Open Access

Multilocus analyses indicate a mosaic distribution of hybrid populations in ground squirrels (genus Ictidomys)

Cody W. Thompson^{1,2}, Faisal Ali Anwarali Khan^{1,2,3}, Frederick B. Stangl Jr⁴, Robert J. Baker^{1,2} & Robert D. Bradlev^{1,2}

¹Department of Biological Sciences, Texas Tech University, Lubbock, Texas 79409-3131

²Natural Science Research Laboratory, Museum of Texas Tech University, Lubbock, Texas 79409-3191

³Department of Zoology, Faculty of Resource Science and Technology, Universiti Malaysia Sarawak, Kota Samarahan, Sarawak 94300, Malaysia ⁴Biology Department, Midwestern State University, Wichita Falls, Texas 76308

Keywords

Amplified fragment length polymorphism, cytochrome-b, Ictidomys, mosaic hybrid zone, secondary contact, Y-linked structural maintenance of chromosomes.

Correspondence

Cody W. Thompson, Museum of Zoology, University of Michigan, 1109 Geddes Avenue, Ann Arbor, MI 48109. Tel: (734) 615-2810; Fax: (734) 763-4080; E-mail: cwthomp@umich.edu

Funding Information

Portions of this research were supported by grants awarded to C. W. Thompson from the American Society of Mammalogists, Association of Biologists at Texas Tech University, Southwestern Association of Naturalists, Texas Academy of Science, and the Michelle C. Knapp Graduate Research Award through Texas Tech University. Additional funds came from the Texas Tech University Biological Database Program. F. A. A. Khan was supported by funds from the Universiti Malaysia Sarawak, Higher Education Ministry of Malaysia, and the Department of Biological Sciences, Texas Tech University.

Received: 29 April 2013; Revised: 26 July 2013; Accepted: 12 August 2013

Ecology and Evolution 2013; 3(13): 4448-4460

doi: 10 1002/ece3 755

Introduction

Hybrid zones are the result of the interaction of genetically distinct groups that produce offspring of mixed ancestry (Barton and Hewitt 1985; Arntzen 1996; Abbott et al. 2013). Hybrid zones often provide a natural laboratory for

addressing major evolutionary concepts (Hewitt 1988; Baker et al. 1989; Harrison 1993). For example, studies of hybrid zones offer the opportunity to examine genetic control of speciation, mechanistic models of maintenance, premating and postmating isolation, direction of introgression, and other processes and patterns of hybridization

Abstract

DNA sequence data from mitochondrial cytochrome-b (Cytb) and Y-linked structural maintenance of chromosomes (SmcY) genes were combined with 478 nuclear loci obtained from amplified fragment length polymorphisms (AFLP) to assess the extent of hybridization and genetic spatial structure of populations in two hybridizing species of ground squirrel (Ictidomys parvidens and Ictidomys tridecemlineatus). Based on AFLP analyses of 134 individuals from 28 populations, 10 populations were identified that possessed hybrid individuals. Overall estimates of F_{ST} values revealed strong support for population structure in the Cytb data set; however, analyses of the SmcY gene and the AFLP data indicated ongoing gene flow between species. Pairwise F_{ST} comparisons of populations were not significant for the SmcY gene; although they were significant for the Cytb gene, indicating that these populations were structured and that gene flow was minimal. Therefore, gene flow between I. parvidens and I. tridecemlineatus appeared to be restricted to populations that exhibited hybridization. In addition, the fragmented nature of the geographic landscape suggested limited gene flow between populations. As a result, the distributional pattern of interspersed parental and hybrid populations were compatible with a mosaic hybrid zone model. Because ground squirrels display female philopatry and male-biased dispersal, the ecology of these species is compatible with this hypothesis.

© 2013 The Authors. Ecology and Evolution published by John Wiley & Sons Ltd. This is an open access article under the terms of the Creative Commons Attribution License, which permits use,

distribution and reproduction in any medium, provided the original work is properly cited.