

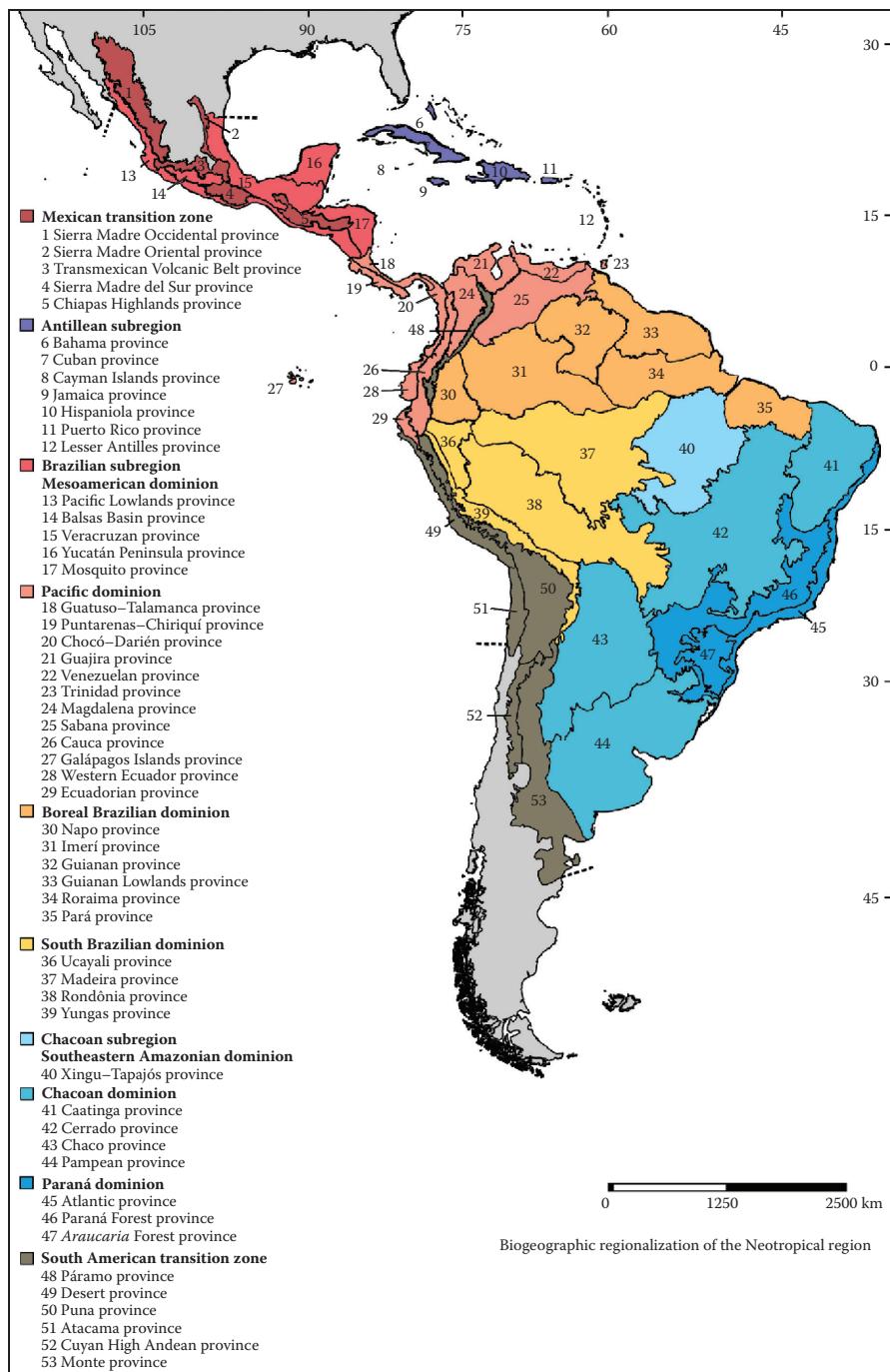
JUAN J. MORRONE

NEOTROPICAL BIOGEOGRAPHY

Regionalization and Evolution



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Regionalization and Evolution

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*I dedicate this book to Chico Mendes (1944–1988),
guardian angel of the Neotropical forests.*



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Preface

The Neotropics comprise the tropical areas of the Americas, from Mexico to Argentina. Hundreds of biogeographic studies of plant and animal taxa from this region have been published for more than 150 years, producing a basic knowledge that is synthesized into regionalization schemes. In this book, I address two central questions of evolutionary biogeography: which areas are recognized within the Neotropical region and how did their biotas evolve? In the last few decades, molecular phylogenetics and parametric model-based biogeography have allowed the postulation of complex biogeographic scenarios for particular taxa, and the search for biotic patterns has been somewhat neglected. However, I feel that biogeographic regionalizations based on the distributional patterns of plant and animal taxa are still relevant in the twenty-first century because they constitute the background knowledge of systematic, ecological, evolutionary, and other kinds of studies.

The biogeographic regionalization of the Neotropical region presented herein consists of three subregions, two transition zones, seven dominions, and 53 provinces. For each unit, I provide the valid name according to the International Code of Area Nomenclature (ICAN) followed by a list of citations and synonyms, a brief characterization, and some endemic and characteristic taxa. In order to deal with biotic evolution, I refer to the identification of biotas through areas of endemism and generalized tracks, their relationships based on track and cladistic biogeographic analyses, and, when possible, the cenocrons or biotic subsets that have been identified within them. This attempt of synthesis is based on a vast bibliography that I have compiled for more than two decades. I feel grateful to many authors who have provided insights on the regionalization and evolution of the Neotropical region. Particularly inspirational for my work were Jorge Artigas, Ángel L. Cabrera, Joel Cracraft, Léon Croizat, Philip Darlington, Gonzalo Halffter, Michael Heads, René Jeannel, Guillermo Kuschel, Emilio Maury, Ernst Mayr, Paul Müller, Eduardo Rapoport, Osvaldo Reig, Raúl Ringuelet, Donn E. Rosen, Jerzy Rzedowski, Jay Savage, George G. Simpson, Arne Takhtajan, Alfred R. Wallace, and Abraham Willink.

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CHAPTER 1

Theoretical Background

Evolutionary biogeography integrates distributional, phylogenetic, molecular, and paleontological data in order to discover biogeographic patterns exhibited by plant and animal taxa, and assess the historical changes that have shaped biotic assembly (Morrone, 2009). Biogeographic regionalizations are hierarchical classifications categorizing geographic areas in terms of their endemic taxa and their relationships. They represent the syntheses of different evolutionary biogeographic analyses and, at the same time, constitute the background knowledge of other studies (e.g., systematic, ecological, or evolutionary).

EVOLUTIONARY BIOGEOGRAPHY

Evolutionary biogeography is the integrative study of distributional, phylogenetic, molecular, and paleontological data, aimed to discover biogeographic patterns and assess the historical changes that have shaped them (Morrone, 2009). It follows a stepwise approach. First, areas of endemism or generalized tracks are identified and considered as hypotheses about biotic identity based on the distributional congruence exhibited by different plant and animal taxa. Second, cladistic biogeographic analyses test these hypotheses, based on the available phylogenetic evidence on the taxa analyzed. Third, biogeographic regionalization is achieved based on cladistic biogeographic hypotheses. Fourth, the molecular dating of divergences between lineages and fossil data allows the identification of cenocrons, which represent subsets of taxa within a biota, identified by their common origin and evolutionary history. Cenocrons incorporate a temporal dimension that implies a time frame of the dispersal of sets of taxa into the biota. Finally, after the biotas and cenocrons have been identified, one may construct a geobiotic scenario by accounting biological and non-biological data to explain the episodes of vicariance/biotic divergence and dispersal/biotic convergence that have shaped the evolution of the biotas analyzed.

The dispersal–vicariance model followed herein assumes that the relationship between earth history and life is more complex than what is assumed in simpler models, because biotic history is reticulate. It treats vicariance as the default explanation for general biogeographic patterns and dispersal as the process that shapes the