

1 **Does biogeographic history shape spatial patterns of intraspecific variation in**  
2 **personality, performance and morphology?**

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4 **Roberta Bisconti<sup>1</sup>, Andrea Chiochio<sup>1,\*</sup>, David Costantini<sup>1</sup>, Claudio Carere<sup>1</sup>, Daniele**  
5 **Canestrelli**

6 <sup>1</sup> Department of Ecological and Biological Sciences, Tuscia University, Largo dell'Università s.n.c., 01100  
7 Viterbo, Italy

8 \*Address corresponding to Andrea Chiochio, E-mail: a.chiochio@unitus.it

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10

11 **Abstract:**

12 Addressing how individual variation within populations drives the evolution of biodiversity patterns is a major  
13 challenge in ecology and evolutionary biology. Historical biogeographic processes have had dramatic  
14 consequences on the structure of biodiversity. However, while the interplay between historical processes and  
15 genotypic diversity within populations has been widely investigated, the effects of such processes on  
16 phenotypic diversity remain poorly explored. Here, we investigate whether dispersal-driven processes of  
17 historical biogeographic relevance, such as late Pleistocene range dynamics, have contributed to shape the  
18 geographic patterns of phenotypic trait variation. We focus on dispersal-related personality, morphological  
19 and performance traits in the Tyrrhenian tree frog, *Hyla sarda*, which underwent a northward range expansion  
20 from the Sardinia Island to the Corsica Island during the Last Glacial Maximum, when a temporary land-bridge  
21 connected these islands. We collected tree frogs from four geographic areas along the past expansion route,  
22 controlling for altitude, local habitat, demographic factors, and bioclimatic differences between geographic  
23 areas. Then, we scored intraspecific variation in two personality traits, two performance traits, along with  
24 morphological traits likely involved in the dispersal process. Tree frogs from the northern area in Corsica were  
25 more prudent in a novel environment, they had significantly larger body size, longer limbs, wider heads, and  
26 displayed stronger take-off and adhesion performances compared to individuals from the source area in  
27 Sardinia. Overall, these results suggest a non-random spatial sorting of the intraspecific variation in multiple

28 phenotypic traits during the range expansion phase. In turn, they also suggest that population differentiation in  
29 phenotypic traits associations might be a legacy of past biogeographic dynamics, identifying an overlooked  
30 driver of the current phenotypic architecture of animal populations.

31

32 KEYWORDS: Pleistocene range expansion, dispersal, personality, locomotory performance, biogeography

33

## 34 1. INTRODUCTION

35

36 Understanding the processes shaping the geographic patterns of intraspecific variation is a major challenge in  
37 ecological studies (Holt 2003; Des Roches et al., 2018; Raffard et al., 2019). In the last decades, considerable  
38 efforts have been focused on investigating the processes shaping the spatial distribution of genetic diversity  
39 within and among populations (Hewitt 2000, 2004; Schmitt, 2007; Weiss & Ferrand, 2007; Fonseca et al.,  
40 2023). Pleistocene glacial–interglacial cycles have been identified as a major factor in this respect, as they  
41 triggered periodic expansions and contractions of populations, which have left multiple imprints on their  
42 current genetic structure (Hewitt, 2004; Schmitt, 2007; Weiss & Ferrand, 2007; Fonseca et al., 2023). Although  
43 the link between historical biogeographic processes and intraspecific genetic diversity has been widely  
44 investigated, the contribution of these processes to the evolution of intraspecific phenotypic diversity is still  
45 virtually unexplored. At the same time, mounting evidence from studies on the ongoing range expansions (*e.g.*  
46 biological invasions) is showing that behavioural, morphological, physiological, and life history traits, play a  
47 crucial role in determining the ability of organisms to face new environmental conditions (Simmons &  
48 Thomas, 2004; Cote et al., 2007; Phillips et al., 2010; Shine et al., 2011). Therefore, investigating the interplay  
49 between biogeographic processes and intraspecific phenotypic variation is of great value to our understanding  
50 of how biodiversity evolves and is maintained.

51 Dispersal plays a main role in shaping the spatial pattern of biodiversity (Clobert et al., 2001; Travis  
52 et al., 2002; Ronce, 2007; Shine et al., 2011; Lowe & McPeck, 2014; Canestrelli et al., 2016a; Bonte & Dohrel,  
53 2017). Recent studies have shown that inter-individual differences in dispersal ability can directly affect the  
54 success of colonisation into new areas and the establishment of new populations (Bowler & Benton, 2005;  
55 Cote et al., 2007; Clobert et al., 2009; Canestrelli et al., 2016a). In particular, literature on invasive species is

56 showing that dispersers could be a non-random sample of individuals characterised by specific phenotypic  
57 profiles (Fraser et al., 2001; Phillips et al., 2008; Cote et al., 2010; Canestrelli et al., 2016a; Louppe et al.,  
58 2017). These studies suggest that evolutionary and ecological processes experienced by any expanding  
59 population drive the directional evolution of phenotypic traits at the range-edge of an expansion (Pintor et al.,  
60 2008; Atwell et al., 2012; Brodin et al., 2013; Myles-Gonzalez et al., 2015; Gruber et al., 2017). Behavioural  
61 traits can influence individual dispersal propensity (Cote et al., 2010; Canestrelli et al., 2016a). Individuals  
62 from the range-edge of an expansion would be characterized by a specific suite of personality traits, showing  
63 a more exploratory, more active, and bolder behaviour than those living in long established areas (Atwell et  
64 al., 2012; Myles-Gonzalez et al., 2015; Canestrelli et al., 2016a; Gruber et al., 2017). These traits are predicted  
65 to influence all dispersal stages (*i.e.*, departure, transience, and settlement; Dingemanse et al., 2003; Cote and  
66 Clobert, 2007; Cote et al., 2010; Canestrelli et al., 2016a), resulting in a rapid evolution of divergent  
67 personalities along an expansion route. Locomotory performance is also a key factor for expansion dynamics.  
68 During an expansion individuals would be spatially sorted by their locomotor abilities (Phillips, et al., 2010;  
69 Shine et al., 2011), and evidence showed how this process has often driven a rapid evolution in the locomotory  
70 traits (Travis & Dytham, 2002; Phillips et al., 2010; Shine et al., 2011) associated with morphological  
71 characteristics that would improve the dispersal rate compared to individuals living in long-invaded areas  
72 (Phillips et al., 2006; Phillips et al., 2008; Llewellyn et al., 2010; Louppe et al., 2017). For example, in the  
73 cane toad (*Rhinella marina*), a fast dispersal rate with straighter paths and longer legs has been documented at  
74 the edge of the invaded Australian range (Phillips et al., 2008; Alford et al., 2009; Llewellyn et al., 2010).  
75 However, while evidence on the impact of differential dispersal profiles and personality during contemporary  
76 expanding populations is growing, how phenotypic polymorphisms linked to dispersal abilities would have  
77 contributed to key past processes structuring biogeographic patterns remains substantially unexplored.

78         In this study we aim to bridge this gap by exploring how dispersal-driven processes may have shaped  
79 the contemporary geographic patterns of intraspecific variation in personality, performance, and  
80 morphological traits along the past expansion route of the Tyrrhenian tree frog (*Hyla sarda*; Bisconti et al.,  
81 2011a, Spadavecchia et al., 2021). Previous phylogeographic investigations on this amphibian have provided  
82 relevant insights into its historical range expansion (Bisconti et al., 2011a; Spadavecchia et al., 2021). During  
83 the last Glacial Maximum, when the sea levels were lower, a temporary land-bridge connected the islands of

84 Sardinia and Corsica (Van Andel & Shackleton, 1982; Shackleton et al., 1984), allowing a northward range  
85 expansion of the Tyrrhenian tree frog from Sardinia to Corsica (Bisconti et al., 2011a; Spadavecchia et al.,  
86 2021). However, after the last Glacial Maximum, the sea levels rose, separating the newly established  
87 populations in Corsica from their source populations in Sardinia, thus impeding the subsequent flow of  
88 individuals from the rear. Recent studies investigating physiological traits of tree frog individuals from these  
89 two islands clearly showed a spatial pattern of variation, likely linked to this historical population expansion.  
90 Specifically, populations of the two islands showed a clear differentiation in rates of physiological ageing  
91 owing to oxidative stress, different telomere dynamics, and differences in anti-predator strategies (Liparoto et  
92 al., 2020; Canestrelli et al., 2021; Spadavecchia et al., 2023).

93 We explored the hypothesis that historical biogeographic events of dispersal contribute to shape  
94 patterns of phenotypic diversity. We expect that Sardinia and Corsica treefrogs would differ in dispersal-related  
95 phenotypic traits, such as personality, morphology, and locomotor performance. We also tested the hypothesis  
96 that this legacy of past events should be detected in the geographic pattern of these dispersal-related phenotypic  
97 traits. Specifically, if dispersers are a non-random sample of individuals, we predict that, compared to Sardinia  
98 tree frogs, Corsica tree frogs should be bolder, and have longer limbs and better jumping performance.

99

## 100 **2. MATERIAL AND METHODS**

101

### 102 **2.1 Sampling and housing**

103 Sample collection of tree frogs was carried out along a latitudinal transect from the northern part of the island  
104 of Corsica to the ancestral area of the glacial refuge in central-eastern Sardinia (Figure 1 and Table 1). A total  
105 of 93 tree frogs were collected in spring 2018 from four areas (Figure 1; Table 1). Individuals were captured  
106 with hand nets during the first hours of the night, after acoustic and visual localization and then transported to  
107 the housing facilities at Tuscia University (Ichthyogenic Experimental Marine Centre, CISMAR:), Italy.  
108 Housing conditions are described in detail in Bisconti et al., 2023.

109 The sampling design was accurately planned to reduce possible confounding effects linked to local  
110 habitat, kinship or inbreeding between individuals: a) we sampled two areas per island, and two breeding sites  
111 for each area, collecting individuals from different ponds within each site (see Table 1 and Figure 1); b) all the

112 sampling areas were located on the coasts on the eastern side of species distribution, to avoid both current and  
113 past bioclimatic differences among sampling sites, as showed by previous species distribution model (Bisconti  
114 et al., 2011a); c) we collected tree frogs from ponds exclusively located to coastal areas to avoid any effect  
115 linked to environmental conditions correlated to altitude; d) finally, to avoid any effect due to ecological and  
116 evolutionary effects of colonisation through jump dispersal rather than through a spatial diffusion process, we  
117 intentionally excluded other islands (Elba and Capraia) inhabited by this species (Bisconti et al., 2011a,b;  
118 Bisconti et al., 2023).

119

## 120 **2.2 Ethical note**

121 All the procedures were performed under the approval of the Institute for Environmental Protection and  
122 Research 'ISPRA' (protocol #5944), Ministry of Environment 'MATTM' (protocol #8275), Regione Sardegna  
123 (#12144) and Prefecture of Corsica (#2A20180206002 and #2B20180206001).

124         Permission to temporarily house amphibians at Tuscia University (Ichthyogenic Experimental Marine  
125 Centre, CISMAR:), Italy. All handling procedures outlined in the present study were approved by the Ethical  
126 Committee of the of the Tuscia University for the use of live animals. During captivity the animals were  
127 monitored daily. No adverse effects on the overall health condition of the tree frogs were observed during the  
128 procedures. The animals were released in the original sampling locations at the end of the experimentation.

129

## 130 **2.3 Behavioural tests**

131 To investigate personality traits variation along the post-glacial range expansion route of *H. sarda* we focused  
132 on two personality traits that are commonly associated to the dispersal propensity: exploration attitude in a  
133 novel environment and boldness (Canestrelli et al., 2016a). Exploratory activity is considered crucial in  
134 promoting dispersal, as individuals with a greater propensity toward exploration were generally found at the  
135 expanding range edge (Dingemanse et al., 2003; Cote et al., 2010). Exploration activity was investigated  
136 measuring the latency in exploring a novel environment, and the time spent in exploration. Propensity to  
137 disperse was also assessed by investigating the individual's boldness, *i.e.* the individual's propensity to take  
138 risks (Réale et al., 2007; Canestrelli et al., 2016a). Bold individuals are considered willing to accept the  
139 intrinsic risks of dispersal (Stamps, 1985; Bonte et al., 2012), thus dispersing earlier than shy conspecific.

140 To assess these personality traits, we performed two distinct behavioural tests as described below and each test  
141 was repeated after 10 days to measure temporal consistency. All the tests were performed in the housing room,  
142 to avoid any changes in humidity and temperature that could affect the analysed behavioural features  
143 (Duellman & Trueb, 1994).

144

### 145 *2.3.1 Exploration in a novel environment*

146 Considering the arboreal lifestyle of *H. sarda*, a cylindric arena developed in both vertical and horizontal  
147 dimensions was employed, enriched with plants (*Epipremnum aureum*) and an oakwood that allows the  
148 individuals to hide, eliminating the possibility that the measured behaviours reflected abnormal fear or anxiety  
149 rather than spontaneous exploratory behaviour (see Bisconti et al., 2023). The test is accurately described in  
150 Bisconti et al., (2023). Briefly: after a 5-minute acclimation period, individuals were left to explore the arena  
151 for 10 minutes, and the individual exploration behaviour was recorded using an HD video camera (Nova  
152 Germany model DVR AHD-7908). We extracted the following variables using Boris 5.1.3 (Friard & Gamba,  
153 2016): 1) latency to explore (s); 2) duration of activity (expressed as percentage of duration of test); 3) duration  
154 of time spent on the arena ground floor; 4) jumping activity, expressed as percentage of time in activity  
155 (moving) spent jumping.

156

### 157 *2.3.2 Boldness*

158 This test was performed to assess the individual behaviour across the boldness-shyness behavioural axis. We  
159 considered the latency to exit from a shelter as a proxy of the individual propensity to take risk. A HD video  
160 camera (Nova Germany model DVR AHD-7908) was used to record a rectangular arena where all animals  
161 were caged into a cylindrical dark shelter for a 6-minute acclimation period and then left free to exit (Bisconti  
162 et al., 2023). The software Boris 5.1.3 was used to analyse the individual behaviour, measuring the time it took  
163 to exit the shelter.

164

## 165 **2.4 Locomotory performance tests**

166 To investigate the variation in locomotory performances along the past range expansion route of *H. sarda* we  
167 assessed two performance traits that could have a crucial role in the species dispersal. Considering jumping as

168 the predominant mode of locomotion in tree frogs, we explored two performance traits linked to jumping  
169 abilities: the jumping force, and the adhesion force. The jumping force was inferred evaluating the force at  
170 take-off, that may be considered a crucial trait in determining the dispersal capacity in a tree frog since the  
171 distance reached with jumps is a function of the propulsion force generated at take-off (*e.g.*, Marsh & John-  
172 Alder 1994; Nauwelaerts & Aerts 2006 and references therein). The adhesion force is also considered a proxy  
173 of dispersal abilities because it plays a main role during the landing phase (Duellman & Trueb, 1994; Bisconti  
174 et al., 2023). Indeed, tree frogs have specialised toe pads used during the landing phase (Emerson & Diehl  
175 1980; Federle et al. 2006; Smith et al. 2006). Therefore, the differential ability to adhere to the substrate after  
176 a jump would improve the safety of the landing, avoiding missing the target (Bijma et al., 2016).  
177 The jumping force was quantified by means of a jumping test, while the adhesion force was quantified by a  
178 stickiness test, as described below. Before each test, all the individuals were weighted using a scale Acculab  
179 model (ATILON ATL-224-I).

180

#### 181 *2.4.1 Jumping test*

182 A rectangular arena was videotaped with a video camera (Panasonic model no. DMC-FZ300) placed laterally  
183 to the setup. Each tree frog was fitted with an accelerometer data logger (two Axy-4 units, Technosmart, Rome,  
184 9.15 x 15 x 4 mm, 1 gr weight including battery) set to record triaxial acceleration (0 - 4 g) at 100 Hz, and the  
185 tree frog was induced to jump by stimulating its caudal region (Mitchell & Bergmann, 2016). Five jumps for  
186 individuals were collected (see Bisconti et al., 2023 for further details). The dynamic body acceleration for  
187 each dimension was extracted from the downloaded data (x, y, z; Wilson et al., 2006; Shepard et al., 2008).  
188 The logger's x axis recorded sway, the y axis recorded surge, and the z axis recorded heave (Halsey et al.,  
189 2008). By using Framework 4 software (version 2.5), we added these values to get the Vector sum of Dynamic  
190 Body Acceleration [ $VeDBA = (Ax^2 + Ay^2 + Az^2)$ ], quantifying five VeDBA values for each individual, one  
191 for each jump, and the highest value of VeDBA was chosen to calculate the individual maximum jumping  
192 force at take-off [VeDBA x mass] and considered for the analysis.

193

#### 194 2.4.2 Stickiness test

195 A smooth plastic rotating wheel with a moderate and constant angular speed (3 revolutions per minute) was  
196 used to test the maximum adhesiveness force. We recorded five falls for each individual using Nova Germany  
197 model DVR AHD-7908. The software Tracker (version 4.11.0) was used to extract the angle of fall (radian)  
198 for each of the five falls, and the largest value was selected for the subsequent analyses. The maximum  
199 adhesion force was calculated [ $\cos(\alpha) \times \text{mass} \times g$ ] using Barnes et al. (2006)'s standard protocol (see Bisconti  
200 et al., 2023, for further details).

201

#### 202 **2.5 Morphological traits**

203 Locomotion in anurans with an arboreal lifestyle is characterised by individuals with morphological  
204 adaptations to better perform during jumps and landings (Kamada et al., 2017; Bijma et al., 2016). Thus, we  
205 investigate those morphological traits that directly play a role in these two phases of locomotion, such as body  
206 size and limb lengths (Phillips et al., 2006). We also collected data on another morphological trait that is  
207 intrinsically linked to jumping performance: the head size. Indeed, there is evidence suggesting that the head  
208 plays a crucial role improving the ability of tree frogs in performing jumps, especially during the jumping  
209 phase by maintaining the posture, and during the landing phase by hitting the substrate first (Reilly et al., 2016;  
210 Kamada et al., 2017).

211 We collected measures of the following morphological traits: (1) snout–vent length (SVL; mm); (2)  
212 head size (mm); (3) forelimb length (mm); (4) hindlimb length (mm). To obtain standardised measures, the  
213 tree frogs were anaesthetised by a two-minute submersion in a solution of MS-222 (0.05% m/v), placed in the  
214 prone position on 1x1 cm grid paper, parallel to the plane of a camera lens, and photographed using a Panasonic  
215 LUMIX (DMC-FZ300) digital camera. All photographs were taken with the same camera settings and under  
216 the same lighting conditions. After the photographs, each individual was placed in a humid box until it fully  
217 recovered, and then reintroduced into its own fauna box. The measures were then extracted by the photographs  
218 and analysed by using ImageJ 1.52 (National Institutes of Health, USA). The measurement of head size,  
219 forelimb length and hindlimb length have been standardised by extracting the residual of the regression  
220 between each single trait and the SVL.

221



## 222 2.6 Data analysis

223 We performed all statistical analyses using R software version 4.3.1 (R Core Team 2023). To evaluate the  
224 repeatability of the behavioural traits, we ran Generalized Linear Mixed effect (GLMM)-based repeatability  
225 models using the “rpt” function of the *rptR* package (Schielzeth & Nakagawa, 2011; Stoffel et al., 2017),  
226 setting as Poisson distribution datatype; duration of activity and duration of time spent on the arena floor level  
227 were treated as Proportion datatype. We entered each behavioural trait as a dependent variable and individual  
228 as a random factor. We considered behavioural variables as personality traits, and used them in the subsequent  
229 analyses, when their repeatability value  $R > 0.2$ , and the lower bound of the CI was  $> 0.0$  (Brodin et al., 2013;  
230 Kelleher et al., 2017; Kelleher et al., 2018 and references therein). Preliminary models showed that entering  
231 sampling location within each island as a random factor did not improve the fit of the model (i.e. the AIC value  
232 was not reduced beyond 2); thus, it was not subsequently considered. Also, running the models with or without  
233 females did not change at all both the model fit and the results, and thus we retained females for all the  
234 subsequent analyses.

235 We tested the hypothesis of a phenotypic change across the past expansion route of *Hyla sarda* from  
236 Sardinia to Corsica. First, we tested the association of phenotypic variation with the distance from the estimated  
237 area of origin of the range expansion from Sardinia to Corsica as estimated in Spadavecchia et al. (2021). We  
238 set a distinct linear regression model between each behavioural, performance and morphological variable and  
239 the distance from the area of origin, measured in km using the Google Earth (Google Inc.) “ruler” function.

240 Then, we assessed the differences in personality, performance, and morphological traits between the  
241 two islands by running Generalized Linear Models (GLMs; “lm” function in the basic package *stats*) entering  
242 each variable singly as dependent variables, and island as fixed factor. We also run the analyses including  
243 sampling location within each island as a random factor in generalized linear mixed-effect models (GLMMs;  
244 “lmer” function in the *lme4* package) and selected the best-fit model using the AIC (lowest AIC). For  
245 repeatable personality traits, we used the mean between the two repetitions; the variables were square-root  
246 (frequency of jumping events) or log10 (all the others) transformed to meet the assumption of residuals  
247 normality. To exclude the effect of the body size on the morphological traits, for each measured trait we  
248 extracted the residuals of the regression with the SVL. Normality and homoscedasticity of model residuals

249 were evaluated to ascertain the goodness of model fit. For the selected best-fit models, we contrasted the  
250 estimated marginal means and standard errors from the models as computed using the *emmeans* r package.

251

### 252 3. RESULTS

253

254 The repeatability coefficients with its CI and the p-value of are reported in Table 2. Three out of the five  
255 investigated behavioural traits resulted significantly repeatable: boldness, measured as the latency to exit from  
256 the shelter ( $r = 0.40$ ), the frequency of jumping events during the arena exploration ( $r = 0.35$ ), and the latency  
257 to explore a novel environment ( $r = 0.31$ ). Conversely, activity and duration of time spent on the arena floor  
258 were not significantly repeatable and thus they were not considered as personality traits and excluded from the  
259 subsequent analyses.

260 Results from all the models showed a phenotypic change across the past expansion route of *Hyla sarda*  
261 from Sardinia to Corsica. We found significant associations between phenotypic variation and the distance  
262 from the estimated area of origin of the range expansion from Sardinia to Corsica for all the analysed traits but  
263 two (hind limb length, and head width; see Supplementary Figure S1). Also, we found substantial and  
264 significant differences between Sardinia and Corsican populations for all the analysed traits but one (maximum  
265 jumping force at take-off). The AIC model comparison (Supplementary Table 1) showed that entering  
266 sampling location within each island as a random factor did not improve the fit of the model (*i.e.* the AIC value  
267 was not reduced) for all the traits but two (performance traits), supporting the independence of the pattern  
268 found from local sampling site conditions.

269 We found significant differences in personality traits between Sardinia and Corsica individuals (Table  
270 3), with Corsica individuals showing a shyer profile and higher jumping frequencies. The latency to exit from  
271 the shelter and the latency to explore were longer for Corsica individuals than those showed by Sardinia  
272 individuals ( $p = 0.032$  and  $p < 0.001$ , respectively). The jumping frequency during exploration was  
273 significantly higher in Corsica than in Sardinia ( $p = 0.001$ ). The estimated marginal means and standard errors  
274 for personality differences between Sardinia-Corsica are shown in Figure 2.

275 Sardinia and Corsica populations differed in both jumping force and stickiness, with tree frogs from  
276 Corsica showing higher performance values than individuals from Sardinia (Table 3 and Figure 3). However,

277 only stickiness showed significant differences ( $p < 0.001$ ), whereas the differences in jumping force resulted  
278 not significant ( $p = 0.094$ ). The estimated marginal means and standard errors for personality differences  
279 between Sardinia-Corsica are shown in Figure 3.

280 All morphological traits linked to dispersal ability showed significant differences between the two  
281 populations (Table 4 and Figure 4). The investigated morphological traits showed that Corsica tree frogs were  
282 larger in body size ( $p < 0.001$ ), with longer forelimbs ( $p = 0.044$ ) and hindlimbs ( $p = 0.026$ ), and wider head  
283 size ( $p = 0.011$ ).

284

#### 285 4. DISCUSSION

286 We found substantial and significant differences between individuals from the two islands in all the  
287 investigated phenotypic traits, with the Corsica tree frogs showing a shyer behavioural profile, longer decision-  
288 making time, a significant attitude to jump during exploration, and more efficient dispersal-related  
289 performance and morphological traits.

290 The spatial phenotypic pattern observed could be explained by several main scenarios. Variation in  
291 bioclimatic conditions between the two islands could contribute to shape the differences founded. Climatic  
292 conditions could, indeed, affect habitat variables and the microhabitat cover, that in turn, could differentially  
293 influence the dispersal and the ability in avoiding predators especially for an arboreal species (Martin et al.  
294 2005). However, previous studies showed a comparable bioclimatic condition between the two islands that,  
295 coupled with a negligible latitudinal effect, contribute to excluding a role of variation of climatic condition in  
296 explaining such a pattern of differentiation (Bisconti et al., 2011a). An alternative scenario could be attribute  
297 to differences in ecological community between the two islands. In particular, the presence of different  
298 interacting species within the two islands could contribute to mould local adaptations, for example in avoiding  
299 predation (Kang et al. 2017; Gavriilidi et al., 2022). Yet, Corsica and Sardinia are continental islands, they  
300 share geological origin and history, and no differences in their animal communities had never been reported,  
301 contributing to rule out a possible role of these ecological forces to explain the pattern emerged. Lastly, a  
302 genetic basis of diversity among populations of the two islands could be invoked to discuss the phenotypic  
303 differences showed. Therefore, divergent populations could show different phenotypic traits even within the  
304 species level (Lipshutz, et al., 2017; Stamp et al., 2020). However, previous phylogeographic and population

305 genetic investigations showed the absence of remarkable differences in the levels of genetic variability and the  
306 lack of spatial genetic structure between Sardinia and Corsica populations, thus excluding a genetic influence  
307 in explaining the pattern observed (Bisconti et al., 2011a,b). Our data may suggest that the past range expansion  
308 has been promoted by a non-random sample of individuals, and the evolution of the observed differences  
309 would be the result of different selection pressure along the colonisation route. Individuals that successfully  
310 colonised Corsica Island crossing the land bridge during the expansion phase would be sorted by their dispersal  
311 propensity, and this spatial sorting processes would have promoted directional changes in the phenotypic  
312 profiles of individuals during this past range expansions. This scenario would be further supported by the  
313 observed non-random spatial pattern in the geographical cline of all phenotypic traits studied. All these traits  
314 would show a clear latitudinal gradient of variation that perfectly mirrors the putative route of range expansion  
315 for this species (see Supplementary Figure 1). In the following sections, we will discuss our findings for each  
316 group of traits studied, evaluating their putative implications during range expansion and the colonization of  
317 new environments.

318         The shy behavioural profile showed by the Corsica individuals could appear counterintuitive, as  
319 dispersers at the range edge have frequently been described as bolder and more exploratory than those living  
320 in long-established areas (*e.g.* Pintor et al., 2008; Liebl & Martin, 2012; Atwell et al., 2012; Myles-Gonzalez  
321 et al., 2015; Canestrelli et al., 2016a; Gruber et al., 2017). However, recent empirical studies have been  
322 showing that a shy and more prudent behavioural profile is an efficient alternative strategy to navigate in a  
323 novel environment (Gavriilidi et al., 2022; Mazza & Eccard, 2023; Eccard et al., 2023). On the contrary, a  
324 shy profile in expanding populations could allow a more careful assessment of the surrounding environment,  
325 thereby reducing risks and ultimately increasing the chances of survival (Groothuis & Carere 2005; Réale et  
326 al. 2010; Mazza & Eccard, 2023). By requiring less energy investment in exploration, this behavioural profile  
327 could favour metabolic resource allocation to other critical functions pivotal for a successful colonisation, such  
328 as foraging, growth, and reproduction (Verbeek et al., 1994; Deerenberg et al., 1998; Cockrem, 2007; Coppens  
329 et al., 2010). Besides, the higher propensity of Corsica tree frogs to jump during their overall activity duration,  
330 compared to individuals from the source island, may have facilitated a cost-benefit optimization for these  
331 dispersers. Despite the energetic costs associated with a jumping locomotory mode, it could allow individuals  
332 to cover longer distances with a single movement, interspersed with pauses for energy recovery and for

333 predatory and environmental context assessment, before exploring again. The higher propensity to jump shown  
334 by tree frogs from Corsica is reflected in their better locomotory performance than those from Sardinia. As  
335 key traits in determining species dispersal ability, locomotory performance traits play a direct role in dispersal  
336 processes and, consequently, in the overall success of expansion (Phillips et al., 2006; Phillips et al., 2008;  
337 Llewellyn et al., 2010; Louppe et al., 2017). The performance trait analysis revealed that Corsica tree frogs  
338 exhibit greater jumping and adhesion forces than those from Sardinia. In such an arboreal species, both traits  
339 are expected to significantly enhance dispersal ability, playing a crucial role during both the initial jumping  
340 phase and the subsequent landing phase. This may contribute to covering longer distances with each jump  
341 while minimizing the risk of missing the target during landing (see material and methods; Duellman and Trueb,  
342 1994; Kosmala et al., 2017). More performant profiles showed by Corsica individuals during dispersal are in  
343 line with findings from several other studies exploring locomotory performance traits in contemporary  
344 expanding populations (Phillips et al., 2008; Alford et al., 2009; Llewellyn et al., 2010). In fact, recent studies  
345 on invasive species have investigated the evolution of locomotory performance traits at the range-edge of  
346 expanding populations, revealing a significant improvement in the locomotor abilities of dispersers, thus  
347 suggesting that, during an expansion, individuals may be spatially sorted based on their dispersal-related  
348 performance ability (Phillips et al., 2008, Alford et al., 2009; Llewellyn et al., 2010; Shine et al., 2011; Louppe  
349 et al., 2017).

350 Morphological analyses showed that Corsica tree frogs have a larger body size, longer limbs, and a  
351 wider head size than those from Sardinia. All these features imply a better performance of these individuals  
352 during the dispersal and the colonisation of new environments. A larger body size would confer several  
353 advantages to individuals during an expansion. Studies on amphibians showed that larger individuals have a  
354 better locomotory performance both on land and in water (Cabrera-Guzman et al., 2013). Moreover, a larger  
355 size plays an important role in other essential functions particularly crucial for an amphibian during  
356 colonisation, such as avoiding dehydration (Kosmala et al., 2020), allowing to be more performant in capturing  
357 prey (Cabrera-Guzman et al., 2013) and also increasing the reproductive success, whereby the larger male  
358 would be more successful both in intraspecific competition and during the amplexus (Bowcock et al., 2013;  
359 Herrel et al., 2012; Clarke et al., 2019). Similarly, longer limbs improve locomotion, and, thus, the dispersal  
360 ability (Phillips et al., 2006; Clarke et al., 2019). Longer legs allow a greater propulsive force in jumping

361 (Kamada et al., 2017; Bijma et al., 2016). For example, in the invasive *Rhinella marina*, individuals at the edge  
362 of the Australian invasion front exhibited longer legs, a trait significantly enhancing invasion success (Phillips  
363 et al., 2006). In the context of jumping amphibians like tree frogs, longer forelimbs could also contribute to  
364 improving landings, while dampening the energy of jumps upon touchdown (Bijma et al., 2016). Lastly,  
365 colonising individuals with a wider head would likely adapt better to landings after jumps. The increased  
366 propulsion resulting from longer legs might lead to more forceful landings and given that the head is the first  
367 part of the body to arrive on a surface, a larger head size could confer better management of shock absorption  
368 during collisions.

369 In conclusion, this study provides novel evidence that historical range expansion could have played a  
370 role in explaining the geographic patterns of intraspecific phenotypic variation in natural populations. Rapid,  
371 substantial, and directional evolution in dispersal-enhancing traits had been already documented in populations  
372 currently undergoing range expansion (Phillips et al., 2006; Liebl and Martin, 2012; Gruber et al., 2017;  
373 Kosmala et al., 2017). However, although studies of ongoing invasion provide an excellent opportunity to  
374 investigate the causes and consequences of dispersal dynamics, they do not allow the investigation of their  
375 long-term legacies, in evolutionary perspective (Canestrelli et al., 2016a, b). The selective advantages  
376 associated with adaptive, dispersal-enhancing, traits during an expansion phase may be transient, as suggested  
377 by several empirical evidence (Travis & Dytham, 2002; Brown et al., 2007; Phillips et al., 2008; Brown et al.,  
378 2014). Recently established populations may indeed restore variation (phenotypic and genotypic) through a  
379 slower flow of individuals from the rear or may evolve novel trait states in response to demographic density  
380 increases in newly colonised areas (Cobben et al., 2015; Canestrelli et al., 2016a; Perkins et al., 2016). To date,  
381 we are not aware of studies addressing if the predicted directional changes in dispersal-enhancing traits may  
382 persist in populations long after the expansion stops, and the demographic conditions promoting spatial sorting  
383 processes vanish (Cobben et al., 2015; Canestrelli et al., 2016a; Perkins et al., 2016). In this frame, the  
384 expansion history of the Tyrrhenian tree frogs could offer the opportunity to learn about the long-term legacy  
385 of non-neutral processes at play during a range expansion. Finally, to investigate the genomic underpinnings  
386 of the phenotypic evolution promoted by historical range expansions appears the next mandatory step to reach  
387 a more complete understanding of the eco-evolutionary processes moulding spatial and temporal patterns of  
388 variation of biological diversity.

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633 **FIGURES AND TABLES**

634 **Table 1.** Geographic coordinates and sample size of the sites sampled to study phenotypic variation within  
 635 *Hyla sarda* populations.

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Island	Sampled site	Latitude	Longitude	N
<b>Corsica</b>	Aleria	42.1132	9.5221	14
	San Giuliano	42.2677	9.5184	10
	Etagn de Canettu	41.4456	9.2035	10
	T10	41.4603	9.2170	13
<b>Sardinia</b>	Stazzo Pulcheddu	41.1639	9.3620	8
	Porto Pollo	41.1844	9.3303	15
	Cala Ginepro	40.4481	9.7920	14
	Siniscola	40.5814	9.7691	9

637

638 **Table 2.** Summary of GLMM-based repeatability (R) estimates from multiplicative model. Parametric  
 639 bootstrapping (number of iterations = 1,000) was used to calculate the CI interval and the likelihood ratio test  
 640 to estimate the P-value of the repeatability distribution. Significant traits are shown in bold.

641

Variable	Estimate	95 % CI	P-value
Latency to explore	0.31	0.09-0.50	<b>0.002</b>
Activity	0.04	0.00-0.10	0.051
Time on floor	0.07	0.00-0.21	0.183
Jumping events	0.35	0.14-0.53	<b>0.001</b>
Latency to exit	0.40	0.20-0.55	<b>&lt;0.001</b>

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650 **Table 3.** Linear models showing differences between Corsica and Sardinia Island populations in behavioural  
651 and performance traits. Coefficient estimates ( $\pm SE$ ) of these models are shown with both behavioural and  
652 performance traits as dependent variable and island as fixed factor. Significant contrasts are shown in bold.

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Dependent variable	Reference	Level	Coefficient	SE	t-Value	P
Latency to explore	C	S	-0.526	0.131	- 4.028	<b>&lt;0.001</b>
Jumping events	C	S	-1.256	0.358	-3.504	<b>0.001</b>
Latency to exit	C	S	-0.396	0.181	-2.185	<b>0.032</b>
Jumping	C	S	-3.050	1.552	-1.965	0.094
Stickiness	C	S	-7.731	1.969	-3.926	<b>&lt;0.001</b>

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657 **Table 4.** Linear models showing differences between Corsica and Sardinia Island populations in  
658 morphological traits. Coefficient estimates ( $\pm SE$ ) of these models are shown with morphological traits as  
659 dependent variable and population as fixed factor. Significant contrasts are shown in bold.

660

Dependent variable	Reference Level	Level	Coefficient	SE	t-Value	P
SVL	S	C	-0.238	0.062	-3.844	<b>&lt;0.001</b>
Fore limb	S	C	-0.062	0.030	-2.059	<b>0.044</b>
Hind limb	S	C	-0.100	0.044	-2.288	<b>0.026</b>
Head width	S	C	-0.032	0.0124	-2.595	<b>0.011</b>

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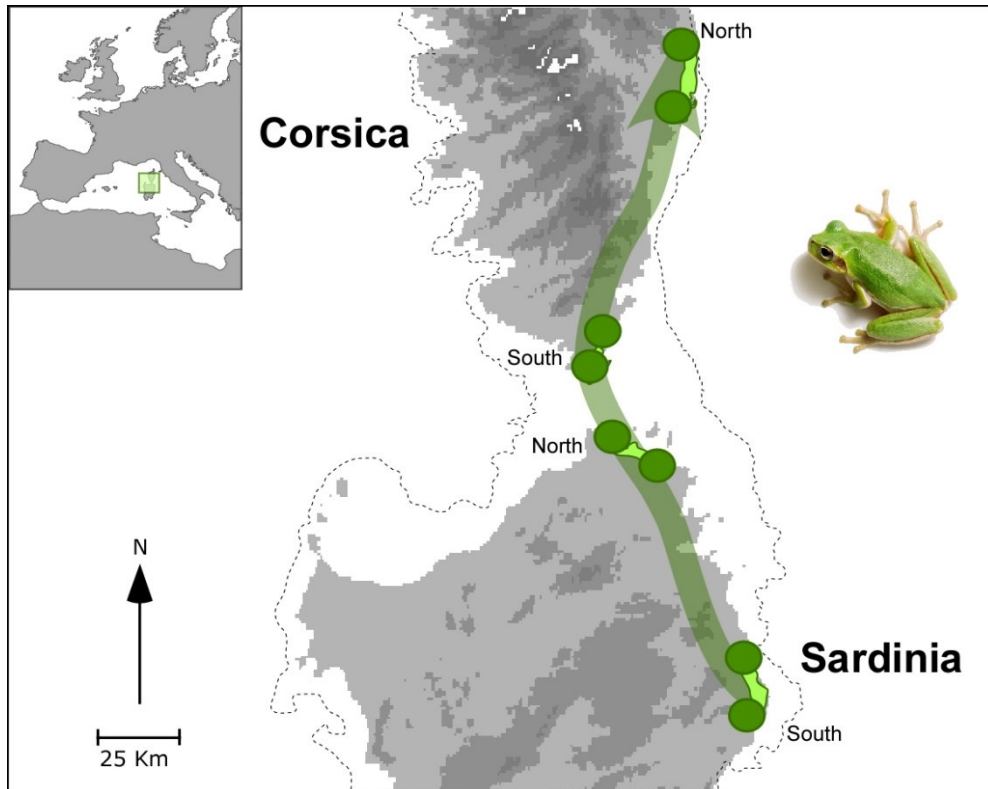
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667 **Figure 1.** Geographical distribution of the 8 sampled populations. The dashed line shows the approximate  
668 location of the coastline during the last glacial maximum (Thiede, 1978).

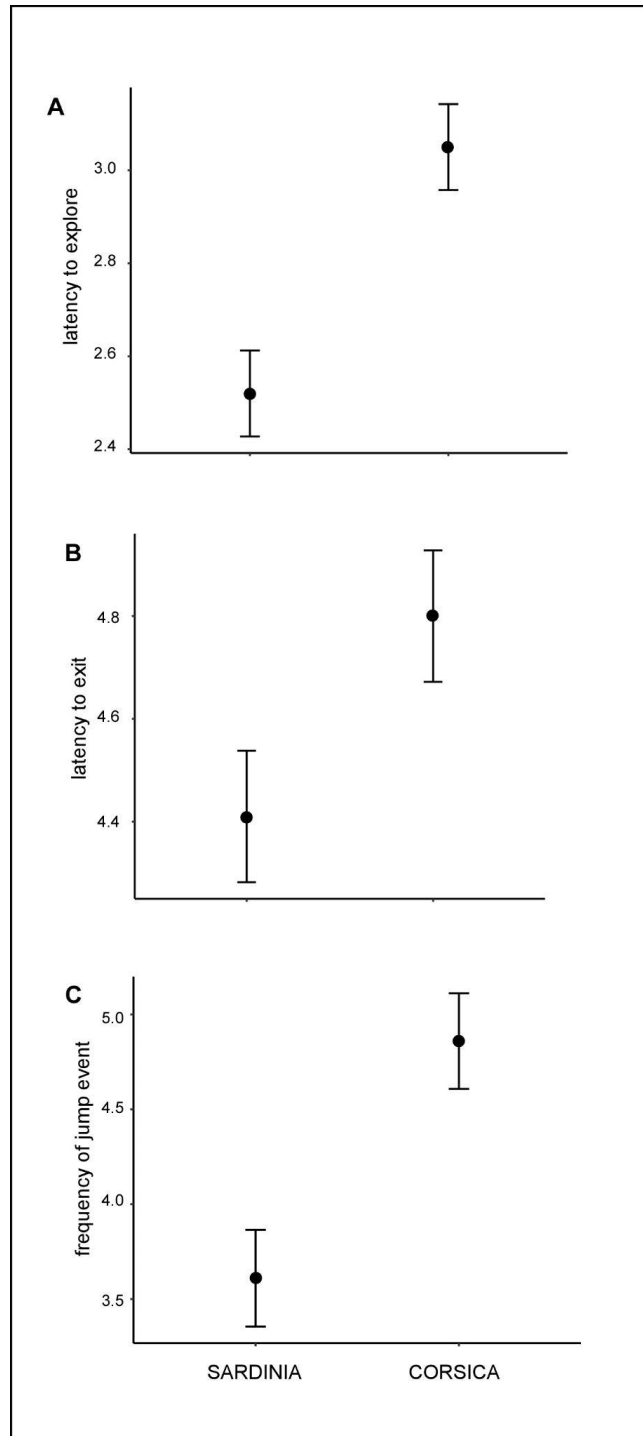
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672 **Figure 2.** Estimated marginal means and standard errors from linear models for personality traits for each  
673 island (A: latency to explore; B: latency to exit; C: frequency of jump event).



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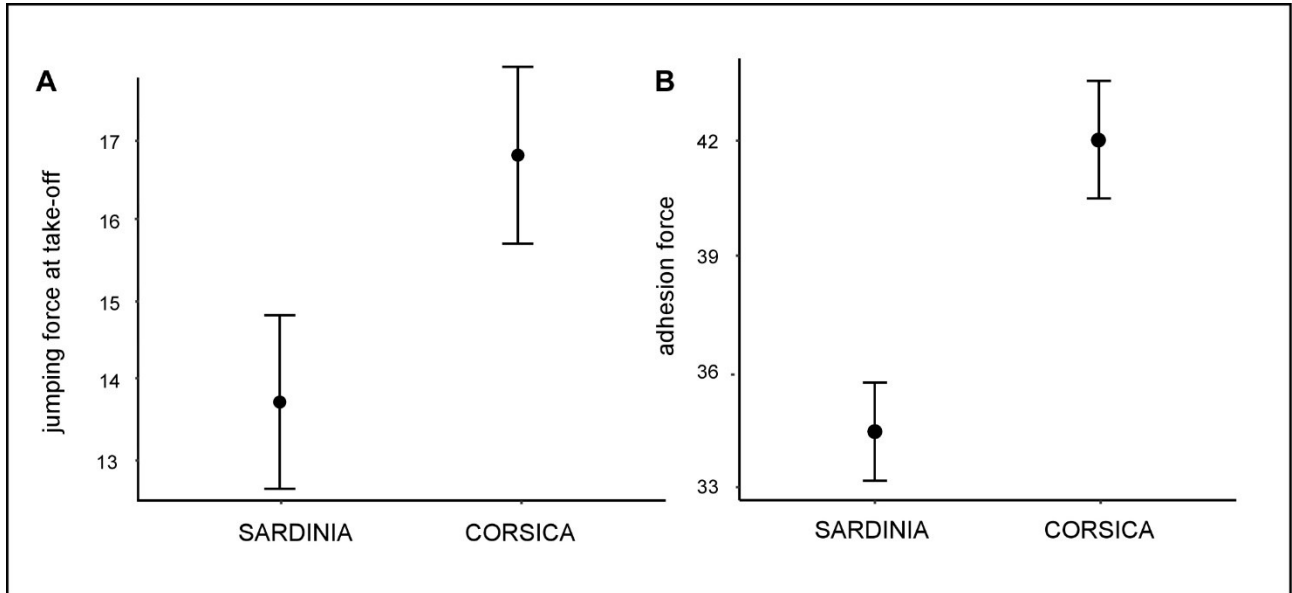
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678 **Figure 3.** Estimated marginal means and standard errors from linear models for performance traits for each  
679 island (A: jumping force at take-off; B: adhesion force).

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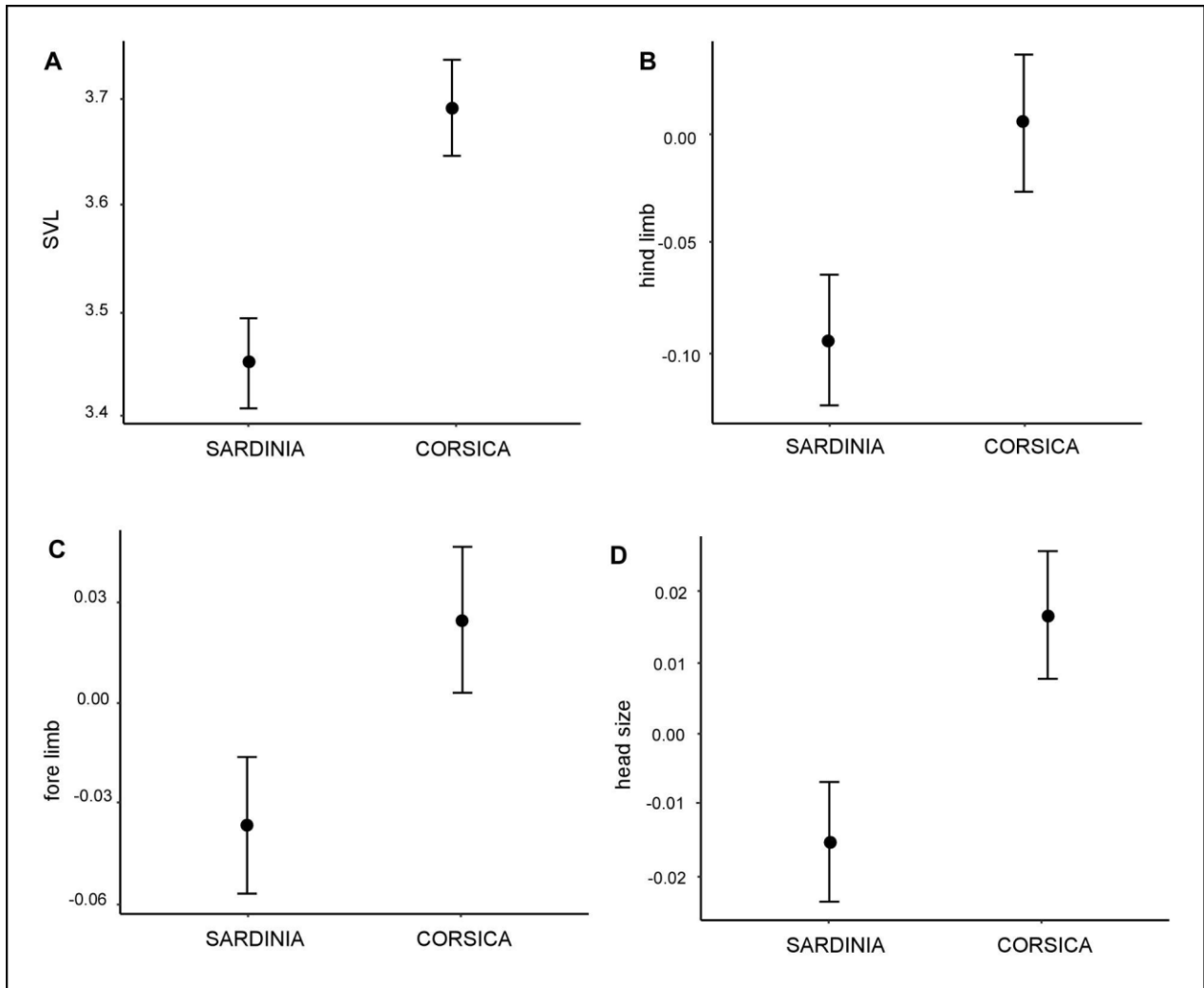
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698 **Figure 4.** Estimated marginal means and standard errors from linear models for morphological traits for each  
699 island (A: SVL; B: hind limb; C: fore limb; D: head size).



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**Table S1.** AIC values of the alternative models run entering each variable singly as dependent variables, and island as fixed factor (Model 1) or including sampling location as a random factor (Model 2); the selected best fit models, i.e. lowest AIC, are reported in bold.

<b>Phenotypic trait</b>	<b>AIC</b>	
<b>Behaviour</b>		
<i>Latency to explore</i>	<b>Model 1</b>	<b>179.06</b>
	Model 2	186.93
<i>Boldness</i>	<b>Model 1</b>	<b>239.34</b>
	Model 2	245.79
<i>Jumping activity</i>	<b>Model 1</b>	<b>369.65</b>
	Model 2	373.49
<hr/>		
<b>Locomotory performance</b>		
<i>Jumping force</i>	Model 1	348.02
	<b>Model 2</b>	<b>343.07</b>
<i>Stickiness</i>	Model 1	384.76
	<b>Model 2</b>	<b>380.61</b>
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<b>Morphology</b>		
<i>Snout–vent length</i>	<b>Model 1</b>	<b>4.42</b>
	Model 2	7.44
<i>Head size</i>	<b>Model 1</b>	<b>-194.23</b>
	Model 2	-177.15
<i>Forelimb length</i>	<b>Model 1</b>	<b>-83.85</b>
	Model 2	-70.10
<i>Hindlimb length</i>	<b>Model 1</b>	<b>-38.92</b>
	Model 2	-26.64
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**Figure S1.** Phenotypic trait variation across the inferred past range expansion route in the Tyrrhenian tree frog, *Hyla sarda*: results from the linear regression model between the distance from the area of origin (in km) and each behavioural, performance and morphological trait investigated in this study.

