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Local Landscape Planning and Management in Rural Areas

Hotspots of stream tadpole diversity in forest and agricultural landscapes in Ranomafana, Madagascar

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Abstract

Logging and human-induced conversion of natural forests into agricultural areas are major drivers of biodiversity loss in the tropics. Anuran larvae can be highly diverse, can reach high biomass and can play important roles in tropical streams; yet, compared to the adult frog communities, relatively little is known about how larval communities respond to disturbance. Information on larvae is highly relevant for amphibian conservation because larvae represent direct evidence of breeding and thus provide a good indicator of species persistence in disturbed habitats. We studied tadpole assemblages in Ranomafana, southeastern part of Madagascar, in streams in a disturbed forest (previously logged forest), at "forest edge" (streams embedded in matrix nearby forest blocks), and compared these to communities in a primary forest. We sampled tadpoles at the microhabitat level ("pools" and "rifes") in 9 streams. We recorded 27 species with a maximum of 17 species/stream recorded at edge. The three habitats harbored diferent assemblages, but, as could be expected, more similarities existed among forest habitats than between forest and non-forest habitats. The most and the least diverse communities were recorded at edge and in the disturbed forest, respectively. Assemblages were dominated by one generalist species, and changes in communities were mostly driven by changes in forest specialists, which either decreased in disturbed forest or were replaced by edge specialists outside forest. Although species richness varied, relative abundances were maintained among habitats, suggesting potential compensatory mechanisms in tadpole biomass. Community structure changed at the microhabitat level: pool environments usually harbored relatively higher species richness and abundance than riffles. Our study highlights the relevance of edge habitats for maintaining amphibian diversity and the pronounced negative efects of past logging activities on tadpole communities. Given the diverse roles of tadpoles in streams, changes in community structure potentially afect critical stream ecosystem processes. The study has strong implications for designing bufer zones around protected areas.

Keywords Logging · Amphibians · Degraded habitats · Edge · Anuran larvae · Streams

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Introduction

As the extent of primary forests is shrinking throughout the tropics (Gibson et al. 2011), and given the insufficient protection provided by reserves (Coad et al. [2019](#page-13-1)), there is increasing interest in quantifying the biodiversity values of disturbed habitats (Edwards et al. [2014;](#page-13-2) Laurance et al. [2014\)](#page-14-0). This has particularly been the case for amphibians (e.g., Cushman [2006](#page-13-3); Kurz et al. [2014](#page-14-1); Riemann et al. [2015](#page-14-2); Ndriantsoa et al. [2017\)](#page-14-3) in lights of their alarming global population declines and considering that at least 30% of species in this taxonomic group are facing extinction (Stuart et al. [2004\)](#page-14-4).

Most studies on amphibian disturbance ecology have typically tended to focus on the adult stage (Ernst and Rödel [2005,](#page-13-4) [2008](#page-13-5); Gardner et al. [2007;](#page-13-6) Riemann et al. [2015;](#page-14-2) Ferreira et al. [2016\)](#page-13-7). Relatively next to nothing is known about the efects of disturbance on larval communities, or, reciprocally, the values of disturbed habitats for amphibian breeding and maintenance. In general, studies on tropical tadpole community ecology are scarce (see review in Borges Júnior and Rocha [2013](#page-13-8)), and still little information is available for the larval stage of many tropical amphibians, even for some purportedly abundant species (Wells [2010\)](#page-14-5). Moreover, there has been an idiosyncratic assumption that tadpole communities would simply match the adult community present at a site; this argument may explain the diference in research pace between these two communities. Although reciprocal infuences on communities of adult and larval stages have been documented (Inger et al. [1986](#page-14-6)), this assumption is not satisfactory because adults can be observed from habitats where no breeding takes place (Skelly and Richardson [2010](#page-14-7)) and not all frog species, in the tropics in particular, have their tadpoles develop in water bodies (Wells [2010](#page-14-5)).

Larvae represent concrete evidence of breeding and provide good indicators of species persistence in modifed landscape. Thus, information on larvae is highly relevant for assessing the quality of disturbed habitats. Larval surveys are less likely to overestimate breeding distribution (Skelly and Richardson [2010](#page-14-7)) and can provide critical information on population trajectories and the factors that may afect abundance, distribution, and assemblages (Skelly and Richardson [2010](#page-14-7)). In contrast to the adults that can be cryptic and for which detection rate can considerably vary with sampling efforts, climate, or calling activities (Vonesh et al. [2010](#page-14-8)), tadpoles' detection rate can be relatively high in a relatively defned small area (Skelly and Richardson [2010](#page-14-7)), making studies on larvae highly pertinent for characterizing amphibian community.

The tropical forests of Madagascar are among the most biologically rich and unique of the world (Harper et al.

[2007](#page-14-9)). More than 90% of Madagascar endemic animal species live exclusively in forest and woodland habitats (Irwin et al. [2010\)](#page-14-10). As for other tropical countries (Burivalova et al. [2014](#page-13-9); Laurance et al. [2014](#page-14-0)), habitat loss, mainly due to logging and forest conversion into agricultural areas (e.g., slash-and-burn agriculture), is a major threat to this biodiversity (Irwin et al. [2010](#page-14-10)).

Ranomafana, in the southeastern part of Madagascar, represents a model system for studies on the effects of habitat disturbance on biological communities (Razafmahaimodison [2004;](#page-14-11) Tecot [2008;](#page-14-12) Herrera et al. [2011;](#page-14-13) Gerber et al. [2012;](#page-13-10) Riemann et al. [2015\)](#page-14-2). One part of Ranomafana National Park was selectively logged approximately 30 years ago but has become a protected area ever since. Selective logging negatively impacted forest structure by reducing basal area (m²/ha) by 53%, mean crown volume (m^3) by 17%, and average tree height by 12% (Ramaharitra [2006](#page-14-14); Tecot [2008\)](#page-14-12). The other parts of the park are relatively less disturbed and could be still considered as primary forests (Tecot [2008\)](#page-14-12). Adjacent to the park are matrix, namely agricultural areas dominated by rice paddy felds, rainfed crops, and banana plantations. Matrix, although it is often highly disturbed, might provide valuable habitat for some amphibian species (Ndriantsoa et al. [2017\)](#page-14-3) and, hence, could be an important component of biodiversity maintenance on a landscape scale. The diferences in ecological conditions over short distances in Ranomafana makes it an ideal study site as small-scale contrasts are more sensitive at detecting ecological determinants than comparisons made on larger scales that were often performed in previous amphibian studies (Parris [2004;](#page-14-15) Ernst et al. [2006\)](#page-13-11).

Ranomafana is characterized by its high amphibian diversity with no less than 112 candidate frog species (Vieites et al. [2009](#page-14-16)), and at least 45% of frogs of Ranomafana National Park reproduce in forest streams to form the world richest stream tadpole assemblages (Strauß et al. [2010\)](#page-14-17), with up to 25 species found within a single stream (Strauß et al. [2013\)](#page-14-18). We took advantage of this established scenario to study the changes in tadpole assemblages in streams in a disturbed forest (formerly logged forest), a habitat matrix (streams at the interface between forest block and agricultural landscape), and compared these to assemblages in streams in a primary forest. Because isolating the efects of logging from other confounding disturbance effects (e.g., tourism, invasive species, cyclone) is difficult, we arbitrary referred this habitat as "disturbed forest" although logging has been known to be the major disturbance recorded in that forest (Tecot [2008\)](#page-14-12).

Small changes in vegetation structure can create signifcant alterations to amphibian communities (Cortés-Gómez et al. [2013](#page-13-12)), and logging, even when conducted selectively, can bear dramatic efects on amphibians, especially on forest specialists for which life-history typically rely on forest habitats (Burivalova et al. [2014](#page-13-9); Ferreira et al. [2016\)](#page-13-7). We asked the following questions: do the three habitats harbor similar assemblages? Do communities change at the microhabitat level within and among habitats? Do assemblages seasonally vary across habitats? We expected to fnd the most diverse community (high species richness and relative abundance) in the least disturbed habitat, i.e., primary forest and the least diverse at edge because of the relative high frequency of disturbance (e.g., frequent slash-and-burn). We also predicted that assemblages would change at the microhabitat level as suggested by earlier habitat-relationship models (Strauß et al. [2013](#page-14-18)). Last, we expected to find different assemblages at diferent sampling periods, as previously recorded for tadpole communities in Ranomafana (Strauß et al. [2016](#page-14-19)).

Materials and methods

Ranomafana National Park (RNP) comprises 43,500 ha of continuous mid-altitude mountain rainforest (500–1300 m a.s.l.). Precipitations are high, with alternating periods of low and heavy rains with an annual precipitation between 1700 and 4300 mm (Wright and Andriamihaja [2003\)](#page-14-20). Periods of heavy rains typically occur between January and April. As a result of slash and burn agriculture, landscapes outside RNP consist of forest fragments embedded in a matrix of cultivated land (e.g., banana and rice paddy felds) and secondary vegetation (i.e., grasslands with bush and shrub vegetation).

Sampling procedures

We sampled tadpoles in 9 streams, 3 at each habitat: primary forest (Vatoharanana), disturbed forest (Talatakely, previously logged forest), and in "matrix" (Ambatolahy) (Fig. [1\)](#page-2-0). "Matrix streams" consisted of streams that crossed nonforested areas, embedded in agricultural areas (rice felds, rainfed crops, banana felds), with adjacent riparian vegetation consisting of small trees, bushes, and strawberry guava (generally less than 5 m on each bank). Nearest forest habitat to these matrix streams was at least 50 m aerial distance. To avoid ambiguity, we refer this type of habitat as "edge" in this study. Note that "edge" here may difer from its classical ecological defnition because these streams were not directly adjacent to forest habitat but were more embedded in matrix habitats. These replicates per habitat were realistically the maximum number of streams that met the objectives of the study and were within the same range of altitude (900–1020 m a.s.l). Samplings were conducted in late October–early November 2014 and in March 2015,

Fig. 1 Map of the study sites within Ranomafana National Park and location of RNP within Madagascar

namely at the beginning and in the later part of the rainy season (Strauß et al. [2016\)](#page-14-19).

The streams were second-order streams and were 2.1–3.05 m width. Each stream was distant of at least 200 m with no direct connection between them; so, tadpoles from one stream could not be washed away to another stream. Mean water temperatures were 18–19 °C during the study.

The general sampling procedure followed the methods of Keller et al. [\(2009\)](#page-14-21) in which we studied assemblages at the microhabitat level (stream section). This was done because amphibian assemblage can strongly vary within few meters in streams (Keller et al. [2009](#page-14-21)), and our feld observations along with previous studies indicated that microhabitat heterogeneity can strongly structure tadpole assemblages (Inger et al. [1986](#page-14-6); Eterovick and Barata [2006;](#page-13-13) Afonso and Eterovick [2007](#page-13-14); Eterovick et al. [2010;](#page-13-15) Borges Júnior and Rocha [2013](#page-13-8); Strauß et al. [2013\)](#page-14-18). In each stream, we sampled tadpoles in 4 pools and 4 riffles, each representing section of 2.5 m with at least 10-m stream distance separating two consecutive "microhabitat-sites". Pools represented sections with debris loading and slow-flowing water; riffles designated habitats with relatively fast fowing stream section with the substrate dominated by pebbles. The "microhabitat-sites"

were not chosen systematically (i.e., fxed distance between microhabitats) but rather at random with irregular intervals to cover habitat heterogeneity (substrate, water velocity, canopy openness, water depth, characteristics of the surrounding vegetation). We sampled tadpoles using dipnets of diferent sizes, adjusted to obtain optimal sampling results for each microhabitat. An important component of the feldwork was to standardize sampling effort that would allow estimating tadpole relative abundance. Sampling consisted of dipnetting tadpoles in microhabitats within 4 min (time was stopped during sample processing), to provide perunit-time density estimates following (Werner et al. [2007](#page-14-22)). We assumed that all tadpoles within the microhabitat were caught because we often did not catch any more individuals at the end of each sampling. Samplings were always conducted in the morning. We ensured that no tadpoles moved from one microhabitat to the next one by always starting sampling downstream.

The tadpoles were kept alive and were brought back to the laboratory. The tadpoles were sorted into series based on morphological differentiation. Because of the high number of species and our inability to distinguish all species, we assigned series provisional numbers. We took specimen of each series and after anesthetization by Tricaine Methanesulfonate (MS-222), took a fragment of tadpole tail that was used in DNA analysis for species identification. DNA barcoding was based on a fragment of the mitochondrial 16SrRNA gene (modified 16Sar (550 bp) (5′-CGCCTGTTTAYCAAAAACAT-3′) and modified 16Sbr (550 bp) (5′-CCGGTYTGAACT CAGATCAYGT-3′) following Bossuyt and Milinkovitch [\(2000](#page-13-16)). PCR products were prepared for sequencing using BigDye Terminator sequencing chemistry (Applied Biosystems, CA, USA).

Environmental characterization

We characterized the adjacent forest and riparian vegetation, representing habitat relevant for the adults. Two 5×10 m plots, with the longer side parallel to the stream, were randomly established on each side of a stream (then 4 plots at each stream). We recorded Diameter at Breast Height (DBH) of trees >5 cm to estimate basal area of riparian vegetation. Within each 5×10 m plot, we had a 5×5 m subplot, in which the number of trees $DBH < 5$ cm (shrubs) was counted. Canopy openness of the habitat was estimated at the center of each plot. Two random 1×1 quadrats were set in each 5×5 plot to measure understory height (3 measurements) and litter depth (3 measurements). Measurements were averaged within each 1 m^2 quadrat. Heights of hanging vegetation were also recorded at 2-m interval along a 5×10 m plot. At the center of each plot, we estimated canopy openness using a fish-eye lens mounted on a digital camera. Canopy openness was estimated using the CanopOn2 software ([http://takenaka-akio.org/etc./canopon2/](http://takenaka-akio.org/etc./canopon2/index.html) [index.html](http://takenaka-akio.org/etc./canopon2/index.html)). These measurements were conducted in October–November 2014.

Data analysis

Species diversity, species richness and relative abundance

We used Shannon's *H*′ and Simpson's indexes to measure species diversity in each habitat. These indexes were computed as follows:

$$
H' = -\Sigma \text{pi} \log{(\text{pi})},
$$

where pi is the proportion of individuals belonging to the *i*th species in the habitat.

$$
D = 1 - \Sigma (n/N)^2,
$$

where $n =$ the total number of individuals of a particular species and N =the total number of individuals of all species. The value of this index also ranges between 0 and 1, and the greater the value, the greater the sample diversity.

We conducted two types of analysis that focused on the stream and on the microhabitat levels for species richness and relative abundance. Species richness corresponded to the maximum number of species found in each stream (stream level) or in each microhabitat (pool or riffle level); relative abundance represented the total number of tadpoles sampled from each stream or from each microhabitat. At the stream level, we analyzed the efects of habitat and time of sampling on species richness and relative abundance using linear mixed-efects models with the function "lmer" in the package "lmerTest" in R (Kuznetsova et al. [2017\)](#page-14-23). In these models, "habitat" and "year" were the factors; "stream" was the random factor. At the microhabitat level, these analyses involved "habitat", "microhabitat", and "year" as explanatory variables and "stream" as random factor. *P* values from these models were obtained by *F* tests based on Satterthwaite's method.

Community analysis

We used non-metric multidimensional scaling (NMDS) to visualize and evaluate patterns of dissimilarity in species composition at the microhabitat level between the three habitats for each sampling period. NMDS can handle data with many zeros, ranked and non-normal data better than classical ordination methods (e.g., PCA, CCA), and is well suited for ecological data. Unlike methods that attempt to maximize the variance or correspondence between objects in an ordination, NMDS represents, as closely as possible, the pairwise dissimilarity between objects in a low-dimensional space. That is, microhabitats that are projected closer to each other on the NMDS coordinate system are more likely to harbor similar species than more distant ones. The number of axis was selected based on the lowest stress. As a rule of thumb, a stress value lower than 0.2 represents a good ft of the data (Clarke [1993](#page-13-17)). The ordination was constructed from a Jaccard dissimilarity matrix using species presence/ absence data. NMDS was performed with function "meta-MDS" from R package "vegan" (Oksanen et al. [2016\)](#page-14-24).

We conducted a three-way perMANOVA with the function "adonis2" from R package "vegan" (Oksanen et al. [2016\)](#page-14-24) to test for differences in species composition at the microhabitat and at the habitat levels across the two sampling periods ("habitat", "microhabitat", and "year"). perMANOVA is a powerful permutation method to detect changes in community structure (Anderson and Walsh [2013](#page-13-18)). The three-way perMANOVA was based on Bray–Curtis dissimilarity and 9999 permutations.

We conducted SIMPER analysis (Clarke [1993\)](#page-13-17) with the presence–absence data to break down the contribution of each species to the observed dissimilarity between the habitats. The function performs pairwise comparisons of groups of sampling units and fnds the average contributions of each species to the average overall Bray–Curtis dissimilarity.

Riparian vegetation structure

We analyzed changes in riparian vegetation structure (understory height, litter cover, vegetation cover, canopy cover, basal area, shrub density, riparian vegetation height) between the three habitats using linear mixed-efects models with the function "lmer" in the package "lmerTest" in R (Kuznetsova et al. [2017\)](#page-14-23). We entered "plot" nested in "stream" as random factors and computed the afore-mentioned environmental parameters as response variables. *P* values from these models were obtained by *F* tests based on Satterthwaite's method. Posthoc tests were conducted using least-square means; results of these tests are directly displayed on the fgures. Data were log-transformed before analysis.

Mixed-effects models, NMDS ordination, and per-MANOVA were performed in R 3.3.3 (R Core Team [2017](#page-14-25)). The SIMPER analysis and the graphs were made on PAST 3.0 (Hammer et al. [2001](#page-14-26)).

Results

We recorded 4444 individuals of 27 species (["Appendix 1](#page-10-0)") and 2764 individuals of 16 species of the family Mantellidae recorded in the beginning (October–November 2014) and in the later part (March 2015) of the rainy season, respectively.

Species diversity and richness

Primary forest and edge were the most diverse communities with edge having higher *H'* value than primary forest. Diversity varied with the period of sampling and in 2015, primary forest harbored the highest species diversity (Table [1\)](#page-4-0). For both diversity indexes, the disturbed forest had the lowest values across the two sampling periods.

In 2014, edge habitats harbored the highest species richness with 21 species (range=14–17 species/stream), followed by primary forest with 17 species (range $=11-15$) species/stream). The lowest species richness was recorded in the disturbed forest with 12 species (range $=8-10$ species) stream). Species richness was lower in 2015 and the highest diversity was then recorded in primary forest with 13 species (range=7–11 species/stream), disturbed forest with 10 species (range=4–7 species/stream). The lowest species richness was recorded at edge with 9 species (range=5–9 species/stream) (Table [1](#page-4-0); Fig. [2](#page-5-0)).

At the stream level, species richness signifcantly difered between the three habitats and sampling period; the interaction between the two factors was marginally signifcant (Table [2](#page-6-0)). At the microhabitat level, habitat, microhabitat, and sampling periods infuenced species richness (Table [3](#page-6-1)). Species richness signifcantly difered between microhabitats within the same habitat (Table [3\)](#page-6-1). Pools generally harbored

Table 1 Species richness, relative abundance, and diversity of stream tadpoles in primary forest, disturbed forest, and at edge

Fig. 2 Tadpole relative abun dances in streams $(n=3$ for each habitat) at edge, in disturbed forest, and in primary forest at the beginning (2014) and in the later part (2015) of the rainy season. Genus names were abbreviated in the graph as fol lows: *Boophis* (B.), *Mantidac tylus* (M.), *Spinomantis* (S.), *Gephyromantis* (Ge.), *Guibe mantis* (Gu.)

Table 2 Results of linear mixed-efects models testing the effects of habitat and time of sampling on tadpole species richness and relative abundance at the stream level

Bold denotes signifcant efect

Table 3 Results of linear mixed-efects models testing the efects of habitat, microhabitat, and time of sampling on tadpole species richness and relative abundance at the microhabitat level

Bold denotes signifcant efect

higher species richness than riffles (average species number per microhabitat, 2014: 7.62 vs 5.83, 2015: 4.75 vs. 4.25).

Relative abundance

At the stream level, tadpole relative abundance did not significantly differ among the three habitats, though there was tendency for primary forest to harbor more tadpole individuals in streams. This was because there was high variation in tadpole relative abundance among streams (Table [1\)](#page-4-0). At the microhabitat level, microhabitat and sampling period influenced tadpole abundance. Pools significantly harbored higher number of tadpoles (average number of tadpole individual per microhabitat, 2014: 87.51 vs. 38.36; 2015: 55.87 vs. 30.5). Significantly higher number of tadpoles was recorded in October 2014 than in March 2015 (Fig. [2;](#page-5-0) Tables [1,](#page-4-0) [2](#page-6-0)).

Tadpoles of the genus *Boophis* dominated the assemblages in all habitats (Fig. [2](#page-5-0)). The genus *Gephyromantis* and *Guibemantis* were represented by one species, respectively. Patterns of species abundance show that by far the most abundant species were *Boophis quasiboehmei*, *B. madagascariensis,* and *B. reticulatus* (Fig. [2\)](#page-5-0). These species were ubiquitous in all streams with *B. quasiboehmei* being the dominant species in all streams and were abundant in all sampling periods (Fig. [2\)](#page-5-0). *B. andohahela*, *B. tasymena,* and *B. sp37* were exclusively recorded at edge where *B. picturatus* was also rare*. Spinomantis perraccae* and *S. aglavei* were absent outside forest and could be considered forest specialists.

Community diversity and structure

The general stress coefficients of NMDS models were 0.119 and 0.105 in 2014 and 2015, respectively, indicating good preservation of ordering relationships of the multidimensional among-microhabitat dissimilarities. Primary forest and disturbed showed overlaps in community structure (Fig. [3\)](#page-7-0).

Results of three-way perMANOVA (Table [4](#page-7-1)) indicated that species assemblages changed at the microhabitat (pools and riffles) and at the habitat levels, and between sampling periods. Tadpole communities significantly changed between microhabitats within each habitat and between sampling periods within each habitat (for all pairwise tests $P < 0.001$, following Bonferoni *P* value corrections, ["Appendix 2](#page-10-1)").

SIMPER analyses revealed that overall dissimilarity between the primary forest and the disturbed forest was 35.43% (vs. 41.55% in 2015), 61.25% (vs. 66.88% in 2015) between the primary forest and the edge, and 54.19% (vs. 62.67% in 2015) between the disturbed forest and the edge. "Specialists" (i.e., species that were only recorded in forest habitats or at edge) mostly explained these dissimilarities (["Appendix 3](#page-10-2)").

Riparian habitat structure

For the parameters we measured, riparian vegetation mainly differed in basal area, litter depth and canopy cover (Table [5](#page-8-0)); we did not fnd signifcant diferences in any other variables. As could be expected, forest habitats had higher basal area, thicker litter, and lower canopy

Fig. 3 Non-metric multidimensional scaling (NMDS) showing differences in species composition between microhabitats in streams at edge (red), disturbed forest (blue), and in primary forest (green) at the beginning $(2014, NMDS stress=0.119)$ and in the later part $(2015,$

NMDS stress=0.105) of the rainy season. Ordination was based on Jaccard dissimilarity using presence–absence data (color fgure online)

Table 4 Results of three-way perMANOVA based on Bray–Curtis dissimilarity analyzing changes in community structure at the microhabitat and at the habitat levels, and between sampling periods

	df	Sum Sq	F	P
Habitat	$\mathfrak{D}_{\mathfrak{p}}$	7.51	40.93	< 0.001
Microhab	1	1.18	12.96	< 0.001
Year	1	1.10	12.02	< 0.001
Habitat: microhab	2	0.49	2.70	0.003
Habitat: year	2	0.45	2.45	0.007
Microhab:year	1	0.13	1.48	0.173
Habitat:microhab:year	2	0.25	1.37	0.176
Residuals	123	11.28		

Bold denotes signifcant efect

openness. Disturbed forest tended to have higher density of shrubs per unit of area compared to the other habitats but this was not signifcantly diferent from the other habitats.

Discussion

As for many other tropical countries, logging and conversion of natural forests to agricultural areas are major threats to biodiversity in Madagascar. Given that current protected areas may not be sufficient in maintaining all extant species in the long term (Coad et al. [2019\)](#page-13-1), it is important to quantify the conservation values of disturbed habitats around protected areas (Irwin et al. [2010\)](#page-14-10). We found that tadpole community structures in disturbed forest and at edge markedly difered from the ones recorded in primary forest.

We expected the highest species diversity in primary forest, but in contrast to our predictions, the highest and the lowest species diversity were recorded at edge and in the disturbed forest, respectively. Logging activities occurred in 1989 and had simplifed forest structure by reducing tree basal area (by 53%) and crown volume (by 17%) in this part of the forest of Ranomafana National Park (Ramaharitra [2006](#page-14-14); Tecot [2008](#page-14-12)). The effects of selective logging on tropical forests are often negative (see review in Burivalova et al. [2014](#page-13-9)) and can halve amphibian richness, especially those forest specialists, at logging intensities of $63 \text{ m}^3/\text{ha}$ (Burivalova et al. [2014\)](#page-13-9). Though it is difficult to compare this value with data available on logging intensity in Ranomafana, the disturbed forest harbored a significantly lesser number of species (species richness 12 vs. 17) and markedly lower abundance of forest specialist species (*Spinomantis* species and *B. picturatus*) than the primary forest, suggesting that logging could be one driver of community dissimilarity between the two habitats.

Spinomantis often call from canopies of large trees and are known to be restricted to undisturbed habitats (Glaw and Vences [2007](#page-14-27)); thus, they are likely very sensitive to logging. Species in this genus partly explained the community dissimilarity between the forest habitats by having **Table 5** Characteristics of riparian vegetation between primary forest, disturbed forest, and forest edge

Bold denotes signifcant diference

lower incidence (here presence or absence in microhabitat) and lower abundance in the disturbed forest. Structure of the riparian vegetation may have little infuence on com-munity dissimilarity (Table [5](#page-8-0)), which could be not surprising because logging majorly targeted the upper part of the forest. It would be misleading, however, to assume that logging is the only source of disturbance in this forest. In fact, high tourism activity and the invasion of strawberry guava *Psidium cattleianum* in this part of Ranomafana National Park are potential factors that may affect frog populations; their effects on amphibians are unknown though.

Edge harbored the highest species richness with up to 21 species in streams. This number is much lower than the 34 species recorded by Ndriantsoa et al. [\(2017\)](#page-14-3) in matrix streams in Ranomafana. Two reasons may explain this difer-ence; first, Ndriantsoa et al. ([2017](#page-14-3)) surveyed more streams (5 streams vs. 3 streams in this study) and focused on the adult populations using call surveys. In this respect, they were likely to detect higher number of species if frogs call from habitats where no breeding occurs (thus no larvae). Second, not all frog species have their tadpoles develop in streams. The question is why relatively more species were detected outside forest. Earlier studies suggested that factors for the maintenance of amphibian diversity in disturbed habitats are vegetation structure and more importantly the availability of breeding habitats (Bickford et al. [2010;](#page-13-19) Riemann et al. [2015\)](#page-14-2). In matrix and fragmented landscapes in Ranomafana, the presence of stream is an important factor of high species richness (Riemann et al. [2015](#page-14-2); Ndriantsoa et al. [2017\)](#page-14-3) independently of the surrounding forest type. Diversity in degraded habitats can be equal (Riemann et al. [2015\)](#page-14-2) or can even be higher than of primary forests (this study). This is interesting because edge efects on amphibians are not always positive. For example, (Schneider-Maunoury et al. [2016\)](#page-14-28) reported decreased abundance in three-quarter of amphibian species with proximity to edge in a neotropical fragmented landscape. However, the authors noted that species-specifc edge efects were not always consistent and some species can have opposite edge responses when measured in diferent landscapes (Schneider-Maunoury et al. [2016\)](#page-14-28). This could be because species have diferent tolerance to modifed habitats (Laurance [1991](#page-14-29)). For example, frog species richness was higher in forest fragments compared to forest block in Amazonia because some species were associated with matrix habitats and many of primaryforest species used these habitats as breeding sites (Gascon et al. [1999](#page-13-20)).

The high species richness at edge is suggested to be result of shared species between forest and edge habitats (increase of generalists), and because of some species that were only recorded at edge (prevalence of edge specialists) (Lövei et al. [2006](#page-14-30)). As forest specialists declined (e.g., *Spinomantis* species, *B*. *picturatus*), other species with niches better suited to the new environmental conditions composed communities at edge, eventually helping diversity to be maintained (Rie-mann et al. [2015](#page-14-2)) or even higher at edge (this study). Species that were only recorded at edge were species in the genus *Boophis*: *B. andohahela*, *B. tasymena, B. elenae, B. luteus, B. luciae, B. periegetes,* and *B. sp37.* Many species in the genus *Boophis* are most abundantly in open areas in altered habitats (Andreone [1994\)](#page-13-21), but probably not all of these afore-mentioned species are edge specialists because an earlier study recorded at least *B. luteus* in continuous forest (Riemann et al. [2015](#page-14-2)). Glaw and Vences ([2007\)](#page-14-27) described *B. andohahela* as a forest specialist, but along with Strauß et al. ([2013](#page-14-18)) we found that this species, at least its larvae, can also adapt to degraded habitats.

The relatively high species richness at edge is intriguing. Edge habitats are characterized by higher temperatures, increased wind speed, and decreased relative humidity (Lehtinen et al. [2003](#page-14-31)), to which amphibians are particularly sensitive. Species at edge could be adapted to open habitats and may even be specialized on disturbed habitats. It is possible that species that were only detected at edge may also occur inside RNP, but some indeed may be restricted to edge habitats. The eastern rainforest belt of Madagascar was originally completely forested and thus, the majority of the species in this study should be forest species. However, frequent natural disturbance such as cyclones infuence forest structure and microclimate and may have favored amphibian adaptation to disturbed habitats. Thus, species adapted to natural disturbance may have better ability to cope with anthropogenic disturbance (Riemann et al. [2015\)](#page-14-2). Andreone [\(1994](#page-13-21)) hypothesized that stream-dwelling species depend less on the microclimatic conditions of the forest foor and may adapt to disturbed environments.

Assemblages changed at the microhabitat level in streams, species richness and abundances were relatively higher in pools than in riffles. The tadpoles of many frog species have affinity to still and slow-flowing stream sections where leaf litter accumulates (Wells [2010\)](#page-14-5). Litter can represent important refuge and food resources for tadpoles (Ramamonjisoa and Natuhara [2018\)](#page-14-32). Even in riffles where gravels represent the main substrate, the tadpoles mainly occupied the slow-running parts of these microhabitats. Indeed, few tadpole species have evolved adaptation to riffle microhabitats; the tadpoles of *B. picturatus* are characterized by an extremely derived oral disc without any keratodonts and with completely reduced jaw sheaths and are known to ingest sand particles (Grosjean et al. [2011\)](#page-14-33). The tadpoles of *B. andohahela* and *B. marojejiensis* have enlarged suctorial mouthparts (nozzle-shaped oral disk) allowing these species attach to rocks and boulders, likely an adaptation to circumventing strong current (Wells [2010](#page-14-5)).

An interesting aspect of the tadpole communities in Ranomafana is the dominance of one species *B. quasiboehmei*. Although species richness differed among habitats, relative abundances were maintained among the three habitats. *B. quasiboehmei* seems to compensate for decline in abundance of other species in forest habitats while outside forest, increases in abundance of other "edge" species allowed abundances to be maintained (Fig. [2](#page-5-0)). These indicate some signals of compensatory mechanisms in which declines in biomass by some species are compensated for by increases in others, eventually allowing distributions of abundance to be maintained (Brown et al. [2001;](#page-13-22) Morgan Ernest and Brown [2001](#page-14-34); Dornelas [2010](#page-13-23)). This has an important implication for ecosystem functioning and stability given that changes in tadpole biomass can have signifcant efects of stream ecosystem processes (Ramamonjisoa and Natuhara [2018](#page-14-32)).

It is unclear from this study whether the populations recorded at edge were simply tadpoles that were fushed downstream after heavy rains from forested parts on higher altitude. However, this might not be the case because earlier studies in the same study site (Riemann et al. [2015](#page-14-2); Ndriantsoa et al. [2017\)](#page-14-3) reported similar diversity outside forest, suggesting that species that were recorded at edge could be already established populations. Moreover, heavy rains typically occur between January and February in Ranomafana (Strauß et al. [2016\)](#page-14-19). Thus, we believe that at least our frst sampling in October 2014 provided a good characterization of tadpole community structure. Communities in the earlier part of the rainy season were more diverse (higher number of species and higher relative abundance) than the ones recorded in the second sampling period. Community at edge exhibited the biggest change in species composition, going from having the richest to the lowest species richness across years among the three habitats. We do not have a clear explanation for this result but it is possible that because samplings were conducted after the period of heavy rains, the tadpoles could have been fushed due to strong currents in streams at edge. Another explanation is that tadpoles at edge may have metamorphosed earlier due to relatively higher water temperature (18 vs. 19 °C) and likely higher resources availability (Ramamonjisoa, unpublished data). Both factors are known to infuence growth and metamorphosis in tadpoles (Alvarez and Nicieza [2002](#page-13-24)).

Conclusions and conservation implications

Primary forests are often labeled "irreplaceable" for sustaining tropical biodiversity (Gibson et al. [2011\)](#page-13-0). However, increasing loss of natural habitats and the limitation of current established protected areas in conserving biodiversity in the tropics have called for the need to assess the values of human-modifed landscapes and evaluate the relevance of degraded habitats for amphibian conservation (Irwin et al. [2010](#page-14-10); Riemann et al. [2015](#page-14-2)). Considerable number of threatened and data-defcient amphibian species is currently outside protected areas for many of which distribution is limited to very small area (Ramamonjisoa et al. [2013](#page-14-35); Nori and Loyola [2015](#page-14-36)). Edge habitats represent typical "hotspots" because of their high species richness and high level of disturbance, and represent priority habitats in conservation planning. The values of edge habitats in maintaining amphibian diversity are nevertheless likely to depend on the distance from forest block due to dispersal limitation of the adults, suggesting that the quality of matrix is of paramount importance for the conservation of amphibians in degraded habitats (Ndriantsoa et al. [2017\)](#page-14-3).

While studies on amphibian disturbance ecology have typically focused on the adult phase, we call for more studies focusing on the larvae. Larvae are good indicators of the quality of disturbed habitats as they represent direct evidence of breeding and species persistence; thus, surveys limited to adult populations may be less informative

for predicting population dynamics. Moreover, given that tadpoles can infuence critical ecosystem processes in tropical streams (Ranvestel et al. [2004](#page-14-37); Colón-Gaud et al. [2008](#page-13-25); Rugenski et al. [2012;](#page-14-38) Ramamonjisoa and Natuhara [2018](#page-14-32)), information on larvae is needed.

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Appendix 1

Species of tadpole recorded in streams in primary forest, disturbed forest (selectively logged forest), and at forest edge in Ranomafana.

D data deficient, *NT* near threatened, *LC* least concern, *VU* vulnerable, *EN* endangered

Appendix 2

Pairwise diferences following per MANOVA on species composition between the three habitats.

2014

Appendix 3

SIMPER analyses on species compositional similarities between primary forest, disturbed forest, and edge.

2014

Appendix 4

Sampling design

In each habitat, we sampled three streams. In each stream, we sampled tadpoles in 4 pools and in 4 riffles.

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