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Short communication

Infection of an invasive frog *Eleutherodactylus coqui* by the chytrid fungus *Batrachochytrium dendrobatidis* in Hawaii

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Abstract

The chytrid fungus *Batrachochytrium dendrobatidis* has contributed to declines and extinctions of amphibians worldwide. *B. dendrobatidis* is known to infect the frog *Eleutherodactylus coqui* in its native Puerto Rico. *E. coqui* was accidentally introduced into Hawaii in the late 1980s, where there are now hundreds of populations. *B. dendrobatidis* was being considered as a biological control agent for *E. coqui* because there are no native amphibians in Hawaii. Using a DNA-based assay, we tested 382 *E. coqui* from Hawaii for *B. dendrobatidis* and found that 2.4% are already infected. We found infected frogs in four of 10 study sites and on both the islands of Hawaii and Maui. This is the first report of *B. dendrobatidis* in wild populations in Hawaii. As the range of *E. coqui* expands, it may become a vector for the transmittance of *B. dendrobatidis* to geographic areas where *B. dendrobatidis* does not yet exist.

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Keywords: Amphibian declines; Biocontrol; Chytridiomycosis; *Eleutherodactylus coqui*; Hawaii; Emerging diseases; Frog; Invasion; Puerto rico

1. Introduction

Nearly one-third of all amphibians are threatened with extinction (Stuart et al., 2004). Chytridiomycosis, a disease caused by the pathogenic fungus *Batrachochytrium dendrobatidis*, has been identified as a causal agent of amphibian declines in the Americas, Europe, and Australia (e.g., Bell et al., 2004; Berger et al., 1998; Bosch et al., 2001; Lips et al., 2004; Muths et al., 2003), and has been found on every continent with amphibians, except Asia (Weldon et al., 2004). *B. dendrobatidis* is a water-borne pathogen that primarily infects keratinized tissues in the epidermis of amphibians and spreads through colonization by motile, aquatic zoospores (Longcore et al.,

1999). Because *B. dendrobatidis* does not survive desiccation (Johnson and Speare, 2003), amphibians are thought to be the primary means by which the disease is transported to new areas (Daszak et al., 2003; Hanselmann et al., 2004; Weldon et al., 2004).

Some invasive amphibians (e.g., *Rana catesbeiana*) are relatively resistant to chytridiomycosis, yet are efficient carriers of the pathogen (Daszak et al., 2004). The Puerto Rican terrestrial frog, *Eleutherodactylus coqui*, is a notable amphibian invader that has not been tested for *B. dendrobatidis* outside of its native range. *E. coqui* has invaded Florida and several islands in the Caribbean, and was accidentally introduced to Hawaii via nursery plants in the late 1980s (Kraus et al., 1999). Direct development and year-round breeding are thought to contribute to its rapid spread. There are now over 250 known populations on the islands of Hawaii and Maui, located mostly in lowland forests on the windward sides (from 0

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51 to 1100 m altitude), with new populations being reported
52 weekly (Kraus and Campbell, 2002).

53 In Hawaii, *E. coqui* appears to establish populations
54 that have greater densities than those in their native
55 range (20,000 frogs/ha on average in Puerto Rico, Stew-
56 art and Woolbright, 1996; K. Beard, unpublished data).
57 The invasion threatens Hawaii's unique ecological com-
58 munities because *E. coqui* predates upon endemic inver-
59 tebrates, which comprise the large majority of Hawaii's
60 endemic fauna (Beard and Pitt, 2005). The invasion also
61 threatens Hawaii's multi-million dollar floriculture and
62 nursery industries due to quarantine restrictions and
63 frog de-infestation measures (Kraus and Campbell,
64 2002). Likewise, property value and tourism are threat-
65 ened because of its loud (80–90 dBA at 0.5 m) mating
66 calls.

67 Numerous methods for managing *E. coqui* popula-
68 tions have been developed in Hawaii; yet, there has been
69 no report of a successfully eliminated population. Bio-
70 logical control based on amphibian diseases is consid-
71 ered an attractive option because Hawaii has no native
72 amphibians. *B. dendrobatidis* has been found to infect *E.*
73 *coqui* in Puerto Rico dating back to 1978 and is thought
74 to contribute to declines at high elevations (Burrowes
75 et al., 2004). Thus, it has been suggested that *B. dendro-*
76 *batidis* could be used to control *E. coqui* (Hawaii State
77 Department of Agriculture, 2004). Our objective was to
78 determine whether *B. dendrobatidis* is already present in
79 *E. coqui* populations in Hawaii.

80 **2. Materials and methods**

81 *E. coqui* were collected from seven locations on the
82 island of Hawaii and three locations on Maui in May
83 and August 2004, respectively (Table 1). Locations were
84 selected to maximize diversity in forest-type, elevation,
85 and geological history. For one night at each location,
86 subadult [snout-vent length (SVL) < 24 mm (Woolbright,

1985)] and/or adult frogs [SVL ≥ 24 mm] were collected 87
by slowly and systematically walking in a 20 × 20 m plot 88
between 2000 and 2200 h. For each frog, SVL and perch 89
height were recorded. Frogs were collected using stan- 90
dard protocols for testing for *B. dendrobatidis* infection 91
[as outlined in O'Neill et al. (in review)] and were pre- 92
served in 70% ethanol. *E. coqui* demonstrated no overt 93
clinical signs of chytridiomycosis when collected, such as 94
unusual sloughing of the skin or mortality. 95

96 We tested 175 subadults and 207 adults for *B. dendro-*
97 *batidis* using the DNA-based assay described by Annis 97
et al. (2004). This assay uses species-specific primers 98
(*B. dendrobatidis*1a and *B. dendrobatidis*2a) located 99
within ITS1 and ITS2 to amplify the 5.8S region of 100
nuclear rDNA. Tissue samples ranged from a whole foot 101
(subadults) to a half toe (adults). DNA was extracted 102
using the protocol from Schizas et al. (1997) with the fol- 103
lowing modifications: the digestion reaction contained 104
20–30 µl Te (10 mM Tris, 0.1 mM EDTA), and 1.0 µl Pro- 105
teinase K (20 mg/ml). Samples were digested and period- 106
ically vortexed for 3 h at 55 °C. PCR protocols were the 107
same as those described in Annis et al. (2004) including 108
the use of Platinum® *Taq* DNA Polymerase (Invitrogen 109
Corporation, Carlsbad, California, USA). 110

111 Positive controls were both pure *B. dendrobatidis*
112 DNA extracted from culture (Joyce Longcore, unpub-
113 lished data) and DNA extracted from *Rana muscosa* that
114 had previously tested positive for *B. dendrobatidis* (Jes-
115 sica Morgan, unpublished data). Negative controls con-
116 sisted of purified water in the PCR reaction and re-
117 analyses of DNA from animals that previously tested
118 negative for *B. dendrobatidis*. PCR products were visual-
119 ized on a standard 1.4% agarose gel. Samples that con-
120 tained a band at 330 base pairs (BP) in length were
121 presumed to be positive for *B. dendrobatidis* infection
122 (Annis et al., 2004). Three samples resulted in a faint
123 band at 330 BP. In these cases, samples were PCR ampli-
124 fied a second time using the first PCR product as the
125 template. To create comparable negative controls in

Table 1
Number of *Eleutherodactylus coqui* examined and diagnosed with the chytrid fungus *Batrachochytrium dendrobatidis* from 10 locations in Hawaii

Location	Island	Coordinates	Elevation (m)	(No. with fungus/No. examined)	
				Subadults	Adults
Hawaiian Paradise Park	Hawaii	N19°36'W154°59'	50	1/42	–
Humane Society	Hawaii	N19°36'W155°01'	135	0/13	–
Kurtistown	Hawaii	N19°36'W154°05'	310	1/18	–
Lava Tree State Park	Hawaii	N19°29'W154°54'	180	4/26	1/75
Manuka Natural Area Reserve	Hawaii	N19°07'W155°50'	560	0/24	–
Puainako/Safeway	Hawaii	N19°42'W155°04'	45	0/4	–
Waipio Overlook	Hawaii	N20°07'W155°35'	305	0/7	–
Kihei Nursery	Maui	N20°44'W156°27'	400	–	0/51
Maliko Gulch	Maui	N20°52'W156°19'	440	2/41	0/75
Miles Makawao	Maui	N20°53'W156°19'	20	–	0/6
Total				8/175	1/207

Frogs were collected in the summer of 2004.

126 these cases, PCR products scored as negative in the first
 127 round were retested with a second PCR amplification.
 128 Tests were conducted blindly and 8% of each run were
 129 controls. To confirm our results, tissues (not DNA) from
 130 eight specimens that tested positive and 10 specimens
 131 that tested negative in our laboratory were analyzed by
 132 Pisces Molecular LLC (Boulder, Colorado, USA).

133 Statistical analyses were conducted using SAS v.9 for
 134 Windows (SAS Institute, Cary, North Carolina, USA).
 135 To determine if there was a difference in the number of
 136 subadult and adults infected, we compared the number
 137 of infected and uninfected individuals using Pearson's χ^2
 138 exact test. To determine if there was a difference between
 139 adult and subadult perch heights, we conducted a *t*-test.
 140 A folded *F*-test suggested that group variances were
 141 unequal; thus, a Satterthwaite approximation was used.
 142 Significant differences were accepted at $p < 0.05$.

143 **3. Results**

144 Of the 382 individuals tested, nine showed positive
 145 bands for *B. dendrobatidis* (Table 1). No bands of other
 146 sizes were observed in any samples. All positive and neg-
 147 ative controls were correctly scored. Sixteen of the 18
 148 specimens that tested either positive or negative in our
 149 laboratory were confirmed by Pisces Molecular LLC
 150 (Table 2). Two samples (one positive and one negative)
 151 were scored differently by the two laboratories.

152 *B. dendrobatidis* was detected at four of the 10 study
 153 sites (Table 1). We detected *B. dendrobatidis* in popula-
 154 tions on both the islands of Hawaii and Maui. *B. dendro-*
 155 *batidis* was found to infect frogs from locations ranging
 156 from 50 to 440 m in elevation (Table 1).

157 Subadults measured 14.50 ± 0.26 mm (SE) SVL and
 158 adults measured 30.52 ± 0.15 mm SVL. We found a
 159 greater infection rate in subadults than adults (4.6% vs.
 160 0.5% tested positive) ($\chi^2 = 6.51$, $df = 1$, $p = 0.013$). Subad-

ults perched closer to the forest floor than adults 161
 (0.45 ± 0.035 m vs. 0.87 ± 0.027 m) ($df = 414$, $t = 9.16$, 162
 $p < 0.0001$). 163

4. Discussion 164

We found that the chytrid fungus *B. dendrobatidis* is 165
 present in Hawaii and infects *E. coqui*. Like other nota- 166
 ble amphibian invaders (Daszak et al., 2003; Hansel- 167
 mann et al., 2004; Pessier et al., 1999; Weldon et al., 168
 2004), *E. coqui* is now known to be a carrier of *B. dend-* 169
robatidis in locations outside of its native range. Because 170
E. coqui is unlikely to be eradicated from Hawaii (Beard 171
 and Pitt, 2005), these populations may represent a stable 172
 source of *B. dendrobatidis* in the Pacific. *E. coqui*, appar- 173
 ently traveling in nursery plants from Hawaii, have 174
 already reached another Pacific island, Guam (Beard 175
 and Pitt, 2005). The potential for *E. coqui* to transmit 176
B. dendrobatidis with future introductions adds to its 177
 capacity to threaten native communities. 178

E. coqui could have transported *B. dendrobatidis* to 179
 Hawaii; however, because the location of source popula- 180
 tion(s) and number of introductions are not known, it is 181
 not presently possible to consider the status of *B. dend-* 182
robatidis in these populations. Alternatively, *E. coqui* 183
 could have acquired *B. dendrobatidis* from non-native 184
 amphibians already in Hawaii, some were purposely 185
 introduced as biological control agents (i.e., *Bufo mari-* 186
nus and *Dendrobates auratus*), while another was 187
 brought in for culinary purposes (*R. catesbeiana*). At 188
 two of the four locations where *E. coqui* was found to be 189
 infected, we observed large *B. marinus* populations. 190
 Instead, *B. dendrobatidis* could have been transported in 191
 infected water (Johnson and Speare, 2003) or been car- 192
 ried there by some mechanical vector. It is interesting 193
 that *B. dendrobatidis* has now been found on an island 194
 with no native amphibians, suggesting that it can survive 195
 in incipient amphibian populations, or that it can arrive 196
 and survive in these locations without amphibians. 197

We found that infection rates of *B. dendrobatidis* were 198
 greater in subadults than adults. This may have occurred 199
 because of one or more of the following hypotheses: (1) 200
 infected subadults have a lower survival rate than unin- 201
 fected subadults, (2) we sampled a greater proportion of 202
 each subadult than of each adult, or (3) subadults are 203
 more vulnerable to infection than adults (assuming they 204
 recover). As has been found in previous studies (Beard 205
 et al., 2003), we found that subadults perch heights were 206
 closer to the forest floor than that of adults. This prefer- 207
 ence is thought to result in part from the greater mois- 208
 ture requirements of subadults (Pough et al., 1983). 209
 Because *B. dendrobatidis* is an aquatic pathogen (Long- 210
 core et al., 1999), the high moisture environment found 211
 closer to the forest floor could contribute to greater 212
 infection in subadult frogs. 213

Table 2
Eleutherodactylus coqui samples that scored positive and negative for
Batrachochytrium dendrobatidis in our laboratory compared to scor-
 ing from Pisces Molecular LLC (Boulder, Colorado, USA)

Population	Island	(No. positive/No. negative)			
		Subadults		Adults	
		Our laboratory	Pisces	Our laboratory	Pisces
Hawaiian Paradise Park	Hawaii	1/1	1/1	–	–
Kurtistown Lava Tree State Park	Hawaii	1/1	1/1	–	–
Maliko Gulch	Hawaii	4/4	4/4 ^a	0/2	0/2
Maliko Gulch	Maui	2/2	2/2	–	–

^a One of our positives was scored negative by Pisces and one of our negatives was scored positive by Pisces.

214 Overall, we found a low infection rate (2.4%, $n = 382$),
 215 even compared to other studies of the same species (7.1%
 216 infected in Puerto Rico, $n = 28$) (Burrowes et al., 2004).
 217 There are several potential explanations for this low
 218 infection rate. Because an infection of *B. dendrobatidis*
 219 may be localized (Berger and Speare, 1998), and different
 220 tissues (sometimes different feet) from the same speci-
 221 men may be analyzed, we believe that DNA tests for *B.*
 222 *dendrobatidis* can produce false negatives. Additionally,
 223 we believe that low-level infections may lead to inconsis-
 224 tent results between different samples from one speci-
 225 men. This is supported by the fact that the one specimen
 226 that we scored positive and Pisces Molecular scored neg-
 227 ative had a light band that was barely detectable with a
 228 single round of PCR in our laboratory. Because we iso-
 229 lated DNA from one tissue sample per specimen and the
 230 test may fail to detect low-level infections, we believe our
 231 estimate of *B. dendrobatidis* infection of *E. coqui* in
 232 Hawaii is conservative.

233 Some studies suggest that *E. coqui* is not particularly
 234 susceptible to *B. dendrobatidis* infection and chytridi-
 235 omycosis. Because *E. coqui* does not congregate in a
 236 breeding chorus or have an aquatic life stage, it would be
 237 expected to have a low prevalence of infection (Lips
 238 et al., 2003). In addition, in contrast to other species (e.g.,
 239 *Bufo boreas*), laboratory tests using different levels of
 240 exposure to *B. dendrobatidis* have shown that *E. coqui*
 241 have no significant response in mortality (Cynthia
 242 Carey, personal communication.). Alternatively, the low
 243 prevalence might simply reflect a recent invasion of the
 244 fungus into these populations. Further research is
 245 needed to determine the susceptibility of *E. coqui* to
 246 *B. dendrobatidis*. We believe that *B. dendrobatidis* should
 247 not be used as a biological control agent because it is not
 248 a species-specific pathogen, many amphibians are highly
 249 susceptible, alternative hosts have not yet been identi-
 250 fied, and it has been shown to be readily dispersed by
 251 human activities.

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