

# Advancing Scientific Research with Digitized Data: *using and reusing biodiversity data*

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Florida Museum of Natural History  
University of Florida



*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. All images used with permission or are free from copyright.*

# Collections: The Library of Life

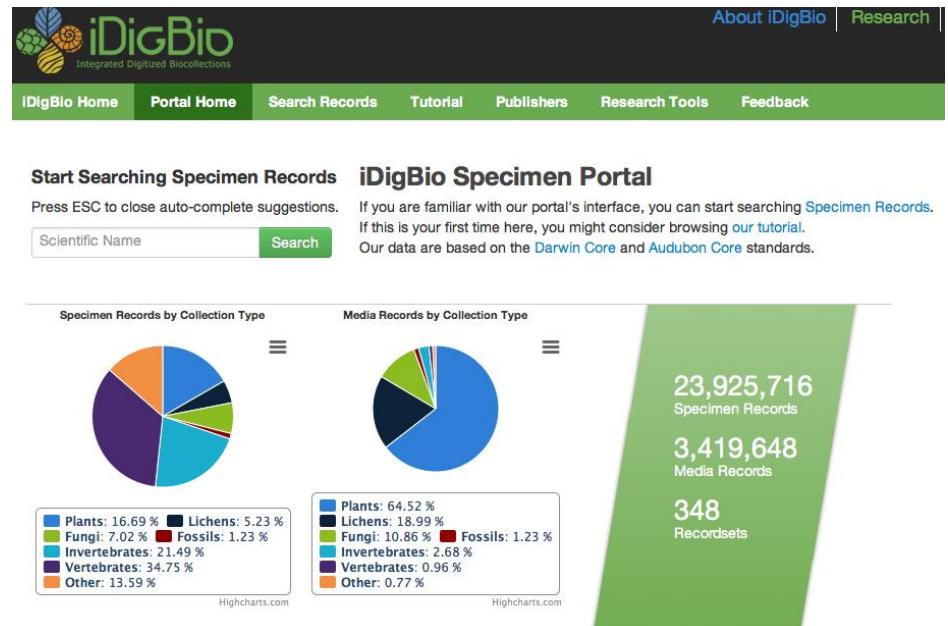


>1600 natural history collections  
in the US alone  
1-2 billion specimens  
in the US  
3-4 billion specimens  
worldwide

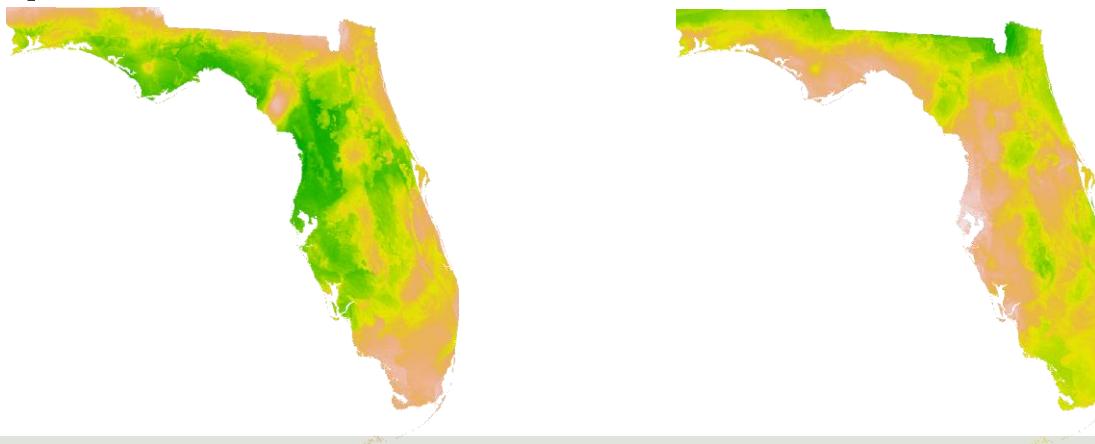


# Research @ iDigBio

- Data portal



- Computational workflows – research use



# Linking Collections to...

- Phylogenies
- Ecology
- Genomics
- Paleontology
- Living Collections
- Other Repositories



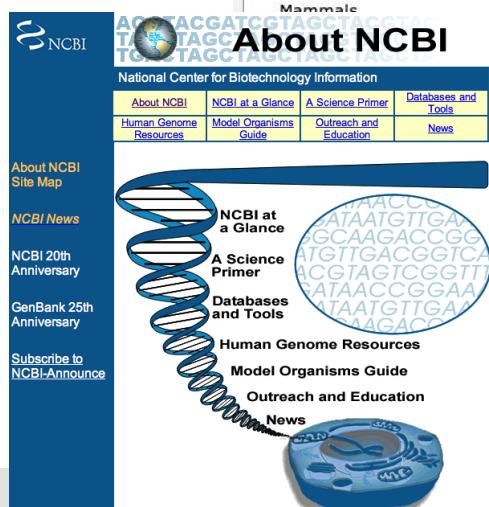

The Paleobiology Database interface features a central search bar with the text "Quick search". Below the search bar is a grid of icons representing various fossil groups: bird, spiral shell, skull, starfish, ammonite, trilobite, and coral. To the left is a circular diagram illustrating the evolution of seed plants, showing the transition from Gymnosperms to Seed Plants (represented by a green circle) and then to Mesangiosperms. To the right is a large, detailed phylogenetic tree of life, with major clades labeled in purple: Raphitoma, Diploria, Duong, UTEX, Actinobacteria, Prochlorococcus, Synechococcus, and others. The tree is densely branched with many smaller labels for specific genera and species.



**Welcome to Morphbank**  
 User: Guest [\[click to login\]](#)



**TreeBASE**  
*A Database of Phylogenetic Knowledge*



**About NCBI**  
 National Center for Biotechnology Information

<a href="#">About NCBI</a>	<a href="#">NCBI at a Glance</a>	<a href="#">A Science Primer</a>	<a href="#">Databases and Tools</a>
<a href="#">Human Genome Resources</a>	<a href="#">Model Organisms Guide</a>	<a href="#">Outreach and Education</a>	<a href="#">News</a>

NCBI at a Glance  
 A Science Primer  
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 News



# Synthetic Research and Complex Data Analysis

- Large data sets
- Heterogeneous data
- Multiple software packages
- Multi-step analytical workflows
- Training

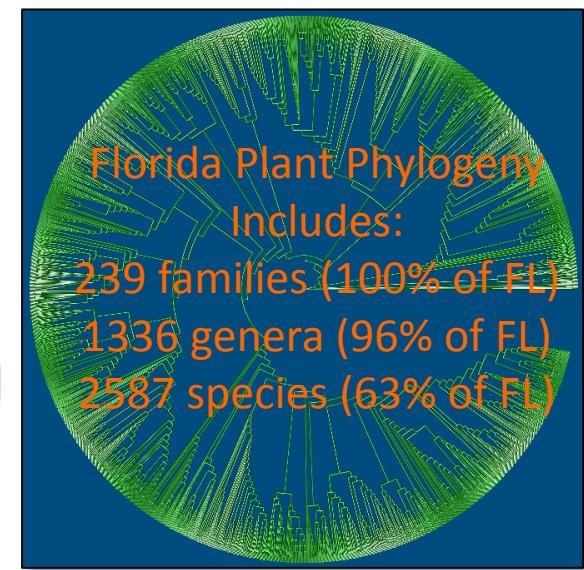
# Synthetic Research: Connections

- Connecting with ecological data
  - Ecological niche modeling
    - Climate change
    - Polyploidy
- Connecting with phylogeny
  - PhyloJive
  - Open Tree of Life
- Connecting with software and tools



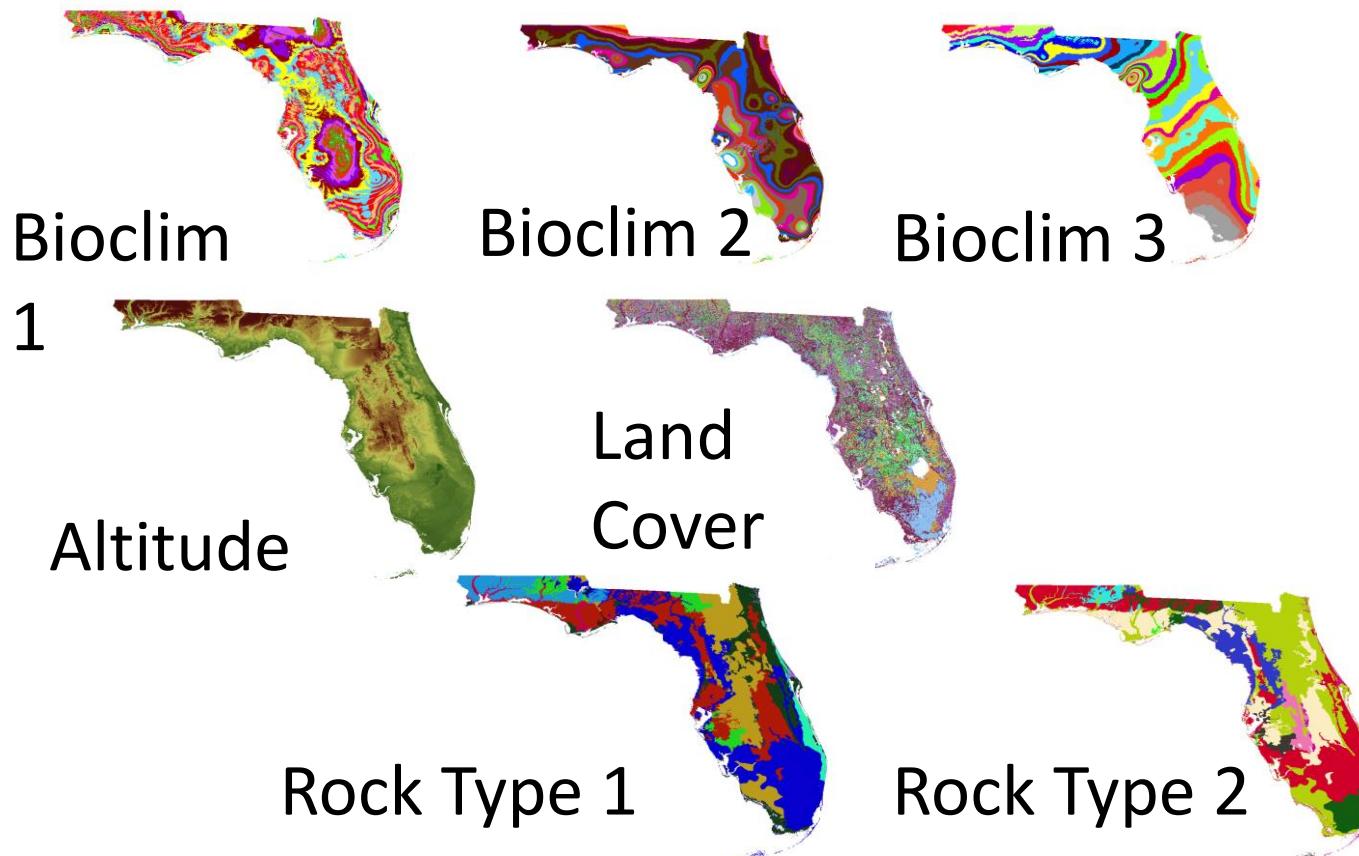
# Florida Plant Diversity in a Changing Climate

Integrating herbarium specimen data, ENM,  
climate change models, and phylogeny



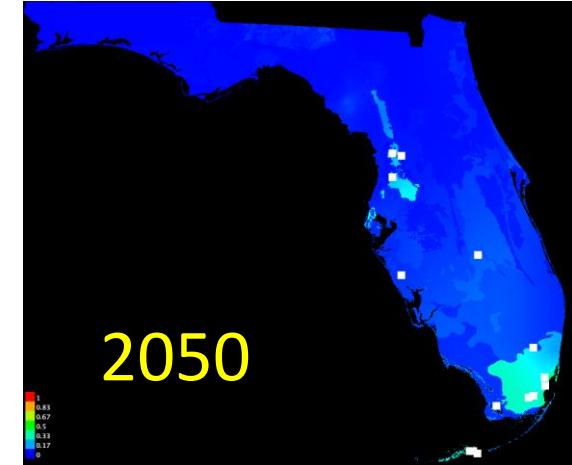
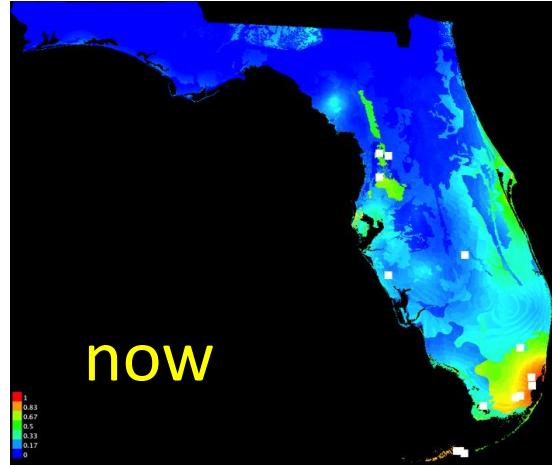
Charlotte Germain-Aubrey

# Ecological Niche Modeling: Environmental Layers

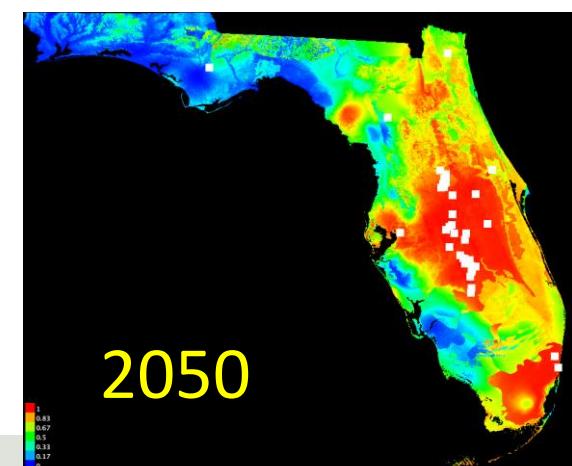
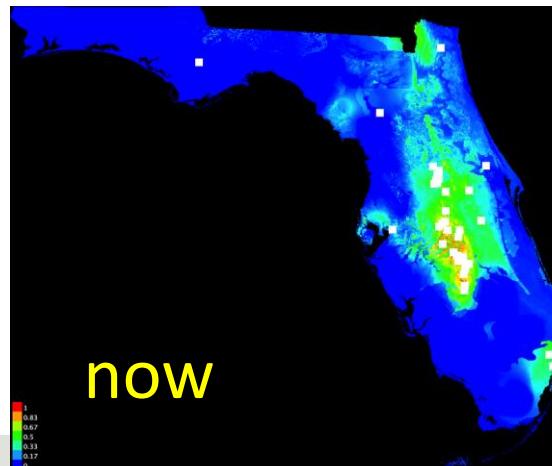


# Responses to Climate Change: Winners & Losers

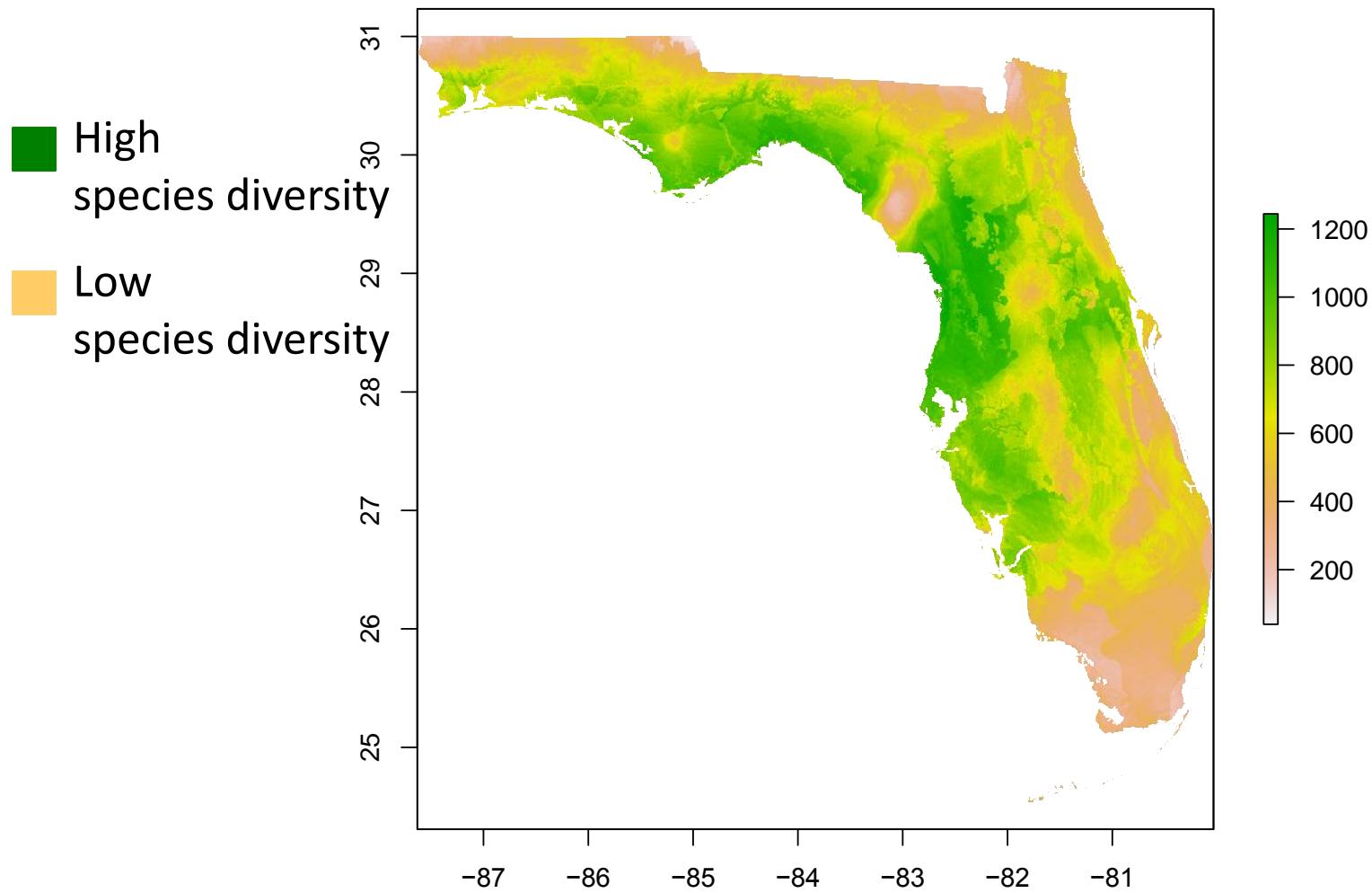
## *Abildgaardia ovata* (flatspike sedge)



## *Prunus geniculata* (scrub plum)



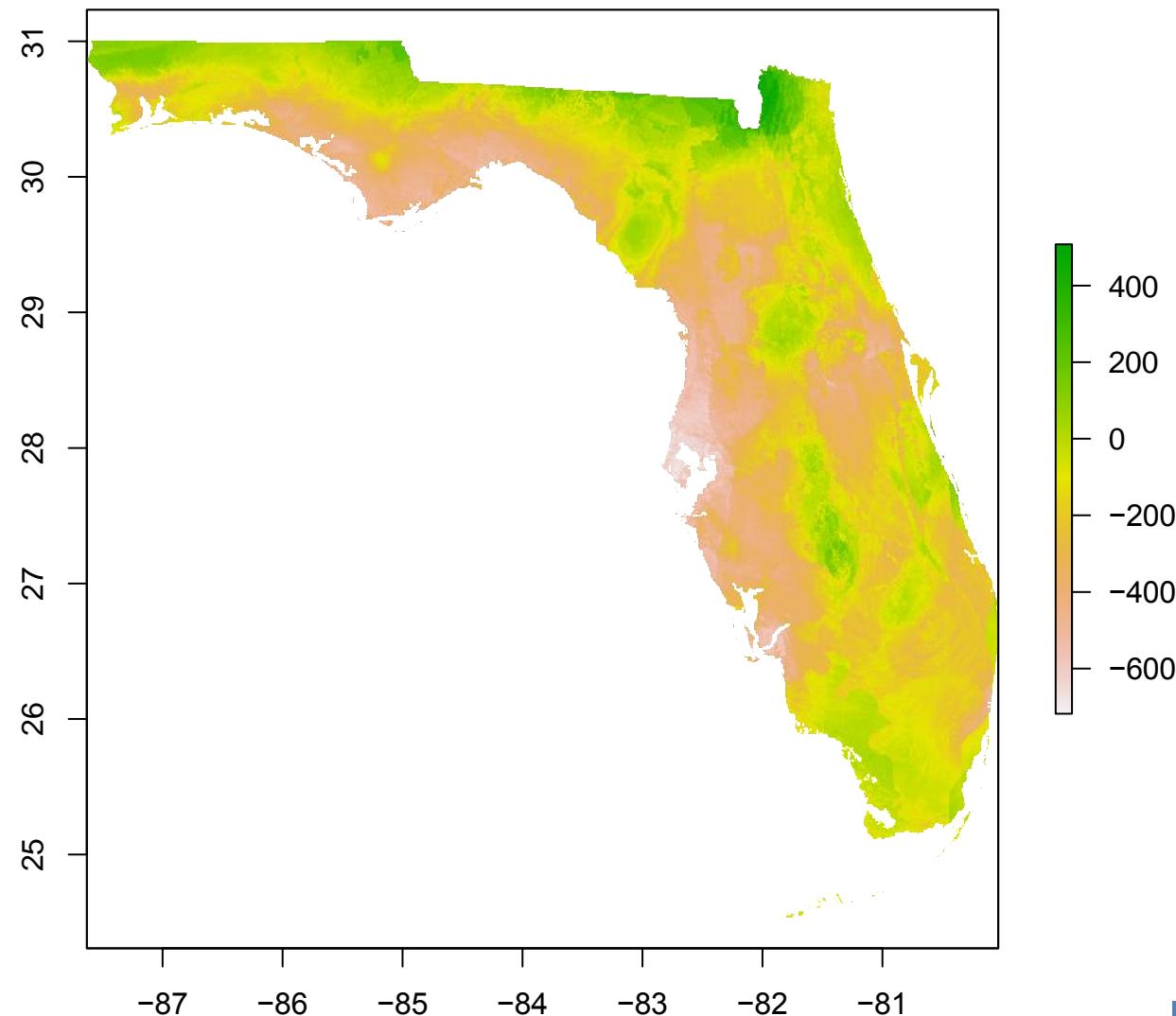
# Florida plant diversity heat map: now



# Between now and 2050...

- Panhandle species moving NORTH!
- Peninsula species moving SOUTH!

# spp 2050 - #spp now



# R package coming soon...

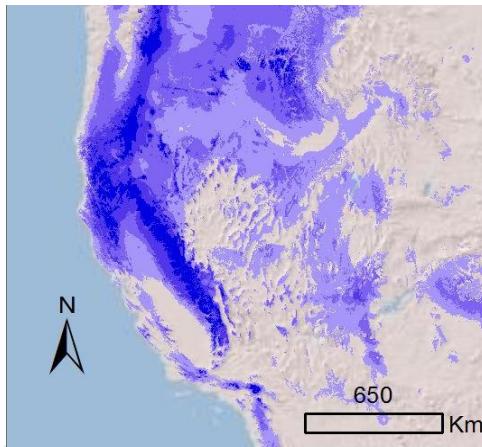
- Data cleaning
  - Lat/long – round decimal points
  - Remove duplicates
  - Bring points back to the coast
  - Removing outliers using a shape file of the desired area
- Extracting climate data from year of collection
  - Prism format
  - Bioclim format
  - If no year given, then default average
- Output formatted for Maxent and Biomod2



c. Germain-Aubrey



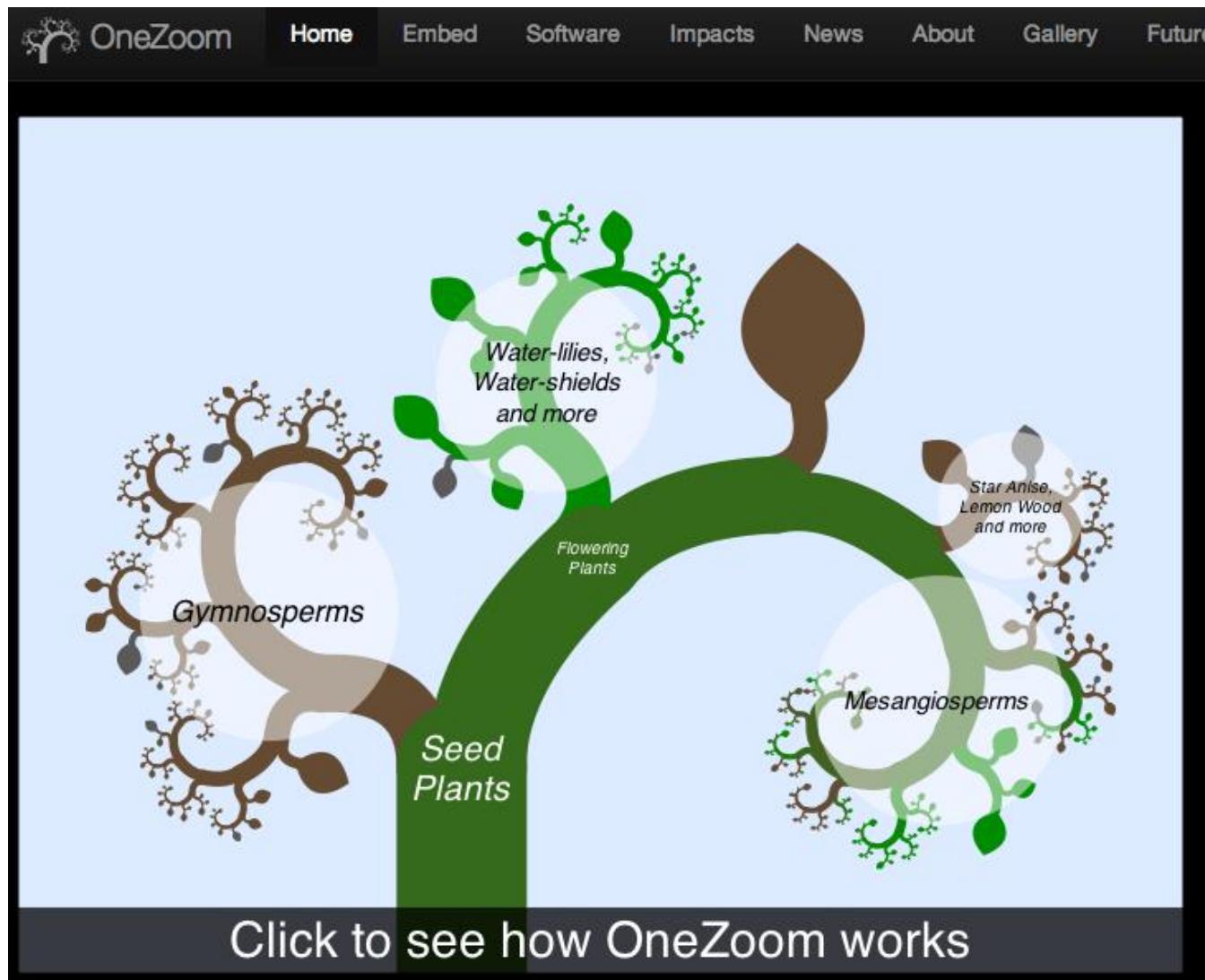
# Distributions of Polyploids and their Diploid Progenitors



- Do polyploids have broader ‘ecological amplitude’ and/or ranges than diploid parents?
- ENM for polyploids and parents
- Predicted distributions under models of climate change
  - Will polyploids and diploids respond in similar ways?

Blaine Marchant

# Linking to Phylogeny



The image shows a phylogenetic tree visualization from OneZoom, illustrating the evolutionary relationships of plants. The tree is rooted at the bottom and branches upwards. Major clades are highlighted in different colors: Gymnosperms (brown), Seed Plants (green), and Flowering Plants (light blue). Within the Seed Plants clade, three groups are labeled: Mesangiosperms (green), Water-lilies, Water-shields and more (light green), and Star Anise, Lemon Wood and more (brown). The tree is composed of numerous small, stylized plant icons representing individual species or genera. A large, dark brown leaf-like shape is positioned in the upper right quadrant of the tree. At the bottom of the visualization, there is a dark banner with the text "Click to see how OneZoom works".

OneZoom

Home Embed Software Impacts News About Gallery Future

Gymnosperms

Flowering Plants

Seed Plants

Water-lilies, Water-shields and more

Star Anise, Lemon Wood and more

Mesangiosperms

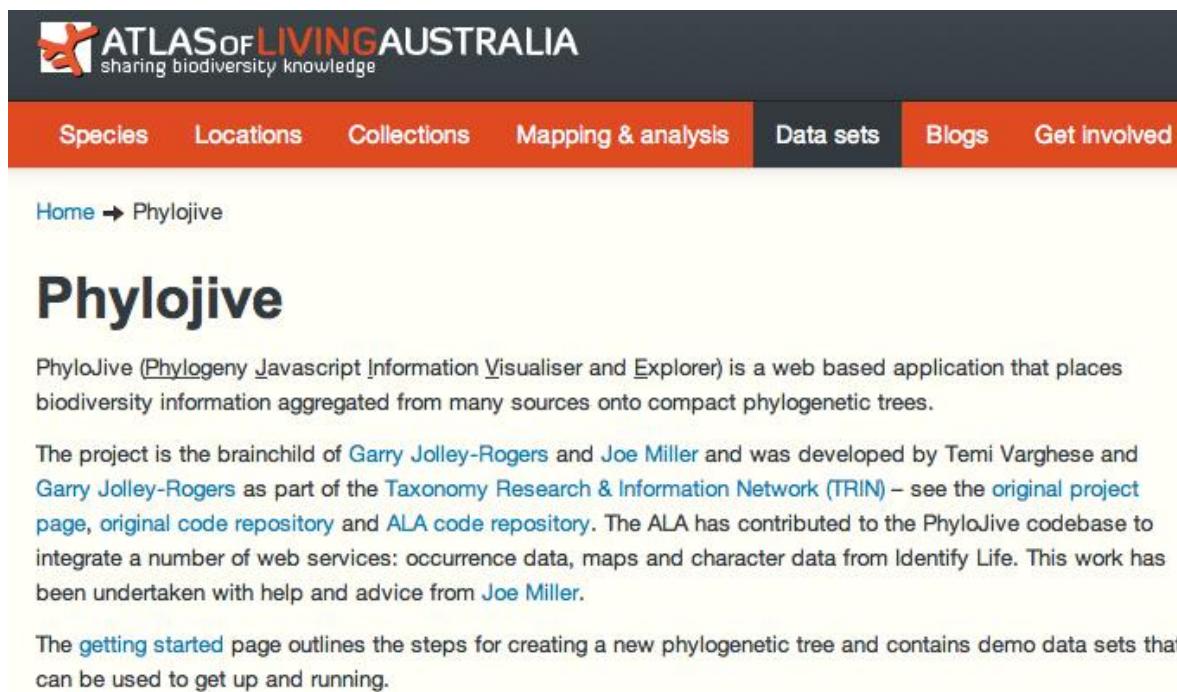
Click to see how OneZoom works

# PhyloJIVE

Links biodiversity data to trees

Joe Miller & Garry Jolley-Rogers

[phylojive.ala.org.au/](http://phylojive.ala.org.au/)



The screenshot shows the ATLAS of LIVING AUSTRALIA homepage with a navigation bar at the top. The navigation bar includes links for Species, Locations, Collections, Mapping & analysis, Data sets, Blogs, and Get involved. Below the navigation bar, a breadcrumb trail indicates the user is at Home → Phylojive. The main content area features a large heading "Phylojive". Below the heading, a paragraph describes PhyloJive as a web-based application that places biodiversity information onto compact phylogenetic trees. Another paragraph details the project's history, mentioning Garry Jolley-Rogers and Joe Miller, and their work with the Taxonomy Research & Information Network (TRIN). It also notes contributions from the ALA and Identify Life. A link to the "getting started" page is provided.



*A. buxifolia*

Source: Australian Plant Image Index Image by: Macd

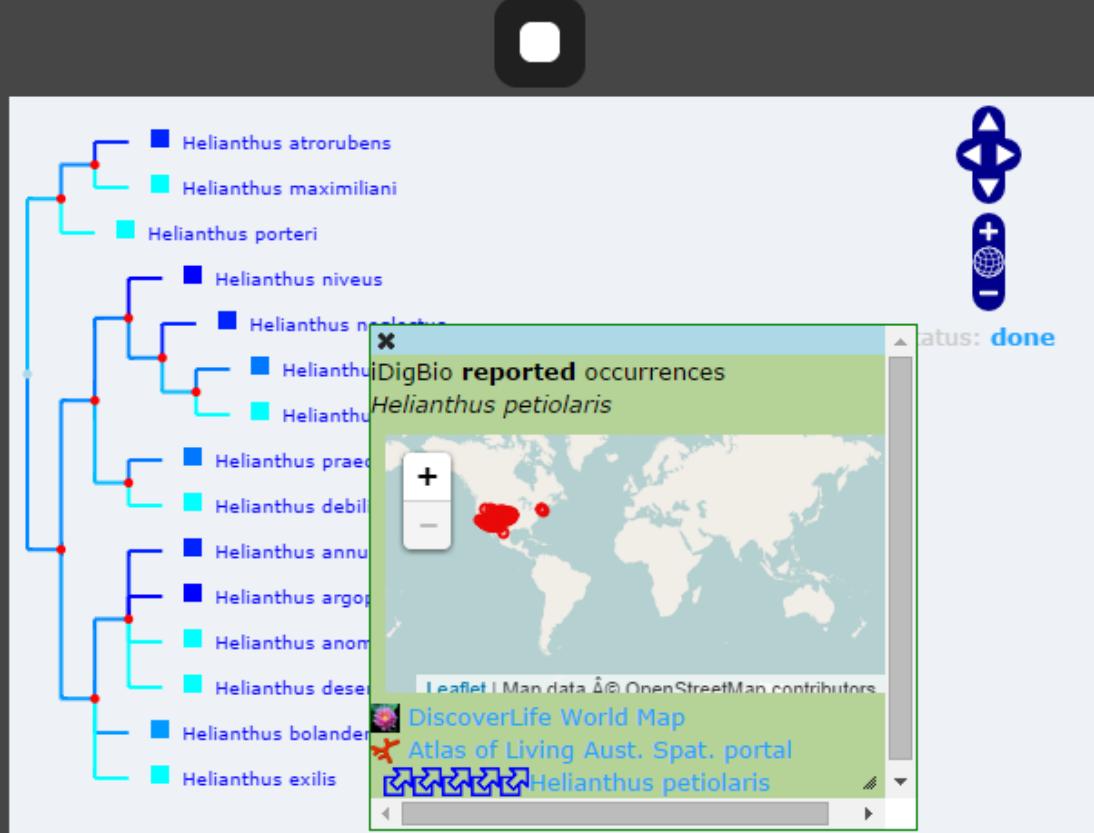
# PhyloJIVE instance in iDigBio

[iDigBio Portal](#)   [PhyloJIVE Home](#)   [OpenTree](#) ▾   [Sample Trees](#)   [Tutorial](#)   [Research Tools](#)

**Existing Tree: Helianthus**  
 Helianthus tree by Joe Miller  
 Select another tree:  
 Helianthus

- Click the top button to get the navigation aid
- Click nodes to get maps and external services
- Try choosing characters (if available) to plot on the tree;
- Align-names feature; search; set-root; rotate, etc.

[Create New Tree](#)



The image shows a phylogenetic tree for the genus *Helianthus*. The tree is rooted at the bottom and branches upwards. Nodes are marked with red dots. Blue squares represent *Helianthus atrorubens*, and cyan squares represent *Helianthus maximiliani*, *Helianthus porteri*, *Helianthus niveus*, *Helianthus petiolaris*, *Helianthus praecox*, *Helianthus debilis*, *Helianthus annuus*, *Helianthus argophyllus*, *Helianthus anomalous*, *Helianthus deserticola*, *Helianthus bolanderi*, and *Helianthus exilis*. A callout window for *Helianthus petiolaris* displays "iDigBio reported occurrences" with a map of the world showing a red dot in North America, and links to "DiscoverLife World Map", "Atlas of Living Aust. Spat. portal", and "Helianthus petiolaris".

- Developed by Garry Jolley-Rogers, Joe Miller, and Temi Varghese
- Integrates biodiversity data with phylogeny
- <http://phylojive.acis.ufl.edu/>

A. Matsunaga

# PhyloJIVE instance in iDigBio



The screenshot shows the PhyloJIVE interface integrated into the iDigBio portal. At the top, there is a green navigation bar with links: iDigBio Portal, PhyloJIVE Home, OpenTree ▾, Sample Trees, Tutorial, and Research Tools. Below the navigation bar, on the left, there is a sidebar with the title "Existing Tree" and a link to "Helianthus tre Miller". A large green leaf icon is displayed. On the right, there is a vertical toolbar with icons for zooming in and out, and a status message "IS: done". In the center, the "OPEN Tree of Life" logo is prominently displayed, followed by the subtitle "A community-assembled ToL".

- Developed by Garry Jolley-Rogers, Joe Miller, and Temi Varghese
- Integrates biodiversity data with phylogeny
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A. Matsunaga

# PhyloJIVE instance in iDigBio



The screenshot shows the PhyloJIVE interface integrated into the iDigBio portal. At the top, there's a navigation bar with links: iDigBio Portal, PhyloJIVE Home, OpenTree ▾, Sample Trees, Tutorial, and Research Tools. Below the navigation, a sidebar on the left displays an "Existing Tree" for "Helianthus tre Miller". It includes a "Select and..." button with "Helianthus" highlighted, a list of instructions for interacting with the tree, and a "Create New T" button. The main area features a large leaf graphic and the "OPEN Tree of Life" logo with the subtitle "A community-assembled ToL". A red box highlights the text "1.8 million described species" located below the logo.

Existing Tree  
Helianthus tre Miller

Select and  
Helianthus

- Click the top of the navigation
- Click nodes and external sources
- Try choosing (if available) a tree;
- Align-names search; set root etc.

Create New T

1.8 million described species

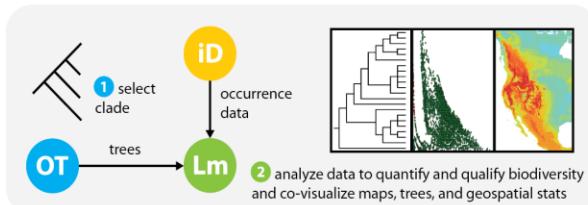
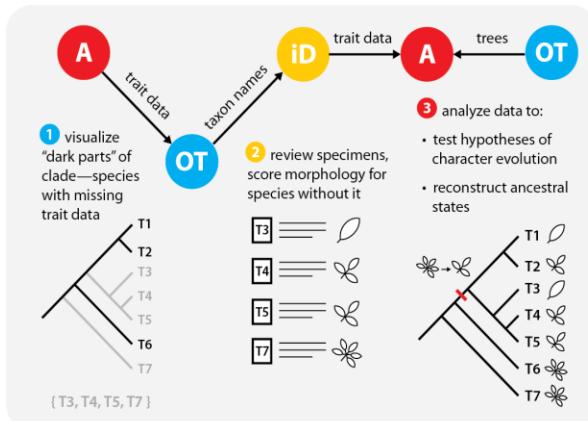
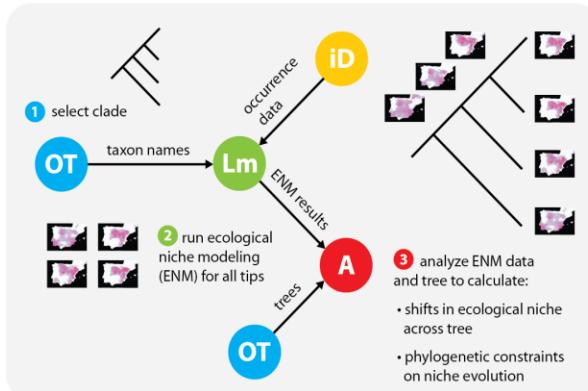
OPEN  
Tree of Life  
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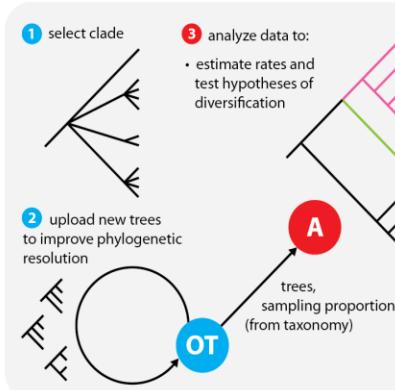
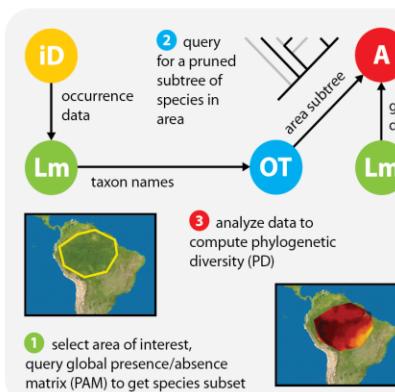
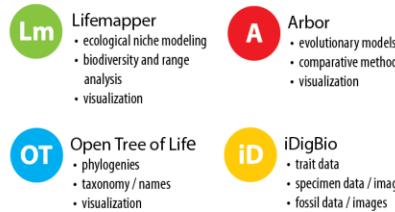
A. Matsunaga

# Connecting Trees, Specimens, Tools

## EXAMPLE WORKFLOWS:



## RESOURCES:

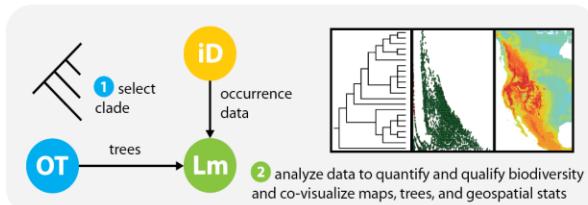
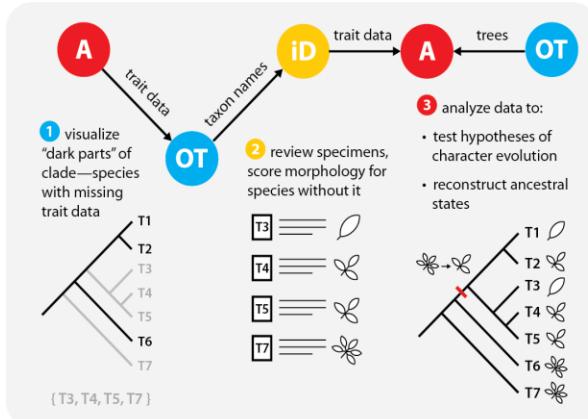
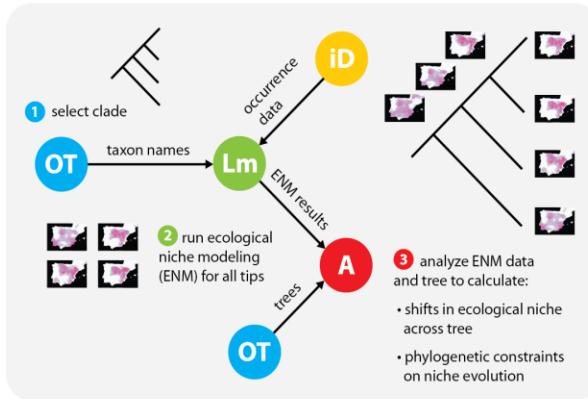


# Connecting Trees, Specimens, Tools

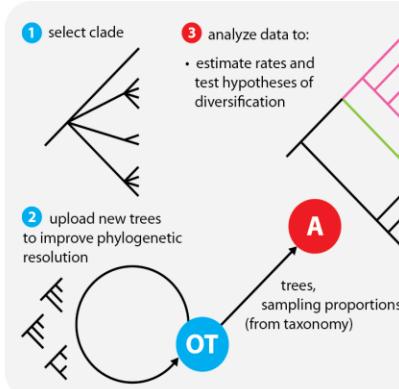
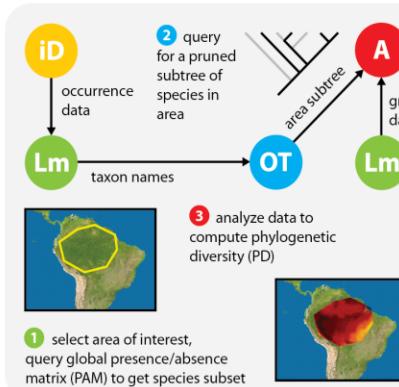
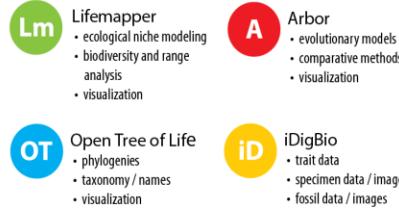


# Connecting Trees, Specimens, Tools

## EXAMPLE WORKFLOWS:

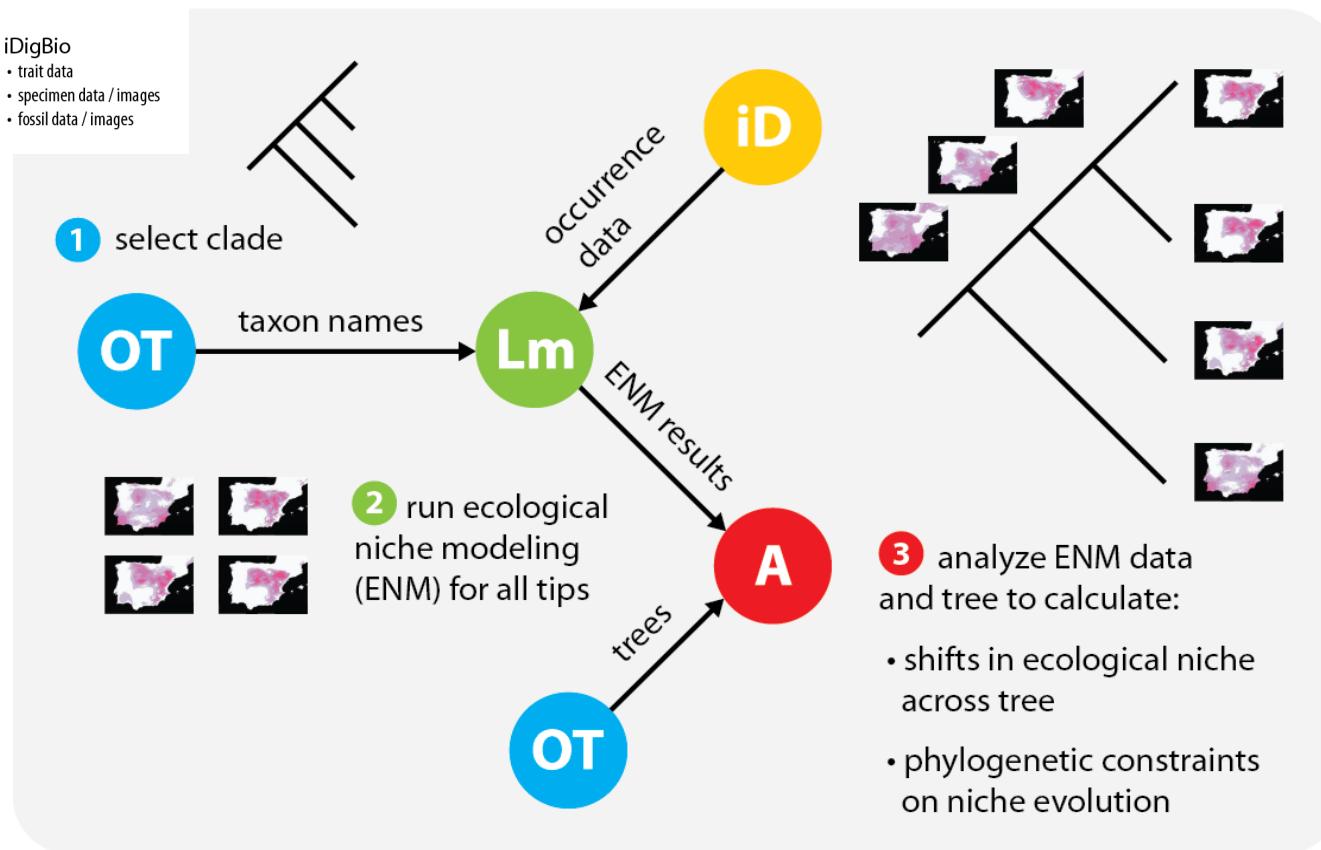
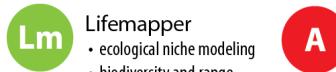


## RESOURCES:



# Connecting Trees, Specimens, Tools

## RESOURCES:



# Connecting Trees, Specimens, Tools



ABI Innovation: Connecting resources  
to enable large-scale biodiversity analyses

D. Soltis, P. Soltis, J. Fortes, A. Matsunaga,  
J. Beach, J. Soberon, S. Smith

## RESOURCES:



Lifemapper  
• ecological niche modeling  
• biodiversity and range  
analysis  
• visualization



Arbor  
• evolutionary models  
• comparative methods  
• visualization



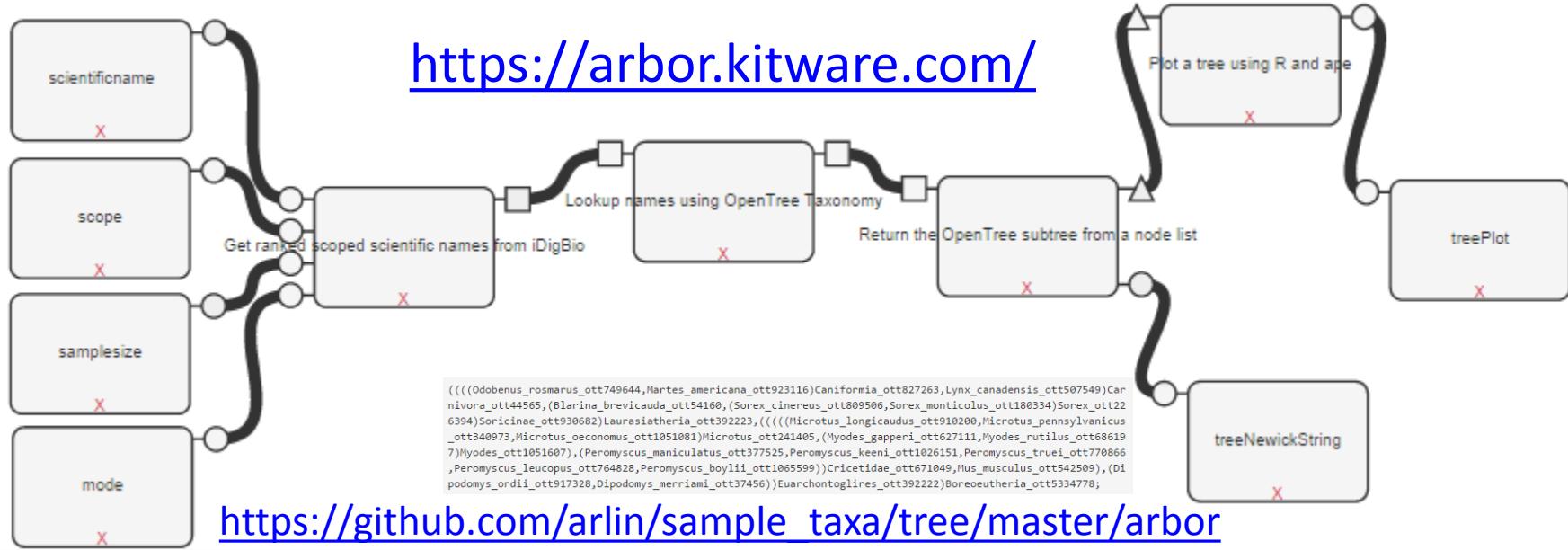
Open Tree of Life  
• phylogenies  
• taxonomy / names  
• visualization



iDigBio  
• trait data  
• specimen data / images  
• fossil data / images

# Arbor, OpenTree, and iDigBio

<https://arbor.kitware.com/>



Workflow to get an induced tree from a configurable iDigBio query >

scientificname  
Mammalia

scope  
\_all

samplesize  
20

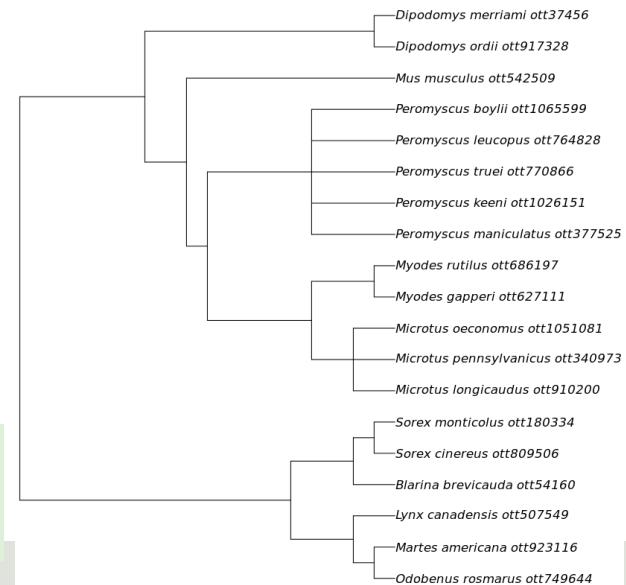
mode  
top

 Run  Close

```
1 {  
2     "query": {  
3         "query_string" : {  
4             "default_field" : "order",  
5             "query" : "rodentia"  
6         }  
7     },  
8     "aggregations": {  
9         "my_agg": {  
10            "terms": {  
11                "field": "scientificname",  
12                "size": 100  
13            }  
14        }  
15    }  
16 }
```

Success! Produced the following outputs:

- Workflow to get an induced tree from a configurable iDigBio query treeNewickString [string]
  - Workflow to get an induced tree from a configurable iDigBio query treePlot [image]



# Specimen Records AND Images

*Callisia graminea*  
grassleaf roseling



29.65, -82.32

number,dwc:preparations,dwc:identificationVerificationStatus,idigbio:subfamily,idigbio:preparationCount,fcc:pickedBy,dwc:eventRemarks,dwc:VerbatimEventDate,dwc:associatedReferences,idigbio:endangeredStatus,dwc:locationAccordingTo,dwc:georeferenceSources,dwc:associatedSequences,dwc:formation,dwc:higherClassification,dwc:catalogNumber,dwc:verbatimSRS,dwc:higherGeography,dwc:individualCount,dwc:decimalLongitude,dwc:datasetName,dwc:month,dwc:georeferencedBy,dwc:eventTime,dwc:identificationQualifier,idigbio:

# Systematics and Taxonomy



*Linnea* (twinflower)

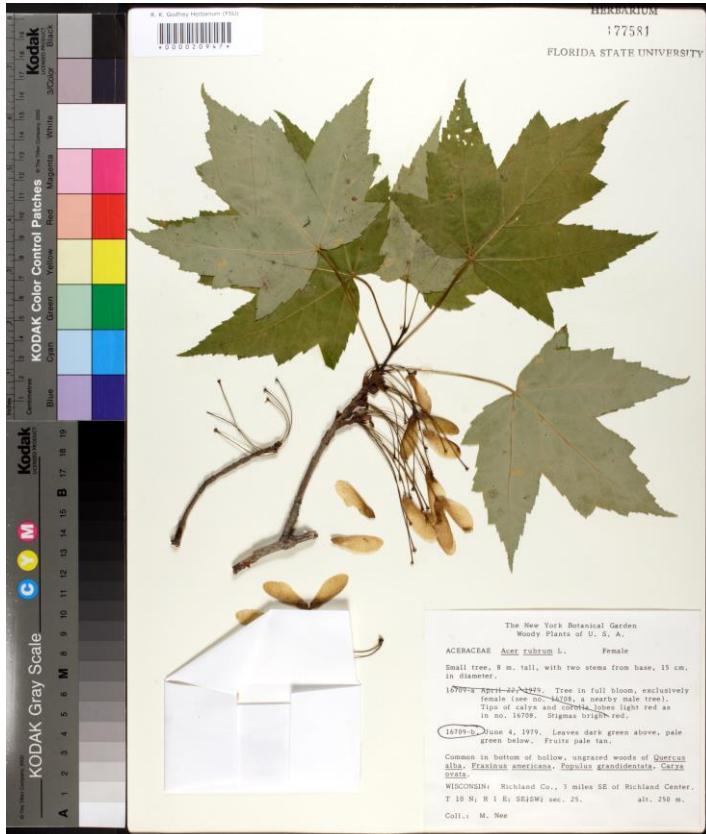
Carl Linné, aka Carolus Linnaeus

TRY

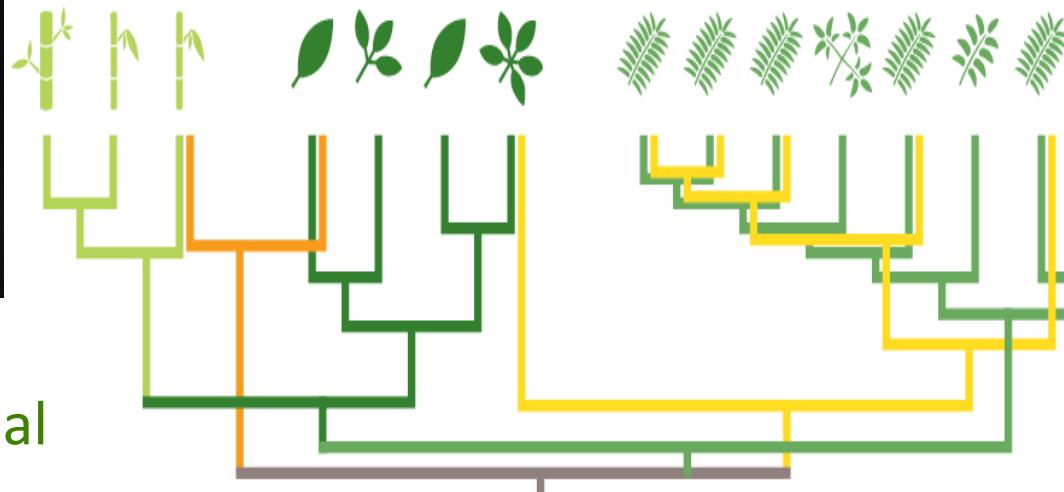
# Plant Trait Database

PhotosyntheticPathway  
Respiration LeafArea NfixationCapacity  
SLA RegenerationCapacity PlantLifespan  
WoodDensity GrowthForm  
PhenologyType LeafN  
LeafP LeafLongevity PhotosyntheticCapacity  
MaxPlantHeight SeedMass

# Using Images to Infer Functional Traits

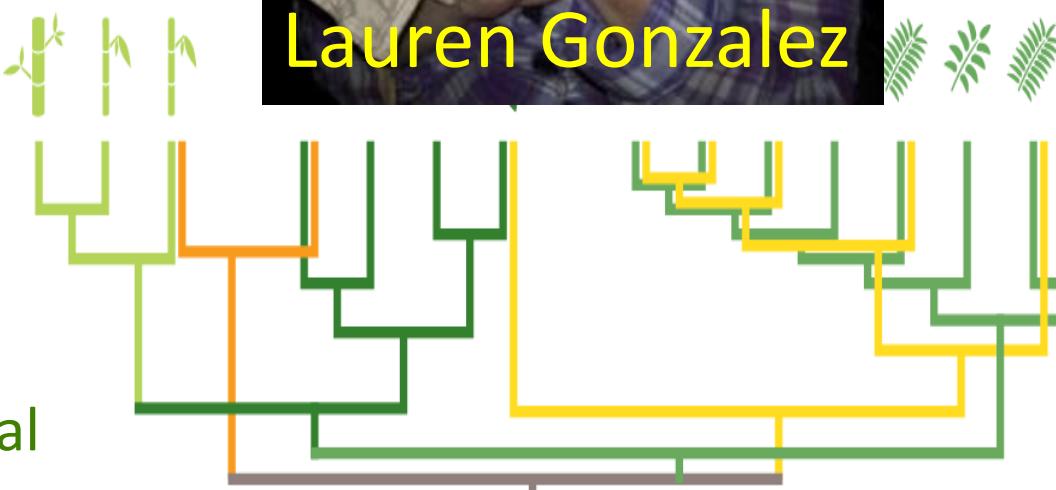


PhotosyntheticPathway  
Respiration LeafArea NfixationCapacity  
SLA RegenerationCapacity PlantLifespan  
WoodDensity GrowthForm  
PhenologyType LeafN  
LeafP LeafLongevity PhotosyntheticCapacity  
MaxPlantHeight SeedMass



Connect to ecology  
Evolution of plant functional  
traits

# Using Images to Infer Functional Traits



Connect to ecology  
Evolution of plant functional  
traits

# Many Research Uses for Specimen Data

- Connections to other resources, e.g. GenBank
- Ecological Niche Modeling
- Integration with phylogeny, e.g. PhyloJIVE
- Complex integration of phylogeny, specimens, ENM, other heterogeneous data
- Images as sources of traits for ecological studies
- Others???
- iDigBio Research Applications Working Group

# Thank you!

**iDigBio colleagues:** J. Fortes, A. Matsunaga  
J. Miller, D. Soltis, S. Smith, J. Beach, J. Soberon, L. Harmon  
**C. Germain-Aubrey, B. Marchant, L. Gonzalez**



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[facebook.com/idigbio](https://facebook.com/idigbio)



[twitter.com/idigbio](https://twitter.com/idigbio)



[vimeo.com/idigbio](https://vimeo.com/idigbio)

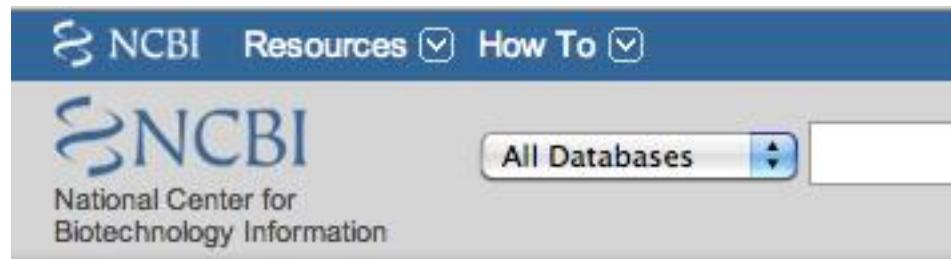


[idigbio.org/rss-feed.xml](https://idigbio.org/rss-feed.xml)



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

# Linking to GenBank



## Welcome to NCBI

The National Center for Biotechnology Information advances science and health by providing access to biomedical and genomic information.

[About the NCBI](#) | [Mission](#) | [Organization](#) | [Research](#) | [NCBI News](#)

- **Examples of the /specimen\_voucher information:**

```
/specimen_voucher="UAM:Mamm:52179"  
/specimen_voucher="AMCC:101706"  
/specimen_voucher="USNM:field series 8798"  
/specimen_voucher="personal:Dan Janzen:99-SRNP-2003"  
/specimen_voucher="99-SRNP-2003"
```



# Connecting ecology to specimens

## Correlation of plant functional traits

