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Remoteness promotes alien species richness on islands

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41

42 **Abstract:**

43 One of the most well-known and general patterns in island biogeography is the decrease in
44 native species richness with isolation, reflecting lower rates of natural dispersal and
45 colonization on remote oceanic islands^{1,2}. During recent centuries, however, a novel, much
46 faster process has increasingly gained importance and altered the composition and richness of
47 island species pools: the human-mediated introduction of alien species³⁻⁷. Analyzing a
48 comprehensive global dataset for alien and native plants, ants, reptiles and mammals on (sub-
49)tropical islands, we found that the number of alien species increases with isolation - a pattern
50 that is opposite to the negative species-isolation relationship (SIR) of native species, and
51 robust across all taxa analyzed. We argue that the reversal of the SIR for alien species is
52 driven by a decrease in the resistance of resident biota to colonization by new species with
53 increasing geographical isolation⁸⁻¹⁰.

54

55 **Main Text:**

56 While the negative SIR for native species is one of the best documented patterns in
57 ecology, it is less clear whether or how the number of alien species on islands is related to
58 isolation. On the one hand, economic theory predicts that fewer commodities are transported
59 to more remote islands¹¹, leading to fewer intentional and accidental alien introductions (i.e.
60 lower propagule pressure), and hence arguably lower colonization rates¹². On the other hand,
61 globalization in trade and transport has considerably reduced the effective isolation worldwide
62 – even of the most remote islands. While natural dispersal to remote islands is extremely rare
63 and has led to the isolation effect in native species, human-aided transport increases the
64 frequency of introduction events by orders of magnitude and the isolation effect might
65 vanish¹. Another line of reasoning suggests that establishment odds of the introduced alien
66 species may be higher on more isolated islands if their impoverished and biologically naïve
67 native biotas provide enhanced ecological opportunities for the introduced species^{8–10}. Taken
68 together, these theories would predict alien species richness on islands to be negatively,
69 positively or uncorrelated with isolation, depending on the trade-offs between colonization
70 pressure and establishment odds. Empirical studies have provided ambiguous results, with
71 negative (for plants and birds¹³), no (for plants¹⁴) or positive (for birds¹⁵, plants¹⁶ and ants
72¹⁷) correlations between alien species and island isolation.

73 Here, we use the most comprehensive datasets currently available of established alien
74 (*sensu* Blackburn et al.¹⁸) and native species numbers on islands to compare the importance of
75 island isolation (i.e., distance to the closest mainland) for native and established alien species
76 richness of vascular plants, ants, reptiles and mammals on subtropical and tropical islands
77 (between 30°N and 30°S latitude; Fig. 1). In our analysis, we account for the effects of other
78 important factors such as island size, climatic and topographic heterogeneity and human
79 impact by using them as additional predictor variables in generalized linear mixed effects
80 models.

81 Across all four taxonomic groups, we found that island isolation has contrasting
82 effects on native and alien species richness. While native species richness decreased with
83 isolation, confirming island-biogeography theory ^{1,2,19}, alien species richness increased with
84 isolation for all four taxonomic groups (only marginally significant for reptiles, Fig. 2 & 3,
85 Table S1, S6). Consequently, when native and alien richness are considered together, we find
86 a marked weakening of the SIRs compared to the pattern for natives only (Fig. 2 & 3, Table
87 S1).

88 The effects of the other predictor variables on species richness were as expected: the
89 numbers of both native and alien species increased with island area (Fig. 3, Table S1).
90 Socioeconomic development (measured as per capita GDP) has a significant positive effect on
91 alien species richness of all taxonomic groups, but it did not affect native species richness
92 (Fig. 3, Table S1). For plants and mammals, per capita GDP was still significant when
93 considering alien and native richness together. Due to the focus on (sub-)tropical islands,
94 climate effects were minor; only native reptile species richness increased with mean annual
95 temperature, and native ant and vascular plant species richness increased with annual
96 precipitation (Fig. 3, Table S1). Finally, alien and native vascular plant and native mammal
97 species richness were positively related to topographic heterogeneity (Fig. 3, Table S1). The
98 robustness of our results was confirmed by a sensitivity analysis that removes potential biases
99 introduced by differences in geographic coverage, sampling intensity and data quality (see
100 Table S2).

101 One possible process behind the positive SIRs for alien species richness is a
102 systematic decrease in the resistance of resident biota to the colonization by new species with
103 increasing geographical isolation. This hypothesis was already formulated by Elton ⁸ and later
104 explicated e.g., by Simberloff ⁹ and Denslow¹⁰. Arguments in favour of this idea emphasize
105 that different resource-use of native and alien species is crucial for successful establishment of
106 the latter ²⁰, and that this divergence likely increases with geographical (and hence commonly

107 evolutionary) isolation. Moreover, particular functional groups, especially large predators and
108 herbivores ²¹, but also pathogens and parasites (e.g., ²²), are generally rarer or absent from
109 remote islands. This leads to reduced predator-escape responses (e.g. island tameness in
110 lizards ²³) and lower resistance to novel parasites in many native island species. As a
111 consequence, introduced predators might have easier access to resident prey, and introduced
112 prey might experience less predation pressure (“enemy release” hypothesis ²⁴). In addition,
113 alien species introduce traits that native island biotas have not been exposed to previously
114 (e.g., allelopathic secondary chemical compounds²⁵) and to which they are naïve (“novel
115 weapons” hypothesis ²⁶), a phenomenon that may increase with isolation as native species
116 become more evolutionarily distinct ²³. Furthermore, as isolated islands usually have a
117 reduced phylogenetic diversity²⁷, the species there might have experienced less competition,
118 and therefore be competitively inferior to alien species from regions with a high phylogenetic
119 diversity (“evolutionary imbalance” hypothesis²⁸). Taken together, these mechanisms may
120 well drive a strong positive correlation between geographical isolation and successful
121 establishment of new arrivals, and hence drive the consistent positive alien species-isolation
122 pattern found in our data.

123 Yet, variation in propagule and colonization pressure might also affect the establishment odds
124 of alien species ¹². In a study on birds ¹⁵, the authors argue that remote islands generally lack
125 native species useful for farming, hunting or aesthetic purposes, which might have led to a
126 greater number of intentional releases of alien birds (i.e., higher colonization pressure),
127 driving a positive SIR. The direct effect of colonization pressure, however, remains difficult
128 to test, as for most taxonomic groups reliable data on introduction events do not exist.
129 Introduction effort is positively correlated with GDP²⁹, and our analyses thus partly corrected
130 for introduction effort by including GDP. Moreover, it seems unlikely that introduction effort
131 (i.e. intentional releases) generally increases with geographic isolation for all tested

132 taxonomic groups (especially for those introduced unintentionally like ants), and thus might
133 have driven the positive SIRs.

134 In conclusion, alien species have markedly changed fundamental biogeographical patterns of
135 island-species richness. The breakdown of biogeographic dispersal barriers, due to human
136 transport, has weakened the classical SIRs. Indeed, the addition of alien species more than
137 halves the effect of isolation on total species numbers. While this pattern has previously been
138 shown for *Anolis* lizards in the Carribean³⁰, we here show that it holds globally for multiple
139 taxonomic groups.

140 Globalization in trade and transport increasingly decouples geographical distance from
141 isolation. As a consequence, immigration rates increase and geographically distant, but no
142 longer isolated islands become packed with species as much as the theory of island
143 biogeography would predict for equal-sized but less isolated islands¹ and may reach a new
144 equilibrium, likely at the expense of many endemic species. However, even if globalization
145 would completely neutralize geographic isolation and natural dispersal barriers, this might
146 explain a weakening of the SIR slopes but not an inversion. Yet, there is a clear congruency of
147 low native diversity and disproportionately high alien species numbers on remote islands. We
148 thus argue that the inverted alien SIR is at least partially driven by a systematic increase in the
149 invasibility due to a decrease in the resistance of resident biota with increasing geographical
150 isolation.

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243 **Methods**

244 **Global island distribution**

245 The dataset comprises a total of 240 islands and island groups (i.e. archipelagos; hereafter
246 also referred to as islands) of oceanic and continental origin with a minimum size of 5 km².
247 We only included subtropical and tropical islands situated between 30°N and 30°S latitudes.
248 Due to the distribution of landmasses across the globe, there are no remote (i.e.
249 geographically isolated) islands in arctic regions. Temperature and isolation are thus closely
250 correlated and would cause a distortion of the isolation effect.

251

252 **Datasets**

253 The dataset differed among taxonomic groups, including 109 islands for vascular plants, 89
254 islands for ants, 129 islands for mammals and 79 islands for reptiles. Species lists of native
255 and established alien species (*sensu* Blackburn¹⁸) were compiled from various sources (Tab.
256 S6). Large data compilations may be affected by biases in data quality and completeness (i.e.
257 varying sampling strategies, differences in taxonomic concepts;^{31,32}). To address these issues,
258 we compiled complete species lists where available based on recent database projects that
259 ensure taxonomic standardization (e.g. using the Plant List for vascular plants;³³).
260 Furthermore, for all other islands where only richness values were available the most up-to-
261 date sources were used, assuming that these sources used a recent taxonomic concept so that
262 biases can largely be excluded.

263 Potential effects of variation in data reliability were tested using a sensitivity analysis
264 (see below). Each island was assigned to a geographic region following the Biodiversity
265 Information Standards (TDWG) classification³⁴ (see Tab. S4). For all islands, we compiled
266 eight predictor variables which represented socio-economic (human population density, per
267 capita gross domestic product), climatic (mean annual temperature, annual precipitation sum)
268 and geographic (island area, elevational range and distance to mainland) variables. Distance

269 to mainland was calculated as the shortest geodesic distance to a continent, excluding
270 Antarctica. The geographical distance is just one metric and ocean currents, winds and the
271 richness of source regions also influence immigration rates for native species ². However,
272 these additional variables are less relevant for aliens as they are introduced through human
273 transport, and so we decided to use geographical distance only. Island area and elevational
274 ranges were calculated for each island and island group. In the case of island groups the
275 cumulative terrestrial surface area of all relevant islands was used. Island area ranged from
276 5.11 km² to 110,730 km², with a median size of 280 km². Data on current climate for each
277 region were derived from WorldClim 2.0 ³⁵). Finally, human population density was derived
278 from the HYDE database ³⁶, and per capita gross domestic product (GDP) from Gennaioli ³⁷,
279 Worldbank ³⁸ and the United Nations ³⁹ (Tab. S5).

280

281 **Statistical analysis**

282 We analyzed the dependence of alien and native species richness (species numbers) on
283 distance to mainland, island area, elevational range, mean annual temperature, annual
284 precipitation sum, GDP and human population density as predictor variables by means of
285 generalized linear mixed effects models (GLMMs) with a Poisson-distributed response
286 (species richness) and the canonical log link function. Human population density, a frequently
287 used surrogate of human impact (e.g. ^{13,14}), was never significant and was thus excluded from
288 the analyses. A random effect intercept term with TDWG 4 region as grouping factor
289 acknowledged political/socio-economic groupings among regions, and a random effect
290 intercept term for island geologic setting (i.e. oceanic islands vs. islands situated on
291 continental shelves ⁴⁰) accounted for possible differences in colonization due to historic
292 connections with continents ². Finally, an additional observation-level random effect term
293 accounted for overdispersion ⁴¹. To improve symmetry, linearity, and to stabilize variances,
294 numerical predictors were subjected to appropriate transformations (natural log for island

295 area, elevational range, distance to mainland; square root for precipitation sum and per capita
296 GDP), and finally standardized. The magnitude of regression coefficients was hence
297 representative of relative effect size. We fitted individual models for alien, native and total
298 (alien plus native) species numbers for every taxonomic group. Model residuals were assessed
299 for spatial autocorrelation by spline (cross-) correlograms, and no spatial-autocorrelation was
300 found (Fig. S1 & S2).

301 All statistical analyses were performed using R (version 3.3.1). For GLMM analyses, we used
302 the function *glmer()* from the package lme4 for fitting⁴² and the function *effect()* from the
303 package effects for partial effect plots. For spline corellograms, we used the function
304 *spline.correlog()* from the package ncf⁴³.

305

306 **Sensitivity analysis**

307 To test the robustness of the assessed relationships between alien species richness and island
308 isolation, we performed a sensitivity analysis. The aim of this analysis was to exclude
309 systematic biases in the data that might stem from heterogeneous sampling intensity or
310 overrepresentation of selected geographical regions, as well as from variable data quality
311 depending on data sources. Therefore, we first systematically excluded islands of a
312 geographic region (based on TDWG level 2 classifications) from the datasets. Then, the
313 number of excluded islands was resampled from the remaining islands to ensure constant
314 sample sizes. Subsequently, we fitted the same GLMMs as were used for the main analysis to
315 the resampled datasets. This procedure was repeated 500 times and confidence intervals were
316 calculated for the regression coefficients and p-values (Table S2). Similarly, we excluded
317 some less reliable data sources, resampled from the remaining islands and recalculated the
318 models.

319

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330 PhyloPic website (www.phylopic.org). Image credit for *Platanus occidentalis* L. goes to
331 Michele M Tobias (<http://phylopic.org/image/806a6ae9-28a0-4dc6-beeb-f6129a44f10e/>). No
332 changes were applied to the pictures. All other images are under no copyright and free to use.

333

334 **List of supplementary material:**

335

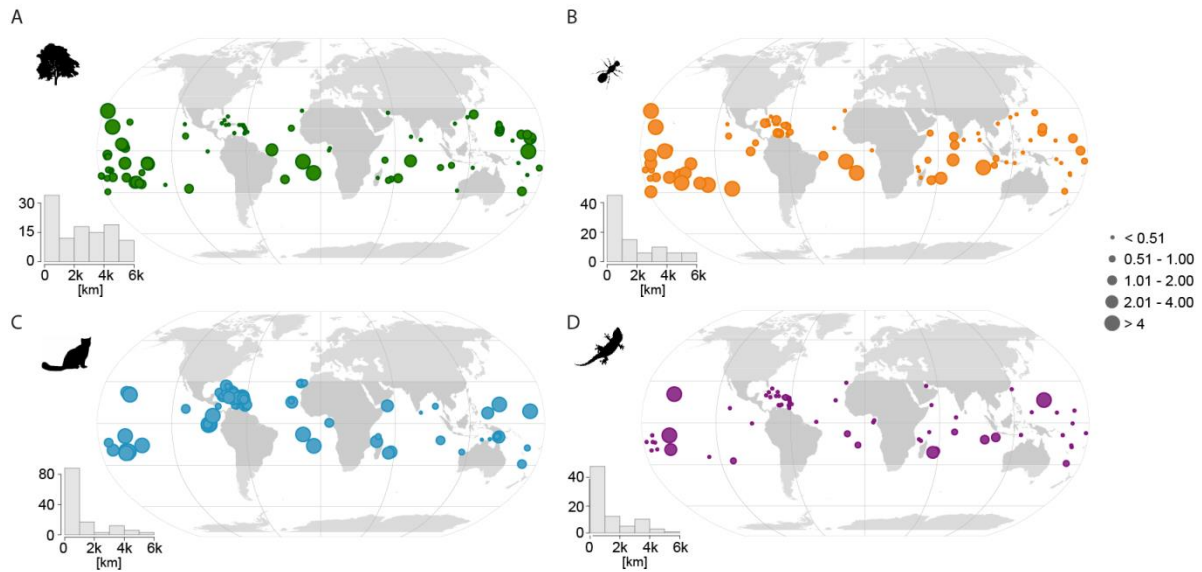
336 Supporting Information

337 Table S1 – S6

338 Fig S1 – S2

339 References

340



341

342 **Fig. 1. Geographic distribution of tropical and subtropical islands used in the study for**

343 **(A) vascular plants, (B) ants, (C) mammals and (D) reptiles.** Symbol size scales with ratios

344 of established aliens in relation to native species. The histograms show the frequency

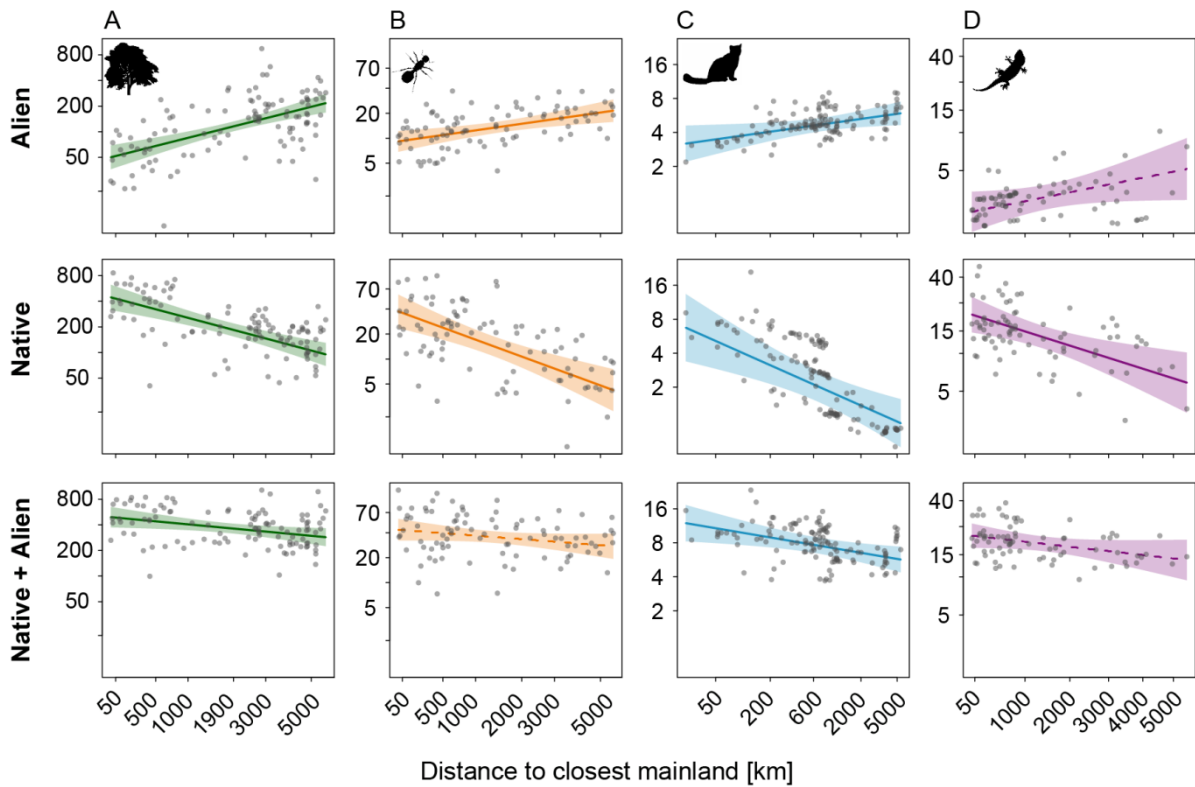
345 distributions of island distance to mainland for the four taxonomic groups. The number of

346 islands included in the analysis differs among the taxonomic groups (vascular plants = 109;

347 ants = 89; mammals = 129; reptiles = 79). Pictograms for the taxonomic groups are taken

348 from www.phylopic.org.

349



351

352 **Fig. 2. Alien and native species richness on islands dependent on island isolation for (A)**353 **vascular plants, (B) ants, (C) mammals and (D) reptiles.** Shown are partial residual plots354 of the species richness-isolation relationships for established alien (1st row), native (2nd row)355 and total (3rd row) species richness (log-log space). Generalized linear mixed effects models

356 with a Poisson-distributed response were applied to additionally account for island size,

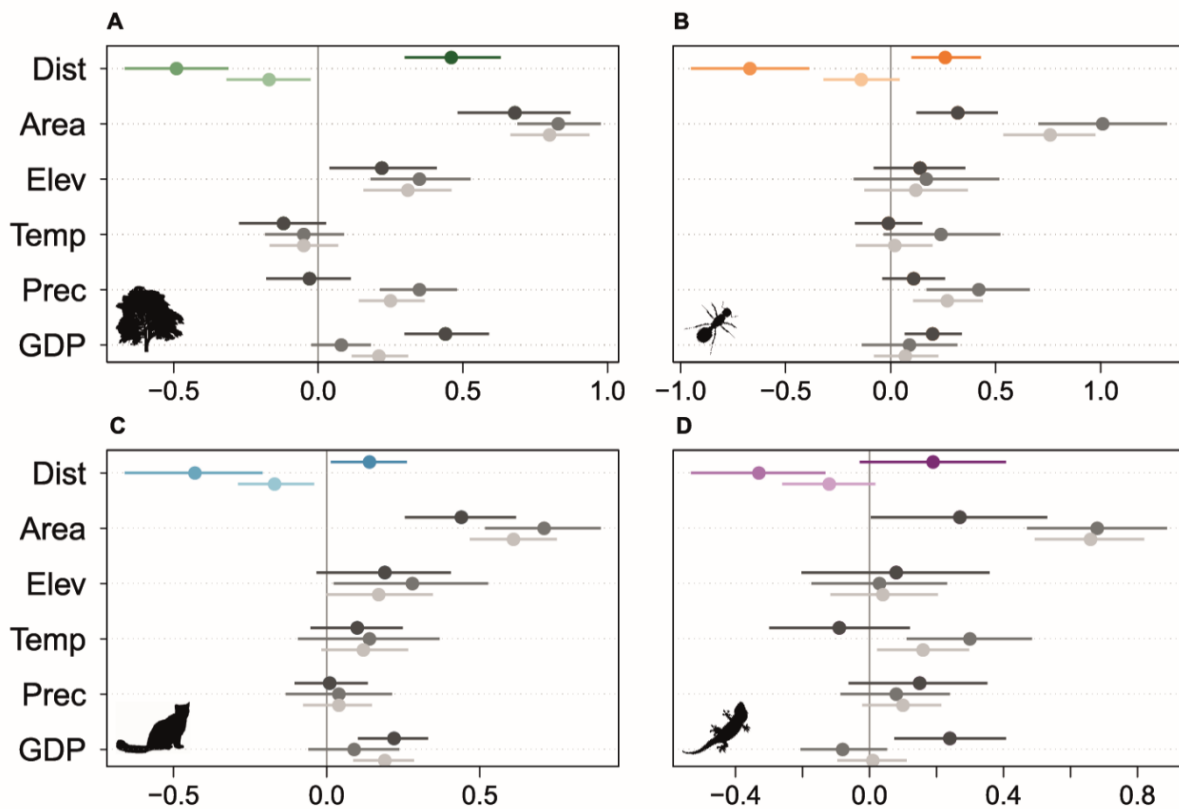
357 heterogeneity (elevational range) climate (temperature, precipitation) and human impact (per

358 capita GDP). Each column represents one taxonomic group. Shading around the regression

359 line indicates its 95% confidence interval. Dashed lines indicate insignificant results.

360 Pictograms for the taxonomic groups are taken from www.phylopic.org.

361



363

364 **Fig. 3. Regression coefficients and 95% confidence limits for the standardized predictor**365 **variables in the generalized linear mixed effects models for (A) vascular plants, (B) ants,**366 **(C) mammals and (D) reptiles.** Dark colors represent the estimates for established alien

367 species, medium colors for native species and light colors for all species. Abbreviations are:

368 Area: island area; Elev: elevational range; Dist: Distance to the closest mainland; Temp: mean

369 annual temperature; Prec: annual precipitation sum; GDP: per capita GDP (for the full model

370 output see Table S1A). Pictograms for the taxonomic groups are taken from

371 www.phylopic.org.

372