

Tracing Disagreement in Taxonomy

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Outline

1. Briefly: the problem
2. Construction of the corpus
3. Feature analysis
4. Preliminary results
5. Future ideas

Take-home message: Disagreement in taxonomy seems to be unevenly distributed; to understand it we'll have to analyze the literature empirically

The problem

Briefly: taxonomy is often taken to be affected by a widespread problem of **rampant uncertainty and disagreement**



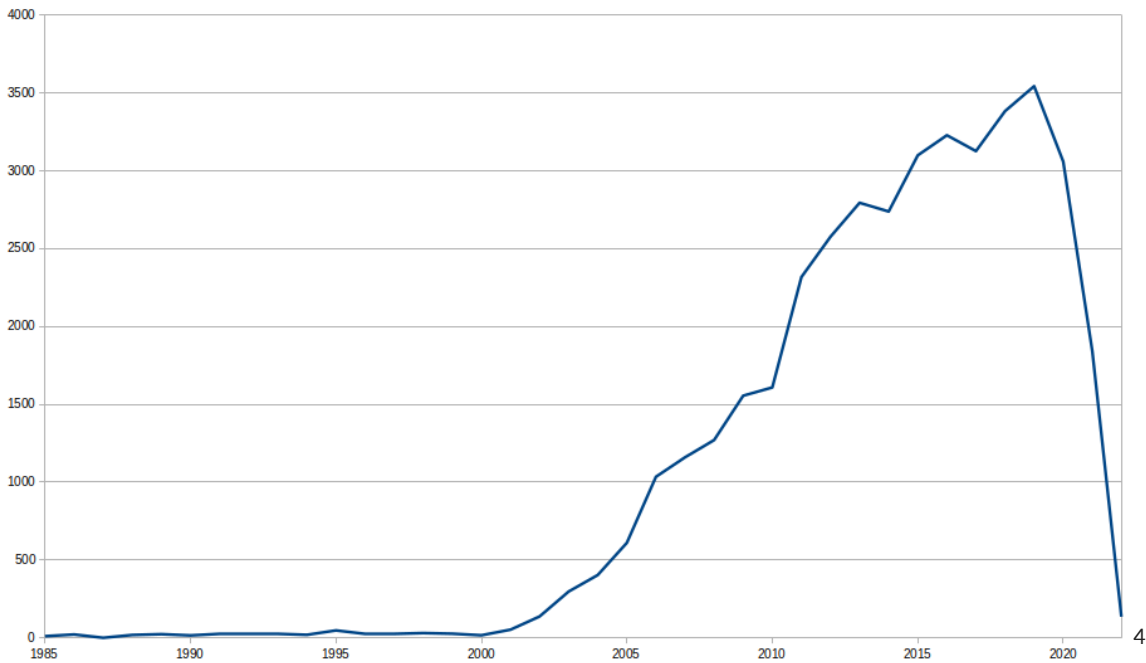
Part of the vast ornithology collection at the American Museum of Natural History.

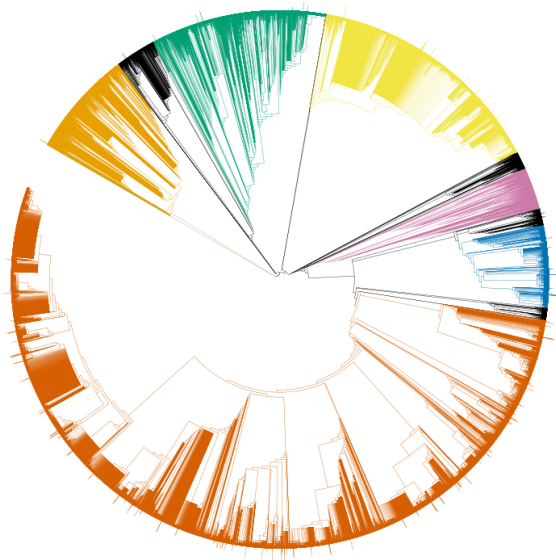
Taxonomy anarchy hampers conservation

The classification of complex organisms is in chaos.
Stephen T. Garnett and **Les Christidis** propose a solution.

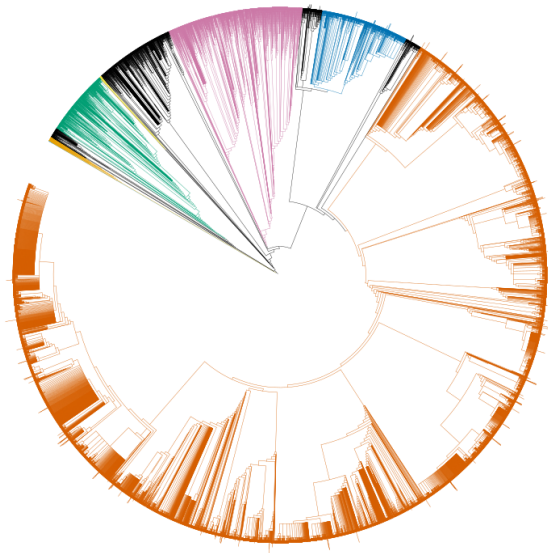
Corpus construction

Journal	Publisher	Size
<i>Zootaxa</i>	Magnolia Press	31,348
<i>ZooKeys</i>	Pensoft	4,940
<i>PhytoKeys</i>	Pensoft	820
<i>Journal of Hymenoptera Research</i>	Pensoft	382
<i>MycoKeys</i>	Pensoft	315
<i>Zoosystematics and Evolution</i>	Pensoft	153
<i>Insecta Mundi</i>	Center for Systematic Entomology	1,367
<i>European Journal of Taxonomy</i>	Museum National d'Histoire Naturelle	1,105





Whole Open Tree of Life



Corpus

Feature Analysis

Global Names Finder (gnfinder): detect the names of species and other groups in text, both by comparison with global lists as well as detection of “probable” names

Global Names Finder (GNfinder)

DOI [10.5281/zenodo.11584025](https://doi.org/10.5281/zenodo.11584025)  reference  go report  A+

Try GNfinder [online](#) or learn about its [API](#).

Very fast finder of scientific names. It uses dictionary and NLP approaches. On modern multiprocessor laptop it is able to process 15 million pages per hour. Works with many file formats and includes names verification against many biological databases. For full functionality it requires an Internet connection.

GNfinder is also available via [web](#) or as a [RESTful API](#).

- [Citing](#)
- [Features](#)
- [Installation](#)
 - [Homebrew on Mac OS X, Linux, and Linux on Windows \(WSL2\)](#)

Locations

Pre-trained model for recognizing locations, organizations, and people in an English-language text (trained by the Bayerische Staatsbibliothek)

The screenshot shows the Hugging Face interface for the model `dbmdz/bert-large-cased-finetuned-conll103-english`. The page header includes the model name, a 'like' button, a 'Follow' button, and the owner's name 'Bayerische Staatsbibliothek' with a follower count of 81. Below this is a row of tags: 'Token Classification', 'Transformers', 'PyTorch', 'TensorFlow', 'JAX', 'Rust', 'Safetensors', and 'bert'. The main navigation bar contains 'Model card', 'Files and versions', 'set', and 'Community'. On the right side of the navigation bar are buttons for 'Train', 'Deploy', and 'Use this model'. The main content area is divided into two columns. The left column contains a message: 'No model card' followed by 'New: Create and edit this model card directly on the website' and a 'Contribute a Model Card' button. The right column contains a 'Downloads last month' section showing '1,372,023' with a line graph. Below this is a 'Safetensors' section with 'Model size: 334M params', 'Tensor type: F32', and a 'Files info' link. The 'Inference Providers' section is active, showing 'HF Inference API' and 'Token Classification' with an 'Examples' dropdown. A text input field contains 'Your sentence here...' and a 'Compute' button is below it. At the bottom of the right column are links for 'View Code Snippets' and 'Maximize'. The footer of the page shows a 'Model tree' link for the model.

Topic Modeling

Convert documents into vectors in a 400-dimensional space (using the doc2vec algorithm), then examine clusters in this space. Normally, each cluster corresponds, more or less, to a subject of discussion.

Topic Modeling

But: **less useful than usual** in this corpus! Often, the clusters indicate how scientists talk about different groups of organisms (“fin, ray, gill, dorsal...”), though some might have a more interesting meaning (“barcoding, biodiversity, DNA...”).

Disagreement

Close-reading of articles where we're sure that taxonomists are disagreeing with each other, to extract lists of keywords.

Désaccord

Example: the *disagreement* list

- critique
- doubt
- opinion
- disagree
- redundant
- reject
- rebuttal
- debate
- invalid
- misunderstanding
- misconception
- allegation
- allegedly
- mistake
- obsolete
- error
- misclassify
- erroneous
- contentious

Disagreement

3 lists: *epistemic values*, *disagreement*, and *pejorative evaluation*

Measure the relative frequency of these words in each article to give them a kind of “disagreement index.”

Methodology

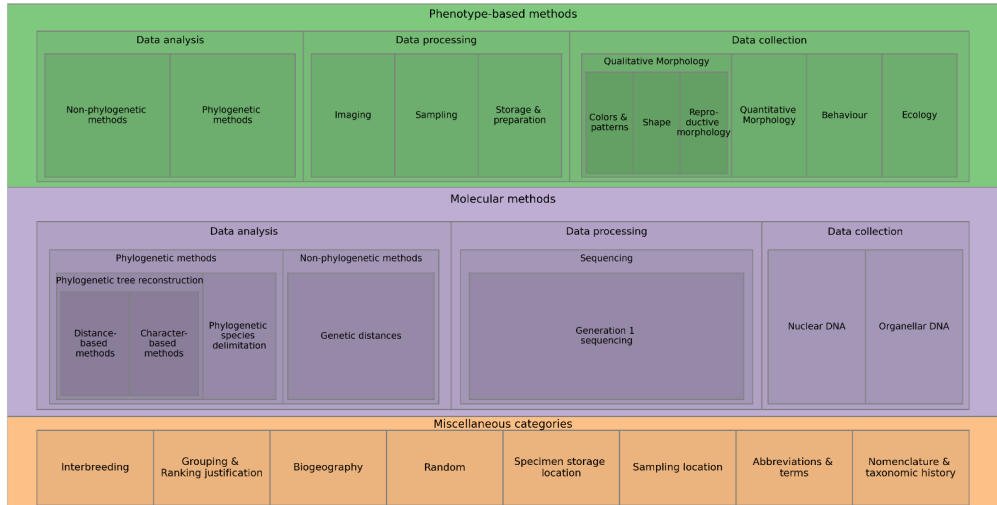
Difficult to detect in taxonomy:

- No “standard” citations for each method
- Different traditions of research (per taxon) = different terminology
- No tradition of describing your methods clearly (exploratory science)
- Lots of amateurs and researchers distributed across the world

Methodology

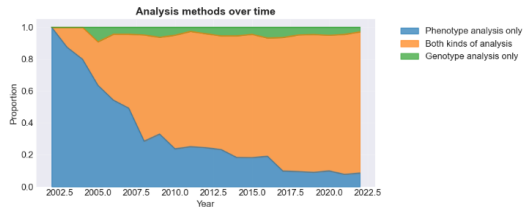
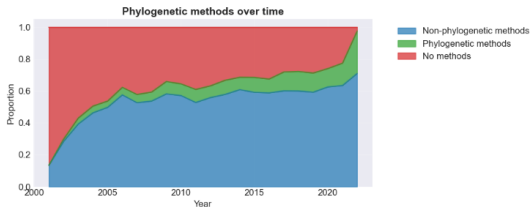
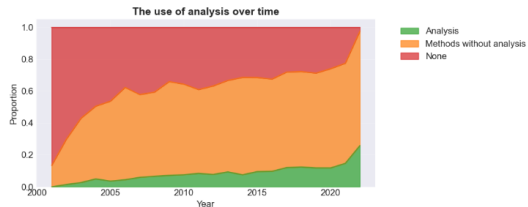
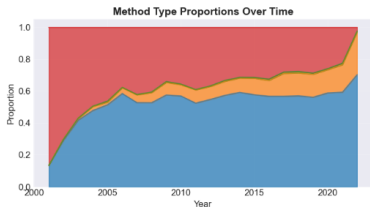
1. Lay out a general, hierarchical structure of methods
2. Isolate the “methods” sections
3. Exploratory analysis with topic modeling of these sections
4. Manual labeling of the paragraphs of these sections
5. Finalize the classification
6. Train classifiers/LLMs to classify the rest of the methods-paragraphs

Methodology

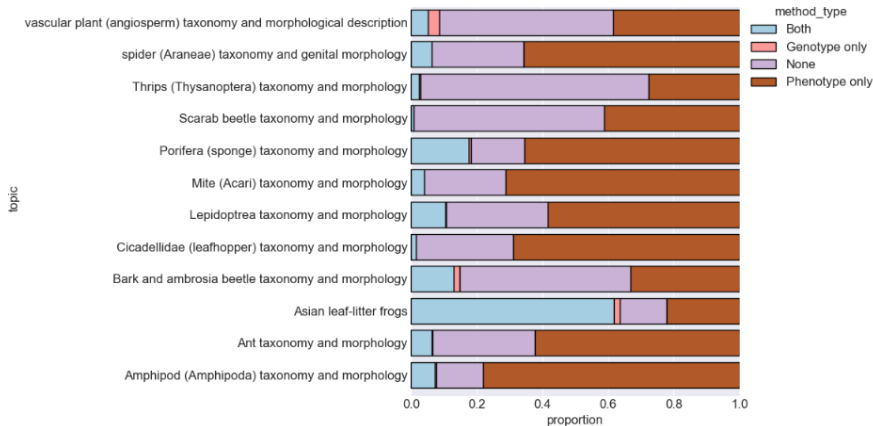


Preliminary results

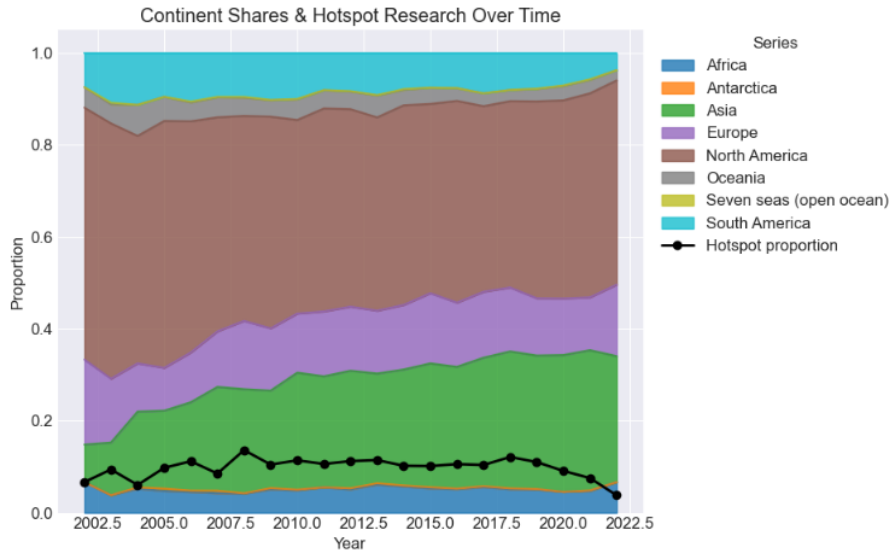
Methodology



Methodology



Taxonomic attention



Disagreement and taxa

Divide organisms into “colloquial” groups (e.g., mammals, fish, birds, ...).

- Lots more disagreement ($> 2\times$): birds ($n = 333$); mollusks ($n = 1064$)
- A bit more ($> 1.25\times$): mammals ($n = 396$)
- A bit less ($< 0.75\times$): fish ($n = 2132$); non-insect arthropods ($n = 7285$)
- A lot less ($< 0.5\times$): prokaryotes (! $n = 13$)

Disagreement and taxa

Second hypothesis: What about the **age** of the group? Test the correlation between the “disagreement index” and the year in which the main genus in the article was described.

We expect a **negative correlation**: the older the group, the more we argue about it.

Disagreement and taxa

Confirmed: **significant negative correlation**

An article on a genus described in 1750 should have a disagreement-index around 0.003 higher than one on a newly described genus (and 0.003 is around the mean disagreement index overall!).

Future ideas

Future ideas

- Correlations with places discussed (and especially eco-regions, biomes, etc.)
- In-depth analysis (close-reading) of changes in methodology with time and across taxa
- Construction of a “high-disagreement” corpus, then analysis of it to detect (maybe?) different senses/kinds of disagreement

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Questions?

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Un problème avec les endroits

On ne peut faire de l'analyse des mots qui décrivent des endroits! Il faut les convertir en coordonnées latitude/longitude. Et faire cette conversion avec les outils normaux (*reverse geocoding*) est **très** cher.

Alors, utilisons un index géographique!

Index géographique

Extrêmement lent:

1. Téléchargez une liste des endroits et leurs coordonnées
2. Mettez le tout dans une base de données
3. Cherchez pour des résultats exacts
4. Sinon, cherchez pour des résultats approximatifs
5. S'il y en a plusieurs, calculez le «centre géographique» de l'article entier et prendre le résultat le plus proche

Concepts d'espèce

Phylo-Phenetic Species Concept

Phylogenetic Species Concept

Genic Species Concept

Cohesion Species Concept

Genealogical Concordance Species
Concept

Genotypic Cluster Species Concept

Genetic Species Concept

Ecological Species Concept

Recognition Species Concept

Genealogical Species Concept

Biological Species Concept

Differential Fitness Species Concept

Compilospecies Concept

Cladistic Species Concept

Hennigian Species Concept

Internodal Species Concept

Mitochondrial Compatibility Species
Concept

Pragmatic Species Concept

Inclusive Species Concept

Biosimilarity Species Concept

