

# Disease reservoirs threaten the recently rediscovered Podocarpus Stubfoot Toad (*Atelopus podocarpus*)

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Abstract.—The Andes have experienced an unprecedented wave of amphibian declines and extinctions that are linked to a combination of habitat reduction and the spread of the fungal pathogen, *Batrachochytrium dendrobatidis* (*Bd*). In the present study, a range of high-altitude habitats in Southern Ecuador were surveyed for the presence of *Bd*. With a particular focus on Yacuri National Park, infection data are presented from across the resident amphibian community. This community contains a once putatively extinct species which was rediscovered in 2016, the Podocarpus Stubfoot Toad (*Atelopus podocarpus*). Across species, local *Bd* prevalence was 73% in tadpoles (n = 41 individuals from three species) and 14% in adults (n = 43 individuals from 14 species). Strikingly, 93% (14/15) of tested tadpoles of the recently described local endemic, *Gastrotheca yacuri*, were infected with a high pathogen load, suggesting that this species likely acts as a reservoir of infection in Yacuri. These findings show that the threat of disease for *A. podocarpus* still exists, and that this species requires urgent action to ensure its survival.

Keywords. Amphibian, Anura, chytrid, conservation, Ecuador, emerging infectious disease, Gastrotheca

Resumen.—Los Andes han experimentado una ola sin precedentes de declinaciones y extinciones de anfibios que están vinculadas a una combinación de factores como la reducción de hábitat y la dispersión del hongo patógeno Batrachochytrium dendrobatidis (Bd). En el presente estudio, muestreamos la presencia de Bd en un rango de hábitats de altura en el sur de Ecuador. Nos enfocamos, particularmente, en el Parque Nacional Yacuri, de donde presentamos datos de infección a través de la comunidad residente de anfibios. Esta comunidad incluye una especie anteriormente considerada como extinta la cual fue redescubierta en 2016, la Rana Arlequín o Jambato de Podocarpus (Atelopus podocarpus). La prevalencia local de Bd fue 73% en renacuajos (n = 41 individuos de tres especies) and 14% en adultos (n = 43 individuos de 14 especies). Sorprendentemente, el 93% (14/15) de los renacuajos examinados de la especie endémica, recientemente descrita, Gastrotheca yacuri, estuvieron infectados con una alta carga del patógeno, sugiriendo que esta especie, probablemente, actúa como un reservorio de infección en Yacuri. Nuestros hallazgos muestran que la amenaza de la enfermedad para A. podocarpus aún existe, y que esta especie requiere acción urgente para asegurar su supervivenciae.

Palabras clave. Anfibios, Anura, quitridio, conservación, Ecuador, enfermedad infecciosa emergente, Gastrotheca

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### Introduction

Amphibians have experienced marked population declines and extinctions across all continents where they occur, and represent a particularly prominent example of the current biodiversity crisis (Hoffmann et al. 2010; Wake and Vredenburg 2008). A main driver for these declines is chytridiomycosis, an infectious disease caused by the fungal pathogen, *Batrachochytrium dendrobatidis* (hereafter *Bd*), which may act synergistically with other



**Fig. 1.** The Podocarpus Stubfoot Toad (*Atelopus podocarpus*) was rediscovered along a single stream in Yacuri National Park (Ecuador) in 2016, after having been presumed extinct in the years following the last previous sighting in 1994. *Photo by Phil Jervis*.

threats such as habitat destruction and climate change (e.g., Hof et al. 2011). *Bd* has affected hundreds of amphibian species globally, and is regarded as the most devastating vertebrate disease ever recorded (Scheele et al. 2019).

The invasion and rapid spread of Bd (global panzootic lineage, BdGPL; Farrer et al. 2011) throughout the Neotropics during the 1980s led to severe population declines in susceptible amphibian species (Pounds et al. 2006; Lips et al. 2006). Amphibian communities at high-elevation sites in the Neotropics, such as the Ecuadorian Andes, occur in the global ecoregions which have been most severely affected by chytridiomycosis (Catenazzi et al. 2011; O'Hanlon et al. 2018). Severe declines have led to range contractions and extinctions of many species, particularly those confined to breeding in high-elevation streams and lakes (Merino et al. 2005; Ron et al. 2003). However, despite the scale of these declines, the epidemiology of chytridiomycosis in the Ecuadorian Andes is not yet well understood. Bd is thought to occur throughout much of the country, and it is generally assumed to currently display an enzootic persistence characterized by low prevalence following the epizootic phase nearly 30 years ago (Catenazzi et al. 2011). Enzootic conditions are, amongst other factors, facilitated by infected reservoir host species which tolerate *Bd* and maintain infection within local amphibian communities after the functional extinction of more susceptible populations (Brannelly et al. 2018; Haydon et al. 2002).

With 64 species currently listed as either Critically Endangered or Extinct on the IUCN Red Lists, the genus *Atelopus* has been disproportionately affected by *Bd* (La Marca et al. 2005; IUCN 2020). In Ecuador, the

Podocarpus Stubfoot Toad, Atelopus podocarpus, was described from museum specimens after the last known individual died shortly after capture in 1994 (Loetters et al. 2011). In line with many other Ecuadorian Atelopus species, anecdotal evidence indicates that before the onset of catastrophic declines, A. podocarpus was relatively common within its range (Ron et al. 2003). In 2016, three individuals of A. podocarpus were rediscovered along a single stream in Yacuri National Park by a field team from the Museum of Zoology (QCAZ), of the Pontificia Universidad Católica del Ecuador (e.g., Fig. 1). However, searches in the surrounding habitat were unsuccessful in finding additional individuals or new populations. The specific aims of the present study are to: (1) survey a range of high-altitude sites in southern Ecuador for Bd presence and prevalence in both tadpoles and adults of all encountered species; (2) revisit the last known site of A. podocarpus in Yacuri National Park, to survey for additional individuals of this species and to establish the Bd infection status of the local amphibian community; and (3) identify potential Bd reservoir species which maintain local infection.

## Methods

Fieldwork was conducted at six sites during 13–21 June 2018, which is in the dry season. Locations were selected at a range of elevations encompassing habitats of previous *Atelopus* occurrences, covering an elevation range of 1,014–3,423 m from eastern foothill forest (Zamora) and eastern montane forest (San Francisco and Loja) to Paramo/Subparamo (Urdaneta, Madrigal del Podocarpus Reserve and Yacuri National Park, see Fig. 2). The ecology of *A. podocarpus* is unknown. However,

Jervis et al.

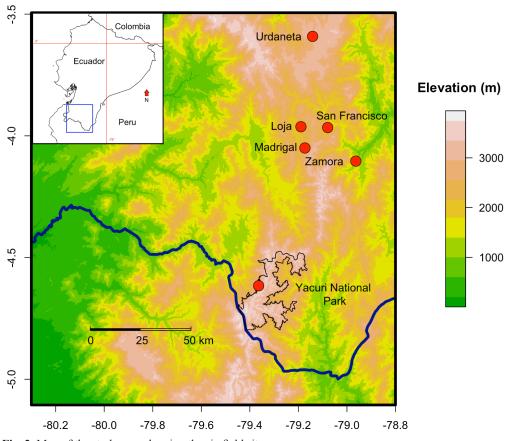


Fig. 2. Map of the study area showing the six field sites.

the closely related *A. ignescens* has been observed breeding in December–January (Peters 1973), so it is likely that sampling occurred outside the breeding season of *A. podocarpus*.

All individuals in this study were captured by hand or with a small fishing net, swabbed with a sterile cotton swab (Medical Wire 100), and immediately released at the site of capture. Tadpoles had their mouthparts swabbed (only the mouthparts of tadpoles can be infected by Bd, Hyatt et al. 2007). Post-metamorphic individuals were swabbed by taking five strokes at the center of the underside, on each flank, the inside of the legs, and the bottom of each of the rear feet. Animals contaminated with soil were washed before swabbing to remove debris, using water purified through mechanical, active carbon, and UV filtration. Swabs were stored in an icebox in the field when possible, until they were returned to the lab and refrigerated below 4 °C until analysis. The coordinates and elevation of each sampled animal was georeferenced using a Garmin GPS. All equipment used in environmental sampling or for handling animals was sterilized in 5% chlorhexidine solution between sites to prevent contamination of potentially disease-naïve sites. Each individual was contained in a new plastic bag and handled with a fresh pair of nitrile gloves to prevent cross-infection. Gloves and bags were disposed of following return from the field.

DNA extractions of swabs were performed using

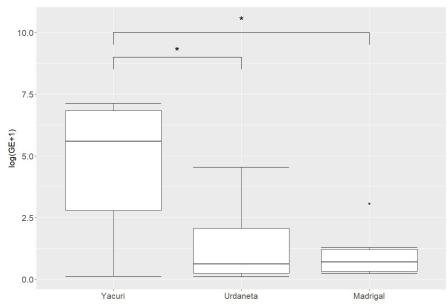
Prep-Man extraction kits (Hyatt et al. 2007), followed by a qPCR-based standard protocol for the quantification of *Bd* prevalence and infection burden (Boyle et al. 2004). Standard curves were generated using 0.1, 1, 10, and 100 *Bd* zoospore standards of BdGPL isolate IA042. To reduce PCR inhibition, samples were diluted 1:10 and the infection burden was multiplied by 10. Infection burden was defined as the number of zoospore genomic equivalents (GE) per swab following Clare et al. (2016). The sample was considered *Bd* positive if both replicates amplified above 0.1 GE.

Differences of infection intensities (GE) between sites were tested using ANOVA with a post-hoc pair-wise Tukey HSD test using R version 3.6.2 (R Development Core Team 2013).

### Results

A total of 41 pre-metamorphic and 43 post-metamorphic individuals representing 18 species from seven sites were swabbed (Table 1). A total of 36 *Bd* positive animals were recorded (30 tadpoles of *Gastrotheca* sp. and six adults from four species in five genera), equating to a total infection rate of 43% (73% for tadpoles and 14% for adults). *Bd* was detected at five of the six sampling sites.

A single male *A. podocarpus* was found along the same stream as the previous expedition at Yacuri National Park. However, further searches along 3.5 km of the



**Fig. 3.** Comparison of infection burdens in *Gastrotheca* tadpoles between three of the six sites. The asterisks denote significant differences. The boxplot was produced in ggplot 2 (Ginestet 2011).

streams surrounding the lake system were unsuccessful at locating any more individuals. Tadpoles of *G. yacuri*, a recently described, locally endemic species (Carvajal-Endara et al. 2019), were characterized by particularly high *Bd* prevalence (93%, n = 15) and infection loads. For *Gastrotheca* species in general, Yacuri had a significantly higher infection burden (P < 0.05) than Madrigal and Urdaneta, the other two sites for which tadpoles were analyzed (Fig. 3).

# amphibians would correspond well with an enzootic disease system (Catenazzi et al. 2017), although the sample size here does not allow unambiguous discrimination between sampling biases related to taxon and life stage. Moreover, an enzootic state does not necessarily equate to stable populations, and further declines are also possible when populations have been affected by *Bd* over significant periods of time (Longo and Burrowes 2010). *Gastrotheca* sp. tadpoles were particularly highly infected by *Bd* (Table 1), suggesting that this genus acts as a reservoir for local disease presence. *Gastrotheca* is a widely-distributed and locally common genus throughout the Andes which can often

and sites. The lower prevalence of Bd amongst adult

### Discussion

This survey found that Bd is widespread in Southern Ecuador, but unevenly distributed between species

Table 1. *Bd* prevalence and mean infection burden by species at each site. The *Atelopus podocarpus* swab was lost in transit from the field site. GE indicates the number of zoospore genomic equivalents as a measure of infection intensity.

Species	n	Life stage	Site	<b>Bd</b> positives	Prevalence	GE (mean)
Boana fasciata	8	Adult	Zamora	1	13% (1–53%)	3.17
Dendropsophus rhodopeplus	6	Adult	Zamora	2	33% (4–78%)	2.52
Dendropsophus sarayacuensis	2	Adult	Zamora	2	100% (16–100%)	1.87
Gastrotheca elicioi	9	Adult	Loja	0	0 (0-34%)	0
Gastrotheca pseustes	10	Tadpole	Urdaneta	6	60% (26-88%)	17.71
Gastrotheca aff. pseustes	16	Tadpole	Madrigal	10	63% (35-85%)	3.08
Gastrotheca yacuri	15	Tadpole	Yacuri	14	93% (68–100%)	465.00
Pristimantis atratus	1	Adult	San Francisco	0	0 (0–97.5%)	0
Pristimantis cf. cajamarcensis	1	Adult	Madrigal	0	0 (0–97.5%)	0
Pristimantis (Huicundomantis) sp.	1	Adult	Madrigal	0	0 (0–97.5%)	0
Pristimantis lymani	1	Adult	San Francisco	1	100% (2.5–100%)	6.22
Pristimantis multicolor	4	Adult	Yacuri	0	0 (0-60%)	0
Pristimantis orestes	1	Adult	Urdaneta	0	0 (0–97.5%)	0
Pristimantis sp. 1	2	Adult	Zamora	0	0 (0-84%)	0
Pristimantis sp. 2	1	Adult	San Francisco	0	0 (0–97.5%)	0
Pristimantis tiktik	4	Adult	Urdaneta	0	0 (0-60%)	0
Rhinella marina	2	Adult	Zamora	0	0 (0-84%)	0

Amphib. Reptile Conserv.

tolerate human modified landscapes. Larval anurans, including *Gastrotheca* tadpoles, are generally tolerant to *Bd* infection (Grogan et al. 2018) as their lack of keratinized skin prevents disease progression until metamorphosis.

As a hypothesis, this situation allows for the proliferation of the pathogen in the habitats under investigation, following the declines and extinctions of more susceptible potential hosts such as Atelopus sp. (Haydon et al. 2002; La Marca et al. 2005; Woodhams et al. 2006). The life history of tadpole-producing Gastrotheca species in high-elevation habitats could allow for the continuous persistence of Bd in breeding pools. Mating takes place on land, and eggs hatch into larvae in the pouch of females which deposit advanced tadpoles into the breeding pool. Many Gastrotheca species do not have fixed reproductive periods and will breed whenever conditions are favorable (Del Pino 1989). Therefore, the combination of slow development and overlapping generations of tadpoles in permanent pools at high elevations creates an ecological system which has previously been characterized by high infection prevalence combined with Bd transmissions to tadpoles of other species (e.g., Alytes obstetricans: Bates et al. 2018 and Rana muscosa: Clare et al. 2016; Rachowicz and Briggs 2007). Although this study represents the first detection of Bd in G. yacuri, tadpoles of the closely related G. riobambae have previously been shown to be capable of maintaining infection across multiple cohorts in a breeding pond monitored over a 9-month period (S. Ron, unpub. data). Future research is needed to determine the potential for Gastrotheca species to act as Bd reservoirs in high-altitude habitats over an extended monitoring period.

Rediscoveries of Atelopus sp. in Bd positive sites are not unusual (Perez et al. 2014; Tapia et al. 2017; Lampo et al. 2011, 2017). However, many of these sites are at lower elevations than Yacuri and possess more diverse amphibian communities, leading to a wider array of options for cross-species infection dynamics. Being home to a smaller number of species, Yacuri National Park could thus be used as an accessible model system to infer the processes which allow for the coexistence of susceptible amphibians within a Bd positive community. An extremely high infection burden in breeding pools was discovered within 50 m of the locality where the four remaining A. podocarpus individuals were found (one individual during this study in 2018, and three individuals discovered in 2016). This suggests that A. podocarpus is still perennially exposed to Bd and therefore at risk of chytridiomycosis, although infection data are unavailable due to the loss of the skin swab from the individual sampled on the second expedition. The high-prevalence, high-intensity infection pattern found here is often seen in epizootic systems, and quantitative population data are required to assess the impact of Bd on

both G. yacuri and A. podocarpus.

A high prevalence of *Bd* in Yacuri could inhibit the proliferation of relic populations of A. podocarpus, and is a major cause for concern for future conservation initiatives. For possible future ex-situ breeding, investigations into the availability of founder individuals are seen as a priority for the species (Conservation Needs Assessment 2012). However, more information is needed on the infection status and size of the population and, until such data are available, we do not recommend the collection of individuals for *ex-situ* conservation. All recent discoveries of this species have been in Yacuri National Park, and all visitors and guides must register in the park office. Hence, dissemination for a citizen-science monitoring project would be relatively straightforward. We also recommend further investigation into the possibilities of improving the habitat for this species, for example through the removal of introduced trout (Martín-Torrijos et al. 2016; Mouillet et al. 2018).

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**Robert Jehle** had a childhood interest in amphibians and their habitats, and feels very fortunate that he could translate that passion into a professional career. Robert is currently a Reader in Population Biology at the University of Salford (United Kingdom), where he teaches in a range of undergraduate and postgraduate programs in Zoology and Wildlife Conservation. His main research area revolves around the ecology, evolution, and behavior of amphibians at the level of populations, often combining evidence from genetic markers with life-history inferences. Robert is a former (2009–2015) Editor of *Herpetological Journal*, and a current (2009–date) Associate Editor of the journal *Animal Conservation*.



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