Copyright © 2007 · Magnolia Press



Molecular systematics of Malagasy poison frogs in the *Mantella betsileo* and *M. laevigata* species groups

FALITIANA C.E. RABEMANANJARA^{1,2}, ANGELICA CROTTINI^{4,5,8}, YLENIA CHIARI³, FRANCO ANDREONE⁵, FRANK GLAW⁶, REMÍ DUGUET⁷, PARFAIT BORA², OLGA RAVOAHANGIMALALA RAMILIJAONA² & MIGUEL VENCES^{8,9}

¹Institute for Biodiversity and Ecosystem Dynamics, Zoological Museum University of Amsterdam, Mauritskade 61, 1092 AD Amsterdam, The Netherlands. E-mail: frabeman@science.uva.nl
²Département de Biologie Animale, Université d'Antananarivo, BP, 906, 101 Antananarivo, Madagascar
³Department of Ecology and Evolutionary Biology, YIBS-Molecular Systematics and Conservation Genetics Lab., Yale University, 21, Sachem Str., New Haven, CT, 06520-8105 USA
E-mail: ylenia.chiari@yale.edu
⁴Universitá degli Studi di Milano, Dipartimento di Biologia, Sezione di Zoologia e Citologia, Via Celoria 26, 20133 Milano, Italy.
⁵Museo Regionale di Scienze Naturali, Sezione di Zoologia, Via G. Giolitti, 36, 10123 Torino, Italy.
⁶Zoologische Staatssammlung München, Münchhausenstr. 21, 81247 München
⁷ Biotope, Agence Océan Indien, 969 ch. cent Gaulettes, 97440 Saint-André, La Réunion

⁸Zoological Institute, Technical University of Braunschweig, Spielmannstr. 8, 38106, Braunschweig, Germany.

E-mail: m.vences@tu-bs.de

⁹Corresponding author

Abstract

Malagasy poison frogs of the genus *Mantella* with its 16 species are currently sub-divided into 5 major groups. Of these, the *Mantella betsileo* group is traditionally understood as containing four species, *Mantella betsileo*, *M. expectata*, *M. viridis* and *M. manery*, while the *M. laevigata* group is considered to be monospecific. A phylogenetic analysis of samples from multiple localities of all species in these two groups, based on sequences of the mitochondrial cytochrome *b* gene, shows the existence of several well-distinct clades in what is currently considered to be *Mantella betsileo*: (1) central-western populations from Kirindy, Isalo, and near Antsirabe close to the Betsileo region, to which the name *M. betsileo* is to be applied, (2) populations of the north-east and north-west, which are closely related to *M. viridis* and to which the name *M. ebenaui* is to be applied, and (3) a clade from southernmost Madagascar and from the Tsingy de Bemaraha, which is sister to *M. expectata* and furthermore includes important intra-clade variation, therefore probably representing one or two undescribed species. Our data also support a large genetic distance of *M. manery* to all other species and its probable sister-group relationship to the sympatric *M. laevigata; M. manery* is consequently transferred from the *M. betsileo* group to the *M. laevigata* group.

Key words: Anura, Mantellidae, Mantella betsileo, Mantella ebenaui, Mantella expectata, Mantella laevigata, Mantella manery, Mantella viridis, Madagascar

Introduction

Vis-à-vis of the world amphibian decline (Kiesecker *et al.* 2001; Pounds *et al.* 2006; Stuart *et al.* 2004), an increased knowledge on genetic data and their interpretations are needed for conservation purposes (e.g. Moritz & Faith 1998; Frankham *et al.* 2002; DeSalle & Amato 2004). Mainly due to fast deforestation, the amphibian biodiversity of the fourth largest island of the world, Madagascar, is heavily threatened (Andreone