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## Deforestation and apparent extinctions of endemic forest beetles in Madagascar

Ilkka Hanski, Helena Koivulehto, Alison Cameron and Pierre Rahagalala

*Biol. Lett.* 2007 **3**, 344-347  
doi: 10.1098/rsbl.2007.0043

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### References

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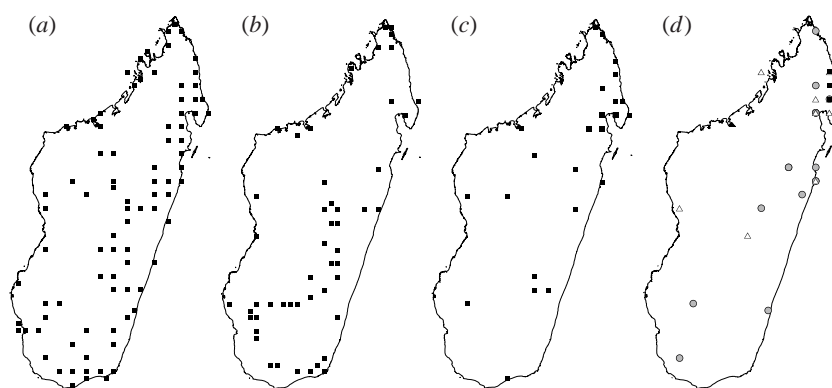


Figure 1. Maps showing (a) the historical sampling localities during 1875–1990, (b) our sampling localities during 2002–2006, (c) the sampling localities of 21 apparently extinct species apart from *H. undatus* and (d) the localities for *H. undatus* prior to 1900 (grey circles), during 1901–1950 (open triangles) and during 1951–1973 (black squares). Localities are mostly shown with resolution of  $0.5^\circ$ .

the range of species  $x$  at time  $t$  (1953 or 2000) as

$$F_{x,t} = (1/n) \sum \sum e^{-\alpha d_{ij}} A_j, \quad (2.1)$$

where  $A_j$  is the percentage of forest cover in cell  $j$ ,  $d_{ij}$  is the distance between the sampling locality  $i$  and cell  $j$  in degrees, and the second summation is over the  $n$  sampling localities for species  $x$ . Thus,  $F_{x,t}$  measures the average amount of forest in the surroundings of the historical sampling localities for species  $x$ , giving decreasing weight to cells with increasing distance from the sampling localities. We assumed  $\alpha=10$ , which gives substantial (greater than 0.05) weight to distances up to 33.6 km. The absolute and relative forest losses were calculated as  $F_{x,1953} - F_{x,2000}$  and  $F_{x,2000}/F_{x,1953}$ , respectively.

As our sampling localities do not evenly cover all of Madagascar, it is possible that we have failed to sample a species because only a few or even none of our sampling localities were within its range. To account for this, we calculated the average distance of the  $n$  historical sampling localities for species  $x$  to our sampling localities as

$$D_x = (1/n) \sum \sum e^{-\alpha d_{ij}}, \quad (2.2)$$

where  $d_{ij}$  is the distance (in degrees) from the  $i$ th historical sampling locality to the  $j$ th locality in our sampling. Since we used the value  $\alpha=1$  in this calculation,  $1^\circ$  (112 km) distance has the weight 0.37.

Other explanatory variables include body size, the last year when the species was sampled prior to our sampling, the (log) number of historical sampling localities, the (log) number of individuals in the Paris collections and the range of the species, defined as the distance between the two most distant historical sampling localities. As the latter three variables are all strongly correlated, we calculated the first principal component as a general measure of past commonness (PC1). PC1 accounted for 90% of variation in the three original variables.

### 3. RESULTS

Out of the 51 species sampled prior to our work and for which locality information is available, we have sampled 29 species but failed to collect 22 other species. We ran stepwise logistic models with the explanatory variables described in §2 to explain apparent extinctions. Relative forest loss ( $F_{x,2000}/F_{x,1953}$ ) entered the model first (table 1). The other two variables that were selected were distance to our sampling localities ( $D_x$ ) and PC1. An equally good model was obtained if PC1 was dropped from the candidate variables, in which case the last year when the species had been sampled was selected at 5% level.

Since the uncollected species cannot be included in our molecular phylogeny (Koivulehto *et al.* in preparation), we cannot critically assess possible phylogenetic bias in apparent extinctions. However, we may use the eight taxonomic species groups of Lebis (1960) as a proxy, as these groups match the clades in the

Table 1. Stepwise logistic regression model explaining whether we have sampled a species of *Helictopleurini* ( $n=29$ ) or not ( $n=22$ ). ( $p$ -value for the full model = 0.38; d.f. = 47.)

explanatory variable	deviance	difference	$p$
constant	69.74		
relative forest loss	61.52	8.21	0.004
distance to our traps	53.24	8.28	0.004
past commonness	49.24	4.00	0.046

molecular phylogeny reasonably well. There is no difference in the fraction of apparently extinct species among the morphological groups ( $p=0.46$ ).

Most of the 22 species that we have not collected have been previously collected from only one ( $n=9$ ) or two ( $n=6$ ) localities, widely scattered across Madagascar (figure 1c). The most striking exception is *Helictopleurus undatus*, which has been collected from 27 localities across much of Madagascar. A closer examination shows that since 1950 this species has been restricted to a small region in the northeast (figure 1d), giving the impression that it gradually disappeared from its former range during the twentieth century.

### 4. DISCUSSION

Helictopleurini are mostly relatively large, many species have colour-patterned elytra and are diurnal and easy to sample with dung and carrion-baited traps. For these reasons, Helictopleurini have been relatively well collected in the past. In our specialized sampling, only four new species were discovered, all of which appear to be very localized and rare. Four species compose only 7% of the described species, which is a small percentage for tropical insects in as large and diverse an area as Madagascar. We conclude that Helictopleurini have been sufficiently well collected in the past to warrant this analysis.

The best predictor of whether a species was collected by us or not was relative forest loss within its past range. The estimated remaining forest cover from 1953 to 2000 ranges from 10 to 60% for different species (figure 2). Species for which less than one-third of the 1953 forest cover remains tended to be apparently extinct. The other factors that had a significant effect on species' occurrence

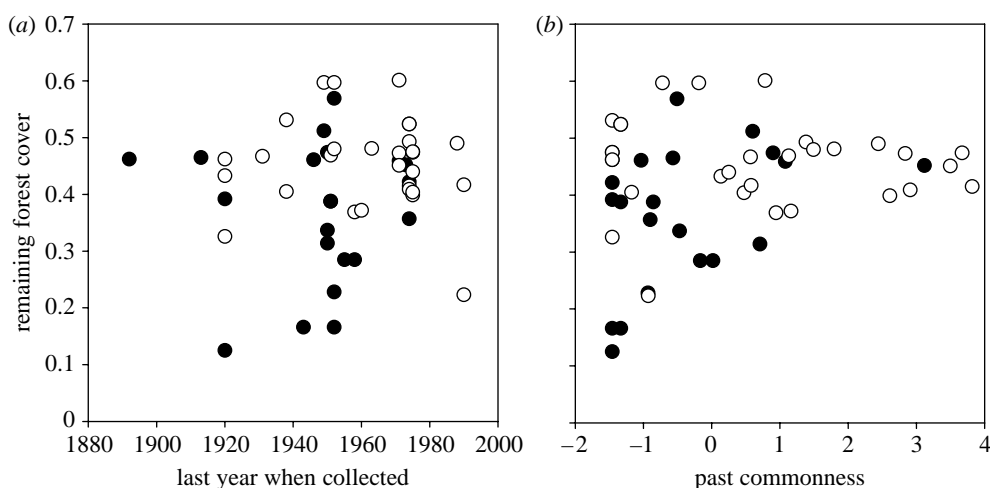


Figure 2. Plots showing whether a species has been collected by us (open symbols) or not (closed symbols, 'apparently extinct' species), depending on relative forest loss within the range of the species plotted (a) against the last year when the species was collected prior to our sampling and (b) against the past commonness of the species (PC1).

during 2002–2006 were PC1, the last year when the species had been collected prior to our sampling and the distances from the past sampling localities to our sampling sites. None of these effects is unexpected. Most of the apparently extinct species have not been collected for 50 years or longer (figure 2).

It is perhaps surprising that the past or current extent of forest within the range of the species had no significant effect on whether we collected the species or not. Admittedly, the forest maps depict a crude picture of forest cover and the past sampling localities provide inaccurate estimates of species' ranges. In this situation, long-term change rather than one-time state of the environment may better reflect species' responses. Furthermore, in many regions forest loss from 1953 to 2000 was a continuation of a longer process. Some species may have been on decline already in 1953 due to deforestation, and the occurrence of species is tracking forest loss and fragmentation with a time lag (Hanski & Ovaskainen 2002).

Forest loss and fragmentation is associated with an increasing pressure on lemurs, the most important dung producers in Madagascar. According to the fossil record, 16 large-bodied lemurs have gone extinct since human colonization *ca* 2300 years ago (Burney *et al.* 2004). The long-term and dramatic decline of *H. undatus* (figure 1d) may be due to high degree of resource specialization. Unfortunately, nothing is known of its biology and by now this once exceptionally widespread species may already be extinct.

Madagascar has large numbers of species with narrow geographical ranges (Wilmé *et al.* 2006). In this situation, and taking into account that Madagascar has already lost most of its forests, very large numbers of insects and other poorly known taxa may already be extinct, effectively extinct or rapidly heading towards extinction due to past and current deforestation. The current plans to expand the protected area network to six million hectares will amount to *ca* 10% of the original forest cover. Species–area considerations suggest that this will protect roughly half of the species (MacArthur & Wilson 1967). Our results are consistent with this prediction.

We thank the personnel of forest reserves and parks in Madagascar for their help in sampling, Evgeniy Meyke for calculations of forest loss, Navjot Sodhi and an anonymous referee for their comments and the Academy of Finland for support (grant nos 64350 and 20286, Finnish Centre of Excellence Programme 2000–2005).

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