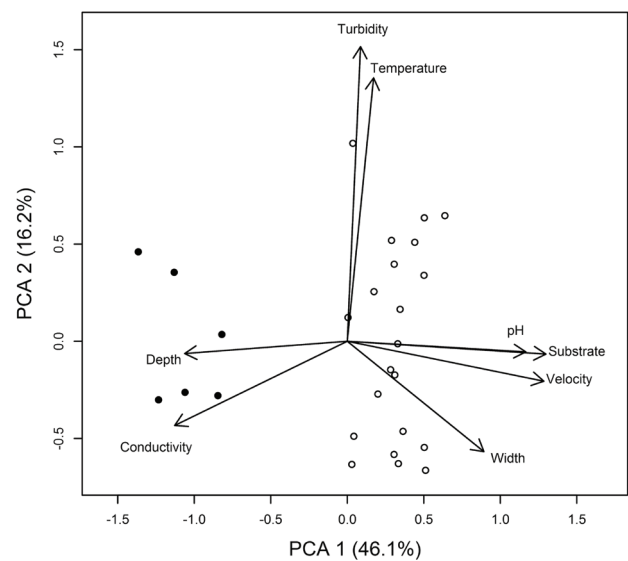
**Fig. 2.** Results of the Principal Component Analysis (PCA) showing the clearwater (open circles) and blackwater (black circles) sites.

Table 2. Species abundances in clear and blackwater streams of Itanhaém River basin.

Species	Abbreviation	Clearwater streams	Blackwater streams	Total	Voucher Number
	Total	1884	1446	3360	
<i>Acentronichthys leptos</i>	Alep	8	115	123	DZSJRP15368
<i>Ancistrus</i> sp.	Asp.	54		54	DZSJRP15384
<i>Atlantirivulus santensis</i>	Asan		23	23	DZSJRP15372
<i>Awaous tajasica</i>	Ataj	14		14	DZSJRP15351
<i>Bryconamericus microcephalus</i>	Bmic	48		48	DZSJRP15374
<i>Callichthys callichthys</i>	Ccal		28	28	DZSJRP15379
<i>Characidium japyhybense</i>	Cjap	1		1	DZSJRP15366
<i>Characidium</i> sp.	Csp.	265	33	298	DZSJRP15357, DZSJRP15365, DZSJRP15363
<i>Crenicichla lacustris</i>	Clac	20		20	DZSJRP15350
<i>Deuterodon iguape</i>	Digu	401		401	DZSJRP15387
<i>Geophagus brasiliensis</i>	Gbra	10	23	33	DZSJRP15361
<i>Gymnotus carapo</i>	Gcar	19	1	20	DZSJRP15378
<i>Gymnotus pantherinus</i>	Gpan	7	49	56	DZSJRP15382
<i>Hollandichthys multifasciatus</i>	Hmul	39	39	78	DZSJRP15377
<i>Hoplias malabaricus</i>	Hmal	1	25	26	-
<i>Hyphessobrycon reticulatus</i>	Hret	2	178	180	DZSJRP15371
<i>Kronichthys heylandi</i>	Khey	295	1	296	DZSJRP15352
<i>Mimagoniates lateralis</i>	Mlat		523	523	DZSJRP15370
<i>Mimagoniates microlepis</i>	Mmic	224		224	DZSJRP15376, DZSJRP15630
<i>Oligosarcus hepsetus</i>	Ohep	4		4	DZSJRP15359
<i>Phalloceros</i> cf. <i>reisi</i>	Prei	136	95	231	DZSJRP15364
<i>Pimelodella transitoria</i>	Ptra	20		20	DZSJRP15390
<i>Pseudotothyris obtusa</i>	Pobt		102	102	DZSJRP15367
<i>Rhamdia quelen</i>	Rque	29	6	35	DZSJRP15388
<i>Rhamdioglanis transfasciatus</i>	Rtra	169		169	DZSJRP15354
<i>Rineloricaria</i> aff. <i>latirostris</i>	Rlat	1		1	DZSJRP15386
<i>Rineloricaria kronei</i>	Rkro	54		54	DZSJRP15383
<i>Schizolecis guntheri</i>	Sgun	1	1	2	DZSJRP12367
<i>Scleromystax barbatus</i>	Sbar	15	29	44	DZSJRP15389
<i>Scleromystax macropterus</i>	Smac		200	200	DZSJRP15356
<i>Synbranchus marmoratus</i>	Smar	4	5	9	DZSJRP15355
<i>Trichomycterus</i> sp. 1	Tsp1	5		5	DZSJRP15362
<i>Trichomycterus</i> sp. 2	Tsp2	12		12	DZSJRP15358
<i>Trichomycterus zonatus</i>	Tzon	26		26	DZSJRP15391

by the physical habitat conditions (Menezes *et al.*, 2007; Gonçalves & Braga, 2012), pointing out a hierarchical pattern of dependence among abiotic features, where higher levels variables related to habitat structure (riparian cover, shade availability, channel morphometry, substrate and water velocity) influence lower levels variables related to water quality and fish community (Cruz *et al.*, 2013; Pulsey & Arthington, 2003; Dias *et al.*, 2010). In the Restinga forest for example, where declivity is lower, the streams have low

water velocities and depositional characteristics, promoting the accumulation and decomposition of humic substances that causes the acid blackwaters (average pH = 4.56) (Janzen, 1974). In addition, the proximity with the sea results in higher levels of water conductivity (Por, 2004). Clearwater streams otherwise drain the mountain ranges of Serra do Mar, coming from higher declivities with water velocities, so high that fine-particulate material does not accumulate as in the blackwater streams.

Table 3. Results of pRDA and variance partitioning for all sites and excluding blackwater streams. Negative fractions were not shown.

	p-value [a]	p-value [c]	Significant variables on pRDA	Limnological [a]	Shared [b]	Structural [c]	Residual
All sites	0.024	< 0.001	pH, width, depth, substrate	0.031	0.238	0.193	0.538
Excluding the blackwater sites		< 0.001	width, depth, substrate			0.347	0.707

Given these differences and the relationship between structural and limnological factors, it is expected a complex interaction between fishes and environment, with a high proportion of variation in community composition (23.8%) assigned to the joint effects of limnology and habitat structure. After removing the blackwater streams from the analyses, it became evident that they were the main source of this joint variability. Considering the isolated effects, a much higher variation was due to structural predictors (19.3%, $p < 0.001$), influenced by depth, width and substrate, stressing the importance of structural complexities to the maintenance of stream fish diversity (Casatti *et al.*, 2009; Dias *et al.*, 2010; Cruz *et al.*, 2013).

Although limnological variables were relatively unimportant in comparison to the structural factors, our results suggest that lower levels of pH may restrict the distribution of some abundant species (e.g., *D. iguape* and *R. transfasciatus*), and favor the occurrence of others (e.g., *M. lateralis*, *S. macropterus*, *H. reticulatus*, *A. leptos*, and *P. obtusa*). It is known that artificial acidification has depletory effects on fishes and aquatic invertebrates (Allan & Castillo, 2007) because several species are incapable to regulate internal pH, which in turn affects oxygen assimilation (Matthews, 1998), reproduction, and calcium and sodium regulation (Fromm, 1980; Freda & Dunson, 1984; Hargeby & Petersen Jr., 1988). Conversely, in natural acidic streams fish richness can be high (Henderson & Walker, 1986; Otto, 2006) because some species are adapted to these harsh conditions (Collier *et al.*, 1990; Menezes *et al.*, 2007). Rahel (1983) suggested that the tolerance to acidic waters results, in part, from physiological adaptations that allow individuals to survive at low electrolytes concentrations. In addition, reproductive physiology and fish behavior may increase the chances of offspring survivor in blackwater streams. For instance, species which are common to blackwaters in southeastern Brazilian coast as *M. lateralis*, *H. multifasciatus*, and *Rachoviscus crassiceps* are internal inseminating, in which the females keep the sperm until finding suitable environmental conditions to lay and fertilize their eggs (Menezes *et al.*, 2007).

The unexplained variation (53.8% with all streams included and 70.6% only with the clearwater streams) was higher than the variation assigned to the environmental factors, and part of this could be attributed to unmeasured

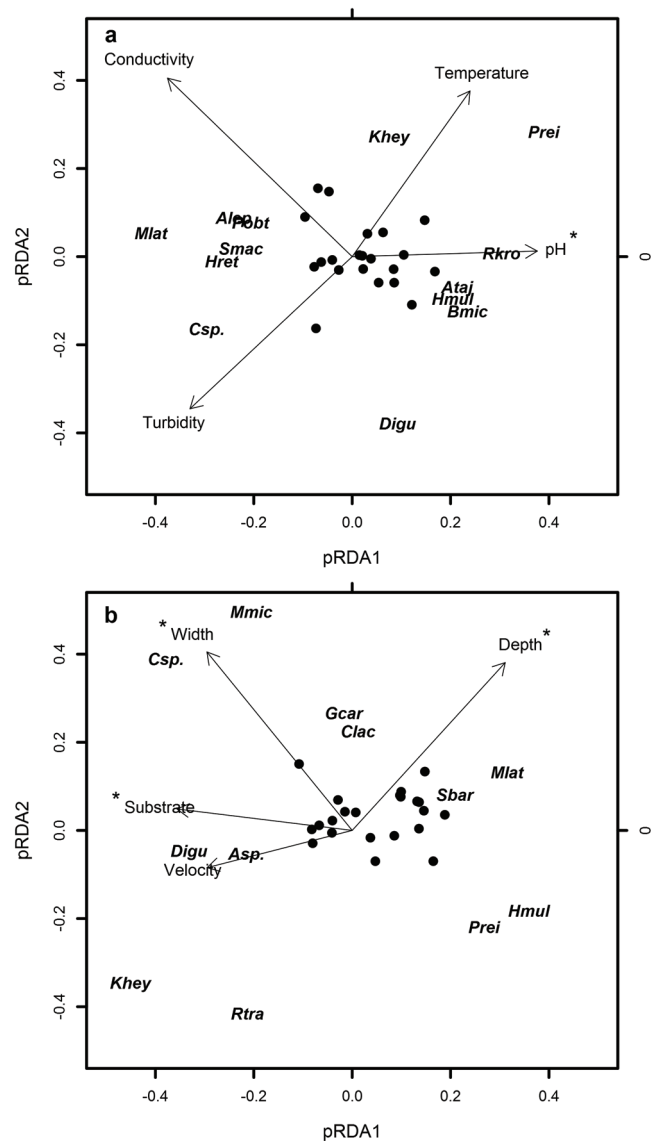


Fig. 3. Results of partial Redundant Analyses (pRDA) showing the patterns of correlation among species abundance with the pure limnological (a), and pure structural (b) predictors. The asterisks indicate environmental variables which were statistically correlated ($p \leq 0.05$) with community composition. Only species with eigenvectors higher than 0.2 were identified by their abbreviated symbols. The others were identified by black dots. This limit was arbitrary but chosen to evidence only the species which were mostly correlated with environmental predictors.

variables (Melo *et al.*, 2011; Landeiro *et al.*, 2012). We did not measure, for example bank slope, riparian vegetation cover, accumulation of wood debris or stream declivity, variables that may influence fish communities (Pulsey & Arthington, 2003; Cruz *et al.*, 2013; Langford *et al.*, 2012). Nonetheless, given the high magnitude of the unexplained variation, it is improbable that they would significantly improve the percent of explanatory variation.

Part of the unexplained variation could also be due to regional factors influencing the organism's dispersion (Falke & Fauch, 2010). In stream habitats, the main rivers could function as dispersal corridors, facilitating the movement of individuals among local communities (Brown & Swan, 2010; Roberts & Hitt, 2010). We found that 19 out of the 34 species captured in the present study (approximately 55%) are present both in streams and the main rivers of the basin (Ferreira & Petrere Jr., 2009). In addition, Mazzoni *et al.* (2004) and Mazzoni & Iglesias-Rios (2007) found evidence of short reproductive migrations for some species of Characidae of Atlantic forest streams, while Mazzoni & Iglesias-Rios (2012) by means of a mark-recapture experiment, found a group of fishes (*Astyanax janaeirensis*, *A. hastatus*, *Parotocinclus maculicauda*, and *Pimelodella lateristriga*), which they called "Long Movement group", that moved at least 6 km within 60 days. Nonetheless, the Mantel test indicated a non-significant influence of site proximities upon assemblage structure, suggesting that including a third spatial matrix would be unhelpful to reduce significantly the unexplained variation of fish-environment association.

Finally, the unexplained variability could be related to our scale of observation. The 50-m long-stretches that we sampled included a variety of mesohabitats of pools, runs and rapids, mainly in the clearwater streams. The influence of mesohabitats within short stretches of Atlantic forest streams was detected by Rezende *et al.* (2010). These authors verified that pool, riffle, and run units varying from 2.10 to 11.7 m-long were significant predictors of fish community structure. Other authors also have found that small patches of mesohabitats (5 m-long) have significant influences on functional fish composition in second and third order streams (Teresa & Casatti, 2012). If mesohabitat heterogeneity within our sampling sites did affect fish composition, which is possible mainly in clearwater streams, averaging the values of habitat predictors and pooling all fishes together might have weakened the observed correlation between fish community composition and local environmental factors and thus, increased the unexplained variation. So far, we cannot test this possibility, but future sampling schemes could be conducted in order to understand how the scale of observation influences our measurement of fish-environment association in these streams.

Despite these possibilities, we verified that high unexplained variances are not unusual in metacommunity studies, and the

reasons for these patterns still need of systematic investigations (Melo *et al.*, 2011). For example, Landeiro *et al.* (2012) partialling out the influences of local environment and space on assemblages of caddisfly larvae in 89 Amazonian streams, found that the percent of unexplained variation ranged from 75.7% to 85.7%, depending on which assemblage subsets were analyzed. Similar fractions, sometimes much higher than 50%, were also found by Falke & Fauch (2010), Siqueira *et al.* (2012), Casatti *et al.* (2012) and Grönroos *et al.* (2013), despite the high number of explanatory variables and the high number of sampling sites included in these studies.

The major limitation of the present study was the small sample size. The low number of blackwater sites did not allow us to investigate the patterns of variation within these streams. This low number of blackwaters in comparison to clearwater streams was partially due to the limited accessibility as they are inside the Restinga forest, and partially because this area is predominantly swampy having few small and shallow streams suitable to electrofishing. The same limitation was found by Gonçalves & Braga (2012) in a nearby basin. On the other hand, the physical habitat structure in clearwater streams (channel width, depth, and substrate) was the main source of variability in community composition.

Despite these limitations, our main conclusions agree with the findings of Gonçalves & Braga (2012) that the lower levels of pH are probably the main environmental filter for blackwater fish communities, and that species associated with these habitats may present some degree of specialization (Menezes *et al.*, 2007). Rocha *et al.* (2007) also stressed that Restinga forests are areas of high endemism threatened by the growing levels of urbanization and fragmentation. Among the five most abundant species, *M. lateralis* and *S. macropterus* are classified as endangered due to their association with Restinga forest streams in the southeastern Brazil (Menezes *et al.*, 2007), and recently, another endangered species (*Rachoviscus crassiceps*) was captured in blackwaters of Itanhaém River basin (Oyakawa & Menezes, 2011). These results are important from the standpoint of conservation planning for that area, since the degree of urbanization in the Itanhaém city has increased from 85.6% in 1970 to 99.59% in 2000 (Pereira, 2002), and because most urbanization pressures are concentrated in the lower stretches of the basin outside from the limits of the Serra do Mar State Park.

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