

RH: disparate views on disparity.

Disparate views on disparity.

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1

Abstract

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3

(Keywords: disparity)

INTRODUCTION

1: disparity is observed everywhere Biodiversity is not a smooth gradient of forms, functions, ecologies and behaviours but it is rather variable and discontinuous. This is at the base of Biological Sciences and is linked to the concept of disparity with the observation that some groups of species are more similar or dissimilar than others. This diversity in morphologies is in fact proper in nature and decoupled from taxonomic diversity: some groups may exhibit low diversity of species by high diversity of shapes (Ruta et al., 2013; Hopkins, 2013, or the other way around).

2: historical evolution of disparity Morphological diversity (hereafter disparity) is based on the observed the (dis)similarities between the morphologies of groups of organisms. In palaeobiology, this concept, as a tool for macroevolutionary studies, stems from several seminal papers from the 90s describing and questioning how the diversity of life forms arose and evolved (Gould, 1989, 1991; Briggs et al., 1992; Wills et al., 1994; Foote, 1994, 1996; Jernvall et al., 1996; Foote, 1997). In parallel, disparity analysis have been extended with modifications to macroevolutionary analysis based on continuous traits or geometric morphometric. The former usage of data was popularised following Harmon et al. (2003) and its implementation in *geiger* (Harmon et al., 2008); while the latter was popularised following (Zelditch et al., 2012) and implementations in *geomorph* (Adams and Otárola-Castillo 2013; Adams et al. 2017 and precursory work from Claude 2008).

3: the great popularity leads it to be used in many methods

Prentice et al. (2011) define disparity as: “a term widely (albeit not always consistently) used to describe the range of forms in a group of organisms, or the difference among different body plans”. Disparity is really popular (Fig. 1):

WHY DO WE NEED DISPARITY

These points should be developed during the meeting

Which macroevolutionary questions are tackled through disparity analysis?

- Bodyplan evolution through time
- Why does some morphologies exist and other not?
- tempo and mode of innovation of genuine novelty
- evolutionary competition
- effects of mass extinctions
- ecological niches

To answer these question, disparity has been used as a proxy for the biological increase/decrease of distinguishable morphological variation. These proximal multidimensional spaces have been described as the morphospace, the eco-space, the functional space, the cladisto space, the developemental space, the allometric space, etc (see Hopkins and Gerber, 2017, and references therein). This multitude of multidimensional spaces arising from the observation of variable and discontinuous

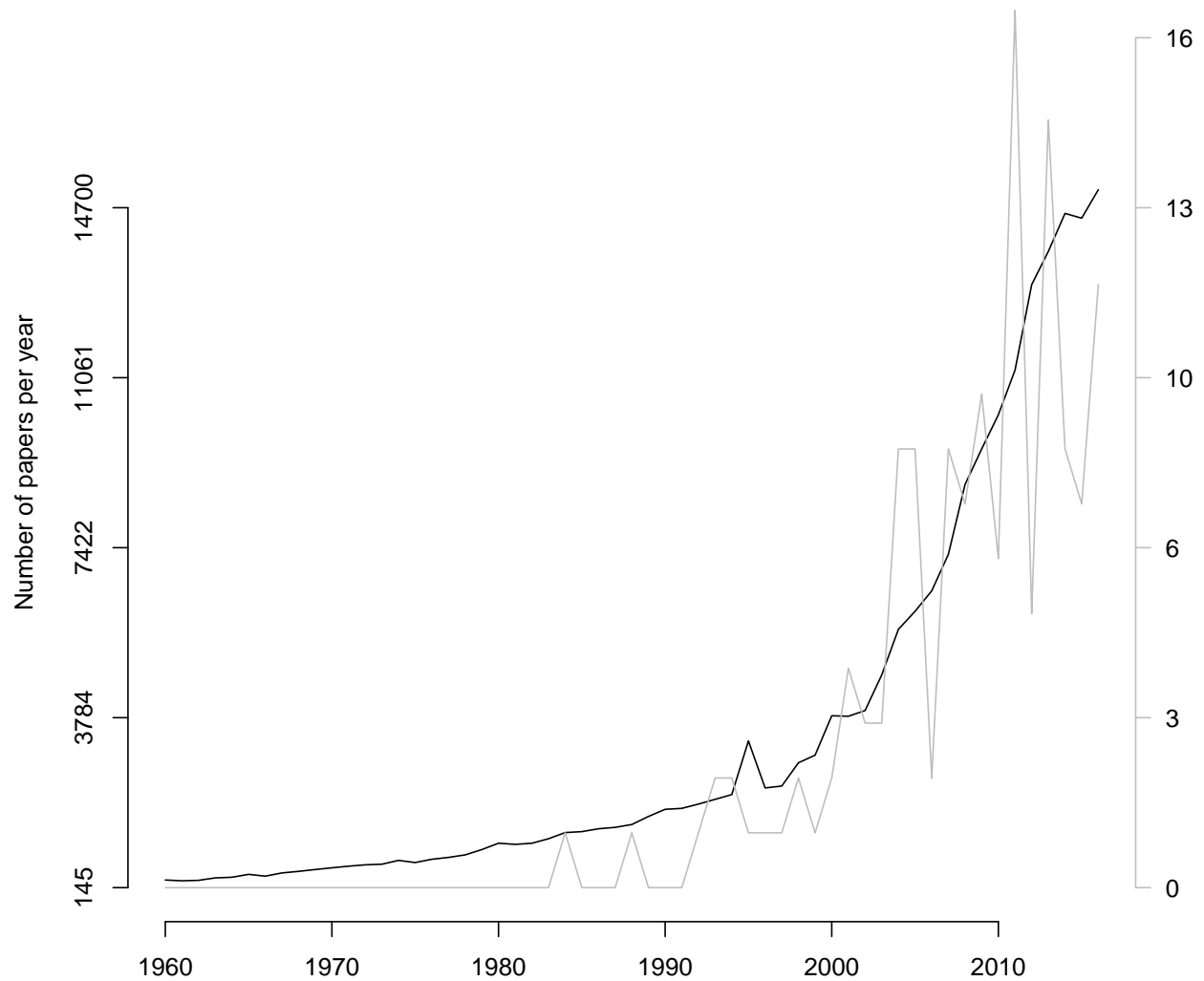


Figure 1: Number of papers on Google Scholar matching the search “morphological disparity” per year. In black, the match is in the paper and in grey, in the title. We collected the number of matches per year from 1960 to 2016 in Google Scholar for the terms "morphological disparity" both in the text (fuzzy matching) or in the title (exact matching). The data was collected on the 1st of November 2017.

43 morphology in Life arises the question of their equivalence and their pertinence. Are
44 the description of these spaces all equivalent? If disparity is the method or the metric
45 describing these spaces then what *is* disparity?

46 WHAT *is* DISPARITY (REALLY)?

47 Disparity can describe either the metric (disparity index (Hopkins and Gerber,
48 2017) or disparity metric (Wills, 2001)) or the whole pipeline (Lloyd, 2015; Zelditch
49 et al., 2012)

50 *Data collection methodology*

51 To assess the usage of disparity in different published studies, we collected
52 methodological data from the 500 first Google Scholar results for the key words
53 “morphological disparity” per order of appearance (accessed on the 1st of November
54 2017). For the 230 relevant papers among the 500 matches, we collected the following
55 methodological data:

- 56 • What was the focal biological group?
- 57 • What kind of data was measured (e.g. landmarks, discrete data, etc.)?
- 58 • Was data collected on the full organism or not?
- 59 • How was the morphospace explicitly defined (e.g. PCA, PCO, MDS, etc.)?

- How was the disparity metric(s) explicitly defined?
- Which statistical test was applied to test the disparity related hypothesis?
- Was phylogeny taken into account or not?

We used only the explicit definition of the morphospace and the disparity metric(s) in this search since a few number of papers had a vague definition of either or both (e.g. a disparity metric was measured but not described anywhere in the paper).

The remaining 270 matches where disparity was mentioned but not measured fell in the following categories: papers out of topic, papers mentioning morphological disparity without measuring it, review papers, papers not accessible (either through a broken link or a paywall) or referenced citation without the paper (as a Google Scholar match).

To reduce the amount of categories for the 230 recorded methods, we concatenated different methods in a smaller number of categories (see supplementary materials and https://github.com/TGuillerme/Disparity_Working_Group/blob/master/Analysis/data_cleaning.Rmd).

[ADD Biological group, full organism and phylogeny in the supplementary]

These points should be developed during the meeting

Disparity data

The data used for disparity methods comes from three main sources:

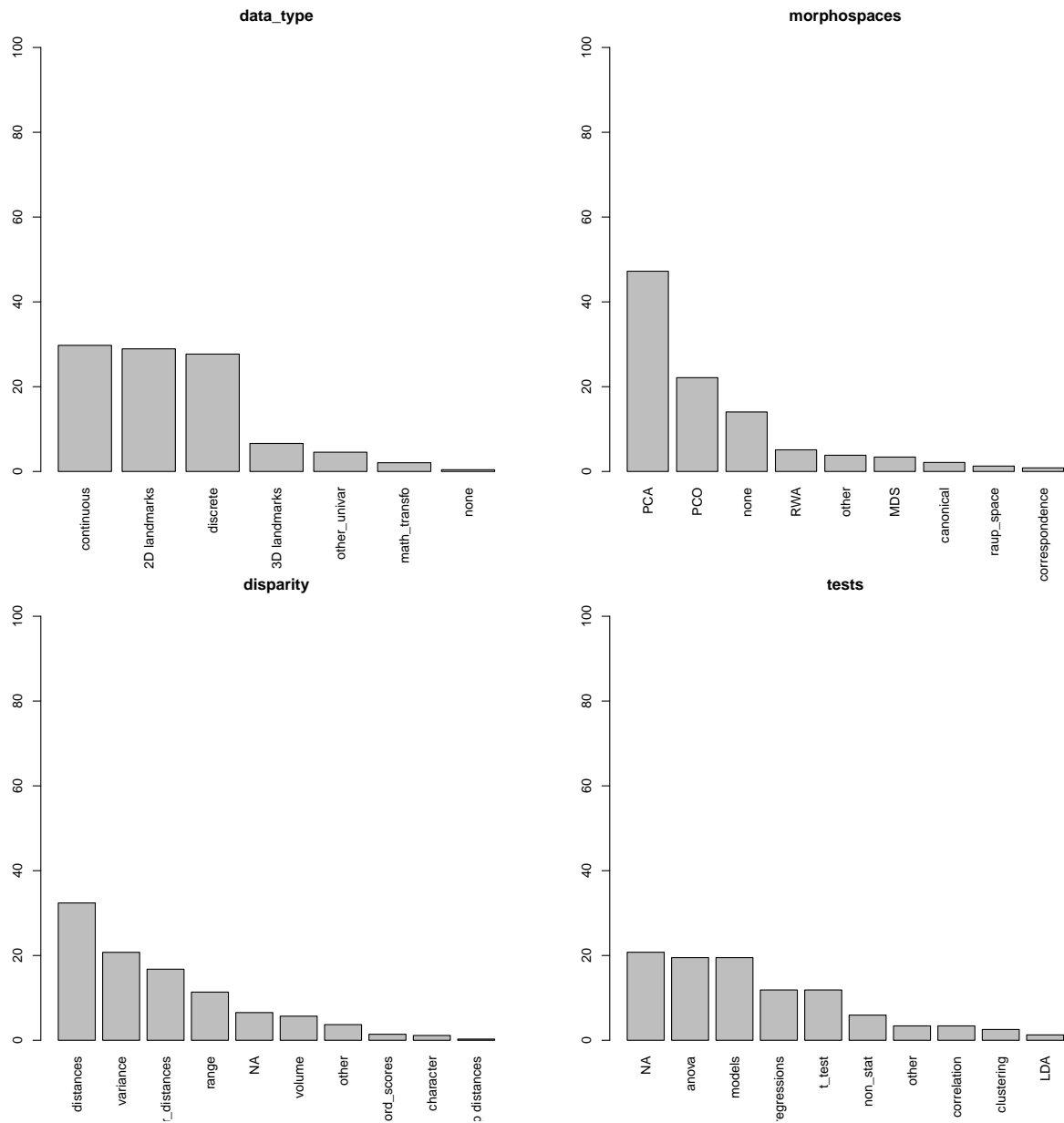


Figure 2: Disparity methods proportional usage: (1) Data type: which input data was used (...); (2) Morphospace: how was the morphospace obtained (...); (3) disparity: what type of disparity metric was calculated (...); (4) test: what type of test was applied (...).

- Continuous data such as limb or body measurements (Slater et al., 2010).
- Discrete morphological characters (sometimes referred to as “Cladistic” characters; Brusatte et al., 2008).
- Geometric morphometric data which is generally the Procrustes transformation of 2D or 3D landmarks (Cooney et al., 2017).

However, data was by far not limited to these three categories (e.g. colours Maia et al. 2013; metabolic rate Nespolo et al. 2016 or chemicals signal García-Roa et al. 2017).

Morphospaces

- PCAs [CITE].
- PCOs [CITE]
- Pairwise distance matrices (no ordination; Harmon et al., 2003).

But also RWA [CITE] or Raup-spaces [CITE].

Disparity metrics

Throughout the 230 analysed papers, we found 103 unique combinations of metrics!

- Distances measurements between species or from a certain point in the morphospace [CITE].

- Ranges and variances of each axis of the morphospace Wills (2001); Ciampaglio et al. (2001)
- Distance between species based on the pairwise distance matrix (not the ordinations) [CITE].

But also metrics based on the volume [CITE], the characters dissimilarity [CITE] or the coordinates of some axis of the ordination [CITE]

Disparity hypothesis

Describing the many outputs (what and how is it tested):

- Variance based (ANOVA, etc.) [CITE].
- Correlation based [CITE].
- Regression based (PGLS, etc.) [CITE].

But also discriminant analysis [CITE] and clustering [CITE].

The three main disparity analysis

From the collected data, we can highlight the use of three main different disparity analysis with their associated data/morphospace/metric/tests and related to specific methodological implementations.

- **The “Claddis” approach:** this group of methods uses discrete morphological data (sometimes referred to as “Cladistic” characters) for the full organisms to build a PCO from the organism’s pairwise distance as a morphospace. Disparity is then often measured as a variation of the ordinated matrix dimensions’ variances or ranges (e.g. the sum of variance or/and the sum of ranges). Hypothesis are often tested using multivariate ANOVAs on the pairwise distance matrix or by simply comparing the confidence intervals overlap of the disparity from different groups.
- **The “geomorph” approach:** this group of method is based on landmark data (2D or 3D) on parts of the organism studied usually the skull) and use a Procrustes transformation of the landmarks that are then directly ordinated using a PCA (but sometimes RWA). Disparity is often measured as a distance metric (e.g. the distance between the species and a point in the morphospace such as its centroid). Hypothesis are then tested using ANOVA type tests with usually no phylogenetic correction (although phylogeny is sometimes used to correct the morphospace).
- **The “dtt” approach:** this method can directly use continuous or discrete data for the full organism without any ordination (but not necessarily), and will measure disparity as the average pairwise distance between species (whether euclidean or any other type of distance). Hypothesis of higher/lower disparity can then be measured using null evolutionary models.

Of course some studies use a combination of these three methods or none of them at all!

Also, among each category XX% of studies use multiple approaches.

[ADD Biological group, full organism and phylogeny in the supplementary]

EXPANDING DISPARITY

How to compare disparity between groups? Is disparity relative?

How to compare disparity between methods?

Can we really say things about competition when looking at disparity in a single group?

Are all characters equal? Character contingency suggests otherwise

What is the relationship between disparity and tree shape?

Do we have a null model for investigating disparity?

Can we really say things about competition when looking at disparity in a single group?

Disparity in other fields

In ecology, disparity bears strong parallels with β -diversity in ecology (a measure of ecological communities (dis)similarity): one biological observation described by a vast array of metrics (Baselga, 2010; Anderson et al., 2011; Donohue et al., 2016).

CONCLUSION

A quick guideline for good disparity analysis:

Maybe we need something like in Parham et al 2012 (Best Practices for Justifying Fossil Calibrations): an easy an identifiable description of the pipeline containing: 1) the type of data, 2) the morphospace and 3) the metric?

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