



# Getting Your Data Published: Sending Data to iDigBio



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TORCH Workshop

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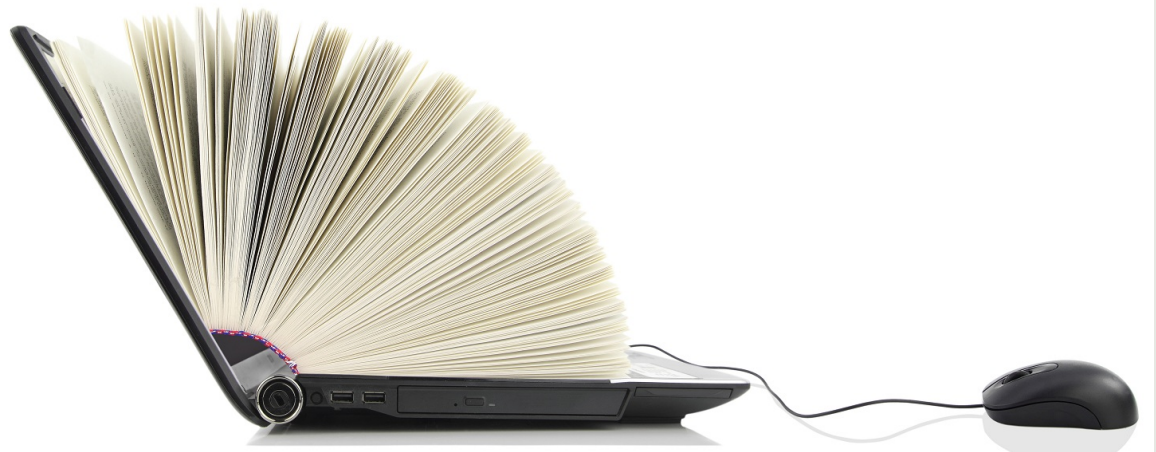
[https://www.idigbio.org/wiki/index.php/Data\\_Ingestion\\_Guidance](https://www.idigbio.org/wiki/index.php/Data_Ingestion_Guidance)



*iDigBio is funded by a grant from the National Science Foundation's Advancing Digitization of Biodiversity Collections Program (Cooperative Agreement EF-1115210). Any opinions, findings, and conclusions or recommendations expressed in this material are those of the author(s) and do not necessarily reflect the views of the National Science Foundation.*

## What do we mean by data publishing?

*making biodiversity data publicly accessible & discoverable, in a standardized form, via a URL.*



# Why publish data? The 4 biggies for data aggregation

# ACCESSIBILITY

# Data Use

# Data Quality

# Attribution



## Data publishing: where to begin with iDigBio?

- Email [data@idigbio.org](mailto:data@idigbio.org)
- There are three ways to share data:

**Least Ideal**

**Most Ideal**



Technical skill vs. time, updatability, data buy-back etc.

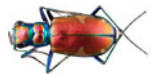
## DATA Method #1 – BEST

- What you already send to GBIF
  - Using Darwin Core field names
  - Packaged in a Darwin Core Archive (DwC-A)
  - On an RSS feed (produced by **IPT**)



## DATA Method #2 – BETTER+

- When you mark your data to publish, all the necessary parts of the package are generated.
  - Custom Darwin Core Archive (DwC-A) on an RSS feed produced by Symbiota
  - almost automatic media
  - <http://symbiota.org>



**Symbiota**

Promoting  
Bio-Collaboration



## DATA #3 – GOOD ENOUGH

- Export your data as CSV/TXT file with DwC fieldnames & let us host it on our IPT or VertNet's

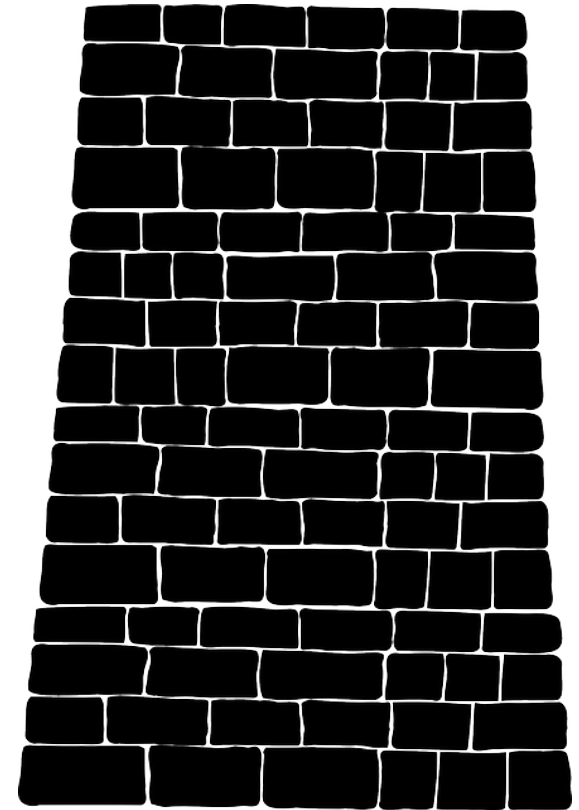
## DATA #4 – Sub Par

- Throw the data over the wall and let us prepare it.
- Has its challenges:
  - data manipulations
  - UUID, data cleansing

↓ buy-back

↓ updates

↓ backlog





## 3 ways to get media to iDigBio:

1. Use Audubon Core extension in IPT

➤ Linked to the specimen

2. Via Symbiota

➤ Linked to the specimen

3. Media appliance

➤ Can be linked to the specimen

## **DATASET INFO: info about the provider (metadata)**

Send your dataset **metadata** with your provider information (eml.xml):

- responsible parties (name, address, email, role)
- institution name, institution code
- URL to the data at your institution
- descriptive paragraph about the institution, collection, and the dataset

## DATASET INFO: rights

Include data rights and rightsHolder information:

- Use Creative Commons standards:

- CC0 for data (not copyrightable)



- CC BY for media (at least)



## **DATASET INFO: update collections lists**

- iDigBio Collections

<https://www.idigbio.org/portal/collections>

- Index Herbariorum

<http://sweetgum.nybg.org/ih/>

- GRBio.org Repositories:

<http://grbio.org/find-biorepositories>

Do you know what your **institutionCode** is?

## Data Quality: Consider searchability in the aggregate

Dates – dwc:eventDate, dwc:day, dwc:month, dwc:year:

- this is not a month: Spring
- this is not a day: 10-18
- this is not a year: 1989? Or [1989]

Taxonomy – fill in dwc:scientificName, parse out the elements, fill in higher taxonomy

- this is not a species: shrimp, daisy

Tics: \* [] {} ?

- Use the verbatim and remarks fields for things that do not fit the definitions.

## Data Quality: Grooming and tics

Your dataset **is no longer just for making labels**, there are other considerations for being digital, and out in the wild:

- 1) Put dates in ISO 8601 format, i.e., YYYY-MM-DD, e.g., 2015-09-17
- 2) Parse apart scientific name
- 3) Conversely, put the piece parts into a scientific name
- 4) Provide as much higher taxonomy as you feel comfortable with, fill in tribe, sub+super family, kingdom, division, class, order) get out of 'family' land.
- 5) Make sure lat and lon coordinates are in decimal, and no N, S, E, W
- 6) Do not export '0' in fields to represent no value, e.g., lat or lon, height
- 7) put elevation in METERS units in the elevation field without the units (e.g., the fields dwc:minimumElevationInMeters and dwc:maximumElevationInMeters already assume the numeric values are in meters, so there no need to include the units with the data)
- 8) And not to get too esoteric, do not use un-escaped newline characters or embedded tabs
- 9) Watch out for diacritics, save in UTF-8

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## When is my work done?

- Digitization is never done
  - Label data
  - Georeferenced
  - Image
- Not until your data are in iDigBio.
  - It is not enough to get to it to Symbiota
    - Publish, re-publish with updates

## Symbiota Notes

- Your dataset
  - Give it a complete name, institution, collection/herbarium
  - Description of the collection – what is in THIS data
  - Good contacts - the person who will respond to requests
- Join the Symbiota working group – community, webinars



# Data Management Plan

- Build a robust DMP – look at DataOne
  - Who will be contributing data (roles and responsibilities)?
    - With what software are they managing their data?
    - Metadata used (Darwin Core?)
  - How will they be mobilizing it
    - Dataset names
    - GUIDs (occurrenceID)
  - Record counts, media counts
  - How they will get it to iDigBio
  - Archiving strategy, backup protocol
  - Responsible parties
  - Other repositories (GBIF, VertNet)
  - Data extensions (e.g., associations)

## Don't Wait – Catch Up!

- Hundreds of herbaria (390+) are ahead of you in their digitization efforts
- Sharing data keeps you relevant
- TORCH: thank you so far !
  - TEX/LL (Tom & George)
  - ASU (Marcy) coming soon
  - BRIT (Jason - Bryophytes & Lichens)
  - CSU – (Clark - Macrofungi)



# Thank you for your attention



[www.idigbio.org](http://www.idigbio.org)



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<webcal://www.idigbio.org/events-calendar/export.ics>



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