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7 Short Title: XXX

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- 20 of this manuscript. HT-C, IM, and DCA performed the analyses. All authors approve of the final product
- 21 and are willingly accountable for any portion of the content.

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

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- Data Archiving: Data are available on DRYAD (doi:10.5061/dryad.xwdbrv1f6 (Tejero-Cicuéndez et al.
- 2021b)). R-scripts are found in the Supplemental Information.

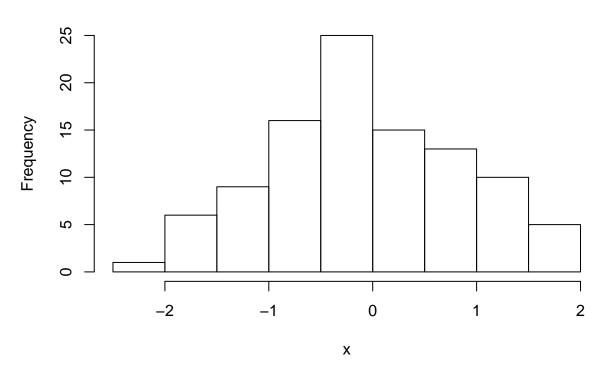
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30 Abstract

asdf

Histogram of x



 $_{32}$ an example

33 Introduction

- 34 some general paragraph on the evolution of phenotypic diversity
- when organisms colonize new and unique habitats, they are subjected to novel ecological selection pressures
- ₃₇ in those habitats. Often these selective pressures elicit changes in body form, as organisms adapt to their
- new habitats (examples: some comment on ecomorphs, etc.). . . . leads to so-called ecomorphs, with such
- well known examples in Anolis lizards, cichlid fishes, etc. It follows that ... Some comment on the fact that
- 40 clades living in diverse ecological conditions often display greater diversity in form and function (REFS).
- 41 However, while the above patterns have been well documented in a variety of vertebrate taxa, what remains
- 42 less known is how allometry plays a role in this phenotypic diversification. We know that XYZPDQ (about
- ⁴³ allometry). Then links to diversity..
- The Afro-Arabian geckos in the genus *Pristurus* afford the opportunity to elucidate the interdigitating effects
- 45 of allometry and habitat specialization on clade-level patterns of phenotypic diversity. Prior work on this
- system (Tejero-Cicuéndez et al. 2021a) has revealed that ... (sentence or 2 about your prior study, getting
- to diversity and ... Importantly, ... something about habitat. What remains unexamined however, is
- 48 XYZPDQ...
- In this study, we ...

50 Materials and Methods

Data

- 52 For this study, we used recently published phylogenetic, phenotypic, and ecological data of the species of
- the genus *Pristurus* including undescribed diversity (Tejero-Cicuéndez et al. 2021a). Briefly, these data
- consisted of: i) a phylogenetic tree of the relationships among all the *Pristurus* species with available genetic
- material; ii) an individual-level phenotypic dataset of linear measurements including body size (snout-vent
- below tended to be length, SVL) and several variables describing body shape: trunk length (TrL), head length (HL), head width
- 57 (HW), head height (HH), humerus length (Lhu), ulna length (Lun), femur length (Lfe), and tibia length
- 58 (Ltb); and iii) ecological data as a discrete character with three states (ground, rock, and tree) reflecting
- 59 the ground-dwelling, rock-climbing, or arboreal habits of each species. For all the analyses, we used only
- those species present in the phylogeny for which the available phenotypic data consisted of five or more

- 51 specimens. This resulted in a curated morphological dataset of 687 individuals from 25 species, with a mean
- 62 of 27 specimens per species, a minimum of nine and a maximum of 56. For more detailed information about
- these data and data collection, especially regarding morphological measurements, refer to the original source
- 64 (Tejero-Cicuéndez et al. 2021a).

65 Statistical Analyses

- To test the hypothesis...
- Mancova body ~ SVL*hab.gp
 - PW of slopes, and inspected reg. coefficients to identify biological trends
- Visualized multivariate regressions via regression scores (sensu Drake and Klingenberg 2008) and
 predicted lines (sensu Adams and Nistri 2010)
- Examine allometry phylogenetically.
- PLS of head vs. SVL and limb vs SVL. Obtained scores on 1st axis for each.
- 73 within-species regressions of Head.sc \sim SVL & limb.sc \sim SVL; obtained regression coefficients (slopes)
- mapped slopes on phylogeny under BM and generated traitgrams to identify changes in allometric relationships across the phylogeny
- Finally, to link allometric patterns with trends in phenotypic diversification we obtained sizestandardized species means, following procedures in H TC paper (residuals from phylo-regressions of
 traits on SVL, residuals). We then performed an ordination to obtain a phylomorphospace, where
 habitat types and species could be observed.

81 Results

82 Discussion

References

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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).

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

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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: rock (beige), ground (dark purple), and tree (magenta).

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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: rock (beige), ground (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.

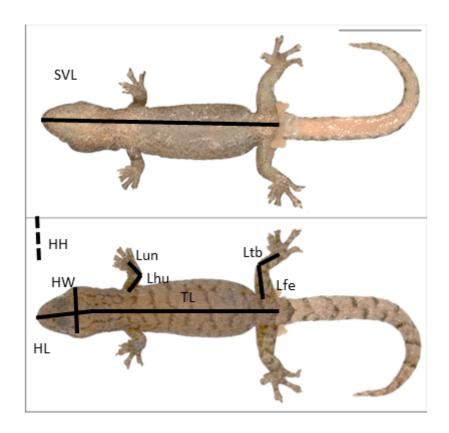


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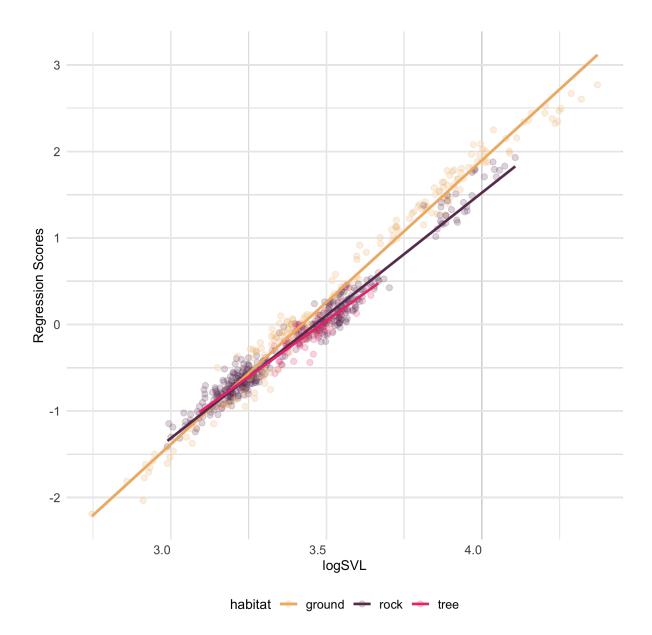


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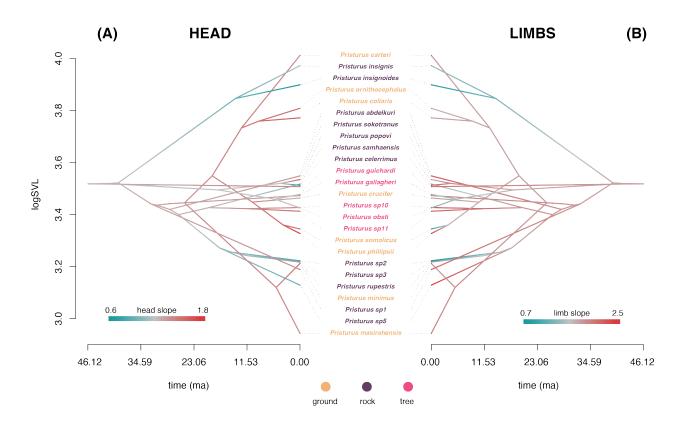


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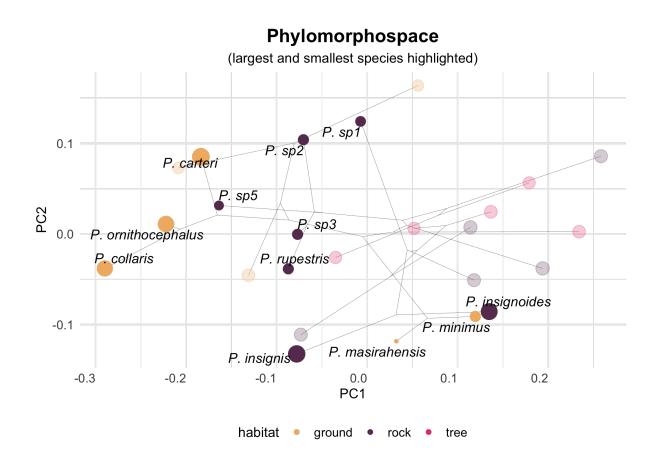


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