# A Review of Biotic Interactions and Taxon Names Found in globalbioticinteractions/ucsb-izc

by Nomer and Elton, two naive review bots review@globalbioticinteractions.org https://globalbioticinteractions.org/contribute https://github.com/globalbioticinteractions/ucsb-izc/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 process of such an openly accessible digital interactions dataset of known origin, and discuss its outcome. The dataset under review, named globalbioticinteractions/ucsbizc, is 6.27MiB in size and contains 2,369 interaction with 8 unique types of associations (e.g., interactsWith) between 395 primary taxa (e.g., Araneidae) and 463 associated taxa (e.g., Lupinus bicolor). The report includes detailed summaries of interactions data as well as a taxonomic review from multiple catalogs.

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## Introduction

## **Data Review**

Data review can be a time consuming process, especially when done manually. This review report aims to help facilitate data review of species interaction claims made in datasets registered with Global Biotic Interactions (Poelen, Simons, and Mungall 2014). The review includes summary statistics of, and observations about, the dataset under review:

For additional metadata related to this dataset, please visit https://github.c om/globalbioticinteractions/ucsb-izc 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, Elton, Nomer combined with third-party tools like grep, mlr, tail and head.

Table 1: Tools used in this review process

tool name	version
elton	0.13.4
nomer	0.5.13
mlr	6.0.0
pandoc	3.1.6.1

The review process can be described in the form of the script below <sup>1</sup>.

# get versioned copy of the dataset (size approx. 6.27MiB) under review elton pull globalbioticinteractions/ucsb-izc

<sup>&</sup>lt;sup>1</sup>Note that you have to first get the data (e.g., via elton pull globalbioticinteractions/ucsb-izc) before being able to generate reviews (e.g., elton review globalbioticinteractions/ucsb-izc), extract interaction claims (e.g., elton interactions globalbioticinteractions/ucsb-izc), or list taxonomic names (e.g., elton names globalbioticinteractions/ucsb-izc)

- # generate review notes
  elton review globalbioticinteractions/ucsb-izc\
  > review.tsv
- # export indexed interaction records
  elton interactions globalbioticinteractions/ucsb-izc\
- > interactions.tsv
- # export names and align them with the Catalogue of Life using Nomer
  elton names globalbioticinteractions/ucsb-izc\
- | nomer append col\
- > name-alignment.tsv

or visually, in a process diagram.

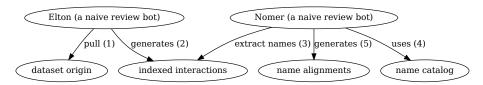


Figure 1: Review Process Overview

You can find a recent copy of the full review script at check-data.sh.

## Results

In the following sections, the results of the review are summarized <sup>2</sup>. Then, links to the detailed review reports are provided.

#### **Biotic Interactions**

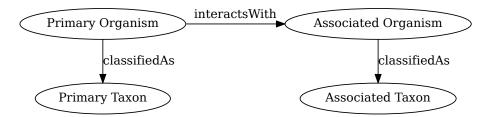


Figure 2: Biotic Interaction Data Model

 $<sup>^2</sup>$ 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.

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/ucsb-izc, is 6.27MiB in size and contains 2,369 interaction with 8 unique types of associations (e.g., interactsWith) between 395 primary taxa (e.g., Araneidae) and 463 associated taxa (e.g., Lupinus bicolor).

An exhaustive list of indexed interaction claims can be found in csv and tsv archives. To facilitate discovery, the first 500 claims available on the html page at indexed-interactions.html are shown below.

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

Table 2: Sample of Indexed Interaction Claims

sourceTaxonName	interactionTypeN	Nam¢argetTaxonName	referenceCitation
Lasioglossum	interactsWith	inside flower of	UCSB-
-		Eschscholzia	IZC00038170
		californica	https://ecdysis.org/collections/individual/index.p
Diptera	adjacentTo	flower of Mimulus	UCSB-
-		auranticus	IZC00038207
			https://ecdysis.org/collections/individual/index.p
Diptera	adjacentTo	flower of Mimulus	UCSB-
	·	auranticus	IZC00038169
			https://ecdysis.org/collections/individual/index.p
Diptera	adjacentTo	flower of	UCSB-
-		Eschscholzia	IZC00038248
		californica	https://ecdysis.org/collections/individual/index.p

Table 3: Most Frequently Mentioned Interaction Types (up to 20 most frequent)

interaction Type Name	count
interactsWith	1574
adjacentTo	573
visits	151
visitsFlowersOf	44
hasHost	18
hostOf	4
eats	3
${\rm coOccursWith}$	2

Table 4: Most Frequently Mentioned Primary Taxa (up to 20 most frequent)

sourceTaxonName	count
Araneidae	133
Lasioglossum	106
Apis mellifera	105
Linepithema humile	87
Cicadellidae	72
Augochlorella pomoniella	63
Agapostemon texanus	47
Solenopsis	46
Hemiptera	39
Heleomyzidae	39
Syrphidae	38
Halictus tripartitus	36
Ceratina acantha	36
Bombus vosnesenskii	35
Temnothorax andrei	35
Colletes	30
Aphididae	27
Lygus	26
Anyphaenidae	25

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

targetTaxonName	count
Lupinus bicolor	282
Lupinus nipomensis	99
Quercus agrifolia	96
Venegasia carpesioides	63
Erigonium	59
Arctostaphylos	49
Marrubium	46
Populus	42
Brassica	40
Salix	39
Encelia californica	37
Foeniculum	36
Atriplex lentiformis	32
Cupressus macrocarpa	31
Heteromeles arbutifolia	28

targetTaxonName	count
flower: Malocothamnus fasciculatus	28
Lupinus succulentus	26
Convolvulus arvensis	26
flower: Calystegia macrostegia	24

Table 6: Most Frequent Interactions between Primary and Associate Taxa (up to 20 most frequent)

sourceTaxonName	interaction Type Name	target Taxon Name	count
Cicadellidae	interactsWith	Lupinus bicolor	52
Solenopsis	interactsWith	Quercus agrifolia	32
Heleomyzidae	interactsWith	Lupinus bicolor	30
Lasioglossum	interactsWith	Venegasia carpesioides	30
Linepithema humile	interactsWith	Populus	29
Lygus	interactsWith	Lupinus bicolor	24
Linepithema humile	interactsWith	Salix	24
Aphididae	interactsWith	Lupinus bicolor	22
Hemiptera	interactsWith	Lupinus bicolor	21
Apis mellifera	interactsWith	Arctostaphylos	19
Temnothorax andrei	interactsWith	Quercus agrifolia	17
Apis mellifera	interactsWith	Marrubium	16
Araneidae	interactsWith	Cylindropuntia prolifera	15
Araneidae	interactsWith	Opuntia cf. littoralis	15
Closterocoris amoenus	adjacentTo	Lupinus nipomensis	15
Apis mellifera	interactsWith	Erigonium	15
Syrphidae	interactsWith	Lupinus bicolor	14
Andrena principalis	interactsWith	Arctostaphylos	13
Thrips	adjacentTo	Lupinus nipomensis	13

#### 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. A tab-separated file can be found at indexed-interactions.tsv  $\,$ 

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



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

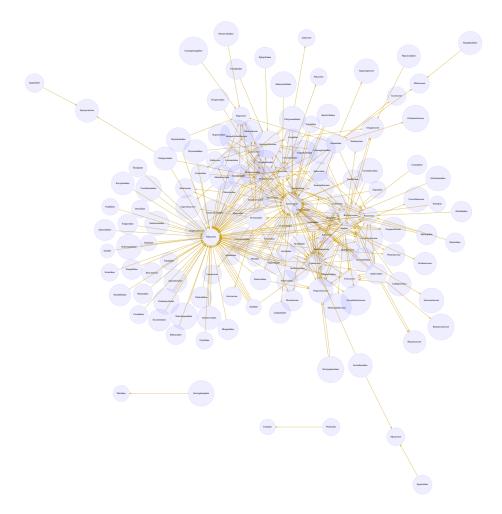


Figure 4: Interactions on the taxonomic family rank as interpreted by the Catalogue of Life. download svg  $\,$ 

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 7: Sample of Name Alignments

providedName	relationName	${\it resolved} {\it CatalogNa}$	mæesolvedName
Distichlis spicata	HAS_ACCEPTED	_ <b>c</b> NoAME	Distichlis spicata
Cryptonevra	HAS_ACCEPTED	_d\AME	Cryptonevra
nigritarsis			nigritarsis
Eulonchus	HAS_ACCEPTED	_d\AME	Eulonchus
Meromyza	HAS_ACCEPTED	_ <b>&amp;A</b> ME	Meromyza

Table 8: 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	280
col	class	1
col	family	105
col	genus	116
col	kingdom	1
col	order	12
col	species	229
col	subfamily	9
col	subgenus	4
col	suborder	2
col	subspecies	8
col	subtribe	1
col	superfamily	4
col	tribe	2
col	variety	1
discoverlife	NA	697
discoverlife	species	64
gbif	NA	252
gbif	class	1

$\overline{\rm resolved Catalog Name}$	${\it resolved} {\it Rank}$	count
gbif	family	108
gbif	genus	133
gbif	kingdom	1
gbif	order	12
gbif	species	250
gbif	subspecies	16
gbif	variety	8
itis	NA	263
itis	class	1
itis	family	105
itis	genus	103
itis	infraorder	2
itis	kingdom	1
itis	order	12
itis	species	236
itis	subclass	1
itis	subfamily	10
itis	suborder	8
itis	subspecies	13
itis	superfamily	4
itis	superorder	1
itis	variety	1
mdd	NA	760
ncbi	NA	279
ncbi	class	1
ncbi	cohort	1
ncbi	family	106
ncbi	genus	115
ncbi	infraorder	3
ncbi	order	12
ncbi	species	217
ncbi	subclass	1
ncbi	subfamily	16
ncbi	subgenus	8
ncbi	suborder	3
ncbi	subspecies	3
ncbi	superfamily	6
ncbi	varietas	1
pbdb	NA	543
pbdb	class	2
pbdb	family	106
pbdb	genus	59
pbdb	infraclass	1
pbdb	infraorder	1

resolvedCatalogName	resolvedRank	count
pbdb	kingdom	1
pbdb	order	13
pbdb	species	10
pbdb	subfamily	15
pbdb	suborder	5
pbdb	superfamily	6
pbdb	unranked clade	3
tpt	NA	756
tpt	genus	1
tpt	species	3
wfo	NA	603
wfo	family	1
wfo	genus	48
wfo	phylum	1
wfo	species	106
wfo	subspecies	2
wfo	variety	1
worms	NA	551
worms	family	83
worms	genus	52
worms	kingdom	1
worms	order	11
worms	species	52
worms	subclass	2
worms	suborder	4
worms	subspecies	1
worms	superfamily	3

Table 9: 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	520
col	SYNONYM OF	79
col	NONE	342
discoverlife	NONE	804
discoverlife	SYNONYM_OF	23

$\overline{\rm resolvedCatalogName}$	relationName	count
discoverlife	HAS_ACCEPTED_NAME	61
discoverlife	HOMONYM_OF	2
gbif	HAS_ACCEPTED_NAME	643
gbif	SYNONYM_OF	139
gbif	NONE	314
itis	HAS_ACCEPTED_NAME	520
itis	NONE	324
itis	SYNONYM_OF	35
mdd	NONE	864
mdd	HAS_ACCEPTED_NAME	1
ncbi	SAME_AS	527
ncbi	SYNONYM_OF	22
ncbi	NONE	344
ncbi	COMMON_NAME_OF	1
pbdb	NONE	622
pbdb	HAS_ACCEPTED_NAME	250
pbdb	SYNONYM_OF	16
tpt	NONE	859
tpt	HAS_ACCEPTED_NAME	6
wfo	HAS_ACCEPTED_NAME	167
wfo	NONE	687
wfo	HAS_UNCHECKED_NAME	9
wfo	SYNONYM_OF	28
worms	HAS_ACCEPTED_NAME	223
worms	NONE	634
worms	SYNONYM_OF	19

Table 10: List of Available Name Alignment Reports

catalog name	alignment results
col	associated names alignments (first 500, full csv/tsv)
ncbi	associated names alignments (first 500, full csv/tsv)
discoverlife	associated names alignments (first 500, full csv/tsv)
gbif	associated names alignments (first 500, full csv/tsv)
itis	associated names alignments (first 500, full csv/tsv)
wfo	associated names alignments (first $500$ , full $csv/tsv$ )

catalog name	alignment results
mdd	associated names alignments (first 500, full csv/tsv)
tpt	associated names alignments (first 500, full csv/tsv)
pbdb	associated names alignments (first 500, full csv/tsv)
worms	associated names alignments (first 500, full csv/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 11: First few lines in the review notes.

reviewDate	${\bf review Comment Type}$	reviewComment
2024-07-23T01:26:39Z	note	source taxon name
		missing: using
		institution-
		Code/collectionCode/collectionId/catalogNumber/occur
		as placeholder
2024-07-23T01:26:39Z	note	found unsupported
		interaction type with
		name: [flying]
2024-07-23T01:26:40Z	note	source taxon name
		missing: using
		institution-
		Code/collectionCode/collectionId/catalogNumber/occur
		as placeholder
2024-07-23T01:26:41Z	note	found unsupported
		interaction type with
		name: [11]

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

Table 12: Most frequently occurring review notes, if any.

reviewComment	count	
source taxon name missing: using	12	
institution-		
Code/collectionCode/collectionId/catalogNumber/occurrenceId		
as placeholder	,	
found unsupported interaction type	3	
with name: [11]		
found unsupported interaction type	3	
with name: [Hovering over]		
found unsupported interaction type	1	
with name: [flying]		

For addition information on review notes, please have a look at the first 500 Review Notes or the download full 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 <sup>3</sup>

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 <sup>4</sup>

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

<sup>&</sup>lt;sup>4</sup>Up-to-date status of the GloBI Index Badge can be retrieved from GloBI's API

If you'd like to keep track of reviews or index status of the dataset under review, please visit [GloBI's dataset index ^[At time of writing (2024-07-23) the version of the GloBI dataset index was available at https://globalbioticinteractions.org/datasets for badge examples.

#### Discussion

This review aims to provide a perspective on the dataset to aid in understanding of species interaction claims discovered. However, it is important to note that this review does *not* assess the quality of the dataset. Instead, it serves as an indication of the open-ness<sup>5</sup> 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 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.

#### Author contributions

Nomer was responsible for name alignments. Elton carried out dataset extraction, and generated the review notes.

#### References

ICZN. 1999. "International Code of Zoological Nomenclature." The International Trust for Zoological Nomenclature, London, UK. https://www.iczn.org/the-code/the-code-online/.

<sup>&</sup>lt;sup>5</sup>According to http://opendefinition.org/: "Open data is data that can be freely used, re-used and redistributed by anyone - subject only, at most, to the requirement to attribute and sharealike."

- Poelen, Jorrit H., James D. Simons, and Chris J. Mungall. 2014. "Global Biotic Interactions: An Open Infrastructure to Share and Analyze Species-Interaction Datasets." *Ecological Informatics* 24 (November): 148–59. https://doi.org/10.1016/j.ecoinf.2014.08.005.
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