A Review of Biotic Interactions and Taxon Names Found in globalbioticinteractions/cu-cuac hash://md5/a2857fbdca8beec53eb9ec81d2077cd7

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https://github.com/globalbioticinteractions/cu-cuac/issues

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Abstract

Life on Earth is sustained by complex interactions between organisms and their environment. These biotic interactions can be captured in datasets and published digitally. We present a review and archiving process for such an openly accessible digital interactions dataset of known origin and discuss its outcome. The dataset under review, named globalbioticinteractions/cu-cuac, has fingerprint hash://md5/a2857fbdca8beec53eb9ec81d2077cd7, is 8.17MiB in size and contains 565 interactions with 3 unique types of associations (e.g., hasHost) between 553 primary taxa (e.g., Arianops) and 73 associated taxa (e.g., ex. Mushrooms). This report includes detailed summaries of interaction data, a taxonomic review from multiple catalogs, and an archived version of the dataset from which the reviews are derived.

Contents

Discussion   **21**

Acknowledgements   **21**

Author contributions   **22**

References   **22**

Introduction

Data Review and Archive

Data review and archiving can be a time-consuming process, especially when done manually. This review report aims to help facilitate both activities. It automates the archiving of datasets, including Darwin Core archives, and is a citable backup of a version of the dataset. Additionally, an automatic review of species interaction claims made in the dataset is generated and registered with Global Biotic Interactions (J. H. Poelen, Simons, and Mungall 2014).

This review includes summary statistics about, and observations about, the dataset under review:

> Clemson University Arthropod Collection https://github.com/globalbioticinteractions/cu-cuac/archive/6cdcbbaa4f7cec8e1eac705be3a999bc5259e00f.zip 2025-04-04T23:36:47.241Z hash://md5/a2857fbdca8beec53eb9ec81d2077cd7

For additional metadata related to this dataset, please visit https://github.com/globalbioticinteractions/cu-cuac and inspect associated metadata files including, but not limited to, *README.md*, *eml.xml*, and/or *globi.json*.

Methods

The review is performed through programmatic scripts that leverage tools like Preston (Elliott et al. 2025), Elton (Kuhn, Poelen, and Leinweber 2025), Nomer (Salim and Poelen 2025), globinizer (J. Poelen, Seltmann, and Mietchen 2024) combined with third-party tools like grep, mlr, tail and head.

Table 1: Tools used in this review process

tool name	version
preston	0.10.1
elton	0.15.9

tool name	version
nomer	0.5.13
globinizer	0.4.0
mlr	6.0.0
jq	1.6
yq	4.25.3
pandoc	3.1.6.1

The review process can be described in the form of the script below [1].

```
# get versioned copy of the dataset (size approx. 8.17MiB) under review
elton pull globalbioticinteractions/cu-cuac

# generate review notes
elton review globalbioticinteractions/cu-cuac\
 > review.tsv

# export indexed interaction records
elton interactions globalbioticinteractions/cu-cuac\
 > interactions.tsv

# export names and align them with the Catalogue of Life using Nomer
elton names globalbioticinteractions/cu-cuac\
 | nomer append col\
 > name-alignment.tsv
```
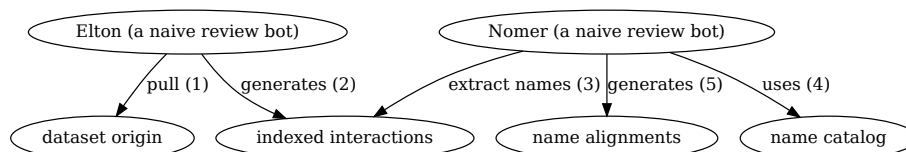
or visually, in a process diagram.

Figure 1: Review Process Overview

You can find a copy of the full review script at check-data.sh. See also GitHub and Codeberg.

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[1]Note that you have to first get the data (e.g., via elton pull globalbioticinteractions/cu-cuac) before being able to generate reviews (e.g., elton review globalbioticinteractions/cu-cuac), extract interaction claims (e.g., elton interactions globalbioticinteractions/cu-cuac), or list taxonomic names (e.g., elton names globalbioticinteractions/cu-cuac)

Results

In the following sections, the results of the review are summarized [2]. Then, links to the detailed review reports are provided.

Files

The following files are produced in this review:

filename	description
biblio.bib	list of bibliographic reference of this review
check-dataset.sh	data review workflow/process as expressed in a bash script
data.zip	a versioned Preston (Elliott et al. 2025) archive of the data under review
HEAD	the digital signature of the data under review
index.docx	review in MS Word format
index.html	review in HTML format
index.md	review in Pandoc markdown format
index.pdf	review in PDF format
indexed-citations.csv.gz	list of distinct reference citations for reviewed species interaction claims in gzipped comma-separated values file format
indexed-citations.html.gz	list of distinct reference citations for reviewed species interactions claims in gzipped html file format
indexed-citations.tsv.gz	list of distinct reference citations for reviewed species interaction claims in gzipped tab-separated values format
indexed-interactions-col-family-col-family.svg	network diagram showing the taxon family to taxon family interaction claims in the dataset under review as interpreted by the Catalogue of Life via Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024)

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[2]Disclaimer: The results in this review should be considered friendly, yet naive, notes from an unsophisticated robot. Please keep that in mind when considering the review results.

filename	description
indexed-interactions-col-kingdom-col-kingdom.svg	network diagram showing the taxon kingdom to taxon kingom interaction claims in the dataset under review as interpreted by the Catalogue of Life via Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024)
indexed-interactions.csv.gz	species interaction claims indexed from the dataset under review in gzipped comma-separated values format
indexed-interactions.html.gz	species interaction claims indexed from the dataset under review in gzipped html format
indexed-interactions.tsv.gz	species interaction claims indexed from the dataset under review in gzipped tab-separated values format
indexed-interactions-sample.csv	list of species interaction claims indexed from the dataset under review in gzipped comma-separated values format
indexed-interactions-sample.html	first 500 species interaction claims indexed from the dataset under review in html format
indexed-interactions-sample.tsv	first 500 species interaction claims indexed from the dataset under review in tab-separated values format
indexed-names.csv.gz	taxonomic names indexed from the dataset under review in gzipped comma-separated values format
indexed-names.html.gz	taxonomic names found in the dataset under review in gzipped html format
indexed-names.tsv.gz	taxonomic names found in the dataset under review in gzipped tab-separated values format
indexed-names-resolved-col.csv.gz	taxonomic names found in the dataset under review aligned with the Catalogue of Life as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format

filename	description
indexed-names-resolved-col.html.gz	taxonomic names found in the dataset under review aligned with the Catalogue of Life as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-col.tsv.gz	taxonomic names found in the dataset under review aligned with the Catalogue of Life as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-resolved-discoverlife.csv.gz	taxonomic names found in the dataset under review aligned with Discover Life bee species checklist as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format
indexed-names-resolved-discoverlife.html.gz	taxonomic names found in the dataset under review aligned with Discover Life bee species checklist as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-discoverlife.tsv.gz	taxonomic names found in the dataset under review aligned with Discover Life bee species checklist as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-resolved-gbif.csv.gz	taxonomic names found in the dataset under review aligned with GBIF Backbone Taxonomy as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format

filename	description
indexed-names-resolved-gbif.html.gz	taxonomic names found in the dataset under review aligned with GBIF Backbone Taxonomy as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-gbif.tsv.gz	taxonomic names found in the dataset under review aligned with GBIF Backbone Taxonomy as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-resolved-itis.csv.gz	taxonomic names found in the dataset under review aligned with Integrated Taxonomic Information System (ITIS) as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format
indexed-names-resolved-itis.html.gz	taxonomic names found in the dataset under review aligned with Integrated Taxonomic Information System (ITIS) as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-itis.tsv.gz	taxonomic names found in the dataset under review aligned with Integrated Taxonomic Information System (ITIS) as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-resolved-mdd.csv.gz	taxonomic names found in the dataset under review aligned with the Mammal Diversity Database as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format

filename	description
indexed-names-resolved-mdd.html.gz	taxonomic names found in the dataset under review aligned with Mammal Diversity Database as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-mdd.tsv.gz	taxonomic names found in the dataset under review aligned with Mammal Diversity Database as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-resolved-ncbi.csv.gz	taxonomic names found in the dataset under review aligned with the NCBI Taxonomy as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format
indexed-names-resolved-ncbi.html.gz	taxonomic names found in the dataset under review aligned with the NCBI Taxonomy as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-ncbi.tsv.gz	taxonomic names found in the dataset under review aligned with the NCBI Taxonomy as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-resolved-pbdb.csv.gz	taxonomic names found in the dataset under review aligned with the Paleobiology Database as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format

filename	description
indexed-names-resolved-pbdb.html.gz	taxonomic names found in the dataset under review aligned with Paleobiology Database as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-pbdb.tsv.gz	taxonomic names found in the dataset under review aligned with Paleobiology Database as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-resolved-tpt.csv.gz	taxonomic names found in the dataset under review aligned with the Terrestrial Parasite Tracker (TPT) Taxonomic Resource as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format
indexed-names-resolved-tpt.html.gz	taxonomic names found in the dataset under review aligned with the Terrestrial Parasite Tracker (TPT) Taxonomic Resource as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-tpt.tsv.gz	taxonomic names found in the dataset under review aligned with the Terrestrial Parasite Tracker (TPT) Taxonomic Resource as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-resolved-wfo.csv.gz	taxonomic names found in the dataset under review aligned with the World of Flora Online as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format

filename	description
indexed-names-resolved-wfo.html.gz	taxonomic names found in the dataset under review aligned with the World of Flora Online as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-wfo.tsv.gz	taxonomic names found in the dataset under review aligned with the World of Flora Online as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-resolved-worms.csv.gz	taxonomic names found in the dataset under review aligned with the World Register of Marine Species (WoRMS) as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped comma-separated values format
indexed-names-resolved-worms.html.gz	taxonomic names found in the dataset under review aligned with the World Register of Marine Species (WoRMS) as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped html format
indexed-names-resolved-worms.tsv.gz	taxonomic names found in the dataset under review aligned with the World Register of Marine Species (WoRMS) as accessed through the Nomer Corpus of Taxonomic Resources (J. H. (ed.). Poelen 2024) in gzipped tab-separated values format
indexed-names-sample.csv	first 500 taxonomic names found in the dataset under review in comma-separated values format
indexed-names-sample.html	first 500 taxonomic names found in the dataset under review in html format
indexed-names-sample.tsv	first 500 taxonomic names found in the dataset under review in tab-separated values format
interaction.svg	diagram summarizing the data model used to index species interaction claims

filename	description
nanopub-sample.trig	first 500 species interaction claims as expressed in the nanopub format (Kuhn and Dumontier 2014)
nanopub.trig.gz	species interaction claims as expressed in the nanopub format (Kuhn and Dumontier 2014)
process.svg	diagram summarizing the data review processing workflow
prov.nq	origin of the dataset under review as expressed in rdf/nquads
review.csv.gz	review notes associated with the dataset under review in gzipped comma-separated values format
review.html.gz	review notes associated with the dataset under review in gzipped html format
review.tsv.gz	review notes associated with the dataset under review in gzipped tab-separated values format
review-sample.csv	first 500 review notes associated with the dataset under review in comma-separated values format
review-sample.html	first 500 review notes associated with the dataset under review in html format
review-sample.tsv	first 500 review notes associated with the dataset under review in tab-separated values format
review.svg	a review badge generated as part of the dataset review process
zenodo.json	metadata of this review expressed in Zenodo record metadata

Archived Dataset

Note that *data.zip* file in this archive contains the complete, unmodified archived dataset under review.

Biotic Interactions

In this review, biotic interactions (or biotic associations) are modeled as a primary (aka subject, source) organism interacting with an associate (aka object, target) organism. The dataset under review classified the primary/associate
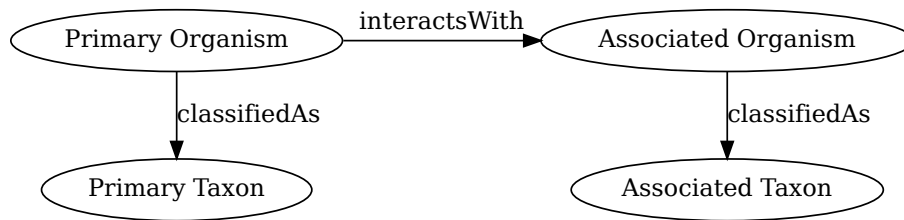
Figure 2: Biotic Interaction Data Model

organisms with specific taxa. The primary and associate organisms The kind of interaction is documented as an interaction type.

The dataset under review, named globalbioticinteractions/cu-cuac, has fingerprint hash://md5/a2857fbdca8beec53eb9ec81d2077cd7, is 8.17MiB in size and contains 565 interactions with 3 unique types of associations (e.g., hasHost) between 553 primary taxa (e.g., Arianops) and 73 associated taxa (e.g., ex. Mushrooms).

An exhaustive list of indexed interaction claims can be found in gzipped csv and tsv archives. To facilitate discovery, a preview of claims available in the gzipped html page at indexed-interactions.html.gz are shown below.

The exhaustive list was used to create the following data summaries below.

Table 3: Sample of Indexed Interaction Claims

sourceTaxonName	interactionTypeName	targetTaxonName	referenceCitation
Trichophaga tapetzella	interactsWith	owl pellets	https://scan-bugs.org:443/portal/collections/individual/index.
Thyridopteryx ephemeraeformis	hasHost	larvae	https://scan-bugs.org:443/portal/collections/individual/index.
Phyllonorycter blancardella	interactsWith	apple foilage	https://scan-bugs.org:443/portal/collections/individual/index.
Cosmopterix attenuatella	hasHost	reared ex Lespedeza	https://scan-bugs.org:443/portal/collections/individual/index.

Table 4: Most Frequently Mentioned Interaction Types (up to 20 most frequent)

interactionTypeName	count
hasHost	276
adjacentTo	273
interactsWith	16

Table 5: Most Frequently Mentioned Primary Taxa (up to 20 most frequent)

sourceTaxonName	count
Arianops	3
Paranthrene simulans	2
CU | CUAC | d198856b-3b84-4738-aa59-97ad8bc5c234 | CUAC000122368 | c25eec7b-5dd7-478a-9ffe-5a6dbe928bf9	2
CU | CUAC | d198856b-3b84-4738-aa59-97ad8bc5c234 | CUAC000122369 | 96511200-6583-4b6e-a6d3-83c549e76e40	2
CU | CUAC | d198856b-3b84-4738-aa59-97ad8bc5c234 | CUAC000122370 | 6bfd0638-dc01-4b42-a3f2-02891077c18b	2
CU | CUAC | d198856b-3b84-4738-aa59-97ad8bc5c234 | CUAC000122371 | ecadbb68-42f3-4051-a970-a077ad93ce3b	2
CU | CUAC | d198856b-3b84-4738-aa59-97ad8bc5c234 | CUAC000122372 | 70da6ace-03ce-4e1c-afe7-6f0eff656014	2
CU | CUAC | d198856b-3b84-4738-aa59-97ad8bc5c234 | CUAC000122373 | 76f91ef0-d4df-45b6-b42b-9dff7f41e44a	2
Stenomorphus californicus	2
Pentacora ligata	2
Aradidae	2
Trichophaga tapetzella	1
Thyridopteryx ephemeraeformis	1
Phyllonorycter blancardella	1
Cosmopterix attenuatella	1
Episimus tyrius	1
Endothenia hebesana	1
Epiblema strenuana	1
Tortrix viridana	1

Table 6: Most Frequently Mentioned Associate Taxa (up to 20 most frequent)

targetTaxonName	count
ex. Mushrooms	84
dying pinus	61
ex. rotten log	59
flowers	55
ex. Polypore	32
ex. Dead turtle	29
ex. Maclura fruit	25
Under Hemlock	17
dead squirrel	14
freshly cut pine stump	12
ex. Asimina fruit	11
dead pine at night	11
bark dead tree	10
exposed rock face	9
mud bank	9
palmetto leaf at night	6
nymph	6
vegetation	6
Ex. Crepe myrtle	5

Table 7: Most Frequent Interactions between Primary and Associate Taxa (up to 20 most frequent)

sourceTaxonName	interactionTypeName	targetTaxonName	count
Arianops	adjacentTo	underside of embedded rock.	2
Pentacora ligata	adjacentTo	bare rock beside rapids	2
Aradidae	adjacentTo	dead wood at night	2
Trichophaga tapetzella	interactsWith	owl pellets	1
Thyridopteryx ephemeraeformis	hasHost	larvae	1
Phyllonorycter blancardella	interactsWith	apple foilage	1
Cosmopterix attenuatella	hasHost	reared ex Lespedeza	1

sourceTaxonName	interactionTypeName	targetTaxonName	count
Paranthrene simulans	hasHost	Japanese Maple	1
Paranthrene simulans	hasHost	emerged 5. May. 1999	1
Episimus tyrius	hasHost	7 Aug. 1989	1
Endothenia hebesana	adjacentTo	Snap-Dragon	1
Epiblema strenuana	interactsWith	ragweed	1
Tortrix viridana	hasHost	ex. Oak	1
Lerema accius	hasHost	pupa inside rolled up leaf of Johnson grass	1
Atalopedes campestris	hasHost	apple tree	1
Megathymus yuccae	hasHost	pupae	1
Erynnis juvenalis	adjacentTo	wing	1
Papilio troilus	adjacentTo	spice bush	1
CU | CUAC | d198856b-3b84-4738-aa59-97ad8bc5c234 | CUAC000055434 | 4d7feb7a-8ea6-4085-9f3a-c071d9737f6f	adjacentTo	dying pinus	1

Interaction Networks

The figures below provide a graph view on the dataset under review. The first shows a summary network on the kingdom level, and the second shows how interactions on the family level. It is important to note that both network graphs were first aligned taxonomically using the Catalogue of Life. Please refer to the original (or verbatim) taxonomic names for a more original view on the interaction data.

Figure 3: Interactions on taxonomic kingdom rank as interpreted by the Catalogue of Life download svg

You can download the indexed dataset under review at indexed-interactions.csv .gz. A tab-separated file can be found at indexed-interactions.tsv.gz

Figure 4: Interactions on the taxonomic family rank as interpreted by the Catalogue of Life. download svg

Learn more about the structure of this download at GloBI website, by opening a GitHub issue, or by sending an email.

Another way to discover the dataset under review is by searching for it on the GloBI website.

Taxonomic Alignment

As part of the review, all names are aligned against various name catalogs (e.g., col, ncbi, discoverlife, gbif, itis, wfo, mdd, tpt, pbdb, and worms). These alignments can help review name usage or aid in selecting of a suitable taxonomic name resource.

Table 8: Sample of Name Alignments

providedName	relationName	resolvedCatalogName	resolvedName
Cosmopterix attenuatella	HAS_ACCEPTED_NAME	col	Cosmopterix attenuatella
Atalopedes campestris	HAS_ACCEPTED_NAME	col	Atalopedes campestris
Erynnis juvenalis	HAS_ACCEPTED_NAME	col	Erynnis juvenalis
Osorius	HAS_ACCEPTED_NAME	col	Osorius

Table 9: Distribution of Taxonomic Ranks of Aligned Names by Catalog. Names that were not aligned with a catalog are counted as NAs. So, the total number of unaligned names for a catalog will be listed in their NA row.

resolvedCatalogName	resolvedRank	count
col	NA	585
col	class	1
col	family	2
col	genus	11
col	kingdom	1
col	order	1
col	species	19
discoverlife	NA	621
gbif	NA	582
gbif	class	1

resolvedCatalogName	resolvedRank	count
gbif	family	2
gbif	genus	12
gbif	kingdom	1
gbif	order	1
gbif	species	21
gbif	subspecies	1
itis	NA	591
itis	family	2
itis	genus	10
itis	kingdom	1
itis	order	2
itis	species	13
itis	subclass	1
mdd	NA	620
ncbi	NA	588
ncbi	family	2
ncbi	genus	10
ncbi	order	2
ncbi	species	18
ncbi	subclass	1
pbdb	NA	606
pbdb	class	1
pbdb	family	2
pbdb	genus	6
pbdb	kingdom	1
pbdb	order	2
pbdb	species	2
tpt	NA	618
tpt	species	2
wfo	NA	613
wfo	genus	4
wfo	phylum	1
wfo	species	2
worms	NA	606
worms	family	1
worms	genus	3
worms	infraorder	1
worms	kingdom	1
worms	order	2
worms	species	6
worms	subclass	1

Table 10: Name relationship types per catalog. Name relationship type "NONE" means that a name was not recognized by the associated catalog. "SAME_AS" indicates either a "HAS_ACCEPTED_NAME" or "SYNONYM_OF" name relationship type. We recognize that "SYNONYM_OF" encompasses many types of nomenclatural synonymies (ICZN 1999) (e.g., junior synonym, senior synonyms).

resolvedCatalogName	relationName	count
col	HAS_ACCEPTED_NAME	36
col	SYNONYM_OF	5
col	NONE	586
discoverlife	NONE	626
gbif	HAS_ACCEPTED_NAME	47
gbif	SYNONYM_OF	9
gbif	NONE	583
itis	HAS_ACCEPTED_NAME	29
itis	NONE	592
itis	SYNONYM_OF	3
mdd	NONE	622
ncbi	NONE	592
ncbi	SAME_AS	34
pbdb	NONE	607
pbdb	HAS_ACCEPTED_NAME	14
pbdb	SYNONYM_OF	7
tpt	NONE	620
tpt	HAS_ACCEPTED_NAME	2
wfo	NONE	615
wfo	HAS_ACCEPTED_NAME	7
wfo	SYNONYM_OF	1
worms	HAS_ACCEPTED_NAME	17
worms	NONE	607

Table 11: List of Available Name Alignment Reports

catalog name	alignment results
col	associated names alignments report in gzipped html, csv, and tsv)
ncbi	associated names alignments report in gzipped html, csv, and tsv)
discoverlife	associated names alignments report in gzipped html, csv, and tsv)

catalog name	alignment results
gbif	associated names alignments report in gzipped html, csv, and tsv)
itis	associated names alignments report in gzipped html, csv, and tsv)
wfo	associated names alignments report in gzipped html, csv, and tsv)
mdd	associated names alignments report in gzipped html, csv, and tsv)
tpt	associated names alignments report in gzipped html, csv, and tsv)
pbdb	associated names alignments report in gzipped html, csv, and tsv)
worms	associated names alignments report in gzipped html, csv, and tsv)

Additional Reviews

Elton, Nomer, and other tools may have difficulties interpreting existing species interaction datasets. Or, they may misbehave, or otherwise show unexpected behavior. As part of the review process, detailed review notes are kept that document possibly misbehaving, or confused, review bots. An sample of review notes associated with this review can be found below.

Table 12: First few lines in the review notes.

reviewDate	reviewCommentType	reviewComment
2025-04-10T11:12:37Z	note	found unsupported interaction type with name: [Division]
2025-04-10T11:12:37Z	note	found unsupported interaction type with name: [Comp]
2025-04-10T11:12:37Z	note	found unsupported interaction type with name: [Stand]
2025-04-10T11:12:37Z	note	found unsupported interaction type with name: [Inkto Collection]

In addition, you can find the most frequently occurring notes in the table below.

Table 13: Most frequently occurring review notes, if any.

reviewComment	count
source taxon name missing: using institution-Code/collectionCode/collectionId/catalogNumber/occurrenceId as placeholder	530
found unsupported interaction type with name: [Inkto Collection]	2
found unsupported interaction type with name: [Division]	1
found unsupported interaction type with name: [Comp]	1

For additional information on review notes, please have a look at the first 500 Review Notes in html format or the download full gzipped csv or tsv archives.

GloBI Review Badge

As part of the review, a review badge is generated. This review badge can be included in webpages to indicate the review status of the dataset under review.

Figure 5: Picture of a GloBI Review Badge [3]

Note that if the badge is green, no review notes were generated. If the badge is yellow, the review bots may need some help with interpreting the species interaction data.

GloBI Index Badge

If the dataset under review has been registered with GloBI, and has been succesfully indexed by GloBI, the GloBI Index Status Badge will turn green. This means that the dataset under review was indexed by GloBI and is available through GloBI services and derived data products.

Figure 6: Picture of a GloBI Index Badge [4]

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[3]Up-to-date status of the GloBI Review Badge can be retrieved from the GloBI Review Depot

[4]Up-to-date status of the GloBI Index Badge can be retrieved from GloBI's API

If you'd like to keep track of reviews or index status of the dataset under review, please visit GloBI's dataset index [5] for badge examples.

Discussion

This review and archive provides a means of creating citable versions of datasets that change frequently. This may be useful for dataset managers, including natural history collection data managers, as a backup archive of a shared Darwin Core archive. It also serves as a means of creating a trackable citation for the dataset in an automated way, while also including some information about the contents of the dataset.

This review aims to provide a perspective on the dataset to aid in understanding of species interaction claims discovered. However, it is important to note that this review does *not* assess the quality of the dataset. Instead, it serves as an indication of the open-ness[6] and FAIRness (Wilkinson et al. 2016; Trekels et al. 2023) of the dataset: to perform this review, the data was likely openly available, **F**indable, **A**ccessible, **I**nteroperable and **R**eusable. The current Open-FAIR assessment is qualitative, and a more quantitative approach can be implemented with specified measurement units.

This report also showcases the reuse of machine-actionable (meta)data, something highly recommended by the FAIR Data Principles (Wilkinson et al. 2016). Making (meta)data machine-actionable enables more precise procesing by computers, enabling even naive review bots like Nomer and Elton to interpret the data effectively. This capability is crucial for not just automating the generation of reports, but also for facilitating seamless data exchanges, promoting interoperability.

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[5]At time of writing (2025-04-10) the version of the GloBI dataset index was available at https://globalbioticinteractions.org/datasets

[6]According to http://opendefinition.org/: "Open data is data that can be freely used, re-used and redistributed by anyone - subject only, at most, to the requirement to attribute and sharealike."

Author contributions

Nomer was responsible for name alignments. Elton carried out dataset extraction, and generated the review notes. Preston tracked, versioned, and packaged, the dataset under review.

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