Lifemapper on SAGE2

Michael Elliott¹, James Beach², C. J. Grady², Aimee Stewart², José A. B. Fortes¹

¹ACIS Lab, University of Florida, USA ²University of Kansas, USA

PRAGMA37 San Diego September 11-14, 2019





Project Overview

- A collaboration between myself and researchers at the University of Kansas
- **The goal**: make Lifemapper accessible from SAGE2



Lifemapper

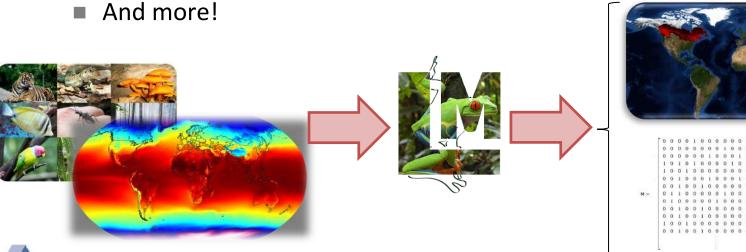






What is Lifemapper?

- Biodiversity modeling software
- Inputs:
 - Species occurrence data
 - **Environmental data**
- **Outputs:**
 - Potential habitats for species
 - Correlations between species occurrences







BiotaPhy

■ The *BiotaPhy* project:

Connect biodiversity projects through workflows with integrated data providers



Lifemapper

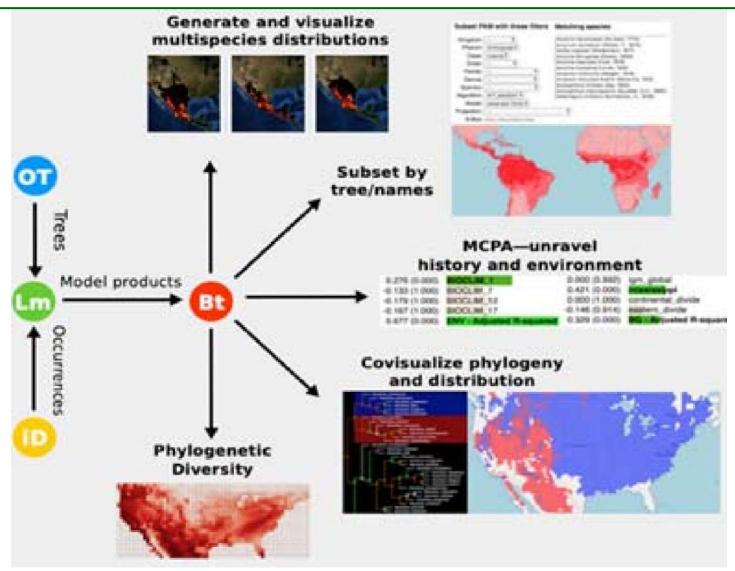








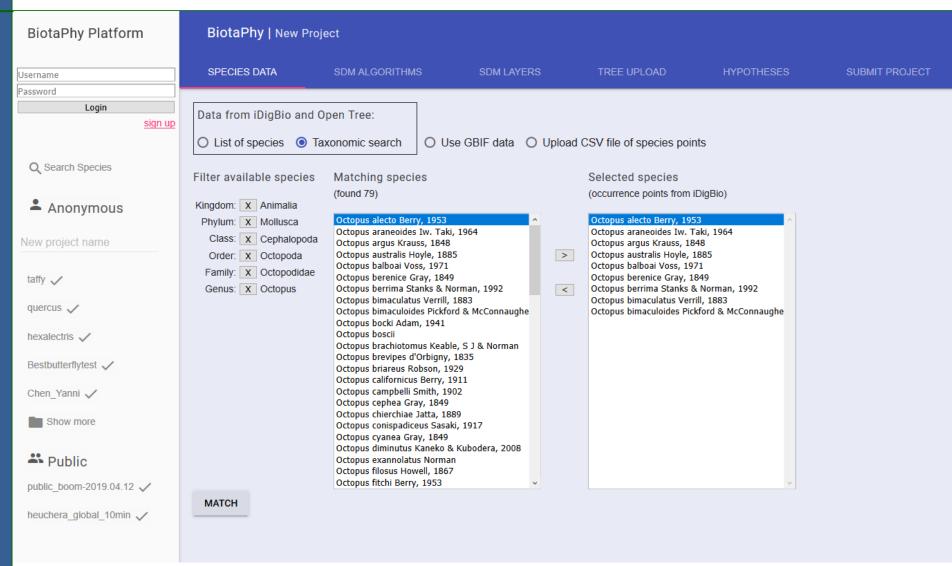
BiotaPhy







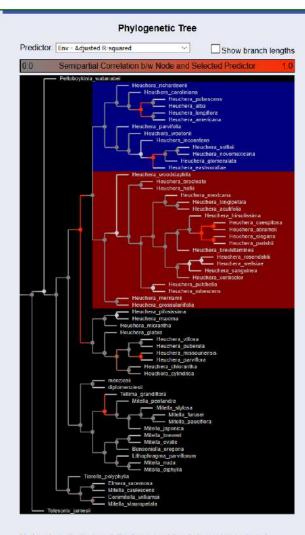
BiotaPhy's Online Interface

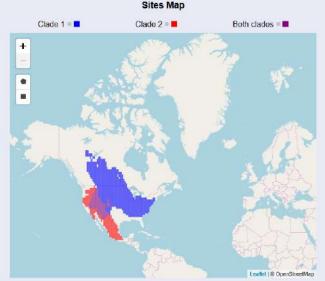




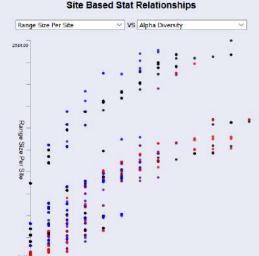


BiotaPhy's Offline Interface





The map shows sites where the selected species are present. Use the select by bounding box or by polygon tools to highlight in the tree which species are present at those selected sites.



The scatter plot shows relationships between site based statistics; each dot represents a single site. If sites in the map are highlighted, the corresponding sites are highlighted in the same color in the scatter plot. You can brush sites in the plot by clicking and dragging to create a selection box. Doing so will highlight the selected points as well as those sites on the map and the species present at those sites in the tree. Use the two drop down boxes to select the metrics to use for the X and Y axes.

Alpha Diversity

Node color indicates correlation between sister clades and the selected predictor. Selecting a node highlights aggregated presence of species of one clade in blue and the other in red. Sites where species of both sides are present are purple.





Lifemapper on SAGE2

10:35 am localhost:9292 Webview: BiotaPhy - MCPA with Modeled Data for Proje Webview: Occurrence Map BiotaPhy | MCPA with Modeled Data for Project ##gridset name## BiotaPhy | MCPA with Modeled Data for Project heuchera global 10min Predictor: Env - Adjusted R-squared Show branch lengths Sites Map Semipartial Correlation b/w Node and Selected Predictor Clade 1 = Clade 2 = Both clades = Heuchera richardsonil Heuchera caroliniana Heuchera_pubescens Heuchera_alba Heuchera_longiflora Heuchera_americana Hauchera wootonii Heuchera_inconstans · Heuchera soltisii Heuchera novomexicana Heuchera glomerulata Heuchera_eastwoodiae chera brevistaminea Heuchera_resendahlii Heuchera sanguinea Leaflet | © OpenStreetMap Heuchera_villosa The map shows sites where the selected species are present. Use the select by Heuchera puberula Heuchera missouriensis bounding box or by polygon tools to highlight in the tree which species are present at Heuchera parviflora those selected sites. Heuchera chlorantha Heuchera_cylindrica Tellima grandiflora Mitella pentandra Mitella_stylosa Mitella furusei Mitella pauciflora Mitella japonica Webview: Lifemapper Package soniella oregona ithophragma parvillorum BiotaPhy | Package Mitella nuda Model Phylogenetic Occurrence Scatter Plot Elmera racemosa Map Projection Tree Mitella_caulescens Conimitella williamsii The BiotaPhy Project is supported by NSF BIO Award #1458422 Mitella stauropetala

Lifemapper on SAGE2

The vision for this app:

 Allow users to easily share visualized Lifemapper results with each other

- Facilitate side-by-side visual comparison between different packages
- Make Lifemapper more accessible







Future Work

- Port the BiotaPhy online interface into SAGE2
 - Requires communication with Lifemapper
- Cross-package features
 - Visualize data from multiple packages on one map
 - Allow users to submit multiple packages back to Lifemapper for correlation analyses
 - Then provide visualizations for these new results





Thank you!

This work is funded in part by grants (NSF ACI 1550126 and NSF ACI 1234983) from the National Science Foundation.



