

Versioned Archive and Review of Biotic
Interactions and Taxon Names Found within
globalbioticinteractions/maarjam
hash://md5/d5a65712ed0d848e01778c8942624c44

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<https://globalbioticinteractions.org/contribute>
<https://github.com/globalbioticinteractions/maarjam/issues>

2026-03-30

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/maarjam, has fingerprint hash://md5/d5a65712ed0d848e01778c8942624c44, is 58.7MiB in size and contains 64,586 interactions with 1 unique type of association (e.g., hasArbuscularMycorrhizalHost) between 1,535 primary taxa (e.g., *Glomus* sp.) and 657 associated taxa (e.g., *Prunella vulgaris*). This report includes detailed summaries of interaction data, a taxonomic review from multiple catalogs, and an archived version of the dataset from which the reviews are derived.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 :

Öpik, M., Vanatoa, A., Vanatoa, E., Moora, M., Davison, J., Kalwij, J.M., Reier, Ü., Zobel, M. 2010. The online database MaarjAM reveals global and ecosystemic distribution patterns in arbuscular mycorrhizal fungi (Glomeromycota). *New Phytologist* 188: 223-241.

<https://github.com/globalbioticinteractions/maarjam/archive/95ec69414a616178c195c5bd1d3973f31fd432026-03-28T03:09:41.300Z> hash://md5/d5a65712ed0d848e01778c8942624c44

For additional metadata related to this dataset, please visit <https://github.com/globalbioticinteractions/maarjam> 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	0.11.1
elton	0.16.7
nomer	0.6.2
globinizer	0.4.0
mlr	6.0.0
jq	1.6
yq	4.25.3
pandoc	3.1.6.1
duckdb	1.3.1
mapserver	7.6.4

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

```
# get versioned copy of the dataset (size approx. 58.7MiB) under review
elton pull globalbioticinteractions/maarjam

# generate review notes
elton review globalbioticinteractions/maarjam\
> review.tsv

# export indexed interaction records
elton interactions globalbioticinteractions/maarjam\
> interactions.tsv

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

or visually, in a process diagram.

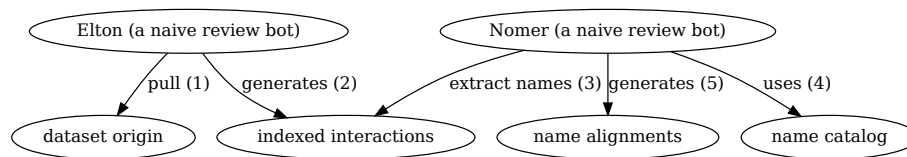


Figure 1: Review Process Overview

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

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

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
<code>biblio.bib</code>	list of bibliographic reference of this review
<code>check-dataset.sh</code>	data review workflow/process as expressed in a bash script
<code>data.zip</code>	a versioned archive of the data under review
HEAD	the digital signature of the data under review
<code>index.docx</code>	review in MS Word format
<code>index.html</code>	review in HTML format
<code>index.md</code>	review in Pandoc markdown format
<code>index.pdf</code>	review in PDF format
<code>indexed-citations.csv.gz</code>	list of distinct reference citations for reviewed species interaction claims in gzipped comma-separated values file format
<code>indexed-citations.html.gz</code>	list of distinct reference citations for reviewed species interactions claims in gzipped html file format
<code>indexed-citations.tsv.gz</code>	list of distinct reference citations for reviewed species interaction claims in gzipped tab-separated values format
<code>indexed-interactions-col-family-col-family.svg</code>	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.parquet	species interaction claims indexed from the dataset under review in Apache Parquet format
indexed-interactions.png	species interaction claims indexed from the dataset under review plotted on a map
indexed-interactions.map	mapserver configuration to plot species interaction claims indexed from the dataset under review on a map
indexed-interactions.gpkg	species interaction claims indexed from the dataset under review in GeoPackage format
indexed-interactions-h3.gpkg	geospatially clustered h3 species interaction claims indexed from the dataset under review in GeoPackage 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

filename	description
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.parquet	taxonomic names found in the dataset under review in Apache Parquet 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
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-col.parquet	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 Apache Parquet 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

filename	description
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-discoverlife.parquet	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 Apache Parquet 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
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

filename	description
indexed-names-resolved-gbif.parquet	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 Apache Parquet 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-itis.parquet	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 Apache Parquet 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-mdd.parquet	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 Apache Parquet 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

filename	description
indexed-names-resolved-ncbi.parquet	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 Apache Parquet 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
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-pbdb.parquet	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 Apache Parquet 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

filename	description
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-tpt.parquet	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 Apache Parquet 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
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

filename	description
indexed-names-resolved-wfo.parquet	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 Apache Parquet 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-resolved-worms.parquet	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 Apache Parquet 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

filename	description
interaction.svg	diagram summarizing the data model used to index species interaction claims
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.

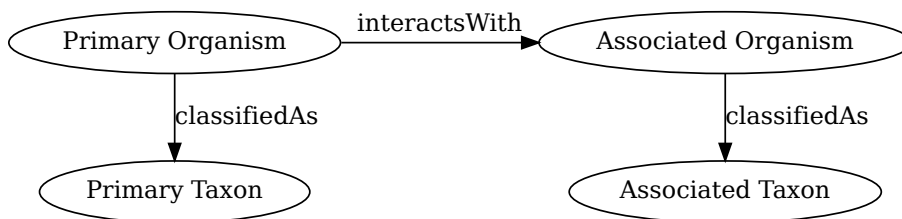


Figure 2: Biotic Interaction Data Model

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 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/maarjam`, has fingerprint hash: `//md5/d5a65712ed0d848e01778c8942624c44`, is 58.7MiB in size and contains 64,586 interactions with 1 unique type of association (e.g., `hasArbuscularMycorrhizalHost`) between 1,535 primary taxa (e.g., `Glomus sp.`) and 657 associated taxa (e.g., `Prunella vulgaris`).

An exhaustive list of indexed interaction claims can be found in gzipped `csv`, `tsv`, `geopackage` and `parquet` 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
Kuklospora PT6	hasArbuscularMycorrhizal	CalzadHostium mucunoides	Oba, H.; Shinozaki, N.; Oyaizu, H.; Tawaraya, K.; Wagatsuma, T.; Barraquio, W.L.; Saito, M., 2004. Arbuscular mycorrhizal fungal communities associated with some pioneer plants in the lahar of Mt. Pinatubo, Philippines. Soil Science and Plant Nutrition.
Kuklospora PT6	hasArbuscularMycorrhizal	CalzadHostium mucunoides	Oba, H.; Shinozaki, N.; Oyaizu, H.; Tawaraya, K.; Wagatsuma, T.; Barraquio, W.L.; Saito, M., 2004. Arbuscular mycorrhizal fungal communities associated with some pioneer plants in the lahar of Mt. Pinatubo, Philippines. Soil Science and Plant Nutrition.

sourceTaxonName	interactionType	targetTaxonName	referenceCitation
Kuklospora PT6	hasArbuscularMycorrhizal	Cladoglossum mucunoides	Oba, H.; Shinozaki, N.; Oyaizu, H.; Tawaraya, K.; Wagatsuma, T.; Barraquio, W.L.; Saito, M., 2004. Arbuscular mycorrhizal fungal communities associated with some pioneer plants in the lahar of Mt. Pinatubo, Philippines. Soil Science and Plant Nutrition.
Kuklospora PT6	hasArbuscularMycorrhizal	Cladoglossum mucunoides	Oba, H.; Shinozaki, N.; Oyaizu, H.; Tawaraya, K.; Wagatsuma, T.; Barraquio, W.L.; Saito, M., 2004. Arbuscular mycorrhizal fungal communities associated with some pioneer plants in the lahar of Mt. Pinatubo, Philippines. Soil Science and Plant Nutrition.

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

interactionTypeName	count
hasArbuscularMycorrhizalHost	64586

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

sourceTaxonName	count
Glomus sp.	46342
Scutellospora sp.	2832
Claroideoglomus sp.	1766
Gigaspora sp.	1144
Acaulospora sp.	1075
Archaeospora sp.	657
Diversispora sp.	601
Paraglomus sp.	501
Scutellospora LH-Sc01	128
Glomus DG9	112
Scutellospora Scut1	96
Glomus DG1	94
Glomus DG2	92
Glomus Glo8	88
Glomus MO-G39	84
Glomus DG3	82
Glomus Glo3	76
Scutellospora Scut2	72
Glomus MO-G3	68

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

targetTaxonName	count
Prunella vulgaris	1665
Hevea brasiliensis	1459
Trachycarpus fortunei	1040
Galium verum	920
Araucaria araucana	861
Potentilla chiloensis	834
Festuca pallescens	824

targetTaxonName	count
Cocos nucifera	813
Rosmarinus officinalis	782
Helictotrichon pratense	767
Plantago lanceolata	765
Manihot esculenta	712
Rubus saxatilis	622
Themeda triandra	621
Zea mays	615
Filipendula vulgaris	594
Hepatica nobilis	579
Convallaria majalis	550
Nephrolepis hirsutula	500

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

sourceTaxonName	interactionType	targetTaxonName	count
Glomus sp.	hasArbuscularMycorrhizalHost	Prunella vulgaris	1376
Glomus sp.	hasArbuscularMycorrhizalHost	Galium aparine	816
Glomus sp.	hasArbuscularMycorrhizalHost	Hevea brasiliensis	774
Glomus sp.	hasArbuscularMycorrhizalHost	Arachis hypogaea	752
Glomus sp.	hasArbuscularMycorrhizalHost	Pinus taeda	736
Glomus sp.	hasArbuscularMycorrhizalHost	Chilodactylus chilensis	730
Glomus sp.	hasArbuscularMycorrhizalHost	Thalictrum fortunei	706
Glomus sp.	hasArbuscularMycorrhizalHost	Festuca pallescens	660
Glomus sp.	hasArbuscularMycorrhizalHost	Helictotrichon pratense	642
Glomus sp.	hasArbuscularMycorrhizalHost	Plantago lanceolata	616
Glomus sp.	hasArbuscularMycorrhizalHost	Rosmarinus officinalis	578
Glomus sp.	hasArbuscularMycorrhizalHost	Themeda triandra	574
Glomus sp.	hasArbuscularMycorrhizalHost	Cocos nucifera	572
Glomus sp.	hasArbuscularMycorrhizalHost	Rubus saxatilis	534
Glomus sp.	hasArbuscularMycorrhizalHost	Filipendula vulgaris	518
Glomus sp.	hasArbuscularMycorrhizalHost	Convallaria majalis	500

sourceTaxonName	interactionType	targetTaxonName	count
Glomus sp.	hasArbuscularMycorrhizalHost	Fraxinus acerensis	444
Glomus sp.	hasArbuscularMycorrhizalHost	Digitalis macroblephara	442
Glomus sp.	hasArbuscularMycorrhizalHost	Phlegelis saxatile	434
Glomus sp.	hasArbuscularMycorrhizalHost	Neophytis hirsutula	418

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](#)

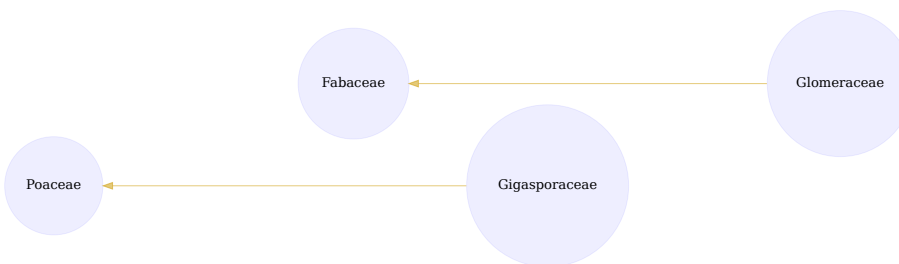


Figure 4: Interactions on the taxonomic family rank as interpreted by the Catalogue of Life. download [svg](#)

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

Geospatial Distribution

If geospatial information was extracted from the dataset under review, the map below will show their distribution. These maps were generated using MapServer (McKenna et al. 2025) tools configured via map configuration indexed-interactions.map :

```
MAP
  SIZE 1600 800
  EXTENT -180 -90 180 90
  PROJECTION
    "init=epsg:4326"
  END
  LAYER # MODIS WMS map from NASA
    NAME "modis_nasa"
    TYPE RASTER
    OFFSITE 0 0 0
    STATUS ON
    CONNECTIONTYPE WMS
    CONNECTION "https://gibs.earthdata.nasa.gov/wms/epsg4326/best/wms.cgi?"

  METADATA
    "wms_srs" "EPSG:4326"
    "wms_name" "OSM_Land_Water_Map"
    "wms_server_version" "1.1.1"
    "wms_format" "image/jpeg"
  END
  CLASS
    STYLE
      COLOR 232 232 232
      OUTLINECOLOR 32 32 32
    END
  END
  LAYER
    NAME "indexed-interactions"
    TYPE POLYGON
    STATUS ON
    CONNECTIONTYPE OGR
    CONNECTION "indexed-interactions-h3.gpkg"
    DATA "indexed-interactions-h3"
    CLASS
      STYLE
        COLORRANGE 253.0 231.0 37.0 32.0 164.0 134.0
        DATARANGE NULL NULL
        RANGEITEM "log_number_of_records"
```

```

        OUTLINECOLOR 0 0 0
      END
    END
  END
END

```

Hexagonal grid cells indicate that interactions claims are available for selected geospatial area: light yellow means relatively fewer claims, dark green relatively more claims.

Figure 5: Hexagonal grid cells indicate that interactions claims are available for selected geospatial area: light yellow means relatively fewer claims, dark green relatively more claims.

Associated data can be found in the geopackage files at indexed-interactions.gpkg for point data and indexed-interactions-h3.gpkg for data clustered in geospatial h3 hexagonals.

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
Acacia erioloba	SYNONYM_OF	col	Vachellia erioloba
Acacia erioloba	SYNONYM_OF	col	Vachellia edgeworthii
Acacia seyal	SYNONYM_OF	col	Vachellia seyal
Acacia seyal	HAS_ACCEPTED_NAME	col	Acacia seyal

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	46
col	family	3
col	genus	38
col	order	1
col	species	573
col	subgenus	1
col	subspecies	32
col	variety	1
discoverlife	NA	675
gbif	NA	38
gbif	family	3
gbif	genus	39
gbif	order	1
gbif	species	582
gbif	subspecies	39
gbif	variety	4
itis	NA	299
itis	family	3
itis	genus	26
itis	order	1
itis	species	335
itis	subspecies	7
itis	variety	6
mdd	NA	674
ncbi	NA	82
ncbi	family	3
ncbi	genus	39
ncbi	order	1
ncbi	species	545
ncbi	subspecies	5
pbdb	NA	644
pbdb	family	3
pbdb	genus	14
pbdb	order	1
pbdb	species	12
tpt	NA	674
wfo	NA	62
wfo	family	3
wfo	genus	24

resolvedCatalogName	resolvedRank	count
wfo	order	1
wfo	species	576
wfo	subspecies	15
worms	NA	538
worms	family	3
worms	genus	12
worms	order	1
worms	species	119
worms	subspecies	3
worms	variety	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	SYNONYM_OF	3245
col	HAS_ACCEPTED_NAME	2702
col	NONE	47
discoverlife	NONE	2829
gbif	SYNONYM_OF	3436
gbif	HAS_ACCEPTED_NAME	2858
gbif	NONE	39
itis	SYNONYM_OF	70
itis	HAS_ACCEPTED_NAME	2003
itis	NONE	779
mdd	NONE	2826
ncbi	SYNONYM_OF	66
ncbi	SAME_AS	2680
ncbi	NONE	85
pbdb	NONE	1268
pbdb	HAS_ACCEPTED_NAME	28
pbdb	SYNONYM_OF	1530
tpt	NONE	2826
wfo	SYNONYM_OF	113
wfo	HAS_ACCEPTED_NAME	564
wfo	NONE	2206
wfo	HAS_UNCHECKED_NAME	43

resolvedCatalogName	relationName	count
worms	NONE	1161
worms	HAS_ACCEPTED_NAME	3203
worms	SYNONYM_OF	1563

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	reviewCommentType	reviewComment
2026-03-30T07:31:29Z	note	target taxon name missing
2026-03-30T07:31:29Z	note	target taxon name missing

reviewDate	reviewCommentType	reviewComment
2026-03-30T07:31:29Z	note	target taxon name missing
2026-03-30T07:31:29Z	note	target taxon name missing

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
target taxon name missing	21431
found unsupported interaction type with name: [cultured spores]	1028
found unsupported interaction type with name: [other]	304
found unsupported interaction type with name: [soil]	93

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 6: 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.

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



Figure 7: 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, **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.

Acknowledgements

We thank the many humans that created us and those who created and maintained the data, software and other intellectual resources that were used for producing this review. In addition, we are grateful for the natural resources providing the basis for these human and bot activities. Also, thanks

⁵At time of writing (2026-03-30) 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."

to <https://github.com/zygoballus> for helping improve the layout of the review tables.

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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