

1 Stability in the Face of Global Decline: A 20-Year Study of Arthropods 2 in an Oceanic Archipelago

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53 Summary

54 Insect declines have been reported globally but whilst island ecosystems are potentially facing
55 exacerbated challenges, no long-term studies (LTER) have confirmed this trend. This study utilises
56 the first available LTER data on island invertebrates, targeting epigeal and canopy arthropods from
57 the Azores Archipelago (Portugal), and covering over 20 years in three distinct sampling events from
58 30 standard sites. We investigate changes in abundance, biomass, and species richness within these
59 arthropod communities, focusing on the proportions of endemic and introduced species, and trends
60 among single-island endemics and forest-dependent endemics.

61 We reveal no significant declines in overall abundance, biomass, or richness of endemic or non-
62 endemic native arthropod populations, although we observe considerable variability between
63 sampling events in both epigeal and canopy communities. Of the species analysed, 26% experienced
64 declines, 15% increased in abundance, and 59% showed no change. We found minimal evidence for
65 the rapid spread of exotics or their increasing influence in the communities, nor any exotic-driven
66 extirpation of endemic species. Forest-dependent endemic species declined at a lower rate than

67 anticipated, suggesting that the extinction debt for these species may be less severe than previously
68 thought. The three-decade-long conservation of Azorean native forests may have contributed to
69 these unexpectedly stable populations.

70

71 Main

72 Collectively, oceanic islands harbour disproportionate amounts of biodiversity and in particular, high
73 proportions of endemics compared to continents¹. However, in addition, oceanic islands also
74 possess some of the world's most threatened biotas¹. Their vulnerability stems from high levels of
75 anthropogenic exploitation, habitat degradation, and the deleterious influence of introduced species
76 relative to their land area^{2,3}, which is further exacerbated by the effects of ongoing climate change⁴
77 and the elevated vulnerability of species evolved in isolation⁵. Consequently, the rate of historical
78 and contemporary anthropogenic extinction on islands surpasses that of the continents⁶.

79 Arthropods are fundamental elements of most ecosystems and several recent studies have reported
80 declines^{7,8}, particularly in relation to insects. Arthropod declines may lead to an accelerating
81 deterioration of ecological functions provided by these organisms⁹, diminishing ecosystem services
82 (e.g. pollination, regulation of pests, decomposition) and, ultimately, leading to ecosystem
83 collapse¹⁰. Despite several studies raising the alarm in terms of insect declines^{7,11,12}, insect
84 populations do not show a consistently declining pattern, particularly as trends have been found to
85 differ between taxa and functional groups and habitat type¹². As true for other taxonomic groups,
86 some species may increase in prevalence⁸ and the proportion of winners often counterbalances that
87 of the losers¹³.

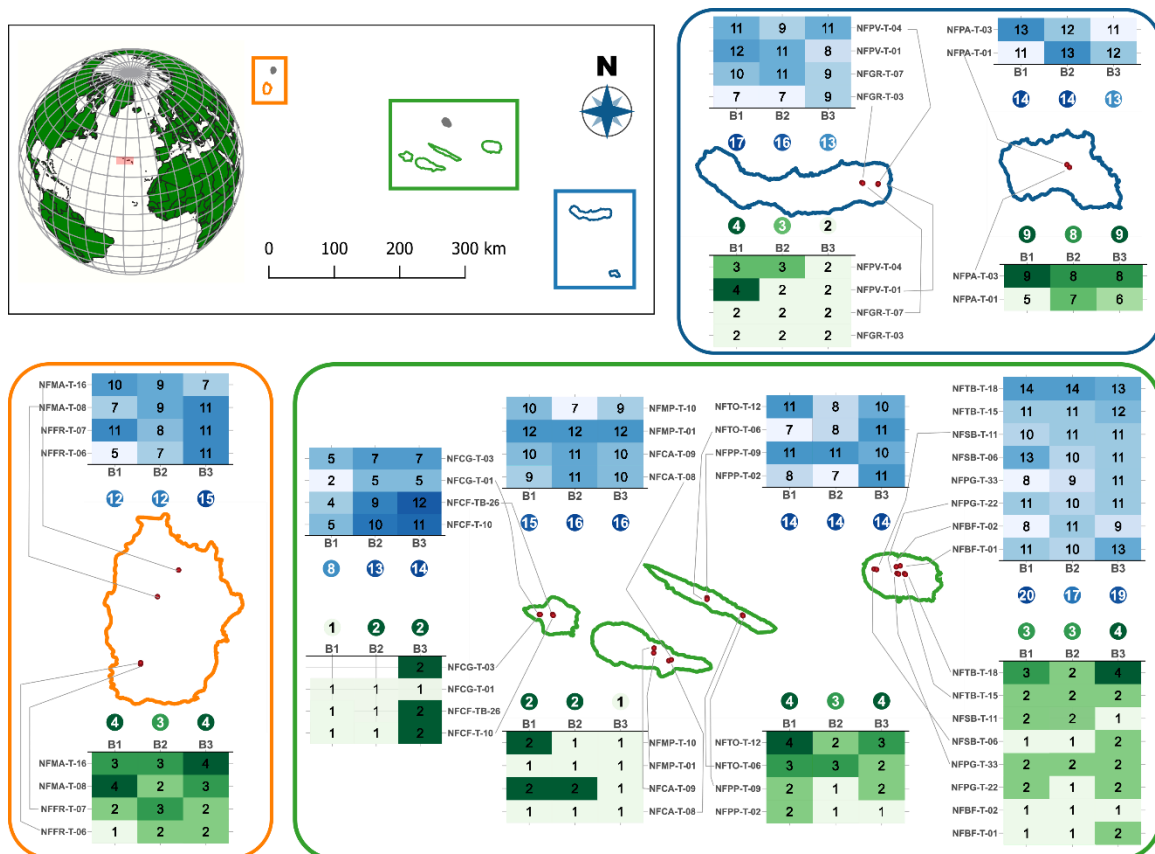
88 While island biotas are known to be particularly threatened, few studies have tested for insular
89 arthropod declines through time. Thus, the contribution of islands to global insect declines remains
90 unknown. Island endemics in particular are expected to decline¹, with many rare endemic species

91 expected to be threatened with extinction. Yet, mostly because of the scarcity of systematic and
92 standardised sampling of, and long-term databases on, insular arthropods, few of these predictions
93 have received sufficient scrutiny to be able to confirm their generality for arthropods of remote
94 islands. Moreover, since the rate of non-native species introduction on islands may exceed the
95 extinction of endemic species¹⁴, it is key to gain understanding into whether population trends of
96 endemic, native but not endemic (henceforth native), and introduced (or exotic) species follow
97 similar trajectories.

98 The humid forests of the Azorean Islands (North Atlantic, Macaronesia) form a distinctive habitat
99 type that once covered ~99% of the archipelago. By the second half of the 20th century its area had
100 shrunk to its current range of only 5% of its original size¹⁵. Due to the legal protection of the most
101 pristine areas, the current area of the forest remnants has not decreased for almost four decades
102 and direct human disturbances have been reduced significantly. Yet, the persisting forest patches
103 are highly fragmented and prone to indirect (mostly anthropogenic) influences, such as the spread of
104 exotic, and potentially invasive, species¹⁶ and climate change^{17–19}. Moreover, it remains unclear
105 whether these forest patches are large enough to sustain viable populations of endemic forest-
106 dependent arthropod species and halt the future extinction processes that have been predicted.
107 Hence, our focus herein is primarily on the endemic species that survive in less-than-ideal
108 circumstances, some of which may not present viable populations, contributing to the so-called
109 extinction debt²⁰. Notwithstanding the apparent urgency of comprehensive conservation strategies
110 for these imperilled species, the lack of in-depth insight into population trends (Prestonian
111 shortfall^{21,22}) of insular arthropods may hamper the development of effective protection measures.
112 In this study, we utilise a dataset of standardised samples collected in 1999–2000, 2010–2011, and
113 2020–2023. This unique data series, compiled as part of the “Biodiversity of Arthropods from the
114 Laurisilva of the Azores” (BALA) project²³, provides an opportunity to gain the first detailed insight
115 into whether arthropods are declining in the native forests of remote oceanic islands. Whilst we pay

116 particular attention to species of outstanding conservation importance, such as single-island
 117 endemics (SIE) and strict forest-dependent Azorean endemics (FDE) considered to be facing
 118 extinction debts^{20,24} (Fig. 1), we also scrutinise how exotics change over time in abundance and
 119 species richness. Moreover, since the overall trends of arthropod abundances may be differentially
 120 influenced by island-wide trends, we also take a simulation approach to investigate the contribution
 121 of each individual island towards the overall changes. Our dataset holds a collection of abundance
 122 data of ground-dwelling (epigeal henceforth, collected with pitfall traps) and canopy-dwelling
 123 (arboreal henceforth, collected by canopy beating) arthropods from 30 repeatedly sampled sites
 124 across seven Azorean islands, collected in three distinct archipelago-wide sampling campaigns (B1,
 125 B2, and B3, henceforth) from 1999 to 2023.

126



127

128 **Figure 1.** The location of the Azorean Archipelago and the seven sampled islands (top left) in the
 129 **three island groups (colour-coded as orange, green, and blue for the Western, Central, and Eastern**

130 groups, respectively). Non-sampled island coloured grey. Sampling sites in the three detailed
131 subfigures are indicated with red dots, and arrows linking them to rows in the blue and orange
132 tables show the number of single island endemic species (SIE) or strict forest-dependent endemics
133 (FDE) (respectively) in each sampling campaign. Numbers in circles represent the number of
134 unique SIE/FDE species caught in each BALA sampling campaign (B1-B3) from the entire island.
135 Abbreviations as: FLO – Flores, FAI – Faial, PIC – Pico, SJG – São Jorge, TER – Terceira, SMG – São
136 Miguel, and SMR – Santa Maria.

137

138 Results

139 *Changes in overall abundance*

140 The data encompass about 17% of the known Azorean arthropod species described to date,
141 including ~33% of known archipelagic endemic taxa from 15 native forest fragments^{25–27}. This
142 corresponds to 30,078 observations of 403 arthropod species (of which 91 or ~23% are archipelagic
143 endemics and 153 or ~38% introduced).

144 To assess the yearly changes in overall arthropod abundances between the sampling events, we first
145 calculated the average differences for the years between consecutive sampling events. Using
146 Cohen's d and bootstrapped confidence intervals, we determined whether standardized effect sizes
147 of transect-wise changes significantly diverged from zero.

148 Our findings indicated no significant overall change in arthropod abundance (i.e. when endemics,
149 natives, and exotics, were all included), between the initial (B1) and final (B3) sampling campaigns
150 conducted from 1999 to 2023, in either ground- or canopy-dwelling communities. However, notable
151 differences were observed between B1 and B2, and B2 and B3. A minor decrease from B1 to B2 was
152 seen for non-endemic native species, which was somewhat offset by increases between B2 and B3.
153 (Supplementary Table S1). However, ground-dwelling species exhibited slight reductions in
154 abundance, with Cohen's d values of -0.47 (CI: [-0.81, -0.12]) for endemics and -0.48 (CI: [-0.82, -

155 0.35]) for natives. The abundance of arboreal native species also declined, with a Cohen's d of -0.32
156 (CI: [-0.57, -0.1]). (Fig 2 A-B). The proportions of endemics and exotics relative to overall number of
157 individuals remained unchanged between B1 and B2. (Supplementary Table S1).

158 *Changes in biomass*

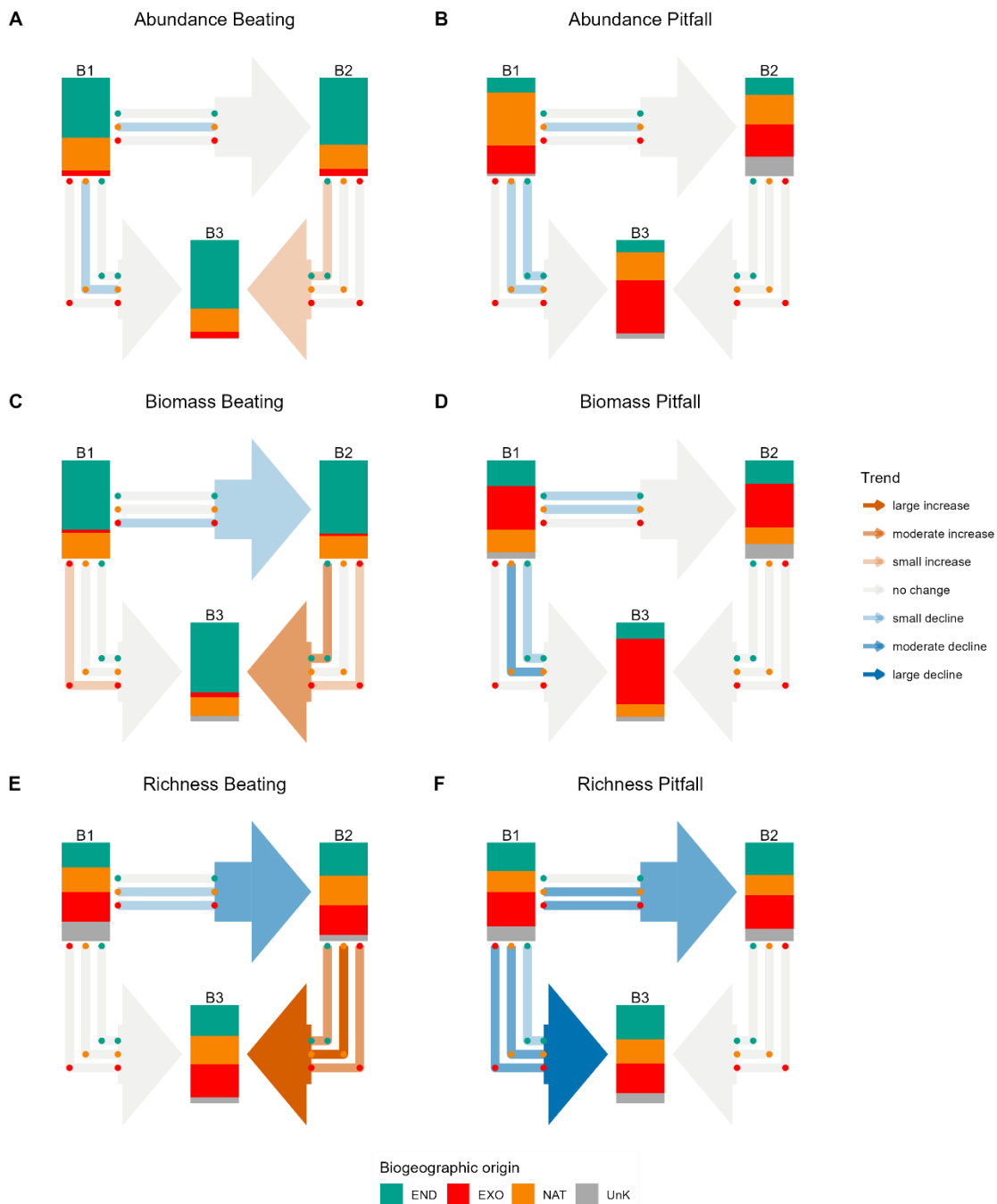
159 Since body mass is a disproportionately important trait driving ecosystem functions and it is often
160 regarded as a superior currency in terrestrial invertebrate community analysis²⁸, we also analysed
161 changes in arthropod biomass. Despite estimating biomass using taxon-specific indices and
162 individual counts (see Methods), the correlation between overall biomass and abundance was not
163 very strong (Pearson's $\rho = 0.82$ for ground-dwellers, 0.68 for canopy-dwellers; $p < 0.001$). Yet,
164 biomass trends closely mirrored abundance patterns, with small-magnitude losses between B1 and
165 B2 offset by gains from B2 to B3, resulting in no net changes in overall biomass across the 25 years.
166 However, when the groups of different biogeographic origin were examined separately, the biomass
167 of both endemic and native ground fauna showed a decline between B1 and B3 ($d = -0.32$, CI: [-0.71,
168 -0.07] and $d = -0.67$, CI: [-1.31, -0.50], respectively), whilst there was an increase in the biomass of
169 arboreal exotics ($d = 0.41$, CI: [0.21, 0.64]) (Fig 2 C-D, Supplementary Table S1). This increase also
170 significantly elevated the biomass-proportion of exotics in the canopy-dwelling community from B1
171 to B3 (Supplementary Table S1).

172 *Changes in species richness*

173 Moderate declines in species richness were noted for both epigeal and arboreal arthropods between
174 B1 and B2, with Cohen's d values of -0.75 (CI: [-1.25, -0.35]) and -0.59 (CI: [-1.32, -0.17]), respectively
175 (Fig 2 E-F, Supplementary Table S1). This was followed by a substantial increase in canopy-dwelling
176 species from B2 to B3 ($d = 1.27$, CI: [0.86, 2.07]), resulting in no net change in arboreal species
177 richness between B1 and B3. Conversely, ground-dwelling species showed no significant increases
178 between B2 and B3, leading to declines in species richness from B1 to B3 ($d = -0.80$, CI: [-1.28, -
179 0.47]).

180 When species groups of different biogeographic origins were analysed separately, ground-dwelling
 181 exotics and natives experienced greater losses from B1 to B3 ($d = -0.68$, CI: $[-1.23, -0.35]$) compared
 182 to Azorean endemics. Arboreal native species showed the largest increase between B2 and B3.
 183 Species-richness proportions of endemics and natives relative to total richness remained unchanged
 184 throughout the study (Supplementary Table S1, Figure S1).

185



186

187 **Figure 2. Summary of changes in abundance (A-B), biomass (C-D), and richness (E-F) between BALA**
188 **sampling campaigns (B1-B3) in arboreal (A, C, E) and epigeal (B, D, F) communities. The proportion**
189 **(from top to bottom) of endemic (END), non-endemic native (NAT), introduced (EXO) species, and**
190 **those with unknown biogeographic origin (UnK) in the communities are shown in the barplots. The**
191 **colours in the barplots, representing biogeographic origins (refer to legend), also correspond to**
192 **the dots along the thin lines that indicate changes for each group (excluding unknown origins). The**
193 **arrows indicate the direction of comparison and their colour indicates the overall changes.**
194 **Changes are standardised to time unit (year) and compared to a ‘zero change’ null model. Change**
195 **was considered statistically not significant when confidence intervals crossed zero. The magnitude**
196 **of change is indicated with colour depth, increasing from small (representing a Cohan’s d value**
197 **between 0.2 and 0.5), through medium (Cohan’s d value between 0.5-0.7) to a large change**
198 **(Cohan’s d > 0.7).**

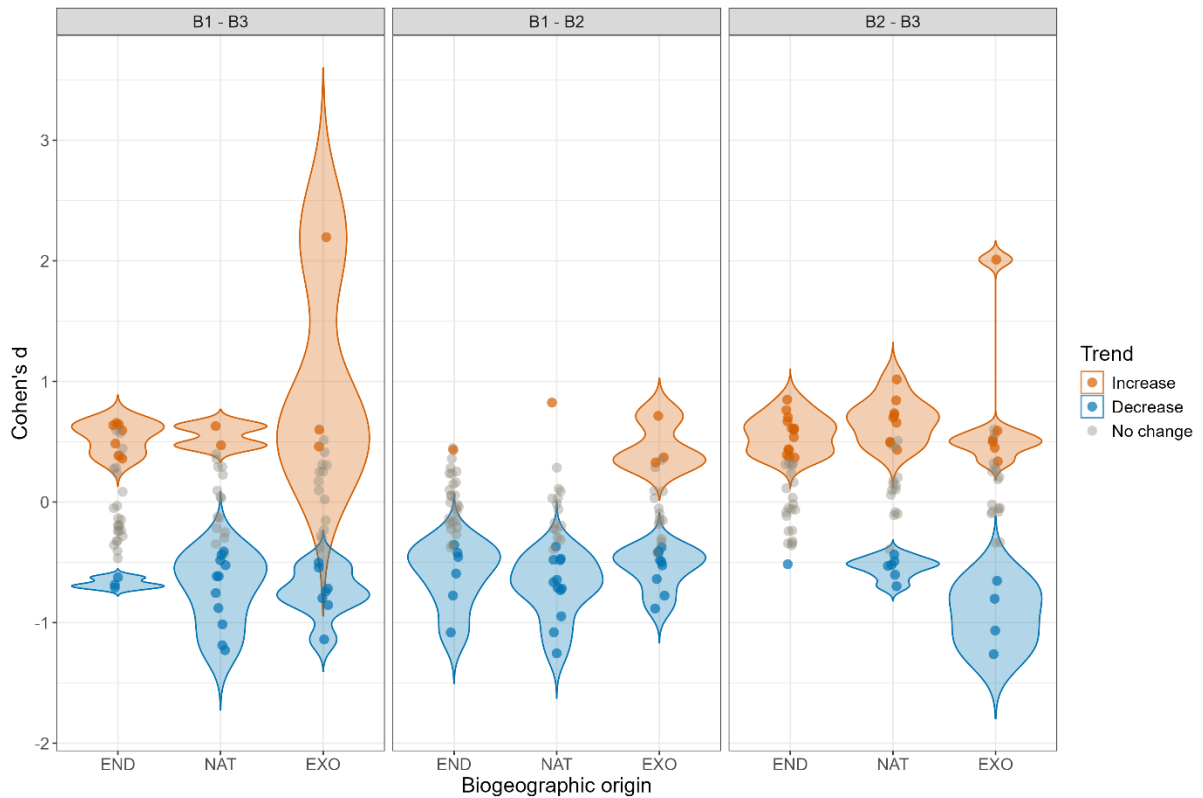
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200 *Changes in individual species abundances*

201 Our second approach focused on the abundance changes of individual species, particularly those of
202 high conservation concern, namely single-island endemics and strict forest-dependent endemics^{20,24}.
203 Of the 403 species present in our dataset, 84 (~21%) were sufficiently abundant to comply with our
204 criteria for formal analysis (see Methods). For these, we used the same methodology as above to
205 investigate how the abundances of individual species changed from B1 to B2 and B3, and from B2 to
206 B3. Of these, 22 species declined significantly from B1 to B3, including 3 with small declines (Cohen’s
207 d between 0.2 and 0.5), 13 with moderate declines (Cohen’s d between 0.5 and 0.7), and 6 with
208 large declines (Cohen’s d > 0.7). The mean Cohen’s d for declines was -0.74 (range: -1.23 to -0.41). In
209 contrast, 5, 7, and 1 species showed small, moderate, and large increases, respectively, with
210 magnitude comparable to that of the declines (mean Cohen’s d = 0.67, range: [0.36, 2.20]).

211 For endemic species, the mean effect size for overall decline (between B1 and B3) was -0.67 (range:
 212 [-0.71, -0.62]), while the increase was recorded at 0.53 (range: [0.36, 0.66]). Introduced species
 213 showed a decline with a mean effect size of -0.76 (range: [-1.14, -0.50]) and an increase of 1.09
 214 (range: [0.46, 2.20]) (Fig 3.).

215



216

217 **Figure 3. Mean change across transects in the abundances of Azorean native forest arthropod**
 218 **species between sampling campaigns. Changes are standardised to time unit (year) and compared**
 219 **to a 'zero change' null model. Species experiencing no significant changes (i.e. confidence intervals**
 220 **cross zero) are marked in grey. Violin plots indicate the first quartile, the median, and the third**
 221 **quartile of the data, as well as the frequency distribution of data points (each dot represents a**
 222 **species). END stands for endemic species, NAT for native but non-endemic, and EXO for**
 223 **introduced species. Only species whose abundance reached the requirements for the statistical**
 224 **analysis are included in the figure.**

225

226 For species whose abundance is too low to allow individual temporal trends to be analysed, we
227 merely compared the sum of collected individuals from each sampling campaign and no statistical
228 tests were conducted to compare the sampling campaigns. Here, 186 (~58%) showed declines, 37
229 (~12%) showed no change (i.e. no more than 20% difference between the first and last sampling
230 events), and 96 (~30%) increased. Endemics and natives accounted for only ~36% of declines, while
231 exotics made up ~38%. No significant differences were found when the proportion of increasing and
232 decreasing species based on biogeographic origin was compared (two-sided Fisher test $p = 0.204$).
233 There were also no differences among the different biogeographic origin in the number of species
234 increasing, being stable, or decreasing in the number of occupied sites (see Methods and
235 Supplementary Table S8).

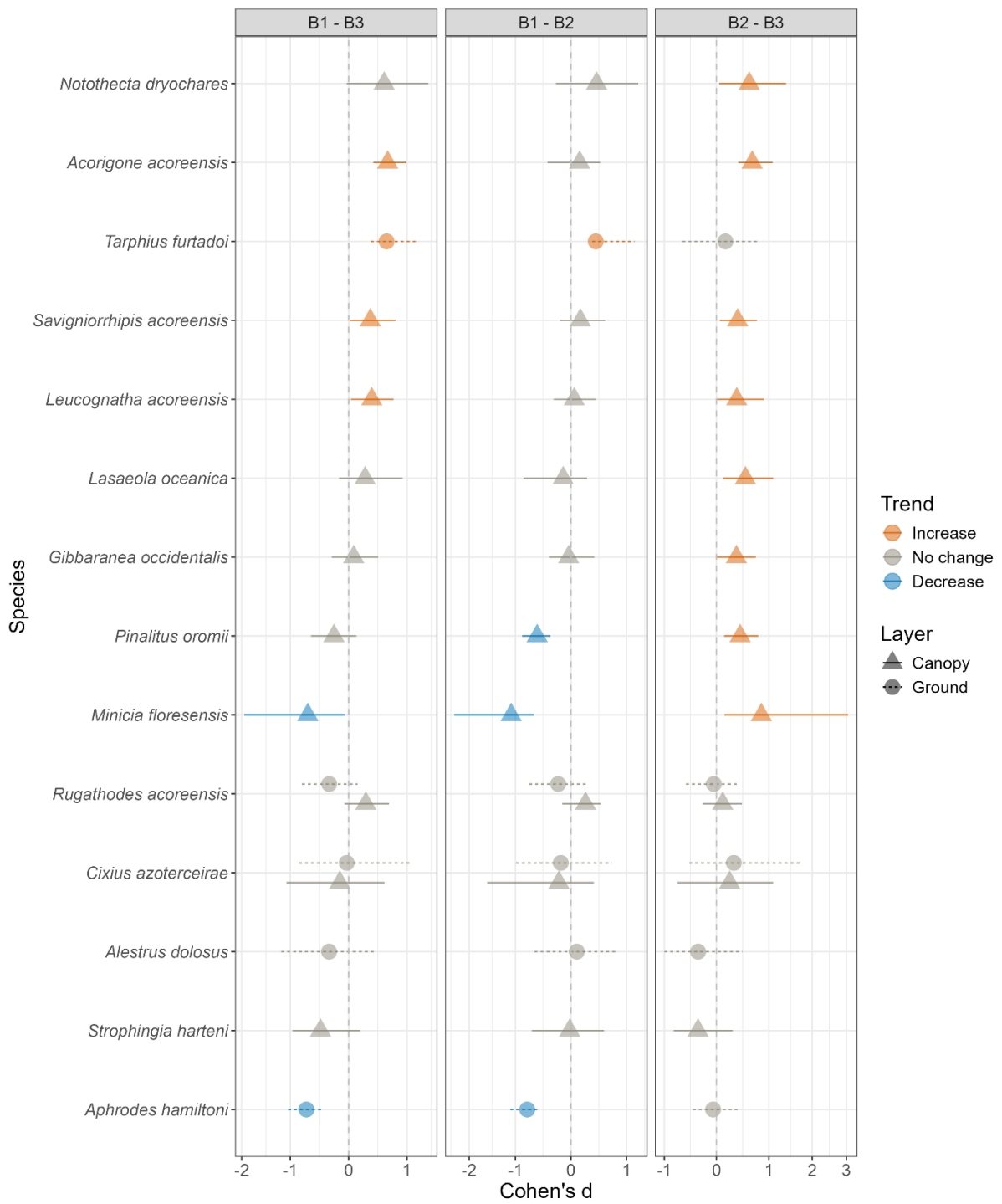
236 *Changes in individual FDE and SIE species abundances*

237 In our samples, 30 SIEs and 34 FDEs were found (Supplementary Table S2), and of these, the trends
238 of 14 FDEs meet our criteria for the calculation of reliable effect sizes and estimation of trends (see
239 Methods). The only SIE with sufficient data for testing, *Cixius azoterceirae*, also appears as an FDE.
240 The abundance patterns for FDE species mirrored overall abundance trends: marked declines
241 between B1 and B2 were followed by increases between B2 and B3. Indeed, while three FDE species
242 declined moderately from B1 to B2 and one increased, seven species showed small to large increases
243 between B2 and B3 without declines. Overall, only two species of FDE, the bug *Aphrodes hamiltoni*
244 and the spider *Minicia florensensis*, showed moderate declines between B1 and B3, whereas four
245 experienced small to large increases in abundance. (Fig. 4, Supplementary Table S2).

246 Of those FDE and SIE species that did not meet our criteria for statistical analysis (see Methods), 25
247 became less numerous (by more than 20% of the original abundance) from B1 to B3, 4 showed no
248 change, and 12 increased (Supplementary Tables S4-S7). The proportions of these outcomes did not
249 significantly differ from non-SIE or non-FDE species (two-sided Fisher test > 0.05). However, the

250 proportion of species decreasing in the number of occupied sites was significantly lower for FDE/SIE
 251 species compared to non-FDE/SIE species and exotics (one-sided Fisher tests, $p = 0.015$ and $p =$
 252 0.048 , respectively, Supplementary Table S8).

253



254

255 **Figure 4. Abundance changes for strict forest-dependent endemics (FDE) from BALA 1 to BALA 3**
256 **(first panel), BALA 1 to BALA 2, and BALA 2 to BALA 3 sampling campaigns. Symbols indicate the**
257 **mean log-transformed Cohen d-value and horizontal lines the corresponding confidence intervals.**
258 **Statistically not significant (i.e. confidence intervals cross zero) changes indicated with grey colour.**
259 **Labels on the x-axis are back-transformed and show mean annual changes in the number of**
260 **individuals.**

261

262 To explore how individual islands contributed to overall trends in abundance, biomass, and richness,
263 we used a simulation approach, removing one island at a time from the dataset or omitting
264 combinations of two or three islands, and recalculating differences between B1 and B3 (see
265 Methods).

266 These analyses showed that Terceira positively contributed to changes in overall abundance and
267 richness of arboreal species, particularly endemics, while negatively contributing to trends in
268 arboreal exotic species. Other islands had varying effects; for example, São Jorge positively
269 contributed to endemic species in epigeal communities but added to declines in canopy arthropods
270 and Santa Maria samples generally made a positive contribution to exotic abundances and richness
271 (Supplementary Figures S2-S10).

272 Discussion

273

274 Understanding population trends of arthropods in insular biotas is essential for effective protection
275 and the preservation of the ecosystem services that they provide as well as to fully answer the
276 challenges of global insect declines. In this study, we found minimal evidence of large-scale
277 depletion in the arthropod fauna of Azorean native forests since the start of sampling (in 1999/2000)
278 and neither did we observe significant declines in either endemic or non-endemic but native
279 arthropod populations. Instead, we found highly variable abundances, biomasses, and species

280 richness between sampling events in both epigeal and arboreal communities, with a balance of gains
281 and losses over time. This finding contrasts with the prevailing narrative of dramatic insect decline
282 worldwide and aligns with other large-scale analyses reporting natural fluctuations or stable insect
283 populations^{29,30}. Since endemics dominate in abundance over natives and exotics in the arboreal
284 communities in the Azores^{15,31}, this unchanging trend was driven by the relative stability of endemic
285 species abundances. Similarly stable populations of endemic flying insects have been reported from
286 Terceira Island over 6 and 10 year study periods^{16,32}. In our study, even the species predicted to
287 decline the most—such as single island endemics and forest-dependent endemics—generally
288 remained stable, with some canopy-specialised endemic species, like the spider *Acorigone*
289 *acorensis*, even experiencing population increases between the first and last sampling campaigns.
290 Moreover, site occupancy for both forest-dependent (FDE) and single island endemics (SIE) did not
291 significantly change and even showed slight increases, indicating at least some degree of habitat
292 stability.

293 Although the current protected remnants of Azorean native forests face some disturbances,
294 including the aggressive spread of the invasive ginger lily (*Hedychium gardnerianum*)³³, which alters
295 the edaphic conditions and microclimate of the forest understorey and inhibits the regeneration of
296 indigenous flora³⁴, these factors do not appear to significantly impact overall native ground or
297 canopy fauna trends. Yet, the invasion by ginger lily may still be linked to negative trends we
298 observed in epigeous species richness and declines in specialist herbivorous species like *Aphrodes*
299 *hamiltoni*. However, excluding São Miguel from our analysis, the largest island where tourism
300 pressure is high and the ginger lily is abundant, did not change the population trends of either the
301 native ground or canopy fauna. Indeed, studies suggest that land use adjacent to native forest
302 remnants has a greater influence on the spread of invasive plants than outdoor activities, such as
303 hiking^{35,36}. Nonetheless, the number of SIE and FDE species decreased only on this island. This
304 discrepancy could be due to the presence of already degraded and small habitat patches that cannot

305 sustain viable populations of specialised species like FDEs and SIEs. Thus, some arthropod species
306 may have already gone extinct due to historical events²⁴ rather than current pressures.

307 Intensive conservation measures implemented over the past three decades in the Azores appear to
308 have mitigated significant direct threats to native habitats and our results allow for cautious
309 optimism regarding the resilience of endemic arthropods to indirect anthropogenic effects. Native
310 forest patches on Terceira Island, which are among the largest in the Azores, seem to exhibit
311 resilience against invasive species, further supporting the idea that habitat protection is crucial for
312 the survival of endemic species. On the other hand, secondary forests of exotic tree species may
313 serve as refuges for some forest specialist endemics, allowing for population stability^{37,38}. However,
314 there is no evidence of spatial rescue effects for isolated populations of rare endemic species³⁹ and
315 no protection fully excludes long-term disturbances from invasive plant encroachment and
316 occasional human or livestock disturbances, nor does it mitigate the ongoing effects of climate
317 change. Indeed, some endemics continue to show declines, with three FDEs—*Eupteryx azorica*,
318 *Cixius azoricus azoricus* leafhoppers, and the *Atlantocis gillerforsi* beetle—not found in recent
319 samples, and others, such as the spider *Minicia floresensis*, becoming increasingly rare.

320 Yet, while Triantis et al.²⁰ predicted that 55-99% of FDEs would eventually face extinction, we found
321 strong statistical evidence of decline in only two species, with untestable indications of decline in 25
322 additional species out of a total of 55 species (SIEs and FDEs combined). Of the 34 FDE species
323 analysed by Triantis et al., only 14 (~41%) showed any signs of decline—far below the predicted
324 range. Moreover, 11 FDE species exhibited population increases, challenging the prevailing narrative
325 of impending biodiversity loss.

326 The limited information on the original arthropod fauna prevents us from ruling out the possibility
327 that the lack of widespread decline among SIE and FDE species indicates that the Azorean fauna had
328 already been impoverished by the time biodiversity research began in this archipelago, with many
329 rare habitat specialists potentially extirpated² or only persisting within relictual sites³⁸. Nonetheless,

330 based on our current knowledge, protecting native habitats appears key to the long-term survival of
331 prospering species, while declining species may benefit from additional focused protection
332 programs⁴⁰.

333 As introduced species often drive endemic extinctions⁴¹, simultaneous increases in exotics and
334 declines in endemics can be anticipated. However, despite reports of a clear increase in non-native
335 species in other studies on Azorean arthropods^{16,32}, we found little evidence that exotics are rapidly
336 outcompeting natives or endemics. On the contrary, declines of comparable magnitudes were noted
337 in the abundances of both endemics and exotics, with slightly (but not significantly) higher increases
338 in exotics. While it is particularly reassuring that neither the abundance nor the species richness
339 proportion of exotics changed in communities over this ca. 25-year timespan, their biomass and
340 biomass-based proportions did increase between the first and last sampling campaigns. While this
341 solely impacted canopy-dwelling communities, the gains exotics exhibited indicate unfavourable
342 changes for indigenous species. Nevertheless, exotics decreased more in site occupancy than SIE and
343 FDE species, suggesting a continuous influx of exotics with low colonization success from disturbed
344 edges of the forest patches⁴². Although these pioneer individuals have the potential to get
345 established and trigger processes that rewire island ecosystems⁴³, in previous studies, neither
346 Whittaker et al.⁴⁴ nor Pozsgai et al.⁴³ found clear evidence for functionally matching exotics
347 outcompeting indigenous species in the Azores. Furthermore, within healthy forest remnants, like
348 those on Terceira Island, the increase of exotics appears limited, while endemic arthropod
349 populations have persisted well. It thus seems possible that long-term environmental shifts,
350 including climate change, provide a greater threat than invasive non-native species.

351

352 *Limitations and future perspectives*

353 While the discontinuous time series presents a limitation, the extensive temporal and spatial scale,
354 combined with significant sampling efforts, make the BALA data a uniquely well-specified basis for

355 uncovering changing diversity patterns. Our sensitivity analysis, which showed minimal divergences
356 in results with up to 20% random sample removal (see Methods and Supplementary Figures S11-
357 S13) confirms the robustness of these patterns. However, the relatively low number of species
358 whose temporal trends could be statistically tested underscores the urgency of establishing a large-
359 scale and continuous monitoring system to assess long-term impacts on arthropod communities.

360 Several key questions remain unanswered. For instance, we know little about nuanced community
361 changes, such as species replacement, or how species traits influence population trends. Future
362 research should delve into the details driving these trends, considering species' traits, environmental
363 constraints, and biotic interactions. Additionally, understanding whether secondary or semi-natural
364 habitats can mitigate biodiversity losses will be essential.

365

366 [Concluding remarks](#)

367 Our findings challenge several studies on global insect declines and the predictions surrounding the
368 demise of insular biotas. They foster moderate optimism for the future of island biotas where
369 habitats remain intact and highlight the importance of long-term monitoring and effective
370 conservation strategies and the need for habitat protection to preserve obscure biodiversity.

371 However, one unfortunate conclusion from our work is that declines may still occur within protected
372 areas, indicating that the quality or size of these preserves may not be sufficient to buffer all
373 negative anthropogenic impacts and prevent biodiversity losses. Thus, our study has some significant
374 implications. First, although habitat protection appears to be effective in at least the short run, some
375 species require additional support from specific protection programs, such as ongoing LIFE
376 Beetles^{40,45} and LIFE Snails⁴⁶ projects. Second, the presence of introduced arthropods does not seem
377 generally to adversely impact indigenous species within Azorean native forest remnants but, instead,
378 the structural health of habitats is crucial for the persistence of indigenous arthropod communities.
379 Therefore, plant invasions may pose a more serious threat than competition from other arthropods.

380 Last but not least, if the natural capital of oceanic islands and the dependent ecosystem services are
381 to be protected, monitoring both indigenous and introduced species, and regularly re-assessing
382 community changes should be a priority⁴⁷. Even simplified, regular monitoring systems utilising
383 advanced technologies and focusing on selected indicator species would be adequate for
384 establishing early warning systems to detect the establishment of potentially competitive species or
385 significant declines in species of high conservation importance. This proactive approach, combined
386 with targeted removal of invasive habitat-structuring plants and competing arthropod exotics, will
387 be key to the success of conservation efforts in the Azores and other insular ecosystems.

388 Methods

389 Study sites and sampling methods

390 We used arthropod data collected from the Azorean archipelago (Fig. 1), using standardised
391 sampling protocols under the umbrella of the ‘Biodiversity of Arthropods the Laurisilva of Azores’
392 (BALA) project^{23,26,27}. In the course of the project, 30 sites of native Azorean humid forests⁴⁸ were
393 chosen for standardised arthropod sampling, repeated in three sampling campaigns: B1 (1999-2002),
394 B2 (2010-2011), and B3 (2021-2023). The sites were spread across 15 native forest remnants, on
395 seven out of the nine Azorean islands (from west to east: Flores, Faial, Pico, São Jorge, Terceira, São
396 Miguel, and Santa Maria), those with remaining native vegetation. Summary data on the transects
397 and remnants are available in Pozsgai et al.²³. All samples were collected between July and
398 September when arthropod activity is at its peak in the Azores.

399 To encompass a broad range of micro-habitats, we employed two primary sampling techniques. The
400 first involved pitfall trapping, to effectively capture soil- and ground-dwelling (epigeal) arthropods.
401 At each site, across a 150 m transect, 30 pitfall traps were evenly distributed, maintaining a
402 separation of 5 m between each. Alternating preservative/attractive solutions of ethylene-glycol and
403 Turquin’s solution⁴⁹ were employed in these traps, maximising catch. Pitfall traps were active for one
404 week in each sampling campaign.

405 The second technique, canopy beating, was employed to target canopy-dwelling (arboreal)
406 arthropods. At each selected site, the beating was performed on the three, locally most abundant
407 tree species, primarily focusing on native *Juniperus brevifolia* (Cupressaceae), *Erica azorica*
408 (Ericaceae), *Ilex azorica* (Aquifoliaceae), *Laurus azorica* (Lauraceae), and *Vaccinium cylindraceum*
409 (Ericaceae). For each of the three species, ten individual trees were randomly selected along the
410 transect and the branches were beaten five times at the height of 1.5–2 m. All visually observable
411 arthropods, except for mites (Acari) and springtails (Collembola), were sampled. Specimens were
412 sorted into morphospecies and, where possible, identified to the biological species. Due to
413 difficulties in their identification, Diptera and Hymenoptera (with the exception of Formicidae, which
414 were included) were excluded from the dataset. Biogeographic origin, categorised as endemic to the
415 Azores, native but not endemic to the Azores (termed as native for brevity), introduced (non-native,
416 also termed as exotic), and of unknown origin, was assigned to each species (Ref.^{23,25} and references
417 therein). Single-island endemics (SIE) were identified based on the dataset described in Ref.²³ and
418 the species list in the Supplementary Material of Ref.²⁰ was used as the basis to identify strict forest-
419 dependent endemics (FDE).

420 We also estimated the species-specific body mass for each non-lepidopteran species in our
421 database. The body length of 3-10 individuals of each (morpho)species in an adult stage was
422 measured and species-specific body mass was calculated based on published body length – body
423 mass equations of the higher taxa (data available from Ref.⁵⁰). Since adult and larval biomasses of
424 Lepidoptera differ significantly, we excluded this group from the biomass-based analysis.
425 Lepidoptera, however, were included in species counts and abundance data. To calculate sample-
426 based biomass, we multiplied the number of collected individuals of each species occurring in the
427 samples by their corresponding species-specific body mass value.

428

429 Data analysis

430 We took two different approaches to explore potential arthropod declines in Azorean native forests.

431 First, we compared the changes at a community level and investigated whether there was a change
432 between the first (B1) and second (B2), second and third/last (B3), and the first and third sampling
433 events in any of three calculated measures: abundances, biomasses, and species richness. We used
434 Spearman's correlation test to estimate the interdependence of biomass and abundance values.

435 Individual samples (i.e. one pitfall trap or the material from beating one tree) were pooled within
436 each sampling transect, for each year when sampling was conducted, and the three measures were
437 calculated for each of these pooled samples. Since the length between two sampling campaigns at
438 the same place varied slightly, we did not directly compare the summarised measures from each
439 sampling round but calculated a standardised yearly change in all three measures. To do so, at each
440 site and for each measure, we subtracted the value of the later sampling from the earlier one and
441 divided it by the number of years that passed between the two sampling events. This gave us the
442 absolute yearly change in species richness, abundance, and biomass for each transect. Although this
443 absolute value was used for further analysis, we also calculated the change as a percentage by
444 dividing the absolute yearly change by the corresponding value (i.e. abundance, biomass, or species
445 richness) of the previous sampling event. To estimate the magnitude of changes between sampling
446 events and to test whether they were significant, we calculated Cohen D-s, using the `cohens_d()`
447 function from the `rstatix` R package⁵¹. In-text categorisation of effect sizes (i.e. "negligible", "small",
448 "moderate", and "large") followed that of Cohen⁵². We estimated the confidence intervals (CI)
449 through a bootstrapping process with 10,000 iterations with the help of the `BootES` R package⁵³ and
450 compared the resulting values to a hypothetical zero change (a 'zero change' null model). We ran
451 this analysis by including all data, and separately for subsets including species of each different
452 biogeographic origin. Moreover, data for this analysis were separated by the sampling method. Thus,
453 each of the three measures resulted in eight trend estimations (pitfall trap – all species, pitfall trap –

454 endemic species, pitfall trap – native species, pitfall trap – introduced species, beating – all species,
455 beating – endemic species, beating – native species, beating – introduced species) between two
456 sampling events.

457 As our second approach, we compared the changes at a species level, paying particular attention to
458 the single-island endemics and strict forest-dependent endemics that may carry an extinction debt²⁰.

459 For this, we separated the species in our samples into two major groups: those represented in at
460 least five samples with four individuals and which were present in at least two sampling campaigns,
461 and those that did not fulfil these criteria. For the first group, we used the same analysis as above to
462 estimate whether abundance differences between two sampling events significantly differed from
463 zero. We considered a species unsuitable for including in the analysis ('rare species' henceforth) if
464 they were represented with fewer than 30 individuals across all samples (i.e. fewer than the median
465 number of individuals per species across all samples, when singletons and doubletons are removed),
466 present in less than 1% of samples, or exhibiting scarcity or low presence (less than four individuals)
467 in two entire sampling rounds. Whilst these values are inherently somewhat arbitrary, lower
468 thresholds resulted in comparisons between extremely low abundances (1-2 individuals), or mean
469 values calculated from unreasonably low sample sizes (three individuals) and lowered the power of
470 our analysis. As these 'rare species' were not abundant nor widespread enough to conduct a fully-
471 fledged analysis, we only categorised the differences between sampling events as "no change",
472 "declines", and "increases" based on how summary abundances differed between samplings. Since
473 we wanted to avoid considering very little variability in abundances as changes, unless at least 20%
474 change between two sampling events was observed, we considered the differences as "no change".
475 We conducted a sensitivity analysis to assess how altering this threshold influences our conclusions
476 but, despite some numerical differences, they held until we reached 30%, and even after minor
477 differences were present only in native species SIE/FDE species and (Supplementary Figure S14).
478 Although we did not conduct formal tests on the differences here, we compared whether the

479 occurrence probability of categories (i.e., no change, declines and increases) differs significantly by
480 undertaking a series of Fisher tests.

481 Fisher tests were also used to compare whether the site occupancy of species groups with a
482 different biogeographic origin or conservation status (as defined by IUCN categorisation), has
483 changed. For this, we recorded at how many sites each species occurred in each sampling event and
484 compared them between sampling events. Changes were only considered when they reached a 20%
485 difference. They then were categorised as “no change”, “declines”, and “increases” and Fisher tests
486 were used to assess if site occupancy changes could be predicted from the biogeographic origin or
487 SIE/FDE categorisation.

488 To assess the impact of island identity on the observed changes we removed one, two, and three
489 islands in every combination from the original dataset and recalculated the changes in community
490 measures (i.e., abundance, biomass, and richness) using the Cohen d calculation above.

491 For the sensitivity analysis, we randomly removed 1, 3, 5, 10, and 15 sites from the analysis and
492 recalculated all community-level changes, using Cohen’s d, as above. We repeated each removal
493 round 1000 times and compared the d values visually. All analyses were conducted in an R
494 environment⁵⁴, with the dplyr⁵⁵, ggplot2⁵⁶, and reshape⁵⁷ packages used for data manipulation and
495 plotting.

496 [Data availability](#)

497 The BALA dataset is available on the Global Biodiversity Facility’s (GBIF) website under the
498 <https://doi.org/10.15468/rpdkx9> identifier. Basic information on the sampling sites and summary
499 datasets used in the analysis are available on GP’s GitHub repository
500 (https://github.com/pozsgaig/BALA_diversity).

501 [Code availability](#)

502 All code and data needed to reproduce the results presented in the article and supplementary
503 material are available on the https://github.com/pozsgaig/BALA_diversity GitHub repository.

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658 Author contributions

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669 Competing interests

670 The authors declare no competing interests.