

Accessing Digital Collections Data Sources for Research: A Tour of iDigBio Data Services

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What's in iDigBio's Repository?

Vouchered specimen records!

- Metadata in Darwin Core, Audubon Core, and userdefined fields
- Images of specimens
- All groups including paleo
- US and international

List	Labels Media Reco	ordsets				Total: 28,164
Family	Scientific Name	Date Collected	Country	Institution Code	Basis of Record	Columns
ACERACEAE	Acer abchasicum Rupr.	no data	Brasil	Rio de Janeiro Botani	PreservedSpecimen	view
ACERACEAE	Acer abchasicum Rupr.	no data	Brasil	Rio de Janeiro Botani	PreservedSpecimen	view
Sapindaceae	Acer accuminata	no data	United States	UCMP	PreservedSpecimen	view
Aceraceae	Negundo aceroides	1850-10-5	no data	MSC	PreservedSpecimen	view
Aceraceae	Negundo aceroides	1887-5-4	United States	MSC	PreservedSpecimen	view
Aceraceae	Acer negundo L. ssp	1891-06-20	USA	UConn	Preserved Specimer	n view
Aceraceae	Negundo aceroides	1895-4-27	United States	MSC	PreservedSpecimen	view
Aceraceae	Negundo aceroides	1895-8-25	United States	MSC	PreservedSpecimen	view
Aceraceae	Negundo aceroides	no data	no data	MSC	PreservedSpecimen	view
ACERACEAE	Acer acinatum Hort.	no data	Brasil	Rio de Janeiro Botani	PreservedSpecimen	view
Sapindaceae	Acer acuminatum Wal	1893-07-20	United States	MO	PreservedSpecimen	view
Sapindaceae	Acer acuminatum	no data	India	UCMP	PreservedSpecimen	view
ACERACEAE	Acer acuminatum Wal	no data	Brasil	Rio de Janeiro Botani	PreservedSpecimen	view







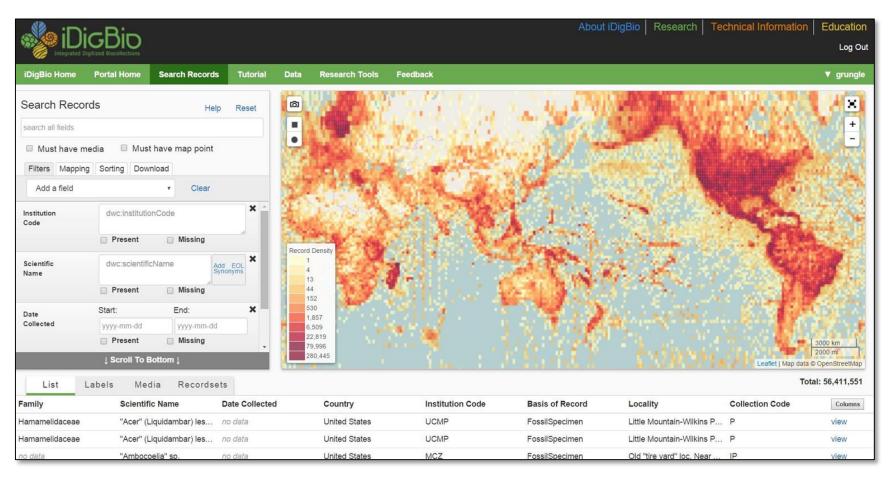
Accessing iDigBio Data

Advantages of the different ways

- 1. Website/portal good to see what's there
- 2. Downloads flexible analysis, permanence of your custom data set
- 3. API exploration of data, building applications
- **4.** Packages easier than direct use of API, literate programming, notebooks, exploration
- 5. Computation big data, cross-dataset analysis



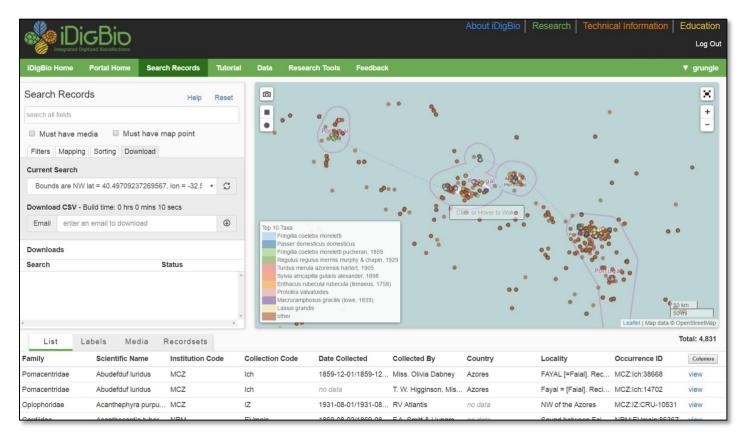
Seeing What's There - The Portal



http://portal.idigbio.org/



Getting What's There - Downloads



Support for "download-and-code" workflow



What's in a Download from iDigBio

Name 📤	Date modified	Туре	Size
meta.xml	2/21/2016 11:25 PM	XML Docu	30 KB
multimedia.csv	2/21/2016 11:25 PM	Microsoft E	1 KB
multimedia_raw.csv	2/21/2016 11:25 PM	Microsoft E	2 KB
occurrence.csv	2/21/2016 11:25 PM	Microsoft E	920 KB
occurrence_raw.csv	2/21/2016 11:25 PM	Microsoft E	737 KB
records.citation.txt	2/21/2016 11:25 PM	Text Docu	1 KB

```
| http://www.idigbio.org/portal (2016),
| 2 Query: {"filtered": {"filter": {"and": [{"term": {"recordset": "ef30a918-b583-41f1-9ac4-4a37591b515a"}}]}},
| 3 1376 records, accessed on 2016-02-21T23:23:08.677528,
| 4 contributed by 1 Recordsets, Recordset identifiers: | http://www.idigbio.org/portal/recordsets/ef30a918-b583-41f1-9ac4-4a37591b515a (1376 records) | 6
```

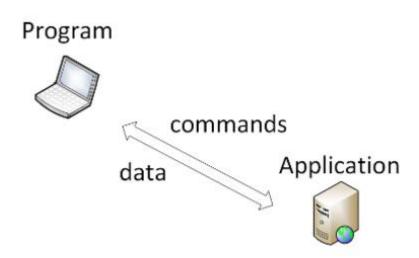
Processed records, raw records, media metadata, record metadata, citation information - everything you need to do reproducible analyses



Getting What's There - API

API: Application Programming Interface

- I want to do some Programming
- using parts of someone else's Application
- and I need an Interface that describes the commands and data to do it.





Getting What's There - API Through a Web Browser

```
search.idigbio.org/v2/sear ×
  ⇒ C search.idigbio.org/v2/search/records?rg={"genus":"acer"}
🛅 Loop 🦲 Personal 🦲 Work 🦲 Reading List 🦲 R 🦲 Learning 🦲 MySQL 🦲 SC/DC
                                                                                Other bookmarks
                                                                             + - View source 🦚
   itemCount: 28164,
   lastModified: "2016-06-27T13:44:46.660Z",
 - items: [
           uuid: "81a6cc35-3180-4c7c-b6b8-951a303a8b18",
           type: "records",
           etag: "61eaf5924d7136f772cce1cec9e5beb7b709fe94",
         - data: {
               dwc:specificEpithet: "medianum",
               dwc:latestPeriodOrHighestSystem: "Tertiary",
               dwc:kingdom: "Plantae",
               dwc:recordedBy: "Webb, S.D.",
               dwc:order: "Sapindales",
               dcterms:references: "http://ucmpdb.berkeley.edu/cgi/ucmp query2?
               spec id=P9078&one=T",
               dwc:individualCount: "1",
```



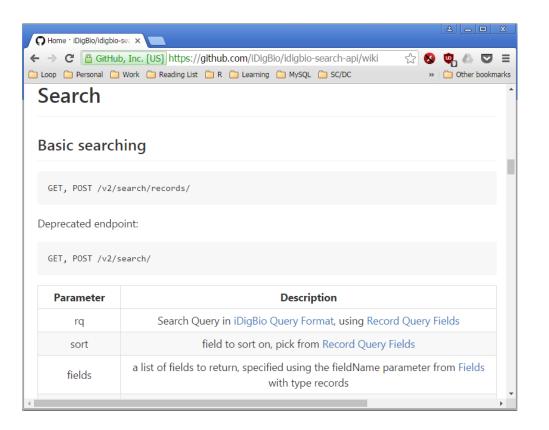
Analyzing What's There - Summary API Through a Web Browser

```
search.idigbio.org/v2/sum ×
          search.idigbio.org/v2/summary/top/records/?top_fields=["basisofrecord"] 5
🗀 Loop 🦲 Personal 🦲 Work 🦲 Reading List 🦲 R 🦲 Learning 🦲 MySQL 🦲 SC/DC
                                                                                            Other bookmarks
                                                                                           + - View source 🙀
  basisofrecord: {
      - preservedspecimen: {
            itemCount: 60536721
      - fossilspecimen: {
            itemCount: 3061794
      - machineobservation: {
            itemCount: 220317
      - livingspecimen: {
            itemCount: 2814
      - humanobservation: {
            itemCount: 889
    itemCount: 64015275
```



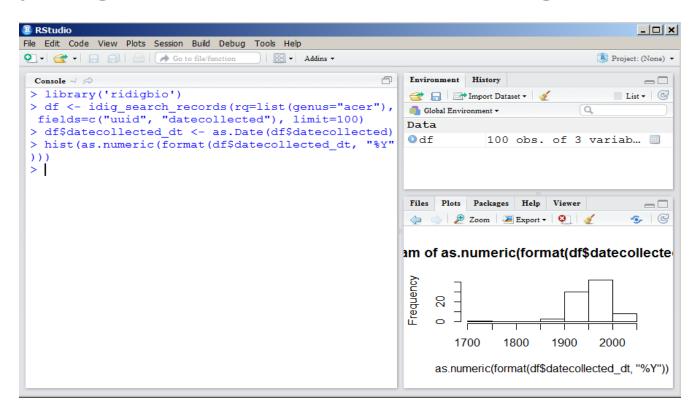
API Documentation

https://www.idigbio.org/wiki/index.php/IDigBio_API





Analyzing What's There - Packages



R ridigbio: https://cran.r-project.org/web/packages/ridigbio/index.html Python idigbio: https://pypi.python.org/pypi/idigbio



Analyzing What's There - Applications



http://lifemapper.org/

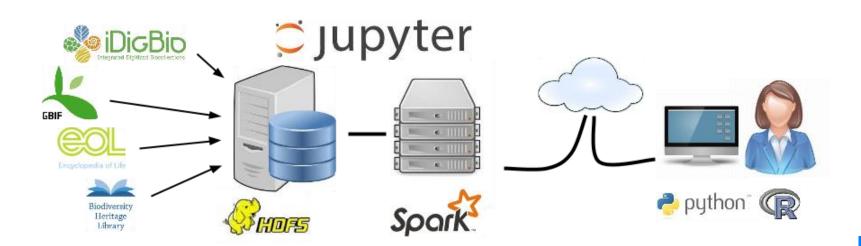


Analyzing What's There - GUODA

Global Unified Open Data Access

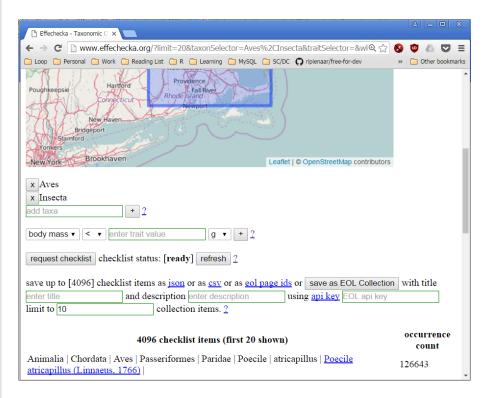
An informal collaboration between technologists from organizations like EOL, ePANDDA, and iDigBio as well as independent biodiversity informaticists.

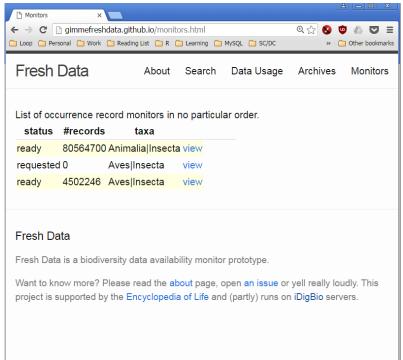
http://guoda.bio





Analyzing What's There - Apps on GUODA





http://www.effechecka.org/



http://gimmefreshdata.github.io/



It's Not Just Us!

Everyone has portals, downloads, APIs, and packages









Where to find some more of them:

https://ropensci.org/packages/

https://www.biodiversitycatalogue.org/services



Challenges - Citation of Data

Easy access can complicate citation of data

- How do you cite 1000's of records in limited space?
- Aggregated datasets, who to cite?
- Permanently recording which records were used

http://www.gbif.org/resource/80761 https://www.idigbio.org/content/idigbio-terms-use-policy



Challenges - Data Quality

We keep the raw data as provided to us but we interpret the data and adjust it to make it searchable such as:

- Flip and sign-correct lat & Ion
- Format dates
- Fill higher taxonomy from GBIF backbone

Data Quality flags are provided as fields

GBIF/TDWG working group for standardizing data changes (with GBIF & ALA)



Opportunities - Reproducible Research Workflows

What is all this for? Why have anything beyond just data dumps? Building efficient, collaborative, and reproducible scientific research!

- Collaborate with peers
- Iterate quickly on ideas
- Re-use and build on the data and code of others
- Publish the complete resources (data and code) to reproduce research



Thank you!





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idigbio.org/rss-feed.xml



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









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