

Elevational distribution patterns differ between exotic and native birds in Reunion Island

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Abstract

Introduction

In this paper, we use past data to document the elevational patterns of bird assemblages on a tropical volcanic island. We specifically address the question whether elevational distribution patterns differ between exotic and native bird species. We analyse patterns along a slope on Reunion Island (Indian Ocean) that presents one of the widest bio-climatic gradients within any oceanic island, *i.e.* from 0 to more than 3000 m on a distance of 20 km. We first characterize species distribution along the gradient and compare distribution parameters across native and exotic species. We then characterize patterns of species richness and compare the observed patterns to two different nul models.

Analysing observed patterns relatively to nul models expectations can inform on assembly processes. Nul models differ in the way they assemble community data, that is species occurrence data, here abundance, in sampled sites. They can include several constraints which consist in preserving some properties of the observed data, such as species overall abundance or local abundance at each sampled site. Here, the two tested models assemble species randomly preserving their frequency along the gradient, in order to retain differences in rarity. The first model is constrained by the total abundance of birds observed locally that is held constant during estimation. The model thus conserves interactions between total abundance and the environment, which can be referred to as the 'site capacity hypothesis'.

This capacity is supposed to be mobilized by birds present to some constant level, which is reflected in the total estimated abundance.

In the second model, these interactions are suppressed supposing that site capacity varies randomly. In this model, the sites are supposed as equal in their capacity to support birds to some random level.

Comparing observed data with nul models allows to explore differences in the elevational patterns of species richness between native and exotic birds under the constraint of observed commonness and site capacity or not.

Studies of this type of oceanic islands may contribute to understand the potentially huge impact of biological invasions (Loope & Mueller-Dombois, 1989).

Material and methods

Studied area

Reunion Island is part of the Mascarene volcanic archipelago, situated in the Western Indian Ocean (55°30' E and 21°05' S)(Fig. 1). This oceanic island appeared three million years ago. In the centre of the island, a plateau (Plaine des Cafres, \approx 2000 m) is flanked by two volcanoes (Piton de la Fournaise, 2631 m; Piton des

Neiges, 3069 m), the sides of which slope steeply towards the sea. There is a very steep climatic gradient: the average minimum ground temperature falls below 0°C above 1500 m in August, above 1800 m from June to October, and above 2300 m all year round; at the highest point, the temperature ranges from -10°C to +26°C (**Cadet1974; Raunet1991**). The studied area, located on the leeward side of the island had a slope of about 0.15 m per m and an annual rainfall ranging from 500 mm at sea level to 2000 mm on the highest mountain slopes **Raunet1991** The land is roughly organized through a land use gradient into a succession of elevational belts (**Cadet1980**) urbanised areas (0-50 m a.s.l.), herbaceous savannah (50-300 m), sugarcane monoculture (300-800 m), rural landscapes with diversified crops (800-1200 m), primary forest mixed with pastures and planted forests (1200-1800 m), heathlands dominated by ericaceous plants (1800-2600 m), and bare soils with a very sparse vegetation of herbaceous species (2600-3069 m) (Fig. ??). Above 1200 m, about 60

Sampling

The transect was sampled through one survey during the breeding season (December, 1997 - April, 1998), using 363 sample points spread along an elevational gradient from 20 m to 2880 m (Fig. ??). Further prospecting between 2880 and 3069 m (Piton des Neiges) revealed no bird presence. Sea birds nest at these high altitudes but were not considered in the study. This was due to of a lack of vegetation, except for scarce amounts of native herbs and weeds. Sampling points were equally distributed all along the elevational gradient, with a mean elevational distance of 8 m between two successive elevations, and a distance of at least 250 m between two successive count points, a mean sampling pressure of 12.7 count points per 100 m elevational band, and covering the overall diversity of vegetation types. Depressions, ridges and mires were avoided in order to avoid extremely wet or dry places. We also avoided places too close to feeding troughs in grasslands as they attract granivorous bird species. We assume that most of the birds were sampled near their nesting sites. In order to limit sampling bias, the different ecosystems found at each 100 m elevational band were surveyed so as not to miss individual bird species specific to individual ecosystems.

Data analysis

We calculated the weighted mid-point of species distribution as the average of all elevations where a species was detected weighted by the number of times it was detected. We also compare species maximal and minimal elevation where observed along the gradient, and amplitude in elevation range. We use the word elevation instead of altitude, as elevation is meters above sea level where altitude is meters above ground. We compared observed species richness in the two groups of bird species (native and exotic) to two different nul models based on random permutations of the original dataset **Gotelli2000** In both models, species overall abundance, that is the total number of records for each species, was preserved in each permutation, in order to account for observed differences in species commonness and detectability. The main difference between the two nul models was in the treatment of site carrying capacity of birds diversity, which under equilibrium hypothesis can be estimated as the total numbers of individuals locally. In the first nul model (N1), sites are considered as equally suitable for birds. In this case, the randomization suppresses any structure that exists in the data due to relationships between carrying capacity and environment (here elevation) in a broad sense. The randomized pattern obtained mimics the distribution of species richness not accounting for those relationships. In the second nul model (N2), the carrying capacity of sites, that is local species richness, was preserved by the randomization procedure. This model holds account for differences in sites to support bird abundance in relation with habitat carrying capacity. In this case, the randomized pattern obtained takes into account both species commonness and site suitability. The total number of individuals observed locally is preserved and species abundance in the dataset are preserved, but individuals and samples are randomized across sites. For each model, we performed a number of permutations ($n = 1999$) to obtain 95-% confidence intervals on simulated species richness.

Results

Twenty-two bird species were recorded over the 363 sampling points, with a mean of 4.3 species per point and 9.4 species per 100-m elevational band along the gradient. Fourteen species were exotic and eight native (Table S1). The number of records varied widely across species between 1 for two species (*Coturnix chinensis*, *Phedina borbonica*) and 268 for *Zosterops borbonicus* (mean: $71.2 \pm \text{s.d.}: 77.8$, Fig. 2).

Amplitudes in elevational range were high. Elevational amplitude exceeded 2000 m for five species and 1000 m for 15 species (Table 1). The largest elevational amplitude measured in native species was for *Zosterops borbonicus* (92 % of the gradient) and for *Acridotheres tristis* in exotic species (83 %). The lowest elevational amplitude measured in native species was registered for *Circus maillardi* (20 %) and for *Perdicula asiatica* in exotic species (8 %). The distribution of weighted mid-points, that is the weighted mean of elevations where a species occurs along the gradient, was highly concentrated between 1000 and 1500 m, with the exception of three exotic species (Fig. 1). The weighted mid-point and minimum elevation were however higher in native species ($p = 0.034$ and $p = 0.024$). By contrast, elevational amplitude and maximum elevation did not differ significantly between the two groups of species ($p = 0.534$ and $p = 0.254$ respectively, Fig. 1). No difference in the variability of species distribution was observed across the two groups: Bartlett test's of variance homogeneity showed that the variability was similar in the two groups for the four tested variables.

The total species richness observed within 100-m elevation bands varied between 2 and 19 (10.4 ± 5.5 , Fig. 3.a). Total richness showed a unimodal mid-elevational peak between 1000 and 1200 m. However, the pattern of species richness differed significantly between exotic and native species with increasing elevation (Fig. 3). Species richness varied between 0 and 7 for native birds (3.5 ± 2.2), and between 1 and 12 (6.9 ± 3.9) for exotic birds (Fig. 3.b). Species richness showed different elevational patterns across the two groups. Species richness of native birds showed a unimodal mid-elevational peak around 1200 m, whereas richness for exotic birds species showed a low-elevation plateau with a small mid-peak also around 1200 m. Exotic species were more numerous than native birds up to mid-elevation, whereas native species were more numerous at higher altitudes (Fig. 3.b). According to richness patterns, three zones could be defined: (1) from 0 to 900 m, there was a significant difference between exotic and native species, both in the number of species contacted and in their variation pattern, as native species increased with the elevational gradient from two to seven while the exotic ones remained at a high level (around 9 species); (2) from 1000 to 2000 m, richness of both exotic and native species presented a peak (around 9 and 6 species, respectively); (3) from 2000 to 3000 m, richness was higher in native species while exotic species were barely present.

In order to explore species co-occurrence according to their status, we compared species richness to random patterns obtained by simulation. Values occurring within the simulated 95% confidence band for the random pattern indicated elevation bands where species richness was not statistically different from random patterns of species assemblage (Fig. 4). In both groups, null patterns showed higher variability at low and high elevation when sites were all considered as equally suitable (randomized total abundance), whereas null patterns were more constrained when the total abundance observed at each site was kept constant (site capacity hypothesis). All models showed mid-elevation peaks however less pronounced under the site capacity hypothesis.

Native species richness was significantly lower than random expectations at low elevation for both null models (Fig. 4.a). This finding indicates depauperate communities where native species occur less frequently than expected by chance. From 900 m above, the observed pattern in native species richness overall supported more the site capacity hypothesis. Regarding exotic birds, species richness was lower than expected by chance above 1500 m high (Fig. 4.b). This result indicates that exotic species in high elevation bird communities are in low numbers compared to the site capacity. At low to mid altitude, the observed pattern in exotic species richness was closer to the confidence interval of the null model without constraint on site capacity (equally suitable hypothesis). The site capacity hypothesis generated higher than observed numbers of exotic species.

Discussion

Tables

Table 1: Summary of species elevation patterns. Occurrence: number of times the species was recorded. Mean : average elevation where the species was observed. Weighted mean: average altitude weighted by the number of times the species was recorded at each site. Min., Max.; minimum and maximum elevation at which the species was observed.

Species	Occurrence	Mean (m)	Weighted mean (m)	Min.	Max.
ACTR	168	1084	1123	20	2400
CIMA	6	1058	1049	800	1370
COCH	1	1010	1010	1010	1010
COCO	33	1267	1307	320	1700
COFR	57	1115	974	185	1660
ESAS	88	1001	987	60	1720
FOMA	268	1051	1033	20	2250
FRPO	2	128	142	70	185
GEST	109	840	804	20	1620
HYBO	67	1405	1410	620	2560
MAMA	5	1528	1615	1240	1705
PADO	72	1128	1202	65	1960
PEAS	16	117	119	50	290
PHBO	1	1495	1495	1495	1495
PLCU	32	580	606	65	1345
PYJO	72	794	805	65	1690
SATE	202	1590	1601	270	2880
STPI	34	1181	1225	100	2150
TEBO	18	1307	1336	900	1680
TUNI	41	832	808	35	1700
ZOBO	229	1371	1347	60	2700
ZOOL	45	1391	1388	620	2600

Figure legends

Figure 2: Box-and-whisker plots indicating the median, first and third quartiles and values outside inter-quartile range of elevations where species were detected. Species are sorted by order of median elevation. Blue indicates native species, , and red exindicate exotic species. Species are labelled with the first two letters of the genus and species names (see Table S1).

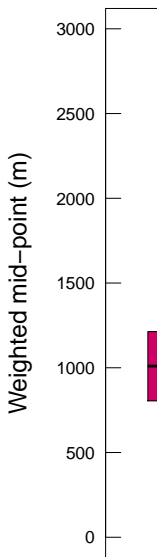
Figure 1: Four features of species distributions with respect to elevation in native and exotic species : (a) weighthd mid elevation point (average elevation where the species was detected weighted by number occurrences), (b) elevation amplitude (difference between highest and lowest elevation of detection), (c) minimum elevation where the species was detected and (d) maximal elevation of detection. Box-and-whisker plots indicate the median, first and third quartiles and values outside inter-quartile range. Symbols indicate individual values ($n = 22$ species), open symbols indicate species with less than 5 records in dataset ($n = 3$ species, see Table 1).

Figure 3: Elevation patterns of species richness : (a) total and (b) according to species status, native or exotic. Richness is calculated here per 100-m elevation band as the number of species detected within the band. Dotted lines indicate generalized additive models fits (GAM) with colored bands showing the 95% confidence interval on estimated richness.

Figure 4: Comparison between observed richness (dots) within 100-m elevation bands for native (a. & c.) and exotic (b. & d.) birds and expected species richness under two different null models that simulate random species assemblages (95%-confidence band). Green lines: null model with fixed site capacity, blue lines: null model with random site capacity (equally suitable sites hypothesis). In both models, species frequencies in the dataset were preserved during simulation ($n = 2000$).

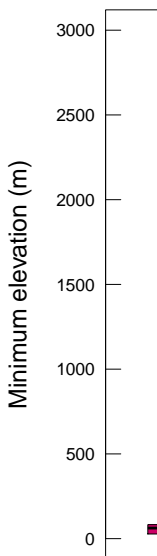
Figures

(a)



```
## Error in eval(expr, envir, enclos): impossible de trouver la fonction "beeswarm"
```

(c)



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## Error in eval(expr, envir, enclos): impossible de trouver la fonction "beeswarm"
```

Figure 1:

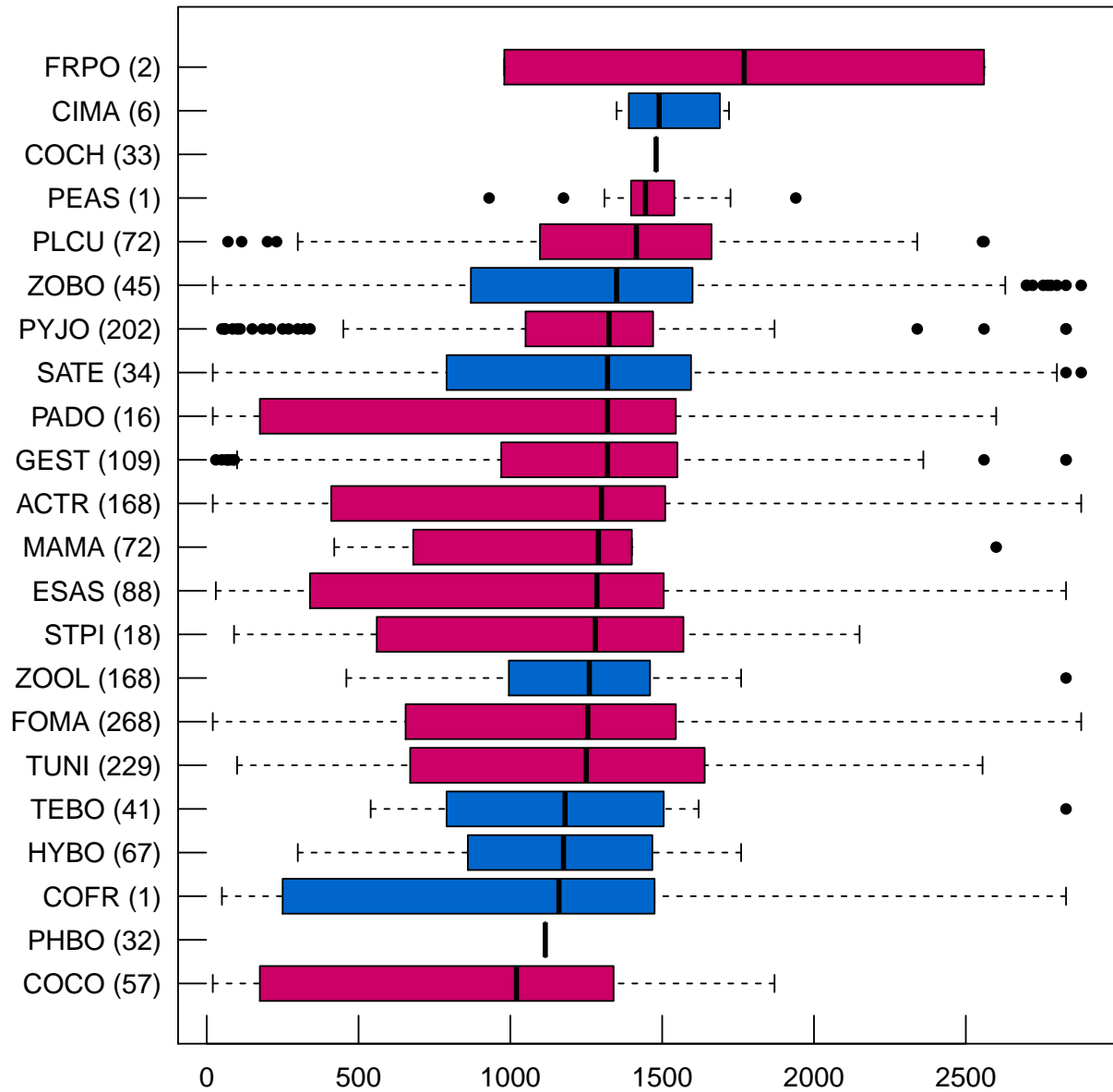
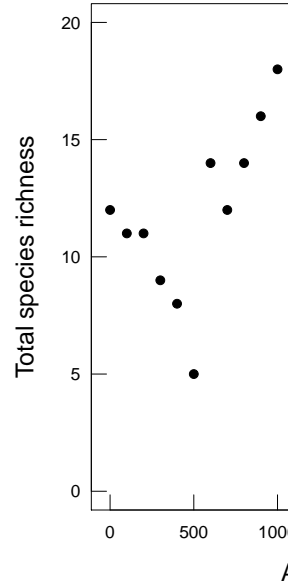


Figure 2:

a



```
## Error in array(np, 1):  'data' doit être de type vecteur, il était 'NULL'
## Error in array(np, 1):  'data' doit être de type vecteur, il était 'NULL'
## Error in eval(expr, envir, enclos):  objet 'fit' introuvable
## Error in eval(expr, envir, enclos):  objet 'fit' introuvable
## Error in eval(expr, envir, enclos):  objet 'ucl' introuvable
## Error in xy.coords(x, y):  objet 'y.polygon' introuvable
## Error in xy.coords(x, y):  objet 'fit' introuvable
## Warning in rm(i.back, i.for, lcl, ucl, se, fit):  objet 'lcl' introuvable
## Warning in rm(i.back, i.for, lcl, ucl, se, fit):  objet 'ucl' introuvable
## Warning in rm(i.back, i.for, lcl, ucl, se, fit):  objet 'se' introuvable
## Warning in rm(i.back, i.for, lcl, ucl, se, fit):  objet 'fit' introuvable
```

Figure 3:

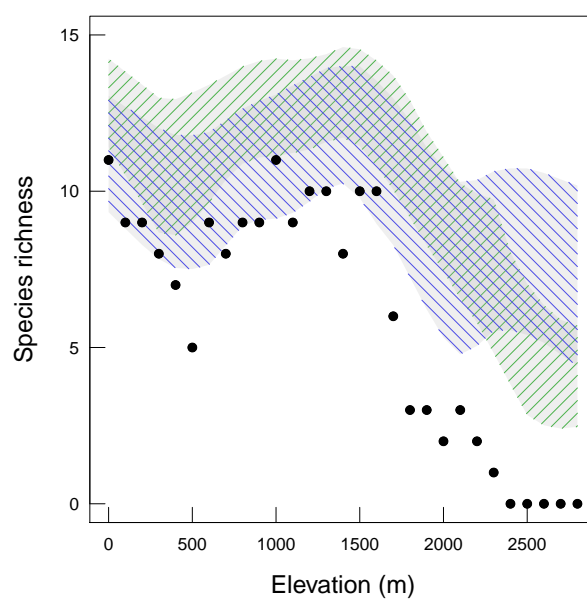
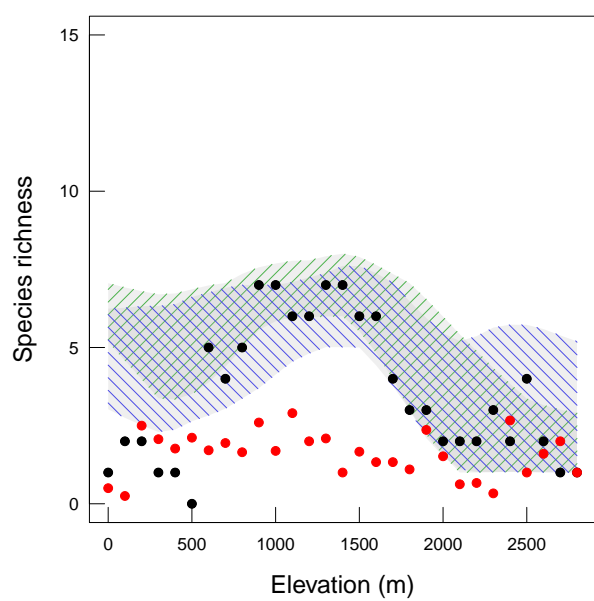


Figure 4:

Table S1: List of bird species detected in the study

Label	Species	Order	Family	Mass (g)	Size (cm)	Status	RLI
ACTR	Acridotheres tristis	Passeriformes	Sturnidae	82 – 134	23 – 29	E	LC
CIMA	Circus maillardi	Accipitriformes	Accipitridae	50	650 – 1000	R	EN
COFR	Collocalia francica	Apodiformes	Apodidae	7 – 13,3		M	NT
COCH	Coturnix chinensis	Galliformes	Phasianidae	31 – 41	12 – 15	E	LC
COCO	Coturnix coturnix	Galliformes	Phasianidae	16	70 – 135	E	LC
ESAS	Estrilda astrild	Passeriformes	Estrildidae	7 – 8	10	E	LC
FOMA	Foudia madagascariensis	Passeriformes	Ploceidae	13	17 – 19	E	LC
FRPO	Francolinus pondicerianus	Galliformes	Phasianidae	200 – 340	30 – 32	E	LC
GEST	Geopelia striata	Columbiformes	Columbidae		20	E	LC
HYBO	Hypsipetes borbonicus	Passeriformes	Pycnonotidae	22	51 – 57	R	LC
MAMA	Margaroperdrix madagascariensis	Galliformes	Phasianidae		24 – 26	N	LC
PADO	Passer domesticus	Passeriformes	Passeridae	26 – 32	14 – 16	E	LC
PEAS	Perdicula asiatica	Galliformes	Phasianidae	57 – 82	17	N	LC
PHBO	Phedina borbonica	Passeriformes	Hirundinidae		13,5	E	LC
PLCU	Ploceus cucullatus	Passeriformes	Ploceidae	17	32 – 45	E	LC
PYJO	Pycnonotus jocosus	Passeriformes	Pycnonotidae	20	23 – 42	E	LC
SATE	Saxicola tectes	Passeriformes	Muscicapidae	12,5	10,5 – 15	R	LC
STPI	Streptopelia picturata	Columbiformes	Columbidae	28	145 – 255	N	LC
TEBO	Terpsiphone bourbonensis bourb.	Passeriformes	Monarchidae	15	10 – 12	M (R)	LC
TUNI	Turnix nigricollis	Turniciformes	Turnicidae		13	N	LC
ZOBO	Zosterops borbonicus borbonicus	Passeriformes	Zosteropidae	10	7 – 8	M (R)	LC
ZOOL	Zosterops olivaceus	Passeriformes	Zosteropidae	10,5	9 – 10	R	LC

Supporting information