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ENVIRONMENTAL PREDATION OF *BATRACHOCHYTRIUM DENDROBATIDIS* DETERMINES INFECTION DYNAMICS

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The occurrence pattern of *Batrachochytrium dendrobatidis* (*Bd*) is not homogeneous at local and regional scales. ^{1,2} Sites which seem appropriate and in vicinity to long-term infected sites, are not always colonized by *Bd*. We hypothesized that the presence of resident microorganisms that prey on *Bd* contributes to explain the observed pattern of *Bd* colonization. The aim of this study was to better understand the interactions of *Bd* with resident microbiotic communities, with a focus on the impact of microorganisms on *Bd* viability, infection probability and infection intensity in amphibian hosts.

Sites in the Pyrenean mountain range with known history of absence or presence of *Bd* were sampled. The decrease in *Bd* zoospore viability after exposure to the water samples was assessed using a method combining ethidium monoazide with real-time PCR³. This method was also used to assess the effect of cultures of several freshwater microorganisms on *Bd* viability. Based on the results of these first experiments several microorganisms were selected for *Bd* zoospore ingestion experiments and for a *Bd* infection experiment with *Discoglossus scovazzi* tadpoles in the absence and presence of microorganisms.

We were able to show a significant correlation between the abundance of microorganisms and Bd zoospore persistence in the Pyrenean water samples. In pure culture, some but not all microorganisms were highly efficient in killing Bd. These highly effective species ingested Bd zoospores and prevented Bd colonization of tadpoles. In conclusion, environmental microorganisms dictate chytrid infection dynamics.

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