

Effects of habitat fragmentation and human disturbance on the population dynamics of the Yunnan snub-nosed monkey from 1994 to 2016

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In this study, we integrate data from field investigations, spatial analysis, genetic analysis, and GLMs to evaluate the effects of habitat fragmentation on the population dynamics, genetic diversity, and range shifts in the endangered Yunnan snub-nosed monkey (*Rhinopithecus bieti*). The results indicate that from 1994 to 2016, *R. bieti* population size increased from less than 2,000 to approximately 3,000 individuals. A primary factor promoting population recovery was the establishment of protected nature reserves. We also found that subpopulation growth rates were uneven, with the groups in some areas, and the formation of new groups. Both the fragmentation index, defined as the ratio of the number of forest patches to the total area of forest patches (e.g. increased fragmentation), and increasing human population size had a negative effect on population growth in *R. bieti*. We recommend that government conservation plans prioritize the protection of particular *R. bieti* populations, such as the Baimei and Jisichang populations, which have uncommon haplotypes. In addition, effective conservation strategies need to include an expansion of migration corridors to enable individuals from larger populations such as Guyoulong (Guilong) to serve as a source population to increase the genetic diversity of smaller *R. bieti* subpopulations. We argue that policies designed to protect endangered primates should not focus solely on total population size but also need to determine the amount of genetic diversity present across different subpopulations and use this information as a measure of the effectiveness of current conservation policies and the basis for new conservation policies.

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Short Title: Population dynamics of *R. bieti*

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ABSTRACT

In this study, we integrate data from field investigations, spatial analysis, genetic analysis, and GLMs to evaluate the effects of habitat fragmentation on the population dynamics, genetic diversity, and range shifts in the endangered Yunnan snub-nosed monkey (*Rhinopithecus bieti*). The results indicate that from 1994 to 2016, *R. bieti* population size increased from less than 2,000 to approximately 3,000 individuals. A primary factor promoting population recovery was the establishment of protected nature reserves. We also found that subpopulation growth rates were uneven, with the groups in some areas, and the formation of new groups. Both the fragmentation index, defined as the ratio of the number of forest patches to the total area of forest patches (e.g. increased fragmentation), and increasing human population size had a negative effect on population growth in *R. bieti*. We recommend that government conservation plans prioritize the protection of particular *R. bieti* populations, such as the Baimei and Jisichang populations, which have uncommon haplotypes. In addition, effective conservation strategies need to include an expansion of migration corridors to enable individuals from larger populations such as Guyoulong (Guilong) to serve as a source population to increase the genetic diversity of smaller *R. bieti* subpopulations. We argue that policies designed to protect endangered primates should not focus solely on total population size but also need to determine the amount of genetic diversity present across different subpopulations and use this information as a measure of the effectiveness of current conservation policies and the basis for new conservation policies.

Key words: habitat fragmentation; human disturbance; *Rhinopithecus bieti*; population size; spatial analyst; range shift; human population density; conservation policies; genetic diversity; land use

INTRODUCTION

Globally, anthropogenic activities resulting in deforestation, habitat loss, the extraction of natural resources, the introduction of invasive species, agricultural expansion, and climate change represent a major driver of the unprecedented decline in biodiversity (*Millennium Ecosystem Assessment, 2006; Magurran & Dornelas, 2010; Newbold et al., 2015*). Quantifying the effects of individual factors and developing solutions to protect environments and endangered is a key scientific challenge (*Mace et al., 2010; Magurran & Dornelas, 2010*). Anthropogenic activities have altered as much as 50% of terrestrial land cover, and land-use patterns, such as the expansion of croplands for industrial agriculture and pastures for grazing cattle and other domesticated animals, has led to a global reduction in the number of species (*Millennium Ecosystem Assessment 2006; McGill 2015*). This has led scientists to refer to the current period as the Anthropocene mass extinction event (*Ceballos, Ehrlich & Dirzo, 2017*), with species extinction rates estimated to be 100–1,000 times greater than during past evolutionary periods (*Raup, 1995; Pimm et al., 1995*). Protecting animal and plant survivorship and biodiversity requires monitoring and managing changes in species population size and genetic diversity (*Dornelas et al., 2014*).

An important form of habitat degradation is fragmentation, which transforms large tracts of continuous habitat into smaller and spatially distinct patches immersed within a dissimilar matrix (*Didham, Kapos & Ewers, 2012; Wilson et al., 2016*). Fragmentation can create detrimental edge effects along the boundaries of habitat patches, alter ecological conditions resulting in the decline of native species, restrict animal movement and gene flow, and sever landscape connectivity (*Crooks & Sanjayan, 2006*) leading to local population extinction in response to insufficient viable “core” habitat (*Ewers & Didham, 2007*). Habitat fragmentation has continued at an alarming rate, and has impaired key ecosystem functions by decreasing biomass, limiting seed dispersal, shifting predator-prey relationships, and altering nutrient cycles (*Haddad et al., 2015*).

Several species of nonhuman primates are particularly affected by habitat fragmentation due to their dependence on intact and biodiverse forested landscapes to obtain a nutritionally balanced diet (*Arroyo-Rodriguez & Fahrig, 2014*). Currently, anthropogenic habitat modifications have resulted in more than half of

the world's primate species listed as Vulnerable, Endangered, or Critically Endangered (*Estrada et al., 2017; Estrada et al., 2018*). Primate populations inhabiting small and isolated forest fragments are especially vulnerable to extinction (*Benchimol & Peres, 2013*). However, given evidence of behavioral plasticity, the benefits of social learning, and the ability to exploit a broad range of different food types, many species of primates are reported to survive, at least over periods of several years, in impacted landscapes by modifying aspects of their activity budget, group size and composition, ranging patterns, and diet (*Onderdonk & Chapman, 2000; Wong & Sicotte, 2007; Boyle & Smith, 2010*).

The endangered Yunnan snub-nosed monkey (*Rhinopithecus bieti*), a large-bodied species of nonhuman primate, that lives in social groups of several hundred individuals (*Kirkpatrick et al., 1998*). A survey in the early 1990's indicated that this species was confined to a narrow region between the Yangtze and Mekong rivers (98°37' to 99°41'E, 26°14' to 29°20'N). Their population size was estimated to be less than 2,000 individuals, distributed across 19 distinct social groups (*Long et al., 1994*). A study by Zhao et al. (2018) integrating data on evolutionary genetics and biogeographical information concluded that both the historical distribution (past 2000 years) and current population structure of the *R. bieti* appears to have been directly impacted by human activities, principally agricultural expansion resulting in severe habitat fragmentation of the Tibetan Plateau and by hunting (*Liu et al., 2009*).

Recent studies modeling habitat change and agricultural expansion across *R. bieti's* range, predict an increase in forest fragmentation and a decrease in habitat quality resulting in range contraction over the next 25-75 years (*Xiao et al., 2003; Wong et al., 2013; Li et al., 2018*). However, some of these studies were based on data collected in the early 1990's and therefore may not accurately represent the current demographic and ecological challenges and conservation pressures faced by *R. bieti*. The main purpose of our study is to: (1) investigate the current population size and distribution of the *R. bieti*, (2) conduct a landscape and spatial analysis of their distribution area, and (3) examine how present day habitat fragmentation influences *R. bieti* population size, genetic diversity, and changes in geographical distribution.

METHODS

Study area and data collection

This field study was approved by State Forestry Administration of China. The study area is a narrow region of 17,000 km² in the northwest of Yunnan Province and the Tibet Autonomous Region. It is bounded by the Mekong River to the west and the Yangtze River to the east (Long *et al.*, 1994). The elevation of the study area varies from 1,300 m to 5,400 m. Our study was conducted across all current distributional areas of *R. bieti* in Yunnan Province and Tibet. Each of the remaining wild groups was surveyed from January to November 2013 and April to September 2016. In order to obtain the most accurate information for this species, we conducted detailed surveys within its main distribution in three national nature reserves (Bamaxueshan, Honglaxueshan and Tianchi), one provincial nature reserve (Yunling), and two sites that are outside of protected reserves (Jinsichang and Bamei) (Fig. 1).

Based on information of group location, a survey was conducted and the number of monkeys per group was censused. Specific locations of *R. bieti* groups were obtained from local forest rangers and officials who regularly patrol the reserves. *R. bieti* were counted directly using auxiliary telescopes to assist in observations (Wu *et al.*, 1988). Each monkey group was observed for 14–21 days, counting all individuals in the group (Wu *et al.*, 1988). The counting of monkeys in each group was based on the following method. Travel in *R. bieti* occurs both on the ground and in trees. However, when crossing open areas, the monkeys often walk slowly in a single or nearly single. A total of 17 groups were censused. *R. bieti* live in groups called multilevel societies (MLS), which are composed of several harem social and reproductive units (Qi *et al.*, 2014). Each harem contains a single adult male, several adult females, and offspring. These harems travel, feed, and range together throughout the day and form a large band. Bands are followed by one or more all male units (AMU) that contain juvenile, sub-adult, and adult bachelor males. Because each band or MLS has been tracked by patrol officers for several years, we were able to obtain an accurate count of its size. This was accomplished by first counting the number of harems in each band and then estimating the number of monkeys per harem and per AMU to obtain a final total. In order to increase census accuracy, at least 3 researchers participated in each

survey and counted/estimated the number of harems and harem sizes.

To monitor changes over time in *R. bieti* demography and distribution, we obtained information on *R. bieti* population size and location in 1990 from Long *et al.* (1994) and compared it with our current data. To assess relationships among suitable habitat, population size, and gene flow, we conducted an analysis of genetic diversity among groups. The genetic diversity for each group was calculated using FSTAT 3.2 (Goudet, 1995) based on microsatellite data from the analyses of 157 individuals' samples (blood, muscle and faecal) in 2007 (Liu *et al.* 2007, 2009). Genetic information was obtained from 11 of the 17 known groups of *R. bieti*.

Landscape data and spatial analysis

We used a land cover map (30"×30" resolution) derived from an assemblage of 13 SPOT5 images across the known distribution of *R. bieti* for 1990 and 2016. This covered an area of 17, 000 km² including the entire *R. bieti* range. Vegetation types suitable for *R. bieti* were defined as dark-coniferous forest dominated by species such as *Abies georgei* and mixed coniferous and broadleaf forest (Kirkpatrick *et al.*, 1998; Xiao *et al.*, 2003; Li *et al.*, 2015). We then verified our habitat classification accuracy using data from the Conservation Information Centre of The Nature Conservancy of China. Fragmentation metrics for *R. bieti* were calculated using FRAGSTATS 4 (McGarigal & Marks, 1995). Patch density (PD, the number of patches divided by total landscape area) was used as indices of fragmentation, with low PD indicating a more connected landscape and high PD indicating a more fragmented landscape (Olsoy *et al.*, 2016). A fragment was considered vegetatively suitable to sustain a population of *R. bieti* if it included dark-coniferous forest and mixed coniferous and broadleaf forest. Given the potential effect that habitat fragmentation can have on whether an area is suitable for *R. bieti*. Supplementary information of forested land, cropland, pasture land, and land reclamation (defined as primary forest converted into a secondary forest, pasture and cropland), as well as information on human population density, was extracted from the History Database of the Global Environment (the pixel size of these data were 0.25 x 0.25 resolution ; HYDE3.1: Goldewijk *et al.*, 2011; He, Li & Zhang, 2015; Li, He & Zhang, 2016). We extracted values of human population density, cropland, and pasture (HYDE3.2; Goldewijk, 2016), all indicators of habitat fragmentation, between the years 1990 and 2016 with the function "extract" in R.

We used Spearman correlations to check for simple linear relationships between the growth rate of the *R. bieti* population and all environmental and landscape variables (fragmentation index, population density, area of cropland, area of pasture, density of households, and density of roads). Then, we applied multiple linear regression analyses to determine the partial and interactive effects of factors affecting *R. bieti* population size. Finally, we applied GLMs to determine the factors affecting *R. bieti* population size.

RESULTS

Based on a census of 17 *R. bieti* subpopulation size isolated from each other and troop size ranged from 40-450 individuals. The mean number of harems per band was 12 ± 3.5 . Across all subpopulations, the total number of *R. bieti* in 2016 was estimated at approximately 3000 individuals. We found that population growth had varied significantly by region. Current population estimates in the central and southwestern zones of the species' range (Fig.1) (Baimaxueshan Nature Reserve and Yunling Nature Reserve) were 147% higher compared to values reported in 1994 (including the formation of three new groups: Gehuaqing, Baijixun(Yongan), Shikuadi). In contrast, size estimates for the southeastern and northern (Honglaxueshan Nature Reserve) areas increased by only 24% and 21% respectively, from earlier estimates. This increase in population size was the result of an increase in the size of bands rather than the number of *R. bieti* bands. We found that 6 groups with populations sizes of less than 100 individuals in 1994 disappeared by 2016. These included those in Bajia, Adong, Houziqin, Heishan, Dapingzi, and Moziping. The population sizes of three groups (Cikatong, Milaka, and Akou (Anyi)) were found to decrease by 20-50% between 1994 and 2016 (Table 1). Four new groups were identified in our census, Zhina, Gehuaqing, Baijixun (Yongan), and Shikuadi. These bands inhabit the northern and southern regions of the species distribution and ranged in size of from 40 to 450 individuals (Table 1).

Our analysis of habitat fragmentation indicates that the southern populations of *R. bieti* (Tianchi nature reserve, Yunling nature reserve and Jinsichang, fragmentation index: 0.99), live in more fragmented habitats and are characterized by smaller band size compared to the Central (N=9) and northern (N=3) populations (Baimaxueshan nature reserve and Honglaxueshan nature reserve, fragmentation index: 0.59). In general, the GLMs indicated that the fragmentation index, human population size, and the area within a group's range devoted to pasture was negatively correlated with *R. bieti* population size. In this case, after model selection based on AIC, only a few variables remained (population size=0.85-3.09 fragmentation index -2.39 population

171 density=0.49 pasture, $R^2=0.52$, $P<0.05$).

172 DISCUSSION

173 The results of our census indicate that the construction of protected nature reserves (87% of the *R. bieti* live in
174 protected areas) has effectively limited human disturbance and reduced hunting, leading to a population
175 increase from less than 2000 to almost 3000 individuals over the past 25 years. During the 1980's, populations
176 of *R. bieti* faced a set of severe anthropogenic challenges. For example, approximately 200,000 m³ of
177 commercial logs were removed annually from their range, an entire band was killed by poachers (*Long et al.*,
178 1994; *Ding, Yang & Liu, 2003*), and the area of suitable habitat decreased by 31% from 1958 to 1997 as large
179 tracts of forests were converted into pastures for cattle grazing (*Xiao et al., 2003*). In addition, in Deqin
180 country alone, approximately 430 *R. bieti* were killed for meat, fur, and medicine from 1970 to 1979, and 68
181 pieces of *R. bieti* fur or bone were sold in local stores (*Bai, 1987*). With the establishment of the Baimaxueshan
182 Nature Reserve, the Honglashan Nature Reserve, the Yunling Nature Reserve and the Tianchi Nature Reserve
183 between 1983 and 2003, *R. bieti* is now protected and this has effectively prevented poaching, the collection of
184 forest products, and livestock grazing within their range. Simultaneously the Chinese government has banned
185 all guns beginning in 1990s, and this has dramatically reduced poaching. Each of these measures has served to
186 protect *R. bieti* and to promote their population recovery over the past 25 years.

187 Despite population increases, the conservation status of *R. bieti* continues to be negatively affected by forest
188 fragmentation and the presence of human settlements in and around their range. Our results indicate that *R.*
189 *bieti* population growth rates varied across its distribution with both increased habitat fragmentation and
190 increased human population density negatively affecting *R. bieti* population growth rates. This was most
191 evident in the southern region (Yunling and Tianchi Nature Reserves), where suitable habitats are highly
192 fragmented and *R. bieti* population growth rates are low. The situation faced by *R. bieti* of Fragmentation-
193 induced reduction in habitat quality and resource availability also has adversely affected the long-term viability
194 of other primate species (*Wahungu et al., 2005; Arroyo-Rodríguez & Mandujano, 2014*) and this has
195 contributed to a primate extinction crisis worldwide (*Estrada et al., 2017, Estrada et al., 2018; Li et al., 2018*). .

196

197 In the case of *R. bieti*, forest fragmentation spatially isolates subpopulations and impeding gene flow.
 198 Moreover, the conversion of natural habitat to agricultural fields and pasture land leads to the local extinction
 199 of small populations, as in the case of the Adong, Houziqin, Heishan, Dapingzi, and Moziping *R. bieti* bands.
 200 Fragmentation also leads to reduction in the availability of large food patches, changes in tree species
 201 composition and diversity, and introduces edge effects leading to changes in soil moisture and local climate,
 202 resulting in a reduction in overall habitat quality (Laurance *et al.*, 2000). Lichen and fruit are important
 203 components of the diet of *R. bieti*, accounting for 50% of total feeding time (Grueter *et al.*, 2009). Across most
 204 forested environments, ripe fruit is characterized by a patchy distribution in space and time. Spatial and
 205 temporal fruit patchiness is exacerbated in fragmented landscapes, limiting the ability of species (Estrada &
 206 Coates-Estrada, 1996). For example, mantled howler monkeys (*Alouatta palliata*) living in more fragmented
 207 landscapes spent more time traveling and searching for food than did groups living in less fragmented
 208 landscapes (Estrada & Coates-Estrada, 1996). We assume that given the large size of *R. bieti* bands,
 209 increased fragmentation results in increased challenges in locating sufficient food. Similarly, lichen is known
 210 to be negatively impacted by human-induced environmental change such as air pollution, and a decline in the
 211 production of lichen is likely to have a severely negative effect on *R. bieti* nutrient intake, reproduction,
 212 population growth rates, and survivorship (Grueter *et al.*, 2008).

213 Our results also indicated that the population growth rate of *R. bieti* was negativity correlated with human
 214 population density and human accessibility to wildlife habitats. Based on our field observations, we noted that
 215 during the *R. bieti* breeding season (month 2 to month 6), local people go into the forest to pick cordyceps,
 216 which frightened and affects *R. bieti* breeding behavior by making *R. bieti* spending more time traveling. In
 217 addition, human accessibility to wildlife habitats has the potential to become a vector for the transmission of
 218 zoonotic diseases between humans, domesticated animals, and wildlife, and rapidly extirpate entire primate
 219 subpopulations from local areas (Patz *et al.*, 2008; Lambin *et al.*, 2010; Murray & Daszak, 2013; Gottdenker *et*
 220 *al.*, 2014; Estrada *et al.*, 2018). We found that in areas with human population densities of greater than 46

human/ km², *R. bieti* growth rates either increased very slowly (rate of 11.4%) or some small bands exhibited negative growth and disappeared from areas such as Bajia, Adong, Houziqin, Heishan, Dapingzi, and Moziping. This reduction has been offset, in part, by the addition of four newly formed bands (Zhina, Gehuaqing, Baijixun (Yongan), Shikuadi) and other bands increasing in size. Moreover, pastures serve as a barrier to migration for *R. bieti* (Kirkpatrick *et al.*, 1998; Grueter *et al.*, 2010) and therefore severely limit the movement of individuals across these highly transformed landscapes and the opportunity to expand into unoccupied habitats and form of new bands.

Our results revealed that *R. bieti* populations in the Yunling Nature Reserve and in Jinshichang are the most isolated, and characterized by certain unique haplotypes (haplotype M1-M4, M6-M11 and M29-30) (Liu *et al.*, 2007). Moreover, despite the fact that total population size has increased dramatically over the past 25 years, the population size of three bands (Milaka, Cikatong, and Akou (Anyi)) has markedly decreased, exacerbating the loss of genetic diversity (Table.1). We suggest that management decisions for endangered species should not focus solely on population size but also consider subpopulation genetic diversity. We recommend that effective conservation policies for this species should prioritize protecting certain targeted populations such as Milaka, Cikatong, and Akou (Anyi) that have high intraspecific genetic diversity but low growth rates (Table 1). In addition, government policies that promote the alleviation of human poverty, especially in rural communities, represent an important conservation tool that indirectly protects *R. bieti*. In this regard, the Chinese government has encouraged local farmers to return land to natural forest, prohibited logging, and relocate from Nature Reserves. This serves to reduce human interference and protect habitats that are critical to the survival of *R. bieti* as well as others threatened taxa.

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Figure legends

375 **Figure 1.** The study area and locations of the bands of *R. bieti* from 1994 to 2016.

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Figure 1

The study area and locations of the bands of *R. bieti* from 1994 to 2016

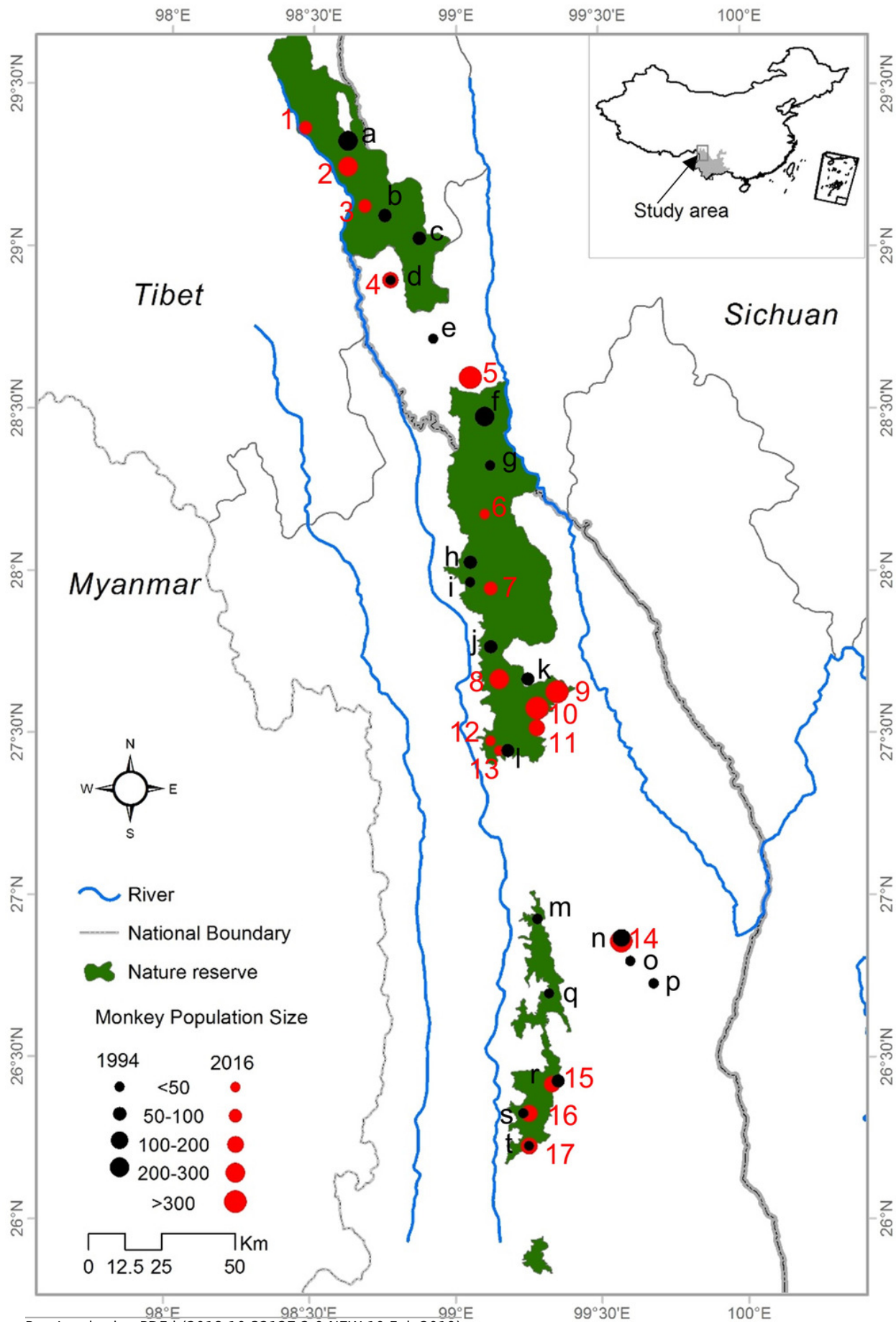


Table 1(on next page)

Changes in the location of natural bands of *R. bieti* between 1994 to 2016

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Table.1 Changes in the location of natural bands of *R. bieti* between 1994 to 2016

Site	Latitude	Longitude	1994	2016	Change group	Intraspecific genetic diversity	
						Unbiased Hz	Obs Hz
Zhina	29.37	98.47	-	80	new	0.54	0.50
Xiaochangdu	29.33	98.62	> 200	280	unchanged	-	-
Milaka	29.13	98.75	<100	60	unchanged	0.58	0.64
Bajia	29.03	98.87	<100	-	disappeared	-	-
Bamei	28.9	98.77	<50	100	unchanged	0.59	0.54
Adong	28.72	98.92	<50	-	disappeared	-	-
Wuyapuya	28.48	99.1	> 200	400	unchanged	0.63	0.64
Cikatong	28.03	99.05	50-100	50	unchanged	0.63	0.65
Guyoulong(guilong)	27.97	99.05	<50	80	unchanged	0.70	0.69
Shiba	27.77	99.12	50-100	200	Unchanged	-	-
Guomorong(Xiangguqibg)	27.67	99.25	50-100	480	unchanged	0.65	0.63
Akou(Anyi)	27.45	99.18	50-100	40	unchanged	0.66	0.59
Jinsichang	26.87	99.57	100-150	310	unchanged	0.57	0.54
Neidaqin(Fuhe)	26.43	99.35	50-100	120	unchanged	0.66	0.71
Lashashan	26.33	99.23	<50	130	unchanged	-	-
Houziqin	26.93	99.28	<50	-	disappeared	-	-
Longma	26.23	99.25	<50	140	unchanged	0.64	0.72
Heishan	26.7	99.32	<50	-	disappeared	-	-
Dapingzi	26.73	99.68	<50	-	disappeared	-	-
Moziping	26.8	99.6	<50	-	disappeared	-	-
Gehuaqing	27.58	99.28	-	450	new	-	-
Shikuadi	27.52	99.28	-	120	new	-	-
Pantiange	27.47	99.15	-	-	unchanged	-	-

Baijixun(Yongan)	27.48	99.12	-	40	new	-	-
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