- The role of habitat and evolutionary allometry in the morphological
- differentiation of *Pristurus* geckos (OR: Evolution along allometric
- lines of least resistance: Morphological differentiation in *Pristurus*
- geckos)
- ORDER TBD: Héctor Tejero-Cicuéndez<sup>1,\*</sup>, Iris Menéndez<sup>2,3</sup>, Adriá n-Talavera,
- Gabriel Riaño, Marc Simó-Riudalbas<sup>1</sup>, Salvador Carranza<sup>1</sup>, and Dean C. Adams<sup>4</sup>
- 8 27 October, 2022
- <sup>9</sup> <sup>1</sup>Institute of Evolutionary Biology (CSIC-Universitat Pompeu Fabra), Passeig Marítim de la
- <sup>10</sup> Barceloneta 37-49, Barcelona <u>08003</u>08002, Spain
- <sup>11</sup> Departamento de Geodinámica, Estratigrafía y Paleontología, Facultad de Ciencias Geológicas,
- 12 Universidad Complutense de Madrid, C/José Antonio Novais 12, Madrid 28040, Spain
- <sup>3</sup>Departamento de Cambio Medioambiental, Instituto de Geociencias (UCM, CSIC), C/Severo
- Ochoa 7, Madrid 28040, Spain
- <sup>4</sup>Department of Ecology, Evolution, and Organismal Biology, Iowa State University, Ames, Iowa,
- 16 50010 USA

18

19

- \*Correspondence: Héctor Tejero-Cicuéndez cicuendez93@gmail.com
- 20 Keywords: Phenotypic Evolution, Morphospace, Allometry, *Pristurus* geckos

22 Short Title: XXX

23

- 24 Author Contributions: All authors collaboratively developed the concept and contributed to all
- 25 portions of this manuscript. HT-C, IM, and DCA performed the analyses. All authors approve of
- 26 the final product and are willingly accountable for any portion of the content.

27

8 Conflicts of Interests: The authors declare no conflicts of interest.

29

- $\textbf{Data Archiving:} \ \ Data \ \ are \ \ available \ on \ \ DRYAD \ (doi:10.5061/dryad.xwdbrv1f6 \ \ (Tejero-Cicu\'endez)$
- et al. 2021b)). R-scripts are available at XXX.

- Acknowledgments: We thank XYZPDQ... This work was sponsored in part by XXX (to SC)
- DCA was funded in part by National Science Foundation Grant DBI-1902511.

# 35 Abstract

36 asdf

### 37 Introduction

Understanding how phenotypic diversity evolves, and elucidating the forces that generate and maintain this diversity, are major goals in evolutionary biology. Because adaptive evolution is the product of natural selection, changes in ecological selection pressures are expected to affect the evolutionary trajectory of phenotypic traits that facilitate an organism's survival in their habitat. Evolutionary theory predicts that differing habitats will exert unique ecological selection pressures on organisms, resulting in associations between ecological and phenotypic traits. Indeed, species inhabiting differing habitats often display functional, behavioral, or phenotypic differences, that have presumably been the result of adaptive diversification in their respective ecological habitats (Collar et al. 2010; Kaliontzopoulou et al. 2015; Price et al. 2015; Martinez et al. 2021; Kolmann et al. 2022).

48

One possible evolutionary outcome of ecological specialization is that organisms inhabiting similar environments display common phenotypic characteristics. When such patterns occur repeatedly (e.g., Losos 1992; Schluter and McPhail 1992), this convergent evolution is treated as strong evidence of adaptation. Indeed the ecomorphological paradigm (sensu Arnold 1983) is predicated, in part, on such cases, which emphasize the strong association between the phenotypic traits that organisms display (morphological, behavioral, or physiological), and the ecological characteristics of their habitat that mediate organismal performance. In vertebrates, ecomorphological trends have been well studied in numerous taxonomic groups, and include the emblematic 'ecomorphs' of Caribbean Anolis lizards that exploit different microhabitats (Losos 1992, 2009; Mahler et al. 2013), differential beak morphology in species of Darwin's finches (Schluter and Grant 1984; Grant and Grant 2006; Reaney et al. 2020), the recurring phenotypes of African lake cichlids across ecological regimes (Albertson and Kocher 2001; Urban et al. 2022), and the distinct body forms of freshwater fishes in benthic benthis and limnetic habitats (Jastrebski and Robinson 2004; Berner et al. 2008; Stuart et al. 2017) among others.

63

However, while the patterns of morphological differences in distinct ecological contexts have

been well documented, less-well understood is how this differentiation has been influenced by the covariance between body parts resulting from body size variation (i.e., allometry). It has long been recognized that the interrelationships among traits can have a strong influence on how phenotypic evolution proceeds, as trait correlations influence the degree to which phenotypic variation is exposed to selection (Wagner and Altenberg 1996). Thus, the integration among traits can constrain phenotypic change in certain directions, or enhance variation along other phenotypic axes (Schluter 1996; Wagner and Altenberg 1996; Wagner and Zhang 2011; Klingenberg and Marugán-Lobón 2013; Goswami et al. 2014, 2016; Felice et al. 2018; Navalón et al. 2020). Further, because nearly all linear traits covary strongly with overall body size (Jolicoeur 1963; Bookstein 2022), allometric trends could be considered the quintessential measure of phenotypic integration. Thus, identifying whether allometric patterns differ across habitats, and how such patterns of trait covariation affect ecomorphological trends among species utilizing those habitats, remains an important question worthy of investigation.

78

The Afro-Arabian geckos in the genus *Pristurus* afford the opportunity to elucidate the interdigitating effects of allometry and habitat specialization on clade-level patterns of phenotypic diversity.

Prior work on this system (Tejero-Cicuéndez et al. 2021a) revealed that the colonization of ground habitats has been a trigger of morphological change, specifically reflected in an increase in body size and shape disparity. Interestingly, some ground-dwelling species are among the largest of the genus and also show increased relative head sizes and limb proportions, while some other species with this ecological specialization have evolved to be among the smallest of the group. Additionally, among the species exploiting rocky habitats (the most common ecological feature in *Pristurus*), there are also species with both considerably large and small body sizes (Tejero-Cicuéndez et al. 2021a). What remains unexplored, however, is how the evolution of body shape is related to differences in body size and whether habitat specialization has an impact in this shape-size relationship.

90

In this study, we employed a combination of multivariate morphometric and phylogenetic comparative analysis to interrogate macroevolutionary patterns of evolutionary allometry in *Pristurus* geckos of Afro-Arabia. Using phenotypic, phylogenetic, and ecological data, we first characterized allometric trends in body form in the group, to discern the extent to which allometric patterns differed across species occupying distinct ecological habitats. We then examined changes in allometric trends across the phylogeny, and linked these patterns to overall phenotypic integration, diversification in morphospace, and habitat utilization among taxa. Overall our results demonstrate that the interplay between ecological specialization and differing allometric trajectories in species with disparate body size may have a determinant role in shaping the phenotypic evolution and hence in adaptive dynamics in this clade.

#### 101 Materials and Methods

### 102 Data

We used a combination of phenotypic, phylogenetic, and ecological data to characterize and evaluate 103 intra- and interspecific allometric trends. The data utilized here were obtained from our prior 104 work on this system (Tejero-Cicuéndez et al. 2021a, 2022), and are briefly described here. First 105 we used a time-dated, molecular phylogeny that included all members of the genus Pristurus, 106 including several currently undescribed taxa. The tree was estimated in a Bayesian framework, 107 using five mitochondrial markers, six nuclear markers, and 21 calibration points (for details see 108 Tejero-Cicuéndez et al. 2022). Next we categorized each species as belonging to one of three 109 ecological groups (ground, rock, or tree), based on descriptions of habitat use found in the literature 110 (see Tejero-Cicuéndez et al. 2021a). Finally, we obtained a phenotypic data set containing body size 111 (snout-vent length: SVL) and eight linear measurements (Figure 1) that described overall body form: 112 trunk length (TrL), head length (HL), head width (HW), head height (HH), humerus length (Lhu), 113 ulna length (Lun), femur length (Lfe), and tibia length (Ltb) (Tejero-Cicuéndez et al. 2021a). We 114 restricted our study to those species represented by nine or more individuals; resulting in a dataset 115 of 687 individuals from 25 species (invidivuals per species:  $\mu = 27$ ; min = 9, max = 56). Species in 116 the phenotypic dataset were then matched to the phylogeny, which was subsequently pruned to 117 arrive at the final topology. All measurements were log-transformed prior to statistical analyses. 118 Additional details regarding data collection and formal descriptions of each linear measurement may 119 be found in the original sources (see Tejero-Cicuéndez et al. 2021a, 2022). The data are found on 120 DRYAD: https://doi.org/10.5061/dryad.xwdbrv1f6 (Tejero-Cicuéndez et al. 2021b). 121

#### 22 Statistical and Comparative Analyses

We conducted a series of analyses to interrogate allometric trends, patterns of integration, and 123 macroevolutionary changes in allometry, relative to differentiation in body form. 124 characterized evolutionary allometry in the genus by performing a phylogenetic multivariate 125 regression of body form on size, using the species means as data. We then performed an analogous 126 procedure at the individual level, regressing body form on size using our entire dataset. From both 127 the species-level (phylogenetic) and the individual-level regression models, we obtained the set of 128 regression coefficients, and calculated the difference between them to describe the extent to which 129 patterns of allometry at the individual level were concordant with evolutionary allometric trends 130 across species.

132

Next we used the individual dataset to determine whether allometric trends in body form differed 133 across habitat groups. This was accomplished by performing a multivariate analysis of covariance, 134 with body size (SVL), habitat, and  $SVL \times habitat$  as model effects. Significance was evaluated 135 using 999 iterations of a permutation procedure, where residuals from a reduced model were 136 randomly permuted in each permutation (RRPP), model statistics were recalculated, and used to 137 generate empirical null sampling distributions to evaluate the observed test statistics (following 138 Freedman and Lane 1983; Collyer and Adams 2007; Collyer et al. 2015). We then compared the 139 multivariate allometric vectors for each habitat group by calculating pairwise differences in their 140 angular direction in morphospace, and evaluating these relative to empirical sampling distributions 141 obtained through RRPP (Collyer and Adams 2007; Adams and Collyer 2009; Collyer and Adams 2013). Patterns of multivariate allometry relative to body size were visualized via regression scores 143 (Drake and Klingenberg 2008) and predicted lines (Adams and Nistri 2010), based on the coefficients 144 and fitted values from the linear model described above.

146

We then examined changes in the size-corrected body shape proportions allometric trends across
the phylogeny. Here we treated the head dimensions and limb dimensions separately, as allometric
trends could potentially differ between these body regions due to differential functional or selective
constraints (Kaliontzopoulou et al. 2010). Because both the head and limb data were multivariate,

we first performed a partial least squares (PLS) analysis (Rohlf and Corti 2000) of the head traits versus SVL, and the limb traits versus SVL, to describe the direction of maximal covaration between each body region and size. Then, we measured the mean residuals of each species to the allometric trend inferred, which show if head and limbs proportions of species are greater or smaller than expected for their body size. The species residuals were then mapped on the phylogeny of Pristurus using a Brownian motion model of evolution, to qualitatively evaluate shifts in head and limbs proportionality across the phylogeny for the group.

Furthermore, we examined the relationship between evolutionary and static allometry. 158 Species-specific PLS scores from each analysis were obtained, and species-specific slopes de-159 scribing the extent of head and limb allometry within each species were extracted from an 160 analysis of covariance modeled as:  $PLS1_{head} \sim SVL * species$  and  $PLS1_{limb} \sim SVL * species$ 161 respectively. This The species-specific allometric slopes (static allometry) were then compared 162 to the evolutionary allometric slopes expected according to their habitat by calculating the 163 correlation and angle among trends. We also compared species static allometry slopes among 164 species and visualized how this slopes changed throughout the were then mapped on the phylogeny 165 of *Pristurus* under using a Brownian motion model of evolution, to qualitatively evaluate shifts 166 in allometry across the phylogeny for the group (for a similar approach see Adams and Nistri 2010). 167

168

Next, because allometry describes the extent to which traits covary with size and with each other 169 (i.e., integration), we conducted an analysis of integration. Here we characterized the extent of 170 morphological integration in body form for individuals within each habitat group. Integration 171 was estimated by summarizing he dispersion of eigenvalues of the trait covariance matrix (sensu 172 Pavlicev et al. 2009). This measure  $(V_{rel})$  was subsequently converted to an effect size (a Z-score), 173 which quantified the strength of morphological integration (see Conaway and Adams 2022). We 174 then performed a series of two-sample tests to compare the strength of morphological integration 175 across habitat groups. Additionally and for comparison, we repeated these analyses on the set of 176 size-standardized trait data, found as a set of shape ratios (sensu Mosimann 1970) where each trait 177 was divided by body size (Supplemental Information). 178

Finally, to relate within-species allometric trends with patterns of phenotypic diversification in the 180 group we generated a phylomorphospace, based on the size-standardized species means obtained from a phylogenetic regression (see Tejero-Cicuéndez et al. 2021a). Here, phenotypic similarities 182 among species, relative to their phylogenetic relationships and habitat affiliations, were observed. 183 A similar phylomorphospace was constructed for species means not corrected for body size, and the phenotypic disparity among species means in each habitat was calculated and subsequently 185 compared (Supplemental Information). All analyses were conducted in R 4.2.1 (R Core Team 2022), 186 using RRPP version 1.3.1 (Collyer and Adams 2018; Collyer and Adams 2022) and geomorph 4.0.4 187 (Baken et al. 2021), and scripts written by the authors (available at XXX). 188

#### $_{\scriptscriptstyle{189}}$ Results

197

Using phylogenetic regression, we found significant evolutionary allometry in body form across species ( $N_{sp} = 25$ ; F = 217.9; Z = 5.53; P < 0.001). Likewise, when allometry in body form was examined across individuals, a similar pattern was observed ( $N_{ind} = 687$ ; F = 7910.8; Z = 9.20; P < 0.001). Further, the vectors of regression coefficients between the two analyses were highly correlated ( $\rho = 0.94$ ) and were oriented in nearly parallel directions in morphospace ( $\theta = 1.49^{\circ}$ ). This revealed that the pattern of multivariate allometry across individuals was concordant with macroevolutionary trends of interspecific allometry among species of *Pristurus* across the phylogeny.

Our analyses also exposed significant differences in the allometry of body form among Pristurus 198 utilizing distinct habitats (Table 1). Here, comparisons of multivariate allometric vectors identified 199 that ground-dwelling *Pristurus* displayed a distinct allometric trend as compared with *Pristurus* 200 occupying both the rock and tree habitats (Table 2). In addition, allometric patterns in both 201 rock and tree habitats were similar to the multivariate line of isometry (Figure 2), while patterns of multivariate allometry in ground-dwelling *Pristurus* was decidely steeper. Inspection of the 203 regression coefficients for each trait (Supplemental Information) confirmed the steeper allometric 204 coefficients for all head and limb traits in ground-dwelling *Pristurus* as compared with rock 205 and tree-dwelling taxa, corroborating this result. Taken together, these findings implied that 206 larger individuals of ground-dwelling *Pristurus* species displayed proportionately larger heads and 207

limbs, as compared with large individuals in taxa utilizing other habitat types. A visualization of multivariate allometric trends (Figure 2) confirmed these statistical findings, and indicated that the allometric trajectory in ground-dwelling animals was more extreme as compared with either rock or tree-dwelling *Pristurus*.

212

Mapping the residuals of species into the phylogeny showed that large ground-dwelling species 213 display greater head proportions than large rock-dwelling species, who displayed smaller heads 214 relative to body size (Figure 3A). Conversely, the opposite pattern was observed when comparing 215 small species utilizing these habitats: ground-dwelling species show small relative head proportions 216 while rock-dwelling species show in general larger head proportions. In contrast, limb shape showed 217 more variable patterns. Although all large ground-dwelling species consistently show large relative 218 limb proportions, large rock-dwelling species are variable in this trait, with P. insignis exhibiting 219 large and P. insignoides small limb proportions. For small species, relative limb proportions seem 220 to be independent from habitat utilization, since there are differences in limb residuals both within 221 rock- and ground-dwelling species (Figure 3B). Inspection of traitgrams thus revealed some degree 222 of discordance in patterns of allometry across body regions; a pattern evidenced by the relatively 223 low correlation between slopes representing head and limb allometry respectively ( $\rho = 0.42$ ). 224 We found that static allometric slopes (i.e., per-species slopes) are in general congruent with 225 evolutionary allometric slopes of each corresponding habitat (Figure XXX static allometry: 226 ADD LINE OF EVOLUTIONARY ALLOMETRY PER HABITAT), although there are some 227 cases of relatively low correlation and wide angles between static and evolutionary allometry 228 (Table XXX cor angles.csv). The comparison of the static allometry among species revealed 229 an important variability even within habitats (Figure XXX static allometry) Table XXX 230 static allometry species comparison.xlsx). When static allometric patterns were mapped on 231 the phylogeny, traitgrams elucidated that changes in allometric trends were not concentrated to 232 specific regions of the phylogeny (Supplemental Information). Rather, increases and decreases in 233 static allometry of both the head traits and the limb traits occurred repeatedly. 234 Examination of patterns of trait covariation revealed strong levels of morphological integration 235 within each habitat type ( $Z_{ground} = 3.97$ ;  $Z_{rock} = 3.72$ ;  $Z_{tree} = 2.15$ ). Further, two-sample tests 236

revealed that the strength of morphological integration was significantly greater in ground-dwelling Pristurus than either those utilizing rock ( $Z_{Groung-Rock} = 6.59$ ; P << 0.001) or tree habitats ( $Z_{Groung-Tree} = 11.17$ ; P << 0.001). Pristurus utilizing tree habitats displayed the lowest levels of integration, which were also significantly less than in the rock habitat ( $Z_{Rock-Tree} = 7.19$ ; P << 0.001). When size was accounted for in the data, levels of integration dropped considerably, though the overall pattern and differences among habitat groups remained the same (Supplemental Information).

244

When allometric patterns were mapped on the phylogeny, traitgrams elucidated that changes in 245 allometric trends were not concentrated to specific regions of the phylogeny (Figure 3). Rather, 246 increases and decreases in allometry of both the head traits and the limb traits occurred repeatedly 247 (see also Supplemental Information). When these patterns were viewed relative to body size, 248 large ground-dwelling species tended to display steeper head allometry as compared to large 249 rock-dwelling species, who displayed shallower head allometry (Figure 3A). Likewise, a similar 250 pattern was observed when comparing small species utilizing these habitats. In contrast, large 251 ground-dwelling species exhibited intermediate patterns of limb allometry, and small species in 252 general were more variable in such patterns (Figure 3B). Inspection of traitgrams thus revealed 253 some degree of discordance in patterns of allometry across body regions; a pattern evidenced by 254 the relatively low correlation between slopes representing head and limb allometry respectively 255  $(\rho = 0.42)$ . 256

257

Finally, when body shape differentiation in *Pristurus* was viewed in phylomorphospace (Figure 258 4), we found broad overlap among habitat groups, though arboreal (tree-dwelling) species were 259 somewhat more separated in morphospace. Rock-dwelling species occupied a slightly larger region of 260 morphospace as compared with the other groups, though this pattern was not statistically significant 261 (Supplemental Information). Intriguingly, when viewed in relation to body size, large Pristurus 262 species were not localized to a particular region of morphospace, nor were smaller species. Instead, 263 the largest rock-dwelling species were found in close proximity to the smallest ground-dwelling 264 species, indicating that they were similar in overall body shape. Likewise, the smaller rock-dwelling 265

species were found close to large ground-dwelling species in morphospace, indicating they displayed similar body shapes as well. Thus, synthesizing the patterns revealed in the phylomorphospace with those identified in our earlier analyses revealed a complex interplay between body shape, body size, habitat use, morphological integration, and multivariate allometry; where species with similar body shapes displayed differing overall size, were found in distinct habitats, and exhibited different allometric trends.

#### 2 Discussion

273

274

275

276

278

279

280

281

282

283

284

285

286

287

288

289

290

291

292

- First paragraph; restate topic/questions (see last par of Intro); summarize general findings . . .
  - something about linking allometry and phenotypic diversification?

The relationship between certain phenotypic traits and the organisms' environment is a central paradigm in evolutionary biology. In this context, disentangling the causes of phenotypic differentiation is essential to understand how natural selection operates. In this study, we evaluated the role of potential drivers of body shape differentiation in the geckos of the genus Pristurus. In particular, we investigated how the interplay of ecological specialization, phenotypic integration and allometric trends have shaped the morphological evolution in this radiation of Afro-Arabian geckos. Our results show that allometric trends and integration patterns are different across habitats, with ground-dwelling species having the steepest multivariate allometric slope and also the strongest morphological integration. These patterns are also different across body parts, with decoupled trends between head and limb proportions. Additionally, we found that changes in static allometric trends are not restricted to specific regions of the phylogeny, but rather they show multiple independent increases and decreases following common dynamics within habitat groups. Overall, these results suggest that the interplay between allometric and integration patterns is a fundamental factor to explain the morphological evolution across a variety of habitats, which is consistent with the theoretical expectation that different ecological contexts impose distinct selective pressures triggering phenotypic change.

• result 1: allometry; overall trend among species nearly identical to that among individuals.

Patterns of multivariate allometry in body form calculated from individuals were found to be nearly identical to those calculated from per-species means in Pristurus geckos. Specifically, the vectors of regression coefficients of the two analyses are virtually parallel ( $\theta = 1.49^{\circ}$ ), indicating that the evolutionary allometry is not substantially different whether measured with individual measurements or with species means in this genus.

293

294

295

296

297

298

299

300

301

302

303

304

305

306

307

308

309

310

311

312

313

314

315

316

317

318

319

320

??We also explored patterns of static allometry to compare them among species and with general trends of evolutionary allometry.

Thus, patterns of static and evolutionary allometry are consistent in *Pristurus*... Something else about this

• result 2: Allometry differs among habitat groups: 'steeper' allometry in Ground-dwelling (implication: proportionately larger heads and longer limbs in species at larger body sizes).

When we compared multivariate allometric slopes of species occupying different habitats, we found that, while rock-dwelling and arboreal species do not significantly differ from the isometric trend, ground-dwelling species have a steeper slope which is statistically different from isometry. This means that large ground-dwelling *Pristurus* present proportionally larger heads and longer limbs relative to other large species, while small species in the ground have proportionally smaller heads and shorter limbs (Figure residuals traitgrams). This is consistent with previous results on the morphological evolution of Pristurus (Tejero-Cicuéndez et al. 2021), where large ground species were indeed found to have also proportionally large heads and long limbs. This suggests that the segregation in body size and shape through differential allometric relationships across habitats responds to adaptive dynamics concerning the colonization of ground habitats, and perhaps with a particular interest of hard ground environments inhabited by the largest ground-dwelling species (including the largest of the genus, P. carteri), which has already been suggested to be the main driver of the morphological evolution in this genus (Tejero-Cicuéndez et al. 2021). This points toward the existence of a specialized form of Pristurus geckos adapted to hard grounds (e.g., some definition of hard ground vs. soft grounds??), illustrating the ecomorphological relationships in the genus with a rather conspicuous 'ecomorph' (see Figure X for an example of the hard-ground

ecomorph, P. carteri).

• result 3: relationship between evolutionary and static allometry, evolution of static allometry mapped in the phylogeny...

There is no general consensus about the relationship between the three types of allometry: ontogenetic (allometry during the development), static (allometry among individuals at the same developmental stage), and evolutionary (allometry across populations or species). This, in turn, is reflected in the broadly ambiguous interpretation of allometric patterns in the literature, with an often uncertain distinction between allometry as an evolutionary constraint and allometry as functional optimization resulting from natural selection (Pélabon et al. 2014, Voje et al. 2014). Even though testing these alternative hypotheses is beyond the scope of this work, our results do suggest that static and evolutionary allometry are very similar in *Pristurus* geckos, which could be explained by low evolvability of allometry, but also by the effect of relatively homogeneous selective pressures at different scales. Further analyses, for instance including a broader phylogenetic context and developmental assessments, are needed to illuminate the relationships between different levels of allometric trends. **EXTEND THIS** 

- result 4: Morphological integration differs among habitat groups. Strongest in ground-dwelling; weakest in tree-dwelling. SOME MEANING (combined with allometric trend implies that patterns of trait covariation are more constrained within ground-dwelling.... Thus, differences in body form are most likely found along this primary axis... (harken to Schluter evolution along lines of least resistance)
  - Additionally, rank-order of magnitude of integration across habitat groups corresponds with the range of body sizes in each: ground-dwelling display the largest size-range, while tree-dwelling the least (Supp. Information). On the one hand this matches the expectation that much of the integration observed in *Pristurus* is the result of allometric trends.... And the fact that levels of integration drop so precipitously when data are size-standardized are in accord with this interpretation. Nevertheless, when size is accounted for, the rank-order of magnitudes of integration remain the same, implying that ground-dwelling *Pristurus* are still relatively constrained in patterns of

trait covariation as compared with the other two groups.

350

351

352

353

354

355

356

357

358

359

360

361

362

363

364

365

366

367

368

369

370

371

372

373

374

375

376

377

349

- This notion was further supported when viewing the phylomorphospace of the species means not adjusted for size (SI). Here (and not surprisingly), PC1 is dominated by size, with small species at one end and larger species at the other. More importantly however, is that the disparity among species utilizing different habitats differed significantly in this space. Here, ground-dwelling displayed significantly greater phenotypic disparity than did the other groups (SI).

Similarly, when analyzing patterns of morphological integration, we found important differences among habitat groups: ground-dwelling species present the strongest integration, which in turn is weakest in arboreal species. Morphological integration occurs when different body parts coevolve, and has been suggested as an evolutionary constraint, since it restricts the specific lines along which integrated structures are allowed to vary (REF). Weaker integration levels (i.e., modularity), on the contrary, might facilitate morphological evolution by allowing a less constrained exploration of the morphospace (REF). However, integration might also be interpreted as a potential driver of morphological change, since it may provide a phenotypic pathway through adaptive lines of least resistance that enable rapid evolutionary processes (Navalón et al. 2020). In this context, our results on allometry and integration suggest that patterns of trait covariation are more constrained in ground-dwelling species, such that their differences in body form are most likely found along this primary axis. The fact that ground species in *Pristurus* have been found to have the widest phenotypic disparity and highest rates of morphological evolution (Tejero-Cicuéndez et al. 2021) is consistent with the idea that integration patterns are acting to facilitate morphological evolution along lines of least resistance.

- result 54: morphospace: Thus there was a reciprocal relationship between body shape and body size across ground-dwelling and rock-dwelling species. SOMEHOW TIE THIS TO integration (DCA pondering this one)
- one interesting... head vs. (correlation of head vs. limb slopes: 0.42. Pretty low. Implies some sort of differential something here, resulting in distinct allometric patterns for these

two body regions. SImilar to Antigoni's work (and refs therein). IMPLICATION: tie this into integration/modularity. Less integrated across the whole organism, and more modular... Future studies should examine this.

Another insightful result is the low correlation between head and limbs in their allometric slopes, which implies different evolutionary trajectories for these two body regions. This is likely to happen when different parts of an organism are subjected to different functional pressures (e.g., head evolution might be mainly influenced by diet while limb evolution might respond more tightly to the substrate used by the species), resulting in a decoupling of their respective morphological change. Ultimately, the combination of selective pressures upon which organisms evolve may lead to differential levels of integration across different body parts, with certain structures coevolving in a similar (i.e., integrated) manner and others in a segregated (i.e., modular) way. This, in turn, may have fundamental implications for the extent of morphological diversification within clades, and can be key to describe the phenotypic divergence observed across the tree of life. Future and more in-depth studies on the evolution of different body parts in *Pristurus* and other lizards, including for instance finer phenotypic data and comprehensive ecological information, may allow for discerning the functional drivers of head and limb evolution.

• In conclusion... -Synthesizing these patterns together ... (summarize: steeper allometry, higher integration, greater disparity in body size and body form all in ground-dwelling species). Together the patterns uncovered in our study imply that phenotypic diversification among ground-dwelling *Pristurus* follows tightly along its allometric trajectory, as evidenced by the higher disparity and stronger morphological integration.... some reference back to Goswami 'fly in a tube' paper. Thus, *Pristurus body forms appear to diversify along* allometric\* lines of least resistance.... (Schluter ref again)

#### 402 References

- Adams, D. C., and M. L. Collyer. 2009. A general framework for the analysis of phenotypic trajectories in evolutionary studies. Evolution 63:1143–1154.
- Adams, D. C., and A. Nistri. 2010. Ontogenetic convergence and evolution of foot morphology
- in european cave salamanders (family: plethodontidae). BMC Evolutionary Biology 10:1–10.
- BioMed Central.
- Albertson, R. C., and T. D. Kocher. 2001. Assessing morphological differences in an adaptive
- trait: A landmark-based morphometric approach. Journal of Experimental Zoology 289:385–
- 410 403.
- Arnold, S. J. 1983. Morphology, performance, fitness. American Zoologist 23:347–361.
- Baken, E. K., M. L. Collyer, A. Kaliontzopoulou, and D. C. Adams. 2021. Geomorph 4.0
- and gmShiny: Enhanced analytics and a new graphical interface for a comprehensive
- morphometric experience. Methods in Ecology and Evolution 12:2355–2363.
- Berner, D., D. C. Adams, A.-C. Grandchamp, and A. P. Hendry. 2008. Natural selection
- drives patterns of lake-stream divergence in stickleback foraging morphology. Journal of
- Evolutionary Biology 21:1653–1665.
- Bookstein, F. L. 2022. Dimensions of morphological integration. Evolutionary Biology 49:342–
- 419 372.
- Collar, D. C., J. A. Schulte, B. C. O'Meara, and J. B. Losos. 2010. Habitat use affects
- morphological diversification in dragon lizards. Journal of Evolutionary Biology 23:1033—
- 1049.
- Collyer, M. L., and D. C. Adams. 2007. Analysis of two-state multivariate phenotypic change in
- ecological studies. Ecology 88:683–692.
- <sup>425</sup> Collyer, M. L., and D. C. Adams. 2013. Phenotypic trajectory analysis: Comparison of shape
- change patterns in evolution and ecology. Hystrix 24:75–83.
- 427 Collyer, M. L., and D. C. Adams. 2022. R: RRPP: Linear model evaluation with randomized
- residuals in a permutation procedure. Vsn. 1.3.1. R Foundation for Statistical Computing,
- Vienna, Austria.
- 430 Collyer, M. L., and D. C. Adams. 2018. RRPP: An r package for fitting linear models to

- high-dimensional data using residual randomization. Methods in Ecology and Evolution 9:1772–1779.
- Collyer, M. L., D. J. Sekora, and D. C. Adams. 2015. A method for analysis of phenotypic change for phenotypes described by high-dimensional data. Heredity 115:357–365.
- Conaway, M. A., and D. C. Adams. 2022. An effect size for comparing the strength of morphological integration across studies. Evolution 76:(In Press).
- Drake, A. G., and C. P. Klingenberg. 2008. The pace of morphological change: Historical transformation of skull shape in st bernard dogs. Proceedings of the Royal Society B:

  Biological Sciences 275:71–76.
- Felice, R. N., M. Randau, and A. Goswami. 2018. A fly in a tube: Macroevolutionary expectations for integrated phenotypes. Evolution 72:2580–2594.
- Freedman, D., and D. Lane. 1983. A nonstochastic interpretation of reported significance levels.

  Journal of Business & Economic Statistics 1:292–298.
- Goswami, A., M. Randau, P. D. Polly, V. Weisbecker, C. Verity Bennett, L. Hautier, and M. R. Sánchez-Villagra. 2016. Do developmental constraints and high integration limit the evolution of the marsupial oral apparatus? Integrative and Comparative Biology 56:404–415.
- Goswami, A., J. B. Smaers, C. Soligo, and P. D. Polly. 2014. The macroevolutionary consequences of phenotypic integration: From development to deep time. Philosophical Transactions of the Royal Society B: Biological Sciences 369:20130254.
- Grant, P. R., and B. R. Grant. 2006. Evolution of character displacement in darwin's finches.

  Science 313:224–226.
- Jastrebski, C. J., and B. W. Robinson. 2004. Natural selection and the evolution of replicated trophic polymorphisms in pumpkinseed sunfish (*Lepomis gibbosus*). Evolutionary Ecology Research 6:285–305.
- Jolicoeur, P. 1963. The multivariate generalization of the allometry equation. Biometrics 19:497–499.
- Kaliontzopoulou, A., M. A. Carretero, and D. C. Adams. 2015. Ecomorphological variation in male and female wall lizards and the macroevolution of sexual dimorphism in relation to habitat use. Journal of Evolutionary Biology 28:80–94.
- Kaliontzopoulou, A., M. A. Carretero, and G. A. Llorente. 2010. Intraspecific ecomorphologi-

- cal variation: Linear and geometric morphometrics reveal habitat-related patterns within

  Podarcis bocagei wall lizards. Journal of Evolutionary Biology 23:1234–1244.
- Klingenberg, C. P., and J. Marugán-Lobón. 2013. Evolutionary covariation in geometric morphometric data: Analyzing integration, modularity, and allometry in a phylogenetic context. Systematic Biology 62:591–610.
- Kolmann, M. A., F. P. L. Marques, J. C. Weaver, M. N. Dean, J. P. Fontenelle, and N. R. Lovejoy. 2022. Ecological and phenotypic diversification after a continental invasion in neotropical freshwater stingrays. Integrative and Comparative Biology 62:424–440.
- Losos, J. B. 2009. Lizards in an evolutionary tree: Ecology and adaptive radiation of anoles.

  University of California Press.
- Losos, J. B. 1992. The evolution of convergent structure in Caribbean *Anolis* communities.

  Systematic Biology 41:403–420.
- Mahler, D. L., T. Ingram, L. J. Revell, and J. B. Losos. 2013. Exceptional convergence on the macroevolutionary landscape in island lizard radiations. Science 341:292–295.
- Martinez, C. M., S. T. Friedman, K. A. Corn, O. Larouche, S. A. Price, and P. C. Wainwright.

  2021. The deep sea is a hot spot of fish body shape evolution. Ecology Letters 24:1788–1799.
- Mosimann, J. E. 1970. Size allometry: Size and shape variables with characterizations of the lognormal and generalized gamma distributions. Journal of the American Statistical
- Association 65:930–945.
  Navalón, G., J. Marugán-Lobón, J. A. Bright, C. R. Cooney, and E. J. Rayfield. 2020. The
  consequences of craniofacial integration for the adaptive radiations of darwin's finches and
  hawaiian honeycreepers. Nature Ecology & Evolution 4:270–278. Nature Publishing Group.

#### pre

- Pavlicev, M., J. M. Cheverud, and G. P. Wagner. 2009. Measuring morphological integration using eigenvalue variance. Evolutionary Biology 36:157–170.
- Price, S. A., S. T. Friedman, and P. C. Wainwright. 2015. How predation shaped fish: The impact of fin spines on body form evolution across teleosts. Proceedings of the Royal Society B: Biological Sciences 282:20151428.
- R Core Team. 2022. R: A language and environment for statistical computing. Version 4.2.1. R

- Foundation for Statistical Computing, Vienna, Austria. 490
- Reaney, A. M., Y. Bouchenak-Khelladi, J. A. Tobias, and A. Abzhanov. 2020. Ecological 491 and morphological determinants of evolutionary diversification in darwin's finches and their 492
- relatives. Ecology and Evolution 10:14020–14032. 493
- Rohlf, F. J., and M. Corti. 2000. Use of two-block partial least-squares to study covariation in 494 shape. Systematic Biology 49:740–753. 495
- Schluter, D. 1996. Adaptive radiation along genetic lines of least resistance. Evolution 50:1766– 496 1774. 497
- Schluter, D., and P. R. Grant. 1984. Determinants of morphological patterns in communities of 498 darwin's finches. The American Naturalist 123:175–196. 499
- Schluter, D., and J. D. McPhail. 1992. Ecological character displacement and speciation in 500 sticklebacks. The American Naturalist 140:85–108. 501
- Stuart, Y. E., T. Veen, J. N. Weber, D. Hanson, M. Ravinet, B. K. Lohman, C. J. Thompson, T. 502
- Tasneem, A. Doggett, R. Izen, N. Ahmed, R. D. H. Barrett, A. P. Hendry, C. L. Peichel, and 503
- D. I. Bolnick. 2017. Contrasting effects of environment and genetics generate a continuum 504 of parallel evolution. Nature Ecology and Evolution 1:158.
- Tejero-Cicuéndez, H., A. H. Patton, D. S. Caetano, J. Šmíd, L. J. Harmon, and S. Carranza. 506
- 2022. Reconstructing squamate biogeography in afro-arabia reveals the influence of a complex 507
- and dynamic geologic past. Systematic Biology 71:261–272. 508

- Tejero-Cicuéndez, H., M. Simó-Riudalbas, I. Menéndez, and S. Carranza. 2021a. Ecological spe-509 cialization, rather than the island effect, explains morphological diversification in an ancient 510 radiation of geckos. Proceedings of the Royal Society B: Biological Sciences 288:20211821. 511
- Tejero-Cicuéndez, H., M. Simó-Riudalbas, I. Menéndez, and S. Carranza. 2021b. Ecological 512 specialization, rather than the island effect, explains morphological diversification in an 513 ancient radiation of geckos. Dryad digital repository. (Doi:10.5061/dryad.xwdbrv1f6). 514
- Urban, S., J. Gerwin, C. D. Hulsey, A. Meyer, and C. F. Kratochwil. 2022. The repeated 515 evolution of stripe patterns is correlated with body morphology in the adaptive radiations of 516 east african cichlid fishes. Ecology and Evolution 12:e8568. 517
- Wagner, G. P., and J. Zhang. 2011. The pleiotropic structure of the genotype—phenotype map: 518 The evolvability of complex organisms. Nature Reviews Genetics 12:204–213. 519

Wagner, G., and L. Altenberg. 1996. Perspective: Complex adaptations and the evolution of evolvability. Evolution 50:967–976.

Table 1: Multivariate analysis of covariance describing variation in body form in Pristurus.

	Df	SS	MS	Rsq	F	Z	Pr(>F)
svl	1	516.036559	516.0365588	0.9203096	10188.69842	9.490057	0.001
habitat	2	6.218510	3.1092552	0.0110902	61.38957	9.322480	0.001
svl:habitat	2	3.974307	1.9871536	0.0070879	39.23464	7.077264	0.001
Residuals	681	34.491245	0.0506479	0.0615124			
Total	686	560.720622					

Table 2: Pairwise comparisons of multivariate allometry vectors. Effect sizes  $(Z_{\theta_{12}})$  based on pairwise differences in angular direction are below the diagonal, and their corresponding significance levels are above diagonal. Significant values in bold.

	Ground	Rock	Tree
Ground	0	0.001	0.001
Rock	6.872	0	0.261
Tree	3.657	0.649	0

## **Figures**

Figure 1. Linear Measurements used in this study. SVL = snout-vent length, TL = trunk length, HL = head length, HW = head width, HH = head height, Lhu = humerus length, Lun = ulna length, Lfe = femur length, Ltb = tibia length (for details see Tejero-Cicuéndez et al. 2021a).

Figure 2. Plot of regression scores and predicted lines representing the relationship between linear body measurements and size (SVL). Individuals are colored by habitat use: ground (beige), rock (dark purple), and tree (magenta). Isometric trend represented by the dashed line.

Figure 3. Traitgrams showing the evolution of body size (SVL) through time based on the
phylogenetic tree of *Pristurus*. Colors represent an evolutionary mapping of regression slopes
describing the relationship of (A) head morphology versus body size, and (B) limb proportions
versus body size (see text for descriptions). Species names are colored by habitat use: ground
(beige), rock (dark purple), and tree (magenta).

Figure 4. Phylomorphospace of *Pristurus*, based on residuals from a phylogenetic regression of body measurements on size (SVL). Species means are colored by habitat use: ground (beige), rock (dark purple), and tree (magenta). Large and small rock-dwelling and ground-dwelling are highlighted with darker colors to highlight their differentiation and relative positions in morphospace.

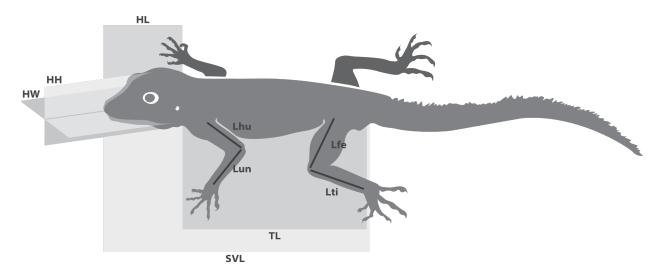


Figure 1: Linear Measurements used in this study. SVL = snout-vent length, TL = trunk length, HL = head length, HW = head width, HH = head height, Lhu = humerus length, Lun = ulna length, Lfe = femur length, Ltb = tibia length (for details see Tejero-Cicu'endez et al. 2021a).

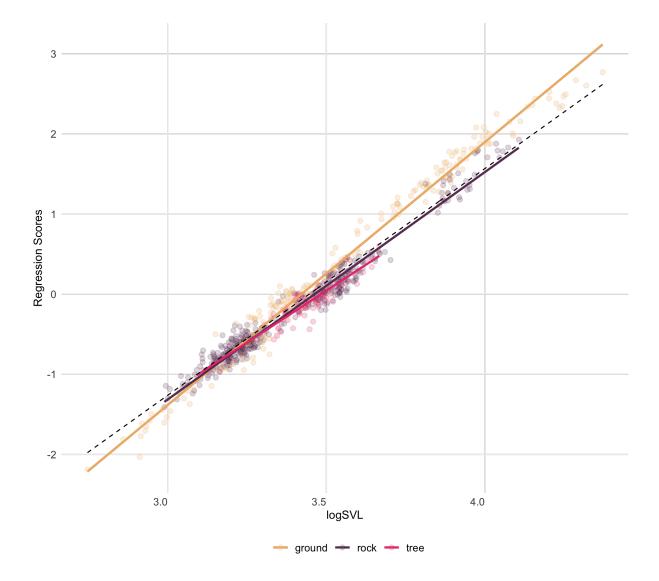


Figure 2: Plot of regression scores and predicted lines representing the relationship between linear body measurements and size (SVL). Individuals are colored by habitat use: ground (beige), rock (dark purple), and tree (magenta). Isometric trend represented by the dashed line.

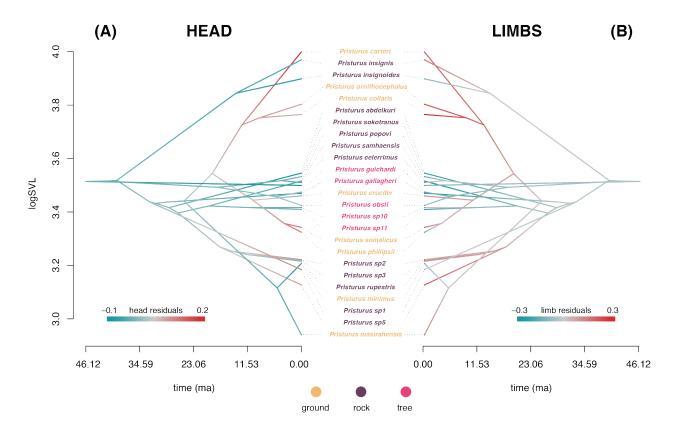


Figure 3: Traitgrams showing the evolution of body size (SVL) through time based on the phylogenetic tree of *Pristurus*. Colors represent an evolutionary mapping of regression slopes describing the relationship of (A) head morphology versus body size, and (B) limb proportions versus body size (see text for descriptions). Species names are colored by habitat use: ground (beige), rock (dark purple), and tree (magenta).

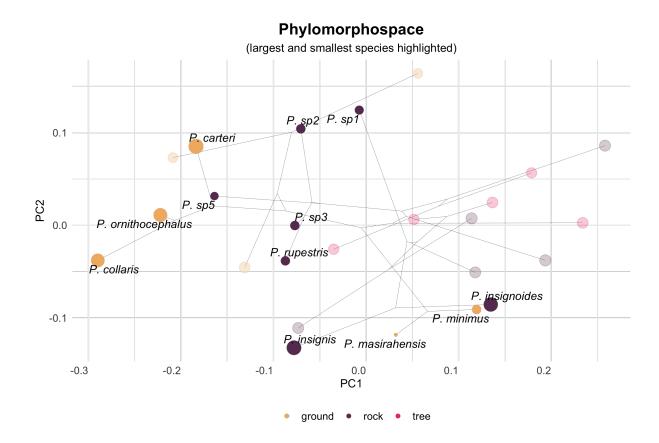


Figure 4: Phylomorphospace of *Pristurus*, based on residuals from a phylogenetic regression of body measurements on size (SVL). Species means are colored by habitat use: ground (beige), rock (dark purple), and tree (magenta). Large and small rock-dwelling and ground-dwelling are highlighted with darker colors to highlight their differentiation and relative positions in morphospace.