ORIGINAL RESEARCH



Invasive black rats menacing endangered lizards

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Abstract

Despite the pernicious impacts that invasive black rats *Rattus rattus* have on island ecosystems, little is known about their effect upon insular reptiles, which are a highly vulnerable but pivotal element of island biota. To bring to light these effects, we evaluated the threat posed by *R. rattus* on the critically endangered Canarian spotted lizard *Gallotia intermedia* by analyzing its frequency of occurrence on rat feces, estimating rat abundance and density, and correlating these parameters with previous lizard censuses. We genetically detected that 14.96% of all rat feces contained *G. intermedia*, with 27.27% of individual *R. rattus* consuming this lizard. Rat density varied from 0.740 ± 0.474 to 2.183 ± 1.137 rats/ha and was correlated with larger declines of *G. intermedia* between past censuses and those of 2019. These results confirm for the first time that *R. rattus* consumes and impacts this endemic and endangered lizard species. From a broader perspective, this is one of the first studies detecting rat impact on a large-sized reptile, which calls for further attention to the interaction between invasive rats and a highly vulnerable but essential component of island ecosystems.

Keywords Canary islands · Conservation genetics · *Gallotia intermedia* · Rat density · Spatially explicit capture-recapture

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Introduction

The invasive black rat *Rattus rattus* is a major threat to global biodiversity (Doherty et al. 2016; Bellard et al. 2017; Dueñas et al. 2021) and one of the most pernicious invaders worldwide (Lowe et al. 2000). On islands, they affect native and endemic flora (Pender et al. 2013; Harper and Bunbury 2015), prey on invertebrates (St Clair 2011), cause native vertebrates' decline and extinction (Towns et al. 2006; Jones et al. 2008; Harris 2009), and eventually trigger cascading ecological effects (Harper and Bunbury 2015; Thoresen et al. 2017). However, experimental evidence on *R. rattus* impacts is still scarce and often relies on its trophic habit studies (Banks and Hughes 2012). That is why *R. rattus* impact upon insular reptiles has been generally assumed to be irrelevant, since its diet is mainly based upon plant material, insects, arachnids, and birds (Banks and Hughes 2012; Shiels et al. 2014), but scarcely on reptiles (Shiels et al. 2014). Nevertheless, it is relevant to highlight that *R. rattus* has been linked to reptile population declines in island ecosystems worldwide (Towns et al. 2006; Smith et al. 2012; Harper and Bunbury 2015), thus understanding the potential impacts of *R. rattus* is needed to properly plan reptile conservation.

More than one-fifth of all reptile species worldwide are threatened by extinction (Cox et al. 2022), with oceanic islands as hotspots of threatened species (Böhm et al. 2013). Invasive mammals have highly contributed to this situation (Böhm et al. 2013; Doherty et al. 2016; Cox et al. 2022), threatening nearly 20% of all reptiles (Bellard et al. 2017). Island reptile conservation is a priority as they have a key ecological role on island ecosystems—e.g., they sustain native plant pollination and seed dispersal interactions (Olesen and Valido 2003) and are cornerstone in trophic networks (e.g., Schoener and Spiller 2010; Piovia-Scott et al. 2019)—and impacts upon them can have other profound effects. For conservation planning purposes, unraveling menaces to island reptiles is essential to define management actions for both reptiles and invasive species. However, this is not a trivial task; threatened reptiles on islands can be relegated to remote areas with difficult access, often with food shortages, where other invasive species can mask the particular impact of *R. rattus*.

The Canarian spotted lizard Gallotia intermedia is a large-sized, critically endangered and endemic reptile of Tenerife (Canary Islands) (Mateo-Miras et al. 2009; BOC 2017), that reaches 15 cm of snout-vent length and over 40 cm of total length (Lopez-Darias et al. 2015; Padilla et al. 2019). Listed among the most endangered reptiles of Europe (Cox and Temple 2009), G. intermedia was believed to be extinct until its rediscovery in 1996 (Hernández et al. 2000)— although its distribution was seemingly larger in the past (Rando 2002; BOC 2017; Padilla et al. 2021). The species currently persists only in small and isolated populations (<500 individuals each) dispersed in two almost inaccessible sea cliffs of the island (Padilla et al. 2019, 2021). Ecological information on the species is mostly lacking, but considering fruits compose over 90% of G. intermedia female diet (Lopez-Darias et al. 2015), it is very likely that it participates in native seed dispersal dynamics similar to other Gallotia species (Pérez-Méndez et al. 2016, 2017). Additionally, as it also consumes invertebrates, it presumably will have an important role in trophic networks, as it has been observed for other giant Gallotia species (Piquet 2022). In 2017, the Canarian Government approved the G. intermedia Recovery Plan, designating six critical areas of conservation and listing a series of priority conservation tasks. Among these tasks, the understanding of the potential effects of rats on G. intermedia population is a priority, as they have been suggested to prey upon eggs and juveniles of endemic lizard species of the archipelago (Rando and López 2001; Traveset et al. 2009; Padilla et al. 2019, 2021). However, no empirical study has yet confirmed this predatory-prey interaction for *G. intermedia*. To fill this gap, we quantified the occurrence of *G. intermedia* in *R. rattus* feces using genetic techniques as a proxy for *R. rattus* consumption (one of the main objectives of *G. intermedia* Recovery Plan; BOC 2017) and estimated rat abundance and density in the cliff area where the species is distributed. We then crosschecked this information with the most recent estimates of *G. intermedia* abundances (Padilla et al. 2019) to further explore the threat posed by *R. rattus* upon this endangered lizard. Our results will directly contribute to design conservation actions for *G. intermedia*, while providing further insights on the effect of invasive rats on island reptiles, an impact that remains largely unexplored worldwide.

Methods

Study area and R. rattus trapping sessions

We designated seven study sites on the two sea cliff areas of Tenerife (Canary Islands, Spain) where *G. intermedia* is present: Teno Rural Park and Natural Monument of Guaza Mountain (Fig. 1) (hereafter Teno and Guaza, respectively). These cliffs are characterized by sparse vegetation from native semi-arid and arid plant communities interspersed with invasive plants (BOC 2017), and present annual mean temperatures of 17.5–20 °C and less than 400 mm rainfall per year (AEMET 2024). To optimize resources and given that most of the sites are only accessible by boat and using climbing techniques, we selected those sites where lizards were most abundant following Padilla et al. (2019), ruling out those excessively difficult to access (Fig. 1).

We trapped *R. rattus* to collect their feces and calculate their density in all sites, by performing three trapping sessions in each site in 2020 (13th July to 3rd August, 31st August to 29th September, and 26th October to 9th November), coinciding with lizards' active period (Rando and Valido 2000; Padilla et al. 2019). We established all trapping sessions on days with favorable weather conditions. Each session consisted of trapping during three consecutive nights, for nine nights per site, using 37 traps distributed among sites (14.89 ± 3.88) traps per site and night). The total number of traps was set to efficiently cover key sites of the species distribution, ensuring manageability given the available human resources and logistics needed to sample these areas. We initially worked with 20 Tomahawk traps (40.64×12.7×12.7 cm) (Tomahawk Live Trap Co., Tomahawk, Wisconsin, USA), 10 FIX-MAN traps $(25 \times 9 \times 9 \text{ cm})$ (FIXMAN Ltd., Bishop's Stortford, UK), and 7 Sherman folding traps $(30.5 \times 9.5 \times 7.5 \text{ cm})$ (H.B. Sherman Traps Inc., Tallahassee, FL, USA). However, since FIXMAN traps produced tail autotomy in two of the first G. intermedia bycatches, we replaced these with other 10 Tomahawk from the first session onwards. Once installed, we georeferenced all traps, which stayed in the same location for the whole session, and baited them with a mixture of flour and canned sardines, previously shown as attractive for *Rat*tus spp. (authors own data). To minimize non-target and maximize R. rattus captures, we activated traps only during night (activating them at dusk and de-activated them at dawn). We always used latex gloves to handle traps and baits to avoid leaving any human odor that could deter rat captures.



Fig. 1 Location of the Canary Islands, the study areas, and sampling sites, showing *Rattus rattus* density (black circle) and *Gallotia intermedia* frequency of occurrence (FO) in rats' diet. FO was calculated as the number of fecal samples in which the species was detected divided by the total number of fecal samples (FO_{exc}; panel A) and the number of rats that consumed *G. intermedia* divided by the total number of rats captured (FO_{ind}; panel B) in each sampling site. Protected natural areas are shown in each panel as dark grey areas. Map created using QGIS 3.36

We checked all traps at dawn and noted down those that were closed, empty, or captured *R. rattus* or non-target species (these latter were immediately released). We transferred captured *R. rattus* to a handling bag (Koprowski 2002), retrieved fresh fecal pellets from the trap using sterile tweezers to increase subsequent DNA detection and minimize contamination (Ando et al. 2020), and dropped samples in tubes with 100% ethanol, labelled with a specific code for each individual (we never simultaneously captured two individuals in a single trap). We preserved all samples at -20 °C once in the lab until later analysis. We identified all captured *R. rattus* using ear-tags (#1005 Size 1 Monel, National Band and Tag Co., Newport, Kentucky, USA) or ear punching (Chen et al. 2016), determined their sex and age (adult vs. juvenile), and released them in the exact same location within a few minutes of handling. We also noted down all individuals that escaped from the trap or during handling and also collected and individually identified any fecal material dropped by them in the trap. We thoroughly cleaned all traps after each capture.

All necessary authorizations to perform this study were available from the competent administration (Cabildo de Tenerife no. AFF 87/20).

Occurrence of G. intermedia in R. rattus feces

After air-drying and homogenizing each fecal sample with a sterile scalpel, we retained 200 mg to extract DNA using the E.Z.N.A. Stool DNA Kit (Omega Bio-tek, Inc. Norcross, Georgia, USA), following manufacturer instructions. We quantified the amount of DNA using NanoDrop ND-1000 spectrophotometer (Thermo Fisher Scientific, Waltham, MA, USA). To detect the occurrence of G. intermedia in each fecal sample, we amplified a 97 bp-sequence of the mitochondrial 12 S rRNA using the primers L1064 5'-TTGACC ACACGAAAGCTTAGA-3' (Maca-Meyer et al. 2003) and Int12SR2 5'-CTGGCGGAAA GTATTGTATTCA-3' (M. Hernández, in litt.), this latter previously designed to differentiate G. intermedia from the sympatric species G. galloti. We performed DNA amplification using 20 µL PCR mixes that contained 4 µL of MyTaq 5X tampon (Bioline, London, UK), 0.15 µL of MyTaq DNA polymerase (Bioline), 0.8 µM of each primer, 0.8 µg/µl of BSA (New England Biolabs), and 10–20 ng of DNA. After optimizing the PCR protocol using small pieces of tails and fecal samples from G. intermedia and G. galloti, PCR conditions consisted of an initial step at 95 °C for 1 min followed by 35 cycles of denaturation for 15 s at 95 °C, annealing for 15 s at 54 °C, and elongation for 10 s at 72 °C, with a final extension step at 72 °C for 5 min. We carried out all reactions in a BioRad T-100 thermocycler (Bio Rad Laboratories, Hercules, California, USA), and included in each of them a positive and a negative control. To minimize contamination, we performed the entire protocol under a laminar-flow bench and using filtered pipette tips. We visualized PCR products in 1.7% agarose gel stained with RealSafe (Real Biotech Corporation, Taiwan), using a 25-700 bp DNA marker (LADD-DN1-500), and subsequently purified and sequenced the sample at the University of La Laguna Genomic Service (SEGAI). We compared DNA sequences with those available in GenBank for G. intermedia, which we extracted using BLAST+2.10.1. (Camacho et al. 2009), and aligned using Clustal W in MEGA X (Kumar et al. 2018). We introduced minor corrections by hand and calculated the frequency of occurrence of G. intermedia as a quotient between the number of feces where G. intermedia was detected and the number of rat feces collected (hereafter FO_{exc}), following previous studies on rat diet (e.g., Caut et al. 2008; Zarzoso-Lacoste et al. 2016). Additionally, to provide a better

description of rat impacts and identify the proportion of individuals contributing to them, we calculated the proportion of *R. rattus* individuals consuming *G. intermedia* from the total of captured rats (including all dead and escaped ones) (hereafter FO_{ind}).

Rattus rattus abundance and density

Following previous research (Fukasawa et al. 2013; Maggs et al. 2015), we calculated R. rattus captures per unit of effort (CPUE) as a proxy of the species abundance in each site using the total number of rats captured divided by the number of active traps per night—i.e., after removing sprung traps and those with by-catches. We calculated CPUE by pooling the data of all traps, as we set a similar number of traps of each type every trapping night in all sites (data not shown). Additionally, we estimated R. rattus density by analyzing capture data and trap location with spatially explicit capture-recapture models performed in secr package (Efford 2023). We divided the capture dataset into seven sessions (one per site) and nine occasions (one per night), and set the detector argument to 'single'. We determined the most appropriate buffer width using 'esa.plot' function and retained its value for subsequent analyses. We modeled density using a half normal detection function, assuming that detection occurred over a constant spatial scale (s) (Piquet and López-Darias 2021), estimating detection probability (g_0) according to trap type and maximizing the conditional likelihood. We used Akaike Information Criterion (AIC) and AIC corrected for small sample sizes (Burnham and Anderson 2002) to evaluate the better fit of an alternative model testing for the occurrence of learned responses affecting g_0 , without finding any support for it (data not shown). We retrieved R. rattus density in each sampling site as Horvitz-Thompson-like estimate following Borchers and Efford (2008) and compared it between Teno and Guaza using Kruskal-Wallis tests. Finally, we evaluated the correlation of rat density with R. rattus abundance, and G. intermedia FOexc and FOind in each site using Spearman rank correlation tests, which are robust to violations of bivariate normality assumptions, non-linear relationships between paired variables and low sample sizes (Bonett and Wright 2000; Quinn and Keough 2002).

Rattus rattus impact on G. Intermedia populations

We performed Spearman rank correlation tests to analyze the relationship of both FO_{exc} and rat density with the following demographic values in each site, extracted from the last available *G. intermedia* census (Padilla et al. 2019) (see Table 1 for values): (1) the total number of *G. intermedia* individuals, (2) *G. intermedia* density, (3) the number of non-reproductive and reproductive *G. intermedia*—i.e., juvenile and average-sized individuals, and sub-adults and adults in Padilla et al. (2019), respectively— (4) and the lizard population trend in each site, calculated as:

$$\frac{\mathrm{N}-\mathrm{P}}{\mathrm{P}}\mathrm{X}\,100\tag{1}$$

where N and P are the number of G. *intermedia* detected in each site in the present and the past, respectively.

 Table 1 Number of Gallotia intermedia individuals [Number], density [Density] (lizards/ha), number of non-reproductive [Non-reproductive] and reproductive [Reproductive] G. intermedia—i.e., juvenile and average-sized individuals, and sub-adults and adults in Padilla et al. (2019), respectively—and lizard population trend (expressed as percentage) [Population trend]—i.e., differential between the current number of lizards and that in the first census performed in each site, divided by the latter—in each sampling site. All data were directly extracted from the last G. intermedia census performed in each site in 2019 or derived from population estimates obtained during these surveys (see Padilla et al. 2019)

Area	Site	Number	Density	Non-reproductive	Reproductive	Population trend
Teno	Punta de Diente de Ajo	16	36.53	7	9	166.67
	Andén de la Amargosa	26	189.23	7	19	550.00
	Andén de las Cañas	31	59.88	9	22	-8.82
	La Galera	51	41.33	16	34	75.86
	Punta del Gigante	21	70.52	1	20	-30.00
Guaza	Andén de los Ñames	23	124.66	7	14	-52.08
	Andén del Parasol	13	75.85	7	5	-71.11

We performed all analyses using R 4.3.3. (R Core Team 2024) (R code file available under https://doi.org/10.6084/m9.figshare.20455530.v1). Results correspond to mean±SD, unless stated otherwise.

Results

Captures and sample collection

We captured *R. rattus* on 123 occasions and collected 127 feces—three belonging to animals that entered the traps but were not captured and another one collected in the vicinity of a trap. We trapped 55 different individuals on all these occasions, three of which deceased during handling and were marked and other three escaped unmarked (Table 2). Therefore, from all trapping occasions, 120 captures corresponded to marked individuals (97.56% of all captures). We captured *R. rattus* on 46, 45, and 29 occasions in the first, second, and third trapping sessions, respectively. From these, we captured adult females, adult males and juveniles on 49, 43, and 28 occasions, respectively. We recaptured marked individuals an average of 2.31 ± 1.58 times, up to a maximum of 7 recaptures for a single individual (Table 2). Recaptures occurred within the same site where rats were originally captured. Non-target captures included one house mouse *Mus musculus domesticus* and 11 *G. intermedia*.

Occurrence of G. intermedia in rodent feces

We detected *G. intermedia* in 19 fecal samples (14.96% of all samples), belonging to 15 different individual rats (27.27% of captured individuals). All detections had 100% similarity with sequences from GenBank for *G. intermedia* (AY154913, AY151923) and with one individual of *G. simonyi* (Z79499). The sequence of 48 bp found between primers allowed us to differentiate between *G. intermedia* and the sympatric *G. galloti* lizard. Both juvenile and adult *R. rattus* consumed *G. intermedia* (seven and nine detections in female and male feces, respectively, and three in juveniles). The number of detections per site and individuals respectively varied from 1 to 6 and 1 to 3 for the total number of positive feces (FO_{exc}: 5.26

Table 2 Total number of *Rattus rattus* capture occasions (including dead and escaped individuals) [C], number of marked (including dead but not escaped individuals), dead and escaped individuals [M/D/E], total number of recaptures in each site [R], and number of feces collected in each site, positive feces and number of *R. rattus* individuals in which *Gallotia intermedia* was detected [F/P/I]. FO_{exc} refers to *G. intermedia* frequency of occurrence in *R. rattus* feces, calculated as a quotient between the number of feces where *G. intermedia* was detected and the number of feces collected. FO_{ind} represents the number of individuals of *R. rattus* that consumed *G. intermedia* from the total of individuals captured (including all dead and escaped ones). Captures per unit of effort (CPUE) (averaged across the three consecutive days of each session) is shown as mean ±SD for each site. Density (rats/ha) estimates correspond to density ±SE and are shown together with the relative standard error (RSE), defined in Efford and Fewster (2013)

Area	Site	С	M/D/E	R	F/P/I	FO _{exc}	FO _{ind}	CPUE	Density	RSE
Teno	Punta de Diente de Ajo	8	5/1/0	5	8/2/2	25.00	40.00	0.040 ± 0.075	1.082 ± 0.662	0.61
	Andén de la Amargosa	7	4/1/0	5	7/3/3	42.86	75.00	0.040 ± 0.065	0.740 ± 0.474	0.64
	Andén de las Cañas	25	7/0/1	23	28/3/3	10.71	37.50	0.159 ± 0.182	1.599 ± 0.880	0.55
	La Galera	18	7/1/0	15	19/1/1	5.26	14.29	0.114 ± 0.136	1.375 ± 0.732	0.53
	Punta del Gigante	16	8/0/1	11	16/2/1	12.50	11.11	0.084 ± 0.123	1.687 ± 0.884	0.52
Guaza	Andén de los Ñames	21	10/0/1	15	21/6/3	28.57	27.27	0.158 ± 0.163	2.183 ± 1.137	0.52
	Andén del Parasol	28	11/0/0	23	28/2/2	7.14	18.18	0.119 ± 0.181	2.147 ± 1.104	0.51

-42.86% [min-max]; FO_{ind}: 11.11 – 75.00% [min-max]) (Table 2). *Rattus rattus* consumed *G. intermedia* all throughout the study season, with seven detections in feces during the first session and six during the second and third sessions, respectively (six, five, and six individuals, respectively). We confirmed that *R. rattus* consumed *G. intermedia* both in Teno (11 positive feces and 10 individuals) and Guaza (8 positive feces and 5 individuals) (Table 2).

Rattus rattus abundance and density

Rattus rattus CPUE ranged from 0.04 ± 0.08 to 0.16 ± 0.18 captures/trap-night (Table 2), with an average of 0.10 ± 0.14 captures/trap-night for all sites and trapping nights together. Average density was 1.54 ± 0.53 rats/ha, with sites located in Teno and Guaza showing similar values (χ_1^2 = 3.75, P=0.053) (Table 2)—the highest value was found in Andén de los Ñames (Guaza) (2.18 rats/ha; Table 2). Relative standard errors in all cases exceeded 0.50 (Table 2). We found no significant correlation between *R. rattus* density and rat abundance, lizard FO_{exe} nor lizard FO_{ind} (P>0.05 in all cases).

Rattus rattus impact on G. Intermedia populations

We found that rat density was negatively correlated to the lizard population trend ($r_S = -0.96$; P = 0.003). No other demographic parameter was correlated to rat density nor FO_{exc} (P > 0.05 in all cases).

Discussion

Our results constitute a clear indication of *R. rattus* consuming the critically endangered *G.* intermedia. Although invasive rats have been suggested as predators of the Canary Islands endemic lizards (Traveset et al. 2009; Padilla et al. 2019), this is the first empirical evidence confirming this predatory interaction. In light of these results and taking into account the ubiquity of *R. rattus* in the archipelago (Nogales et al. 2006) and its broad feeding spectrum (Shiels et al. 2014; Harper and Bunbury 2015; Pomeda-Gutiérrez et al. 2021), the consumption of other endemic lizards within the *Gallotia* genus, including other highly endangered ones (Cox and Temple 2009), seems highly plausible. The critical status of some endemic lizards of the Canary Islands has been frequently attributed to the impact of feral cats (García-Márquez et al. 1997; Nogales et al. 2006; Medina and Nogales 2009); consequently, management efforts within lizards' distribution areas typically target this invasive species (e.g., Rando and López 2001; Rando 2005). However, our results indicate that R. rattus may have also contributed to the demise of these lizards, and might currently represent an additional obstacle to their recovery. Implementing control actions without considering species interactions can lead to unforeseen and highly detrimental consequences (Rayner et al. 2007; Bergstrom et al. 2009; Ringler et al. 2015). For instance, the initial eradication of cats in Little Barrier Island led to a surge in predation pressure by Pacific rats (R. exulans) and the decline in endemic bird breeding success (Rayner et al. 2007). In the particular case of G. intermedia, and given its critical conservation status (BOC 2017), we suggest promptly undertaking conservation actions to control both feral cats and R. rattus simultaneously. From a broader perspective, this research represents one of the few empirical examples worldwide of *R. rattus* potentially impacting a large-sized, endemic reptile (Towns et al. 2006; Hayes et al. 2012; Harper and Bunbury 2015), thus calling for further attention to these vulnerable but essential components of island ecosystems.

We detected a high consumption rate of G. intermedia by R. rattus, as the endemic lizard occurred in 14.96% of all samples collected and in all seasons, and in 27.27% of all individual rats captured. Black rats are usually reported to mainly feed on plants and invertebrates, whereas lizard consumption is often low or even completely lacking (Caut et al. 2008; Shiels et al. 2014; Riofrío-Lazo and Páez-Rosas 2015; Pomeda-Gutiérrez et al. 2021). Caut et al. (2008), however, reported the consumption of small-sized Strand litter skink Caledoniscincus haplorhinus, which appeared seasonally in 13% of all gut samples on Surprise Island, a lower FO than that observed in Tenerife for G. intermedia. However, R. rattus diet studies have been mainly performed through gut or intestinal analysis and stable isotopes (Riofrío-Lazo and Páez-Rosas 2015; Clapperton et al. 2019; Nascimento et al. 2019), so a better understanding of reptile consumption might be reached by using genetic techniques, as they increase the likelihood of prey detection compared to other methods (Farrell et al. 2000; Egeter et al. 2015; Zarzoso-Lacoste et al. 2016; Monterroso et al. 2019). Although the use of these techniques has allowed us to confirm G. intermedia consumption, a real quantification of this interaction is difficult as environmental, biological, and methodological factors can affect DNA quantity or quality in samples (Monterroso et al. 2019)—e.g., prey material and DNA can also degrade within hours, preventing accurate prey detection in fecal samples (Egeter et al. 2015). Our estimates do not allow to precisely extrapolate the number of G. intermedia individuals actually consumed, as occurrence could inform about the consumption of one or multiple individuals by the same or different rats (Egeter et al. 2015; Zarzoso-Lacoste et al. 2016). The next steps of this research should try to elucidate whether the detected consumption is reflecting direct predation or could be due to a potential scavenging trophic behavior of *R. rattus* upon lizard carrion or autotomized tails. However, in the context of these results and the critical conservation status of *G. intermedia* (BOC 2017), the precautionary principle (IUCN 2018) dictates that *R. rattus* must be treated as an additional threat to this endemic lizard.

The population density of R. rattus in areas occupied by G. intermedia, reaching 1.54 rats/ha (ranging 0.74–2.18 rats/ha), is lower than that observed in more productive ecosystems like tropical islands (Uchida 1966; Shiels et al. 2014; Harper and Bunbury 2015; Russell et al. 2018) but comparable to, or even exceeds, that found in other areas of low productivity (Clark 1980; Shenbrot et al. 2010; Harper et al. 2011). These low estimates are consistent with what is expected (Clark 1980; Russell et al. 2011; Harper and Bunbury 2015; Carpenter et al. 2022) for an area receiving limited rainfall (AEMET 2024), presenting sparse vegetation, and hosting populations of rat predators (feral cats F. catus) and competitors (M. m. domesticus) (Rando and Valido 2000; BOC 2017; Padilla et al. 2019; Flores Ravelo and Rando Reyes 2021). However, our estimates should be interpreted as the lowest density attained by R. rattus in these areas, as this parameter may substantially increase with greater resource availability after seasonal rains (Clark 1980; Harper et al. 2015; Goedert et al. 2020; Carpenter et al. 2022). Rattus rattus persisting or even thriving in these areas points to the existence of alternative food resources, like invertebrates and seabird chicks and eggs, both of which are commonly consumed elsewhere (Jones et al. 2008; St Clair 2011) (we detected some algae in some R. rattus feces). The high G. intermedia consumption reported here also suggests this species could contribute to sustaining rat populations, although the fact that lizard consumption is unrelated to rat density could be indicative of G. intermedia not being the primary food source. On the other hand, since our density estimates had a degree of uncertainty exceeding the threshold proposed by Efford (2011), these should be interpreted as a proxy of R. rattus abundance—a cost-efficient and useful alternative to precise density calculations (Stephens et al. 2015) that would be extremely labor-intensive, prohibitive, and even dangerous given our study system. Large density uncertainty could explain the lack of relationship between rat density and CPUE, although this finding could also be a consequence of spatially-explicit density estimates considering trap spacing and rat movements (Efford 2004) or variable capture probability affecting CPUE indices (Wiewel et al. 2009). Sites with higher rat density tended however to present greater rat abundance, so that a positive relationship between both parameters may arise under larger sample sizes. Overall, this study provides the first estimation of rat density in Tenerife and the Canary Islands, although rat abundances have already been reported in other habitats of the island (Hernández Martín 1997; Godoy Betancor 2000). More importantly, our results provide baseline estimates of rat abundance and density in areas occupied by G. intermedia, which are essential to managing the impact of invasive species (IUCN 2018).

We found a correlation between the density of *R. rattus* and the decline of *G. intermedia* populations, suggesting the existence of complex rat-lizard interactions in areas occupied by the endangered lizard. Our results could be explained by direct *R. rattus* predation (expected under the high FO detected) or resource competition between the invasive species and *G. intermedia*, as both species are mainly omnivorous (Rando 2002; Harper and Bunbury 2015; Pomeda-Gutiérrez et al. 2021). Alternatively, the impact here detected could derive from *R. rattus* subsidizing cat populations in the area (Rando and López 2001; Padilla et al.

2019), which would increase predation pressure on the endangered reptile (Courchamp et al. 2000, 2003). This situation could be aggravated by the presence of the congener G. galloti (found in rat feces; data not shown) and the invasive M. musculus, both of which may provide accessible and abundant food resources for F. catus and R. rattus. Indirect interactions involving cats could be particularly relevant considering that G. intermedia occurs in up to 71% of cat feces in certain locations of Teno (Rando and López 2001) and its consumption in Montaña de Guaza has increased in the past years (Flores Ravelo and Rando Reves 2021 and references therein). In spite of that, G. intermedia FO in rats' diet exceeds that of cats in most sites (see Rando and López 2001; Flores Ravelo and Rando Reves 2021), while the only locations holding cat populations in Teno-i.e., La Galera (Rando and López 2001), Punta de Diente de Ajo and Andén de la Amargosa (Padilla et al. 2019)-are those with the highest population recovery of G. intermedia compared to past censuses. Although lizard decline is more pronounced in Montaña de Guaza, where cats are also present (authors' own data), this area also holds the highest rat density according to our results. Trophic relationships are governed by complex predator-prey cycles (Roemer et al. 2002; Ringler et al. 2015; Cerri et al. 2017), thus further research is needed to fully uncover R. rattus impact and understand how trophic relationships are affecting G. intermedia. However, this research allows considering *R. rattus* as a potentially serious threat to *G. intermedia*, whose role in the demise of the endemic lizard may have been masked by multispecies interactions.

Overall, the results presented in this study constitute a call for further attention on *R*. *rattus* as a highly detrimental invader potentially impacting endangered lizards on islands worldwide, and represent the first step in unveiling the impact of an interaction long assumed but never proven and whose consequences could be grave for *G. intermedia*.

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Author contributions MLD conceptualized the study, designed the methodology, was in charge of funding acquisition, resources and supervision. DPP and JMC performed fieldwork. MLG carried out all genetic procedures. MLD, MLG and JCP analyzed the data and wrote the original draft. All authors reviewed and approved the final manuscript.

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Data availability R code file available under https://doi.org/10.6084/m9.figshare.20455530.v1.

Declarations

Competing interests The authors declare no competing interests.

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