

A Review of Biotic Interactions and Taxon Names Found in globalbioticinteractions/pu-perc hash://md5/dee3de1502050be8eee34059dbd03870

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<https://github.com/globalbioticinteractions/pu-perc/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/pu-perc, has fingerprint hash://md5/dee3de1502050be8eee34059dbd03870, is 3.78MiB in size and contains 10,570 interaction with 2 unique types of associations (e.g., interactsWith) between 562 primary taxa (e.g., Acari) and 592 associated taxa (e.g., Ictidomys tridecemlineatus). This report includes detailed summaries of interaction data, a taxonomic review from multiple catalogs, and an archived version of the dataset from which the reviews are derived.

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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:

Purdue Entomological Research Collection <https://github.com/globalbioticinteractions/pu-perc/archive/e0909a7ca0a8df5effccb288ba64b28141e388ba.zip> 2025-04-12T10:16:22.765Z hash://md5/dee3de1502050be8eee34059dbd03870

For additional metadata related to this dataset, please visit <https://github.com/globalbioticinteractions/pu-perc> 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. 3.78MiB) under review
elton pull globalbioticinteractions/pu-perc

# generate review notes
elton review globalbioticinteractions/pu-perc\
> review.tsv

# export indexed interaction records
elton interactions globalbioticinteractions/pu-perc\
> interactions.tsv

# export names and align them with the Catalogue of Life using Nomer
elton names globalbioticinteractions/pu-perc\
| nomer append col\
> name-alignment.tsv
```

or visually, in a process diagram.

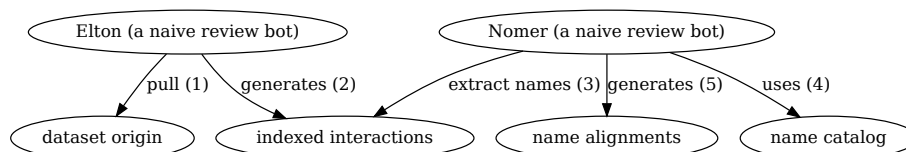


Figure 1: Review Process Overview

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

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

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)

²Disclaimer: 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 kingdom 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

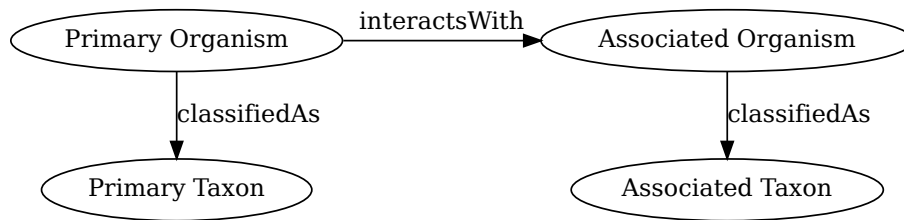


Figure 2: Biotic Interaction Data Model

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/put-perc, has fingerprint hash://md5/dee3de1502050be8eee34059dbd03870, is 3.78MiB in size and contains 10,570 interaction with 2 unique types of associations (e.g., interactsWith) between 562 primary taxa (e.g., Acari) and 592 associated taxa (e.g., Ictidomys tridecemlineatus).

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
Totype velleda	interactsWith	Wild Cherry	https://scan-bugs.org:443/portal/collections/individual/index.
Totype velleda	interactsWith	Apple	https://scan-bugs.org:443/portal/collections/individual/index.
Malacosoma americana	interactsWith	Crabapple	https://scan-bugs.org:443/portal/collections/individual/index.
Malacosoma americana	interactsWith	Crabapple	https://scan-bugs.org:443/portal/collections/individual/index.

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

interactionTypeName	count
interactsWith	10560
killedBy	10

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

sourceTaxonName	count
Acari	742
Oropsylla bruneri	499
Halictus ligatus	368
Halictus confusus	348
Lasioglossum imitatum	316
Andrena wilkella	290
Andrena miserabilis	276
Ctenophthalmus pseudagyrtis	233
Melissodes bimaculata	212
Aug_ella aurata	204
Lasioglossum admirandum	200
Svastra obliqua	150
Nalepella tsugafoliae	136
Haemolaelaps glasgowi	124
Orchopeas leucopus	119
Andrena imitatrix	116
Enderleinellus suturalis	115
Aug_ella persimilis	114
Bombus griseocollis	110

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

targetTaxonName	count
Ictidomys tridecemlineatus	568
Salix	323
Taraxacum	311
Eptesicus fuscus	272
Peromyscus leucopus	267
Salvia sage	211
Salix discolor	207
Melilotus	194
Asclepias incarnata	180
Microtus pennsylvanicus	165
Blarina brevicauda	161
Trifolium pratense	154
Helianthus annuus	153
Cotoneaster	140
Microtus ochrogaster	139

targetTaxonName	count
Asclepias verticillata	136
Alcea hollyhock	131
Helianthus	131
Prunus	128

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

sourceTaxonName	interactionTypeNam	targetTaxonName	count
Oropsylla bruneri	interactsWith	Ictidomys tridecemlineatus	492
Andrena wilkella	interactsWith	Cotoneaster	140
Acari	interactsWith	Eptesicus fuscus	138
Enderleinellus suturalis	interactsWith	Citellus tridecemlineatus	95
Aug_ella persimilis	interactsWith	Salvia sage	92
Lasioglossum admirandum	interactsWith	Taraxacum	92
Acari	interactsWith	Peromyscus leucopus	91
Ctenophthalmus pseudagyrtis	interactsWith	Scalopus aquaticus machrinus	84
Halictus ligatus	interactsWith	Helianthus annuus	80
Orchopeas leucopus	interactsWith	Peromyscus leucopus noveboracensis	74
Halictus confusus	interactsWith	Asclepias verticillata	72
Lasioglossum imitatum	interactsWith	Anise hyssop	68
Nalepella tsugafoliae	interactsWith	Tsuga canadensis	66
Halictus confusus	interactsWith	Asclepias incarnata	64
Andrena miserabilis	interactsWith	Salix	64
Myodopsylla insignis	interactsWith	Myotis lucifugus	57
Halictus ligatus	interactsWith	Helianthus	56

sourceTaxonName	interactionTypeName	targetTaxonName	count
Aug_ella aurata	interactsWith	Salvia sage	54
Chelostoma philadelphi	interactsWith	Philadelphus	50

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](#)

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
Brevipalpus californicus	HAS_ACCEPTED_NAME	col	Brevipalpus californicus

providedName	relationName	resolvedCatalogName	resolvedName
Brevipalpus cardinalis	HAS_ACCEPTED_BY	NA	Brevipalpus cardinalis
Brevipalpus essigi	HAS_ACCEPTED_BY	NA	Brevipalpus essigi
Brevipalpus lewisi	HAS_ACCEPTED_BY	NA	Brevipalpus lewisi

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	272
col	class	2
col	family	20
col	genus	148
col	kingdom	1
col	order	8
col	parvorder	1
col	species	650
col	subfamily	2
col	subgenus	11
col	suborder	2
col	subspecies	27
col	superfamily	2
col	variety	7
discoverlife	NA	925
discoverlife	species	205
gbif	NA	192
gbif	class	2
gbif	family	22
gbif	form	1
gbif	genus	178
gbif	kingdom	1
gbif	order	8
gbif	species	701
gbif	subspecies	35
gbif	variety	14
itis	NA	389
itis	class	2
itis	family	22
itis	genus	139

resolvedCatalogName	resolvedRank	count
itis	kingdom	1
itis	order	10
itis	species	548
itis	subclass	1
itis	subfamily	2
itis	suborder	2
itis	subspecies	13
itis	superfamily	2
itis	variety	1
mdd	NA	1130
ncbi	NA	346
ncbi	class	2
ncbi	family	22
ncbi	genus	161
ncbi	order	8
ncbi	species	570
ncbi	subclass	1
ncbi	subfamily	2
ncbi	subgenus	1
ncbi	suborder	2
ncbi	subspecies	12
ncbi	superfamily	2
ncbi	superorder	1
ncbi	varietas	1
pbdb	NA	906
pbdb	class	3
pbdb	family	13
pbdb	genus	85
pbdb	infraorder	1
pbdb	kingdom	1
pbdb	order	9
pbdb	species	108
pbdb	subfamily	1
pbdb	suborder	3
pbdb	superfamily	2
pbdb	unranked clade	1
tpt	NA	841
tpt	family	2
tpt	genus	11
tpt	order	3
tpt	species	272
tpt	subspecific epithet	3
wfo	NA	806
wfo	family	1

resolvedCatalogName	resolvedRank	count
wfo	genus	114
wfo	species	209
wfo	subspecies	2
wfo	variety	1
worms	NA	912
worms	class	1
worms	family	15
worms	genus	93
worms	kingdom	1
worms	order	8
worms	species	94
worms	subclass	2
worms	suborder	2
worms	subspecies	1
worms	superfamily	2

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	837
col	NONE	283
col	SYNONYM_OF	190
discoverlife	NONE	944
discoverlife	HAS_ACCEPTED_NAME	199
discoverlife	SYNONYM_OF	64
discoverlife	HOMONYM_OF	13
gbif	HAS_ACCEPTED_NAME	1033
gbif	SYNONYM_OF	319
gbif	NONE	202
itis	HAS_ACCEPTED_NAME	718
itis	NONE	400
itis	SYNONYM_OF	61
mdd	NONE	1084
mdd	HAS_ACCEPTED_NAME	67
mdd	SYNONYM_OF	5
ncbi	SAME_AS	761

resolvedCatalogName	relationName	count
ncbi	NONE	357
ncbi	SYNONYM_OF	42
ncbi	COMMON_NAME_OF	1
pdbb	NONE	926
pdbb	SYNONYM_OF	21
pdbb	HAS_ACCEPTED_NAME	229
tpt	HAS_ACCEPTED_NAME	328
tpt	NONE	863
tpt	SYNONYM_OF	127
wfo	NONE	829
wfo	HAS_ACCEPTED_NAME	313
wfo	SYNONYM_OF	55
wfo	HAS_UNCHECKED_NAME	36
worms	NONE	933
worms	HAS_ACCEPTED_NAME	234
worms	SYNONYM_OF	24

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)
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)
pdbb	associated names alignments report in gzipped html, csv, and tsv)
worms	associated names alignments report in gzipped html, csv, and tsv)

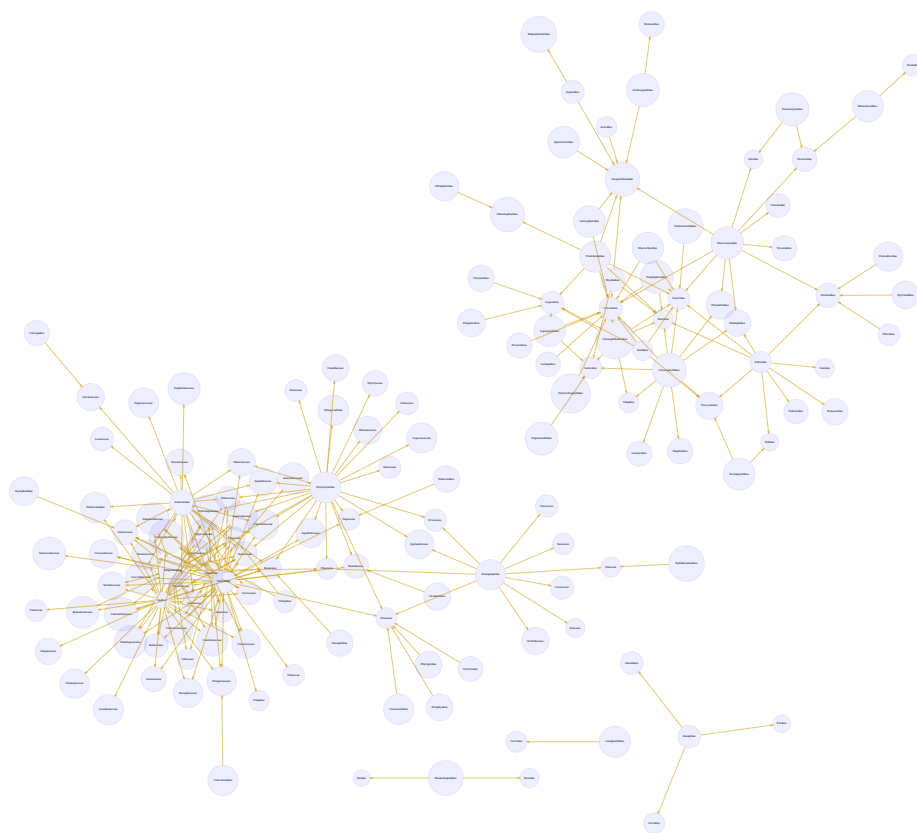


Figure 4: Interactions on the taxonomic family rank as interpreted by the Catalogue of Life. [download svg](#)

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
2025-04-13T08:13:58Z	note	found unsupported interaction type with name: [attacks]
2025-04-13T08:13:59Z	note	source taxon name missing: using institution-Code/collectionCode/collectionId/catalogNumber/occurrenceId as placeholder
2025-04-13T08:13:59Z	note	source taxon name missing: using institution-Code/collectionCode/collectionId/catalogNumber/occurrenceId as placeholder
2025-04-13T08:13:59Z	note	source taxon name missing: using institution-Code/collectionCode/collectionId/catalogNumber/occurrenceId as placeholder

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	8
found unsupported interaction type with name: [attacks]	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 successfully 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

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

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

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

⁵At time of writing (2025-04-13) the version of the GloBI dataset index was available at <https://globalbioticinteractions.org/datasets>

indication of the open-ness⁶ 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 processing 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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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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⁶According 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.”

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