A Review of Biotic Interactions and Taxon Names Found in globalbioticinteractions/inhs-insects hash://md5/26813d12f7b15c1c32cafad666535e2a

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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/inhs-insects, has fingerprint hash://md5/26813d12f7b15c1c32cafad666535e2a, is 183MiB in size and contains 8,835 interaction with 4 unique types of associations (e.g., adjacentTo) between 1,561 primary taxon (e.g., Mezira granulata (Say, 1832)) and 1,750 associated taxon (e.g., under). This report includes detailed summaries of interaction data, a taxonomic review from multiple catalogs, and an archived version of the dataset from which the reviews are derived.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:

Illinois Natural History Survey Insect Collection https://github.com/globalbioticinteractions/inhs-insects/archive/38692496f590577074c7cecf8ea37f85d0594ae1.zip 2025-04-05T00:33:18.083Z hash://md5/26813d12f7b15c1c32cafad666535e2a

For additional metadata related to this dataset, please visit https://github.c om/globalbioticinteractions/inhs-insects 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

tool name	version	
elton	0.15.9	
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. 183MiB) under review
 elton pull globalbioticinteractions/inhs-insects
- # export indexed interaction records
 elton interactions globalbioticinteractions/inhs-insects\
- > interactions.tsv
- # export names and align them with the Catalogue of Life using Nomer
 elton names globalbioticinteractions/inhs-insects\
- | nomer append col\
- > name-alignment.tsv

or visually, in a process diagram.

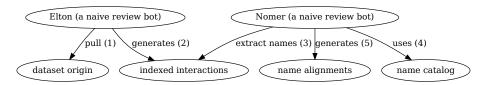


Figure 1: Review Process Overview

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/inhsinsects) before being able to generate reviews (e.g., elton review globalbioticinteractions/inhsinsects), extract interaction claims (e.g., elton interactions globalbioticinteractions/inhs-insects), or list taxonomic names (e.g., elton names globalbioticinteractions/inhs-insects)

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
$indexed\hbox{-}citations.html.gz$	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-	network diagram showing the taxon
family.svg	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)

 $[\]overline{^2 \text{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.cs v	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-it is.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-it is.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.).
indexed-names-resolved-tpt.tsv.gz	Poelen 2024) in gzipped html format 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-w fo.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
$\overline{\text{indexed-names-resolved-wfo.html.gz}}$	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-w fo.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
indexed-names-sample.csv	tab-separated values format first 500 taxonomic names found in the dataset under review in
indexed-names-sample.html	comma-separated values format first 500 taxonomic names found in the
indexed-names-sample.tsv	dataset under review in html format first 500 taxonomic names found in the dataset under review in
interaction.svg	tab-separated values format 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

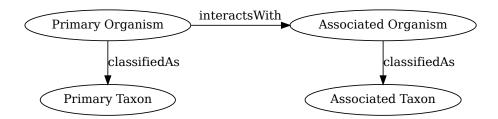


Figure 2: Biotic Interaction Data Model

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/inhs-insects, has fingerprint hash://md5/26813d12f7b15c1c32cafad666535e2a, is 183MiB in size and contains 8,835 interaction with 4 unique types of associations (e.g., adjacentTo) between 1,561 primary taxon (e.g., Mezira granulata (Say, 1832)) and 1,750 associated taxon (e.g., under).

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

$\overline{\text{sourceTaxonName}}$	interaction Type Nam @ arget Taxon Name		${\it reference Citation}$
Sitticus barnesi Cutler, 1965	adjacentTo	table in office	bd2fc9f8-848f- 46fc-9f11- c27ecddb31b6
Hentzia mitrata (Hentz, 1846)	interacts With	herbs + shrubs	0c958a87-6372- 4572-80af- dcda3013df67
INHS INHS Insect Collection 895621 f7af707f- 4742-40a3-96f4- 7df1d13e9b0d	$\operatorname{adjacent} \operatorname{To}$	tree trunks at night	f7af707f-4742- 40a3-96f4- 7df1d13e9b0d
Araneus patagiatus Clerck, 1757	adjacentTo	deciduous trees and shrubs	a474317b-a3da- 4ac3-9905- a94c0dd7bbd2

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

interaction Type Name	count
adjacentTo hasHost interactsWith	5991 2054 758
eats	32

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

source Taxon Name	count
Mezira granulata (Say, 1832)	258
Psammotettix	202
Psyllidae Latreille, 1807	138
Pulicidae Billberg, 1820	125
Empoasca	122
Orchopeas howardi (Baker, 1895)	115
Ptinidae Latreille, 1802	98
Siphonaptera	96
Tetraloniella (Tetraloniella) donata (Cresson, 1878)	94
Myrsidea interrupta (Osborn, 1896)	90
Braconidae Nees, 1811	84
Cynipidae	81
Chalcidoidea	72
Phthiraptera Haeckel, 1896	71
Ichneumonidae	70
Flexamia inflata (Osborn & Ball, 1897)	69
Curculionidae Latreille, 1802	69
Hymenoptera	68
Epitrix cucumeris Harris, 1851	63

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

targetTaxonName	count
under	574
Poa pratensis	335
flowers	217
ex. log	149

targetTaxonName	count
mud	94
flowers of Leguminosae	91
Crow	90
grasses	89
flower	82
tree	78
under oak	74
Cane	73
cottontail	72
Alfalfa	69
pear	65
hips ofRosa blanda	58
bluff	52
wild grape	52
willow	51

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

source Taxon Name	interaction Type N	ameargetTaxonName	count
Mezira granulata (Say, 1832)	adjacentTo	under	258
Psammotettix	adjacentTo	Poa pratensis	202
Ptinidae Latreille, 1802	hasHost	ex. log	98
Myrsidea interrupta (Osborn, 1896)	adjacentTo	Crow	89
Tetraloniella (Tetraloniella) donata (Cresson, 1878)	adjacentTo	flowers of Leguminosae	78
Cynipidae	adjacentTo	tree	69
Flexamia inflata (Osborn & Ball, 1897)	adjacentTo	Poa pratensis	69
Psyllidae Latreille, 1807	adjacentTo	pear	63
Chalcidoidea	interactsWith	hips ofRosa blanda	58

sourceTaxonName	interactionTypeNa	m&argetTaxonName	count
Typocerus confluens Casey, 1913	adjacentTo	flowers	57
Trigonotylus tarsalis (Reuter, 1876)	adjacentTo	grasses	54
Empoasca	adjacentTo	Alfalfa	53
Psyllidae Latreille, 1807	adjacentTo	screens	50
Temnostoma alternans Loew, 1864	adjacentTo	flower	49
Corimelaena (Corimelaena) pulicaria (Germar, 1839)	adjacentTo	flowers	47
Bruchophagus Ashmead, 1888	adjacentTo	Corn	45
Stenocranus similis Crawford, 1914	adjacentTo	Cane	42
Lasioglossum (Dialictus) illinoense (Robertson, 1892)	adjacentTo	Monarda sp.	41
Pycnoscelus surinamensis (Linnaeus, 1758)	interactsWith	rose bench	40

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.

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



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

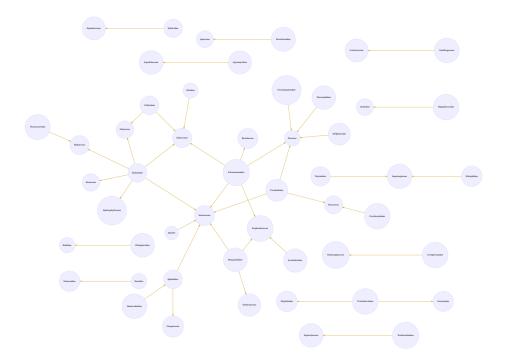


Figure 4: Interactions on the taxonomic family rank as interpreted by the Catalogue of Life. download svg $\,$

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	resolvedCatalogNar	meresolvedName
Lined ground squirrel	NONE	col	Lined ground squirrel
Burn	NONE	col	Burn
Zonotrichia	NONE	col	Zonotrichia
albicollis white			albicollis white
throated sparrow			throated sparrow
Ablabesmyia	HAS_ACCEPTED	_d\AME	Ablabesmyia

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.

$\overline{\rm resolvedCatalogName}$	${\it resolved} {\it Rank}$	count
col	NA	1381
col	class	5
col	family	143
col	genus	362
col	infraorder	1
col	kingdom	1
col	nanorder	1
col	order	22
col	parvorder	1
col	section	1
col	species	780
col	subfamily	9
col	subgenus	32
col	suborder	3
col	subspecies	25
col	subtribe	1
col	superfamily	5
col	superorder	1
	-	

$\underline{{\rm resolvedCatalogName}}$	$\operatorname{resolvedRank}$	count
col	tribe	4
col	variety	2
discoverlife	NA	2721
discoverlife	species	11
gbif	NA	1222
gbif	class	5
gbif	family	145
gbif	genus	394
gbif	kingdom	1
gbif	order	22
gbif	species	925
gbif	subspecies	45
gbif	variety	3
itis	NA	1629
itis	class	5
itis	family	144
itis	genus	290
itis	kingdom	1
itis	order	25
itis	species	591
itis	subclass	1
itis	subfamily	13
itis	suborder	7
itis	subspecies	20
itis	superfamily	6
itis	superorder	1
itis	tribe	1
itis	variety	1
mdd	NA	2731
ncbi	NA	1493
ncbi	class	5
ncbi	family	141
ncbi	genus	353
ncbi	order	22
ncbi	series	1
ncbi	species	666
ncbi	species group	1
ncbi	subclass	1
ncbi	subfamily	16
ncbi	subgenus	13
ncbi	suborder	6
ncbi	subspecies	17
ncbi	superfamily	4
ncbi	superorder	2
	_	

${\bf resolved Catalog Name}$	${\it resolved} {\it Rank}$	count
ncbi	tribe	4
ncbi	varietas	1
pbdb	NA	2302
pbdb	class	5
pbdb	family	131
pbdb	genus	184
pbdb	informal	1
pbdb	infraorder	4
pbdb	kingdom	1
pbdb	order	26
pbdb	species	50
pbdb	subclass	1
pbdb	subfamily	19
pbdb	suborder	9
pbdb	superfamily	5
pbdb	superorder	1
pbdb	tribe	5
pbdb	unranked clade	2
tpt	NA	2545
tpt	family	4
tpt	genus	18
tpt	order	1
tpt	species	163
tpt	subspecificepithet	1
wfo	NA	2543
wfo	family	3
wfo	genus	99
wfo	phylum	1
wfo	species	83
wfo	subspecies	2
wfo	variety	2
worms	NA	2335
worms	class	3
worms	family	98
worms	genus	153
worms	kingdom	1
worms	order	20
worms	species	108
worms	subclass	2
worms	subfamily	3
worms	suborder	6
worms	subspecies	1
worms	superfamily	1

 $\qquad \qquad \text{superorder} \qquad \qquad 1$

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).

$\overline{\rm resolved Catalog Name}$	relationName	count
col	NONE	1772
col	HAS_ACCEPTED_NAME	1482
col	SYNONYM_OF	483
discoverlife	NONE	3300
discoverlife	SYNONYM_OF	1
discoverlife	HAS_ACCEPTED_NAME	10
gbif	NONE	1611
gbif	HAS_ACCEPTED_NAME	1834
gbif	SYNONYM_OF	570
itis	NONE	2024
itis	HAS_ACCEPTED_NAME	1228
itis	SYNONYM_OF	82
mdd	NONE	3240
mdd	HAS_ACCEPTED_NAME	54
mdd	SYNONYM_OF	1
ncbi	NONE	1902
ncbi	SAME_AS	1410
ncbi	SYNONYM_OF	54
ncbi	COMMON_NAME_OF	8
pbdb	NONE	2745
pbdb	HAS_ACCEPTED_NAME	547
pbdb	SYNONYM_OF	45
tpt	NONE	3071
tpt	SYNONYM_OF	87
tpt	HAS_ACCEPTED_NAME	217
wfo	NONE	3067
wfo	HAS_ACCEPTED_NAME	207
wfo	SYNONYM_OF	42
wfo	HAS_UNCHECKED_NAME	23
worms	NONE	2793
worms	HAS_ACCEPTED_NAME	504
worms	SYNONYM_OF	41

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)
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	${\bf review Comment Type}$	${\bf review Comment}$
2025-04-11T01:07:18Z	note	found unsupported interaction type with name: [Pond labeled Section D]
2025-04-11T01:07:18Z	note	found unsupported interaction type with name: [Pond labeled Section E]

reviewDate	${\bf review Comment Type}$	reviewComment
2025-04-11T01:07:18Z	note	source taxon name missing: using institution- Code/collectionCode/collectionId/catalogNumber/occur as placeholder
2025-04-11T01:07:18Z	note	source taxon name missing: using institution- Code/collectionCode/collectionId/catalogNumber/occur 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
found unsupported interaction type with name: [HostSpecies]	623
found unsupported interaction type with name: [PreviousCrop]	578
found unsupported interaction type with name: [A-C)]	339
found unsupported interaction type with name: [Soybeans Variety]	203

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.

 $^{^3\}mathrm{Up}\text{-to-date}$ status of the GloBI Review Badge can be retrieved from the GloBI Review Depot

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 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, Findable, Accessible, Interoperable and Reusable. 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.

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

 $^{^5\}mathrm{At}$ time of writing (2025-04-11) the version of the GloBI dataset index was available at https://globalbioticinteractions.org/datasets

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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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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