

Using Digitized Data in Scientific Research *using and reusing biodiversity data*

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Collections: The Library of Life

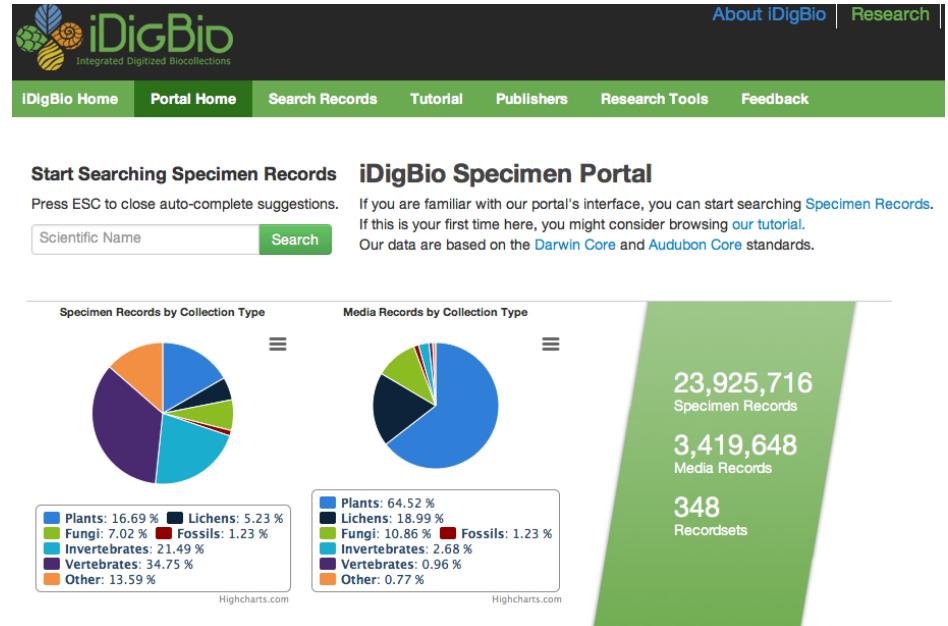


>1600 natural history collections
in the US alone
1-2 billion specimens
in the US
3-4 billion specimens
worldwide

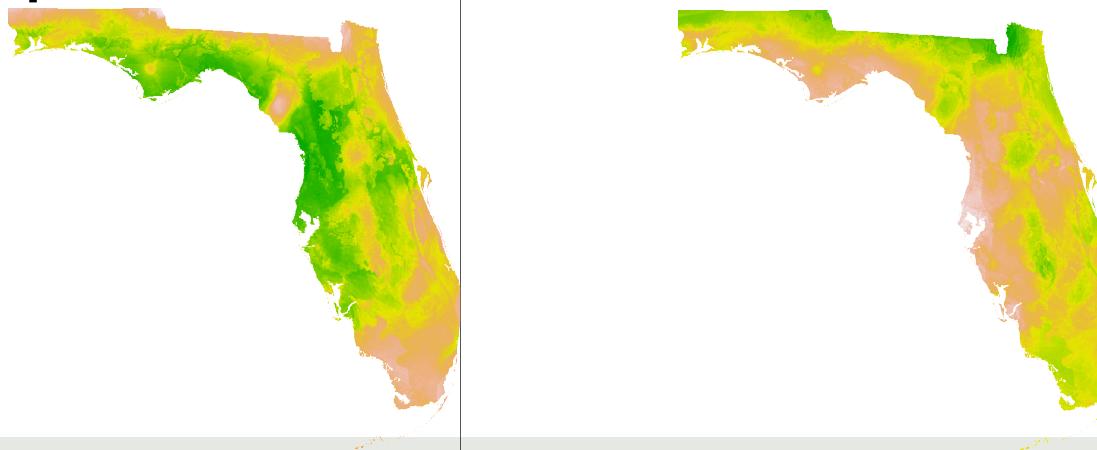


Research @ iDigBio

- Data portal

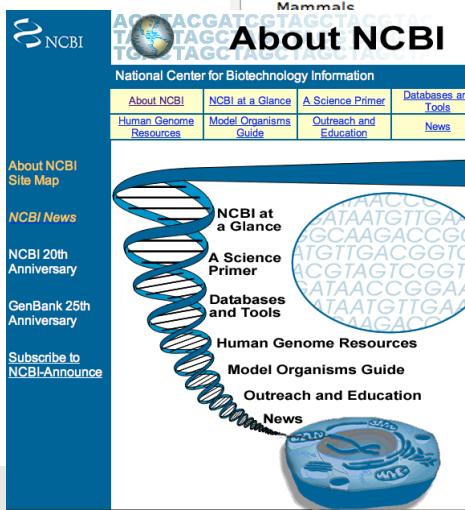
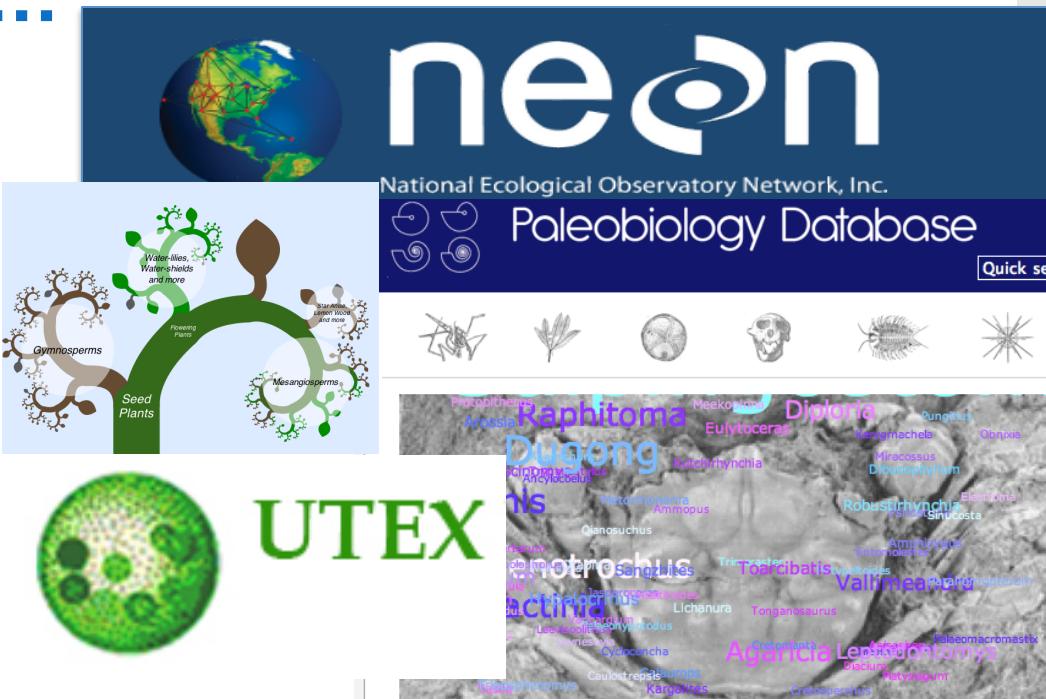


- Computational workflows and environment

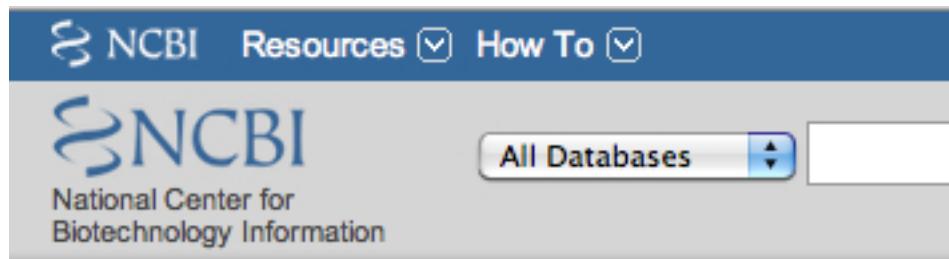


Linking Collections to...

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



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"  
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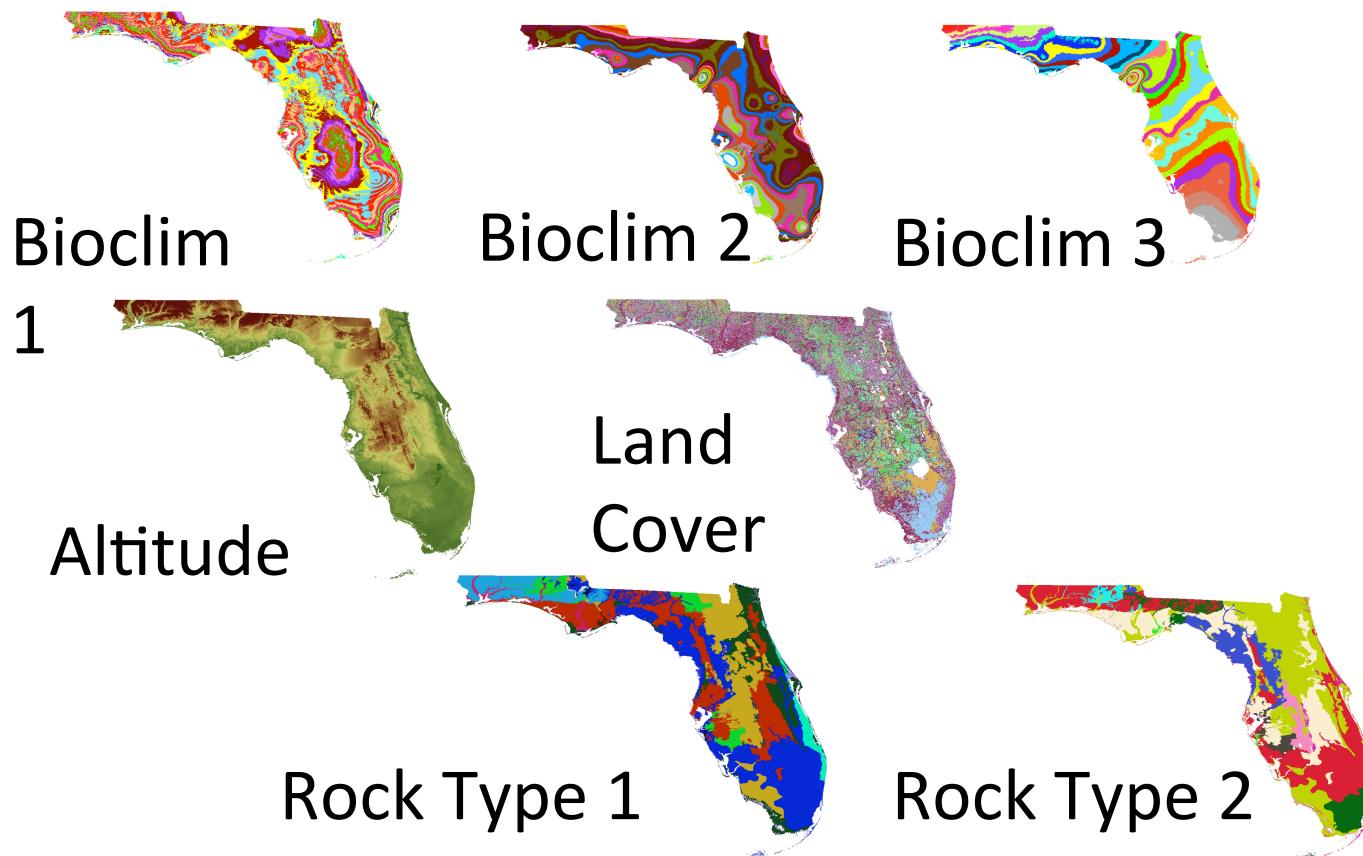
Ecological Niche Modeling: locality information



29.65, -82.32

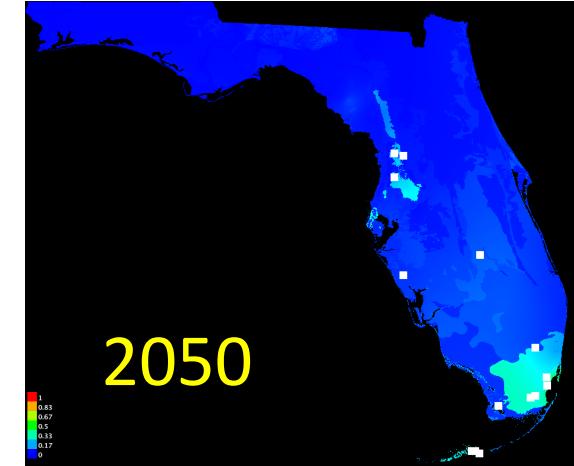
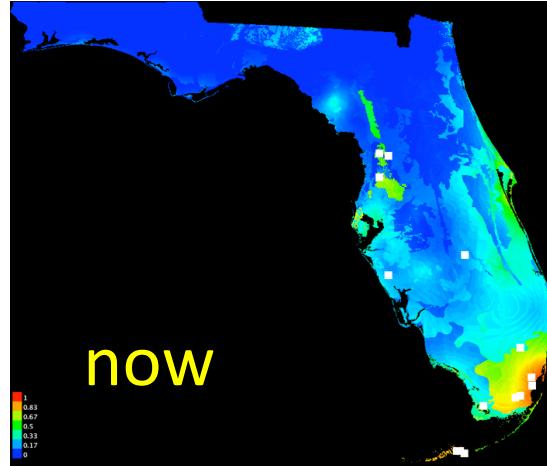
number,dwc:preparations,dwc:identificationVerificationStatus,idigbio:subfamily,idigbio:preparationCount,fcc:pickedBy,dwc:eventRemarks,dwc:VerbatimEventDate,dwc:associatedReferences,idigbio:endangeredStatus,dwc:locationAccordingTo,dwc:georeferenceSources,dwc:associatedSequences,dwc:formation,dwc:higherClassification,dwc:catalogNumber,dwc:verbatimSRS,dwc:higherGeo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

Ecological Niche Modeling: Environmental Layers

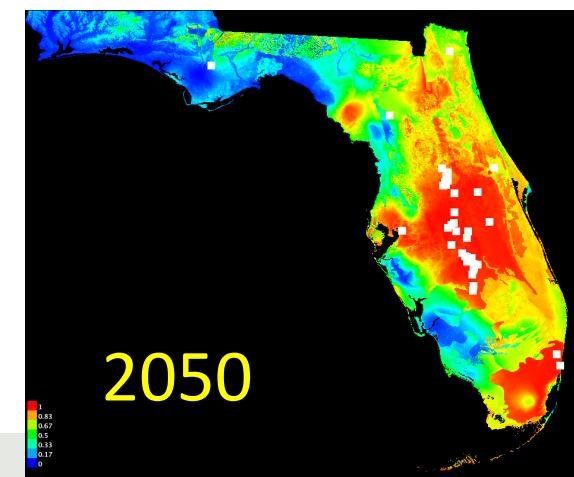
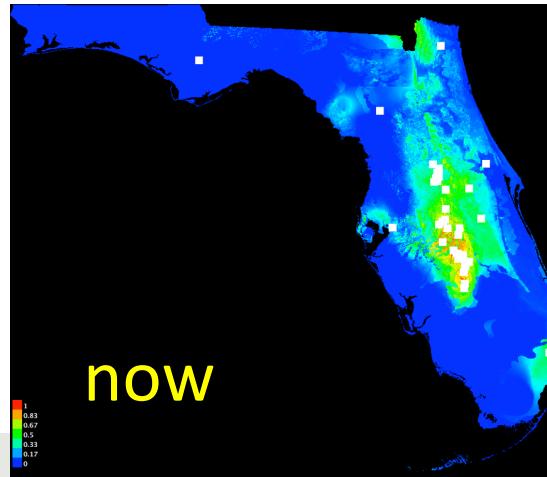


Responses to Climate Change: Winners & Losers

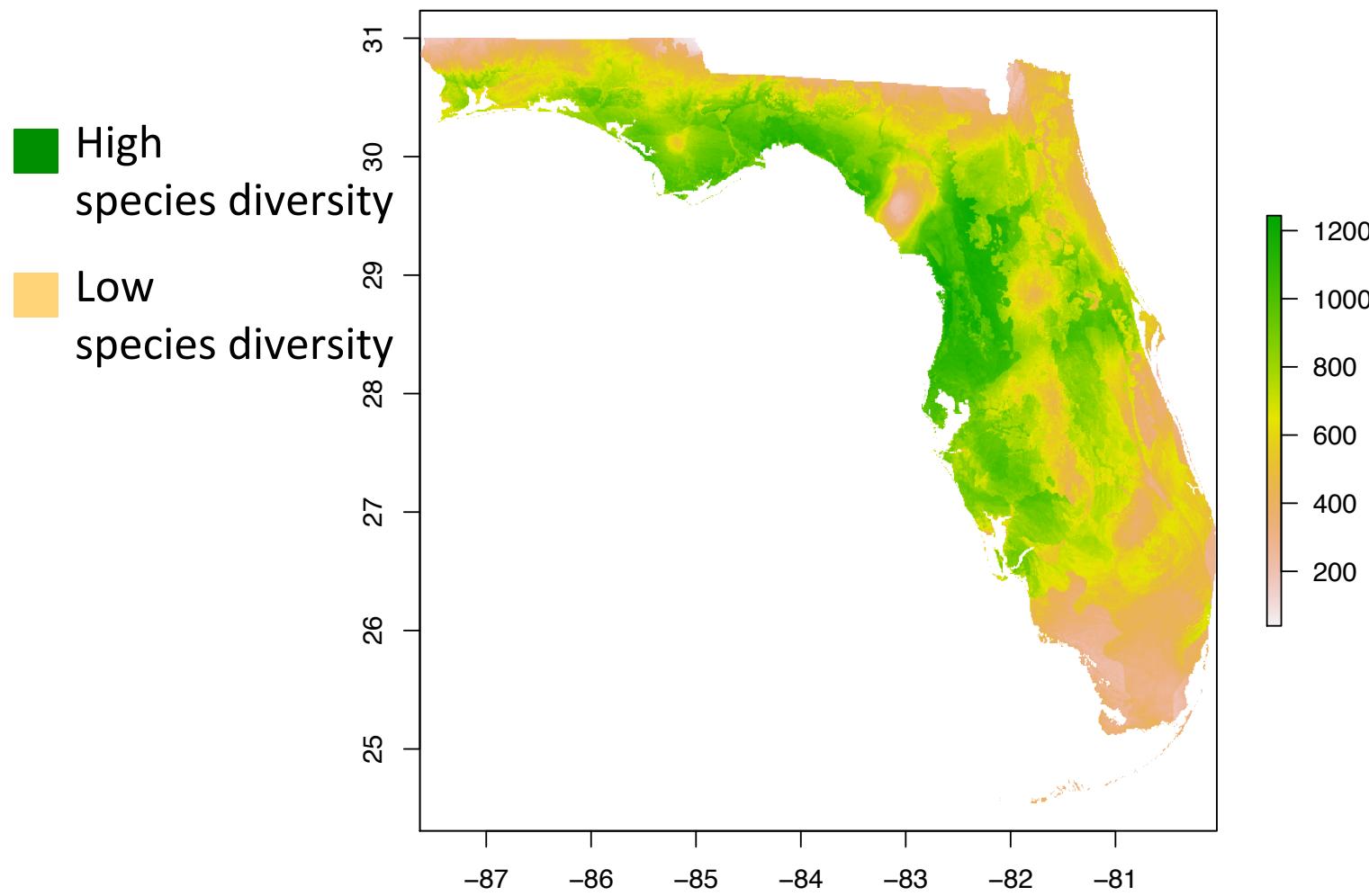
Abildgaardia ovata (flatspike sedge)



Prunus geniculata (scrub plum)



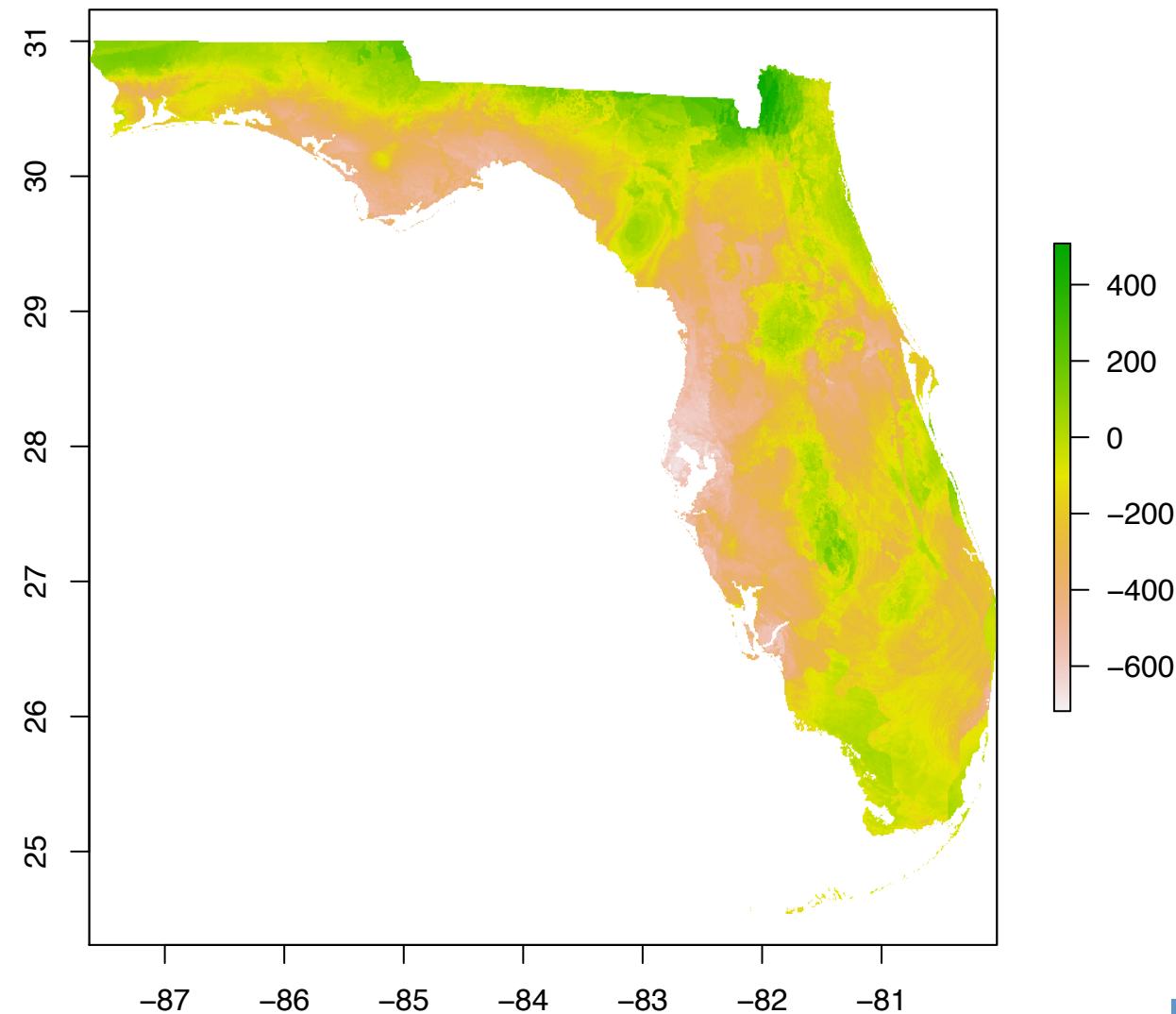
Florida plant diversity heat map: now



Between now and 2050...

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

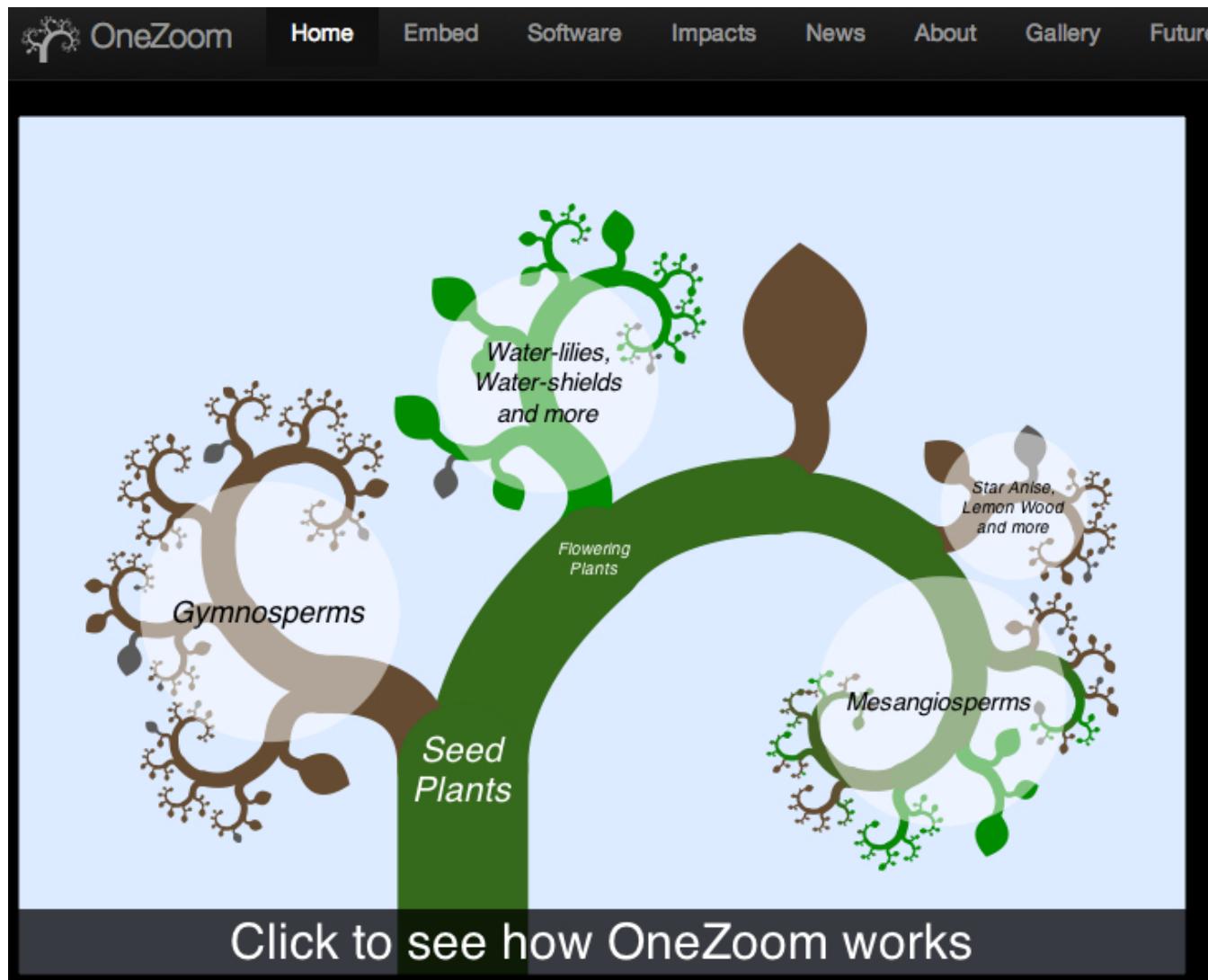
spp 2050 - #spp now



Linking to Phylogeny

OneZoom

Home Embed Software Impacts News About Gallery Future



Water-lilies,
Water-shields
and more

Gymnosperms

Flowering Plants

Seed Plants

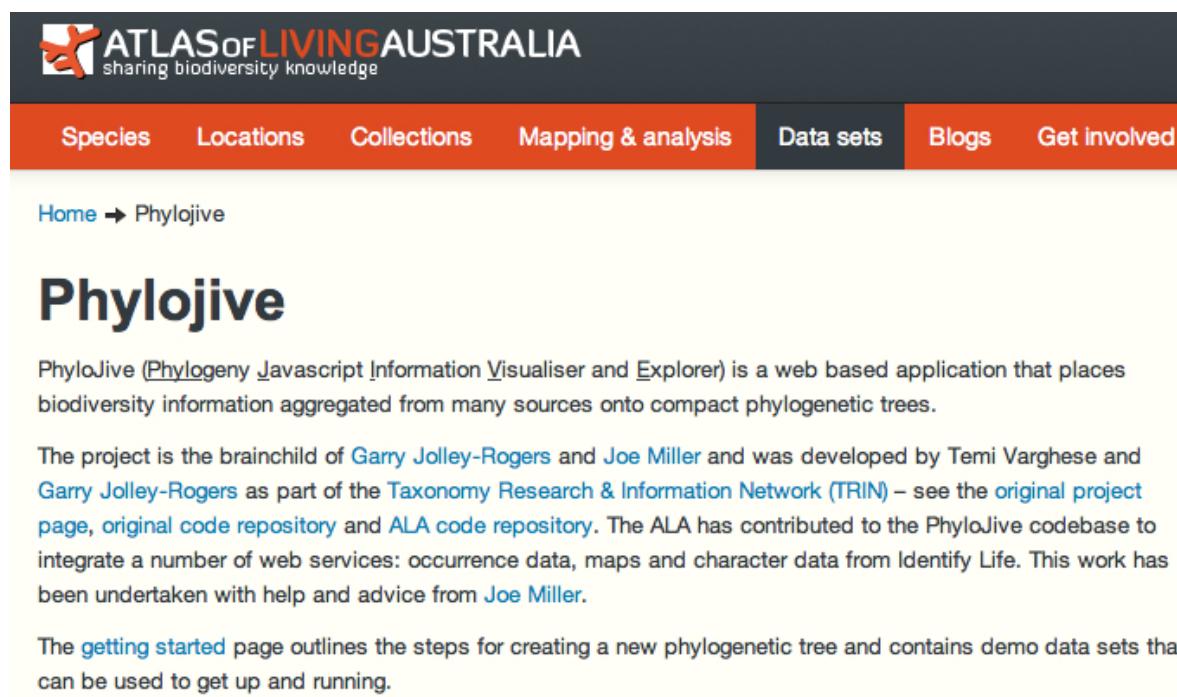
Mesangiosperms

Star Anise,
Lemon Wood
and more

Click to see how OneZoom works

PhyloJIVE

Links biodiversity data to trees
Joe Miller & Garry Jolley-Rogers
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 shows Home → Phylojive. The main content area features a large heading "Phylojive" and a descriptive paragraph about the project. It also contains a block of text detailing the project's history and development, mentioning Garry Jolley-Rogers, Joe Miller, Temi Varghese, and the Taxonomy Research & Information Network (TRIN). A link to the "getting started" page is provided at the bottom.



A. buxifolia
Source: Australian Plant
Image Index
Image by: Macdonald, C.

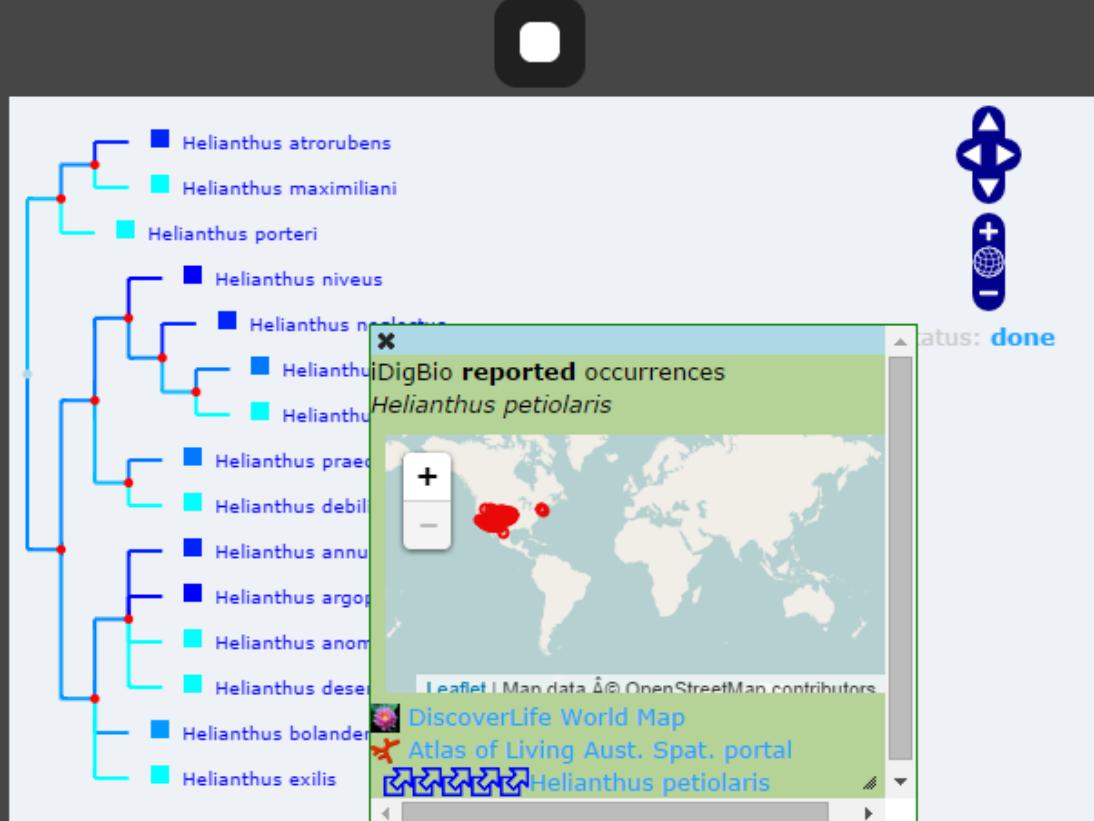
PhyloJIVE instance in iDigBio

[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](#)



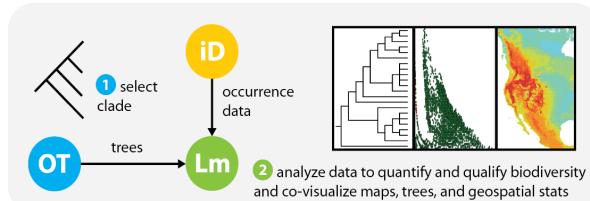
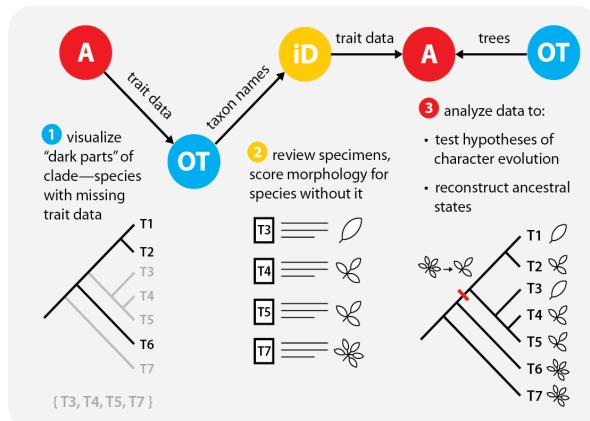
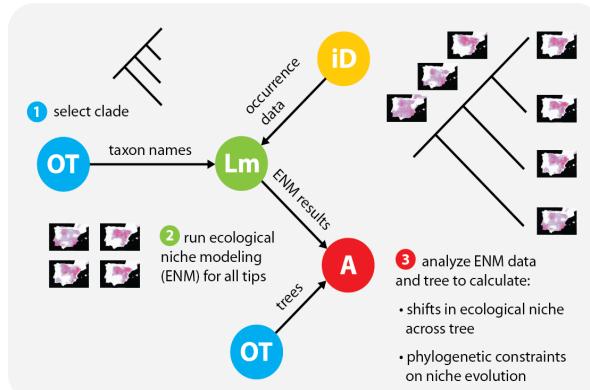
The image shows a phylogenetic tree for the genus *Helianthus*. The tree is rooted on the left and branches to the right. Nodes are marked with red dots. Colored squares next to node labels indicate specific species: blue for *Helianthus atrorubens*, cyan for *Helianthus maximiliani*, and light blue for *Helianthus porteri*. Other species listed include *Helianthus niveus*, *Helianthus petiolaris*, *Helianthus praecox*, *Helianthus debilis*, *Helianthus annuus*, *Helianthus argophyllus*, *Helianthus anomalous*, *Helianthus deserticola*, *Helianthus bolanderi*, and *Helianthus exilis*. A callout window titled "iDigBio reported occurrences" for *Helianthus petiolaris* displays a world map with a red dot over North America, zoom controls, and links to "DiscoverLife World Map", "Atlas of Living Aust. Spat. portal", and "Helianthus petiolaris". The status of the callout is "done".

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

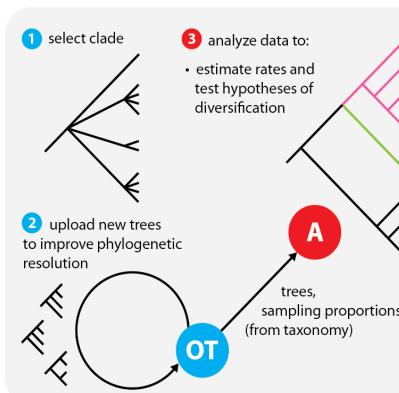
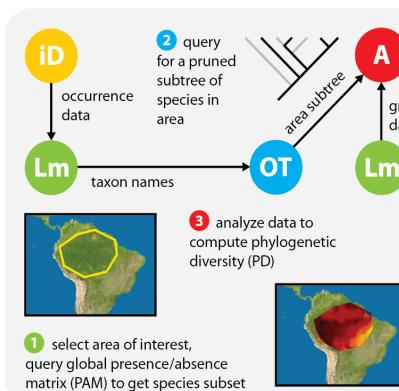
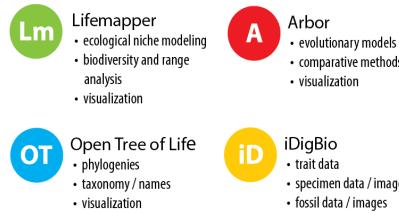
A. Matsunaga

Connecting Trees, Specimens, Tools

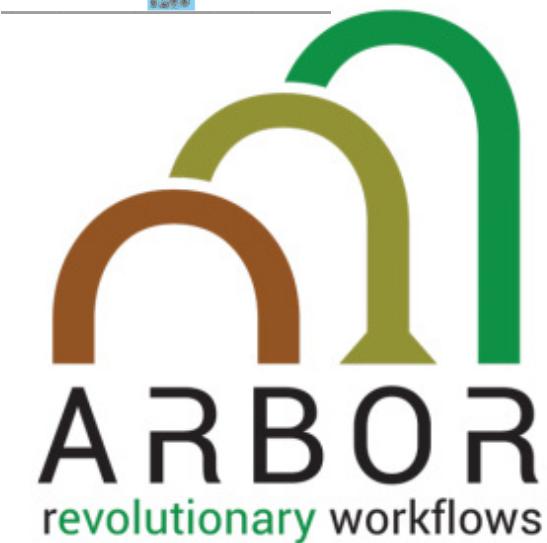
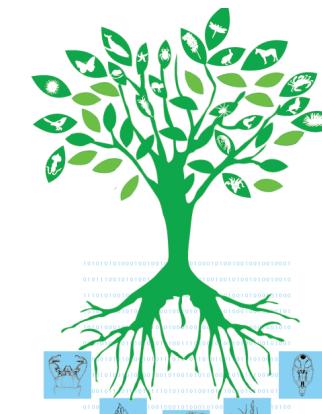
EXAMPLE WORKFLOWS:



RESOURCES:

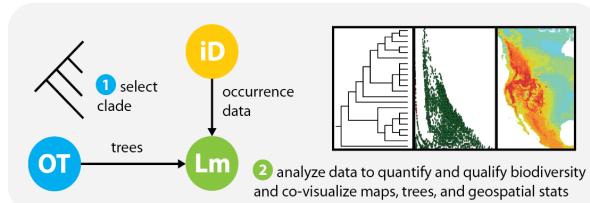
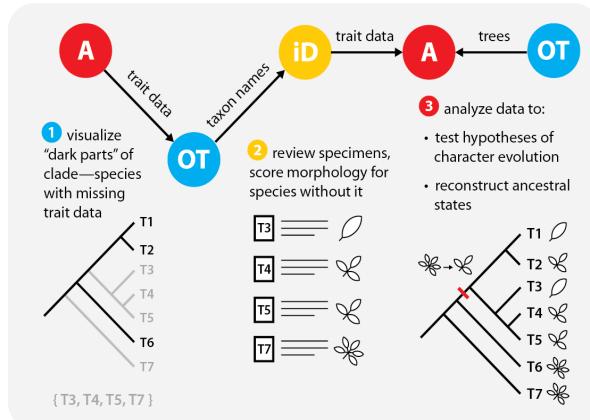
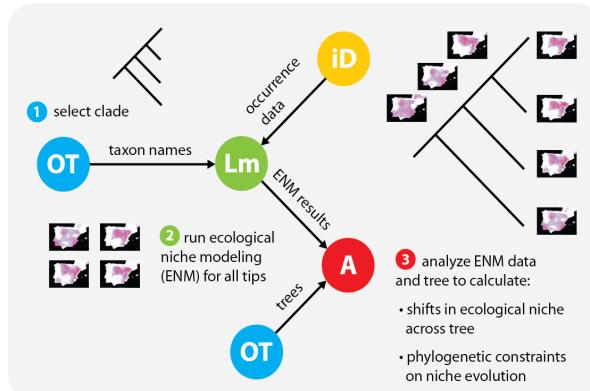


Connecting Trees, Specimens, Tools

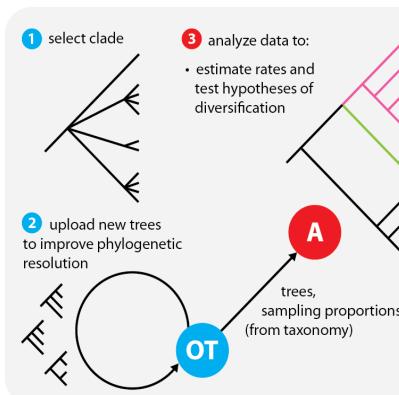
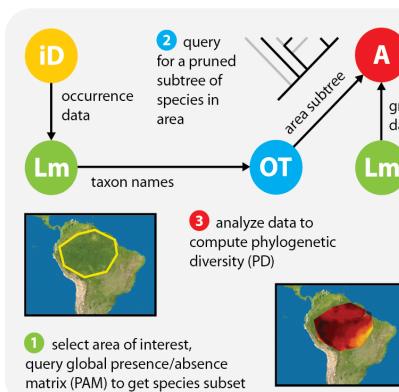
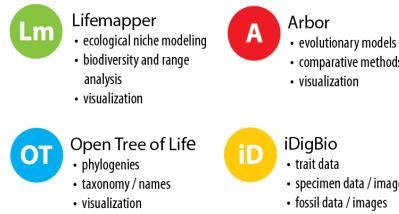


Connecting Trees, Specimens, Tools

EXAMPLE WORKFLOWS:

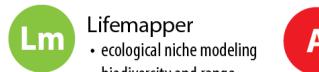


RESOURCES:



Connecting Trees, Specimens, Tools

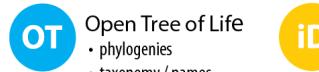
RESOURCES:



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



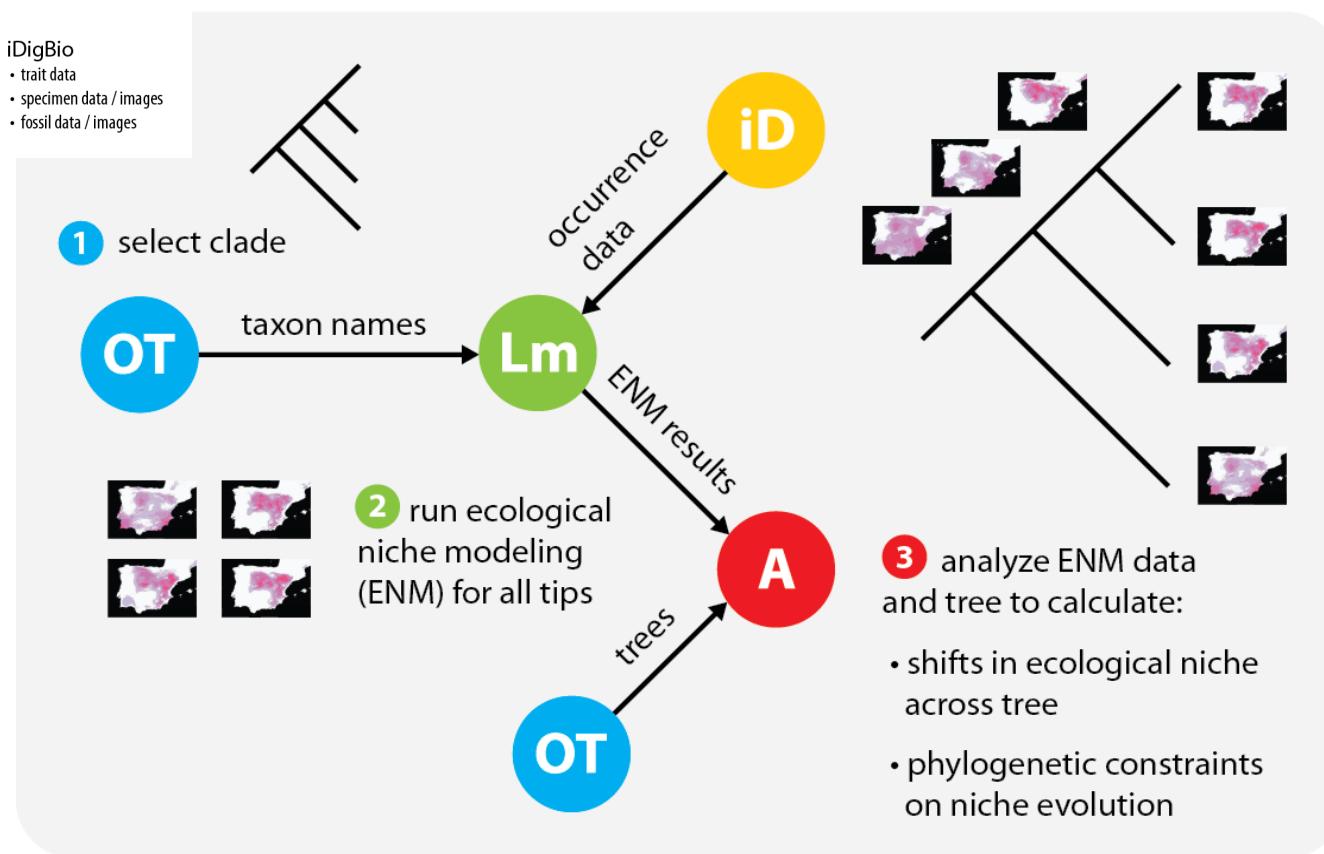
- Arbor
- evolutionary models
 - comparative methods
 - visualization



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

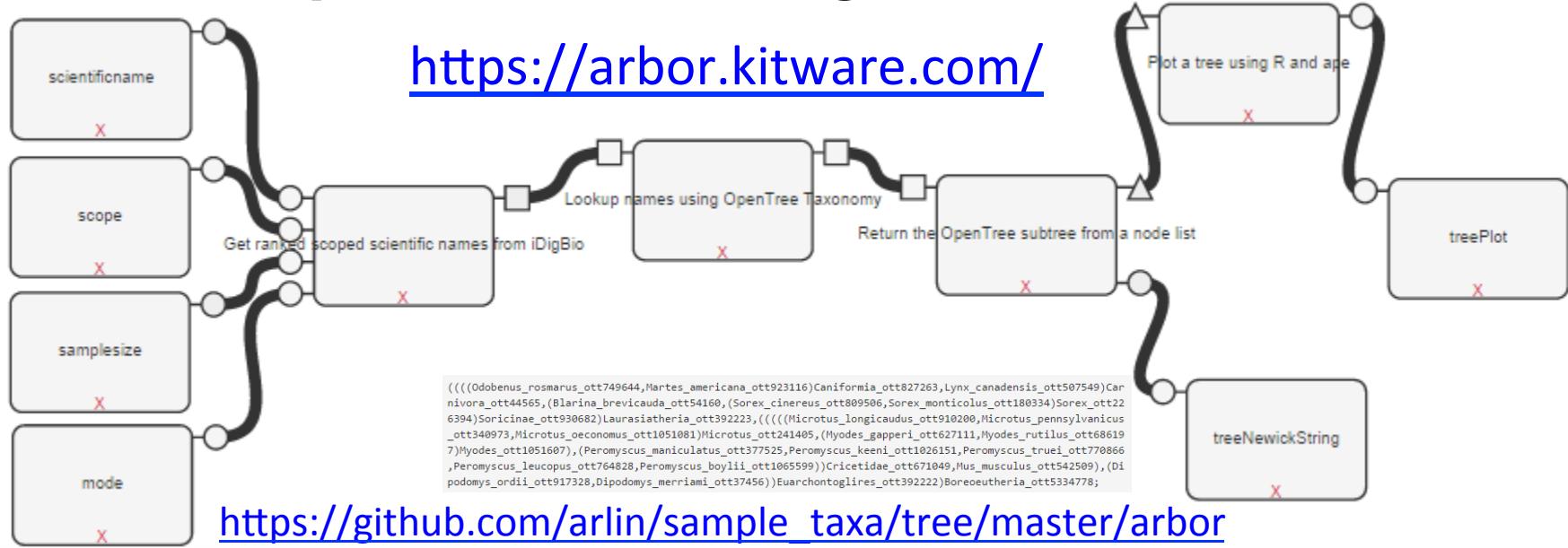


- iDigBio
- trait data
 - specimen data / images
 - fossil data / images



Arbor, OpenTree, and iDigBio

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



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

scientificname

Mammalia

scope

_all

samplesize

20

mode

top

▶ Run Close

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]

tree from a configurable iDigBio query treeNewickString

tree from a configurable iDigBio query treePlot [image]

Phylogenetic tree illustrating the relationships between various rodent species based on mitochondrial DNA sequence data. The tree is rooted on the left and branches to the right, with species names and their corresponding GenBank accession numbers listed at the tips.

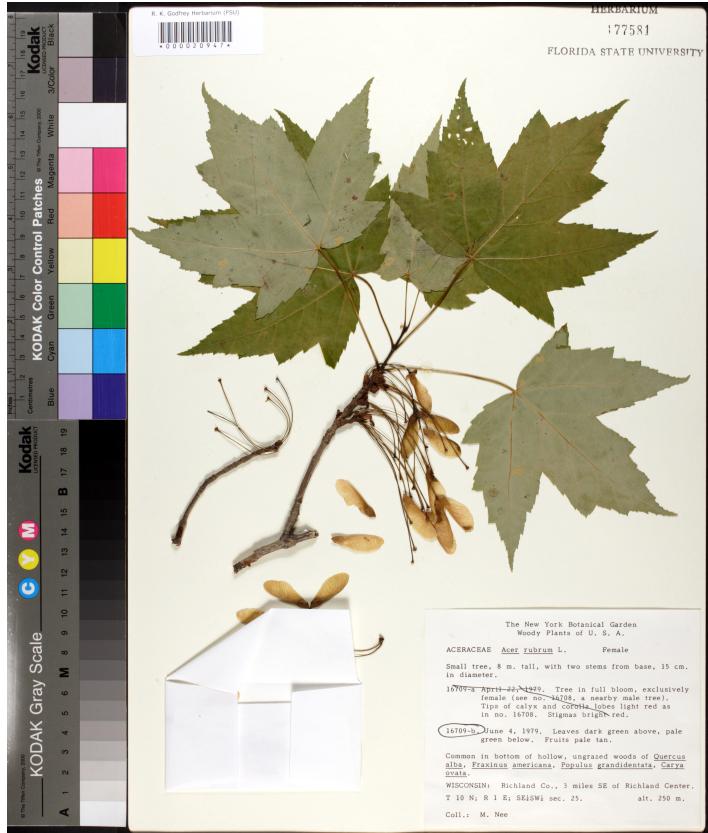
- Dipodomys merriami* ott37456
- Dipodomys ordii* ott917328
- Mus musculus* ott542509
- Peromyscus boylii* ott1065599
- Peromyscus leucopus* ott764828
- Peromyscus truei* ott770866
- Peromyscus keeni* ott1026151
- Peromyscus maniculatus* ott377525
- Myodes rutilus* ott686197
- Myodes gapperi* ott627111
- Microtus oeconomus* ott1051081
- Microtus pennsylvanicus* ott340973
- Microtus longicaudus* ott910200
- Sorex monticolus* ott180334
- Sorex cinereus* ott809506
- Blarina brevicauda* ott54160
- Lynx canadensis* ott507549
- Martes americana* ott923116
- Odobenus rosmarus* ott740644

TRY

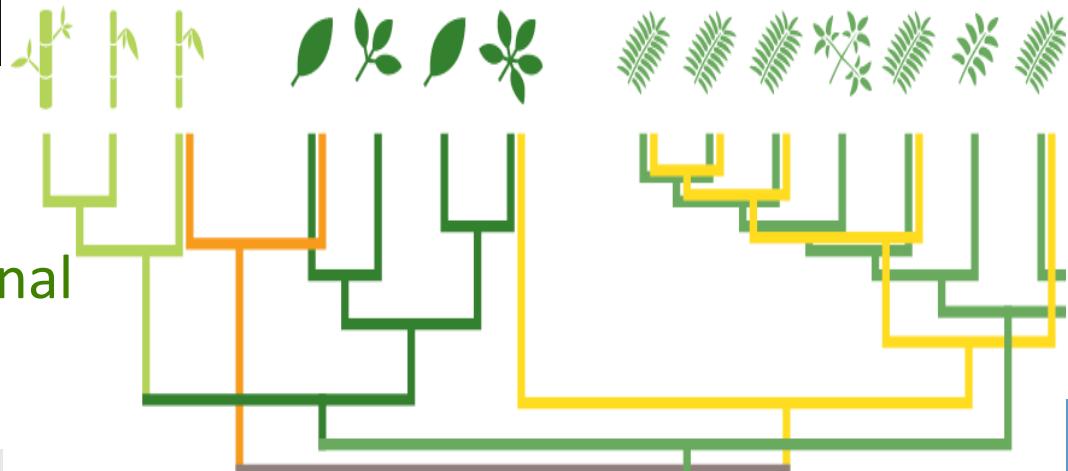
Plant Trait Database



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



Connecting ecology to specimens
Correlation of plant functional traits



Many 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???
- Research Applications Working Group

Thank you!



www.idigbio.org

psoltis@flmnh.ufl.edu



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