



Enabling Biodiversity Research with Open Source Workflow, GIS and Metadata Tools

CJ Grady,

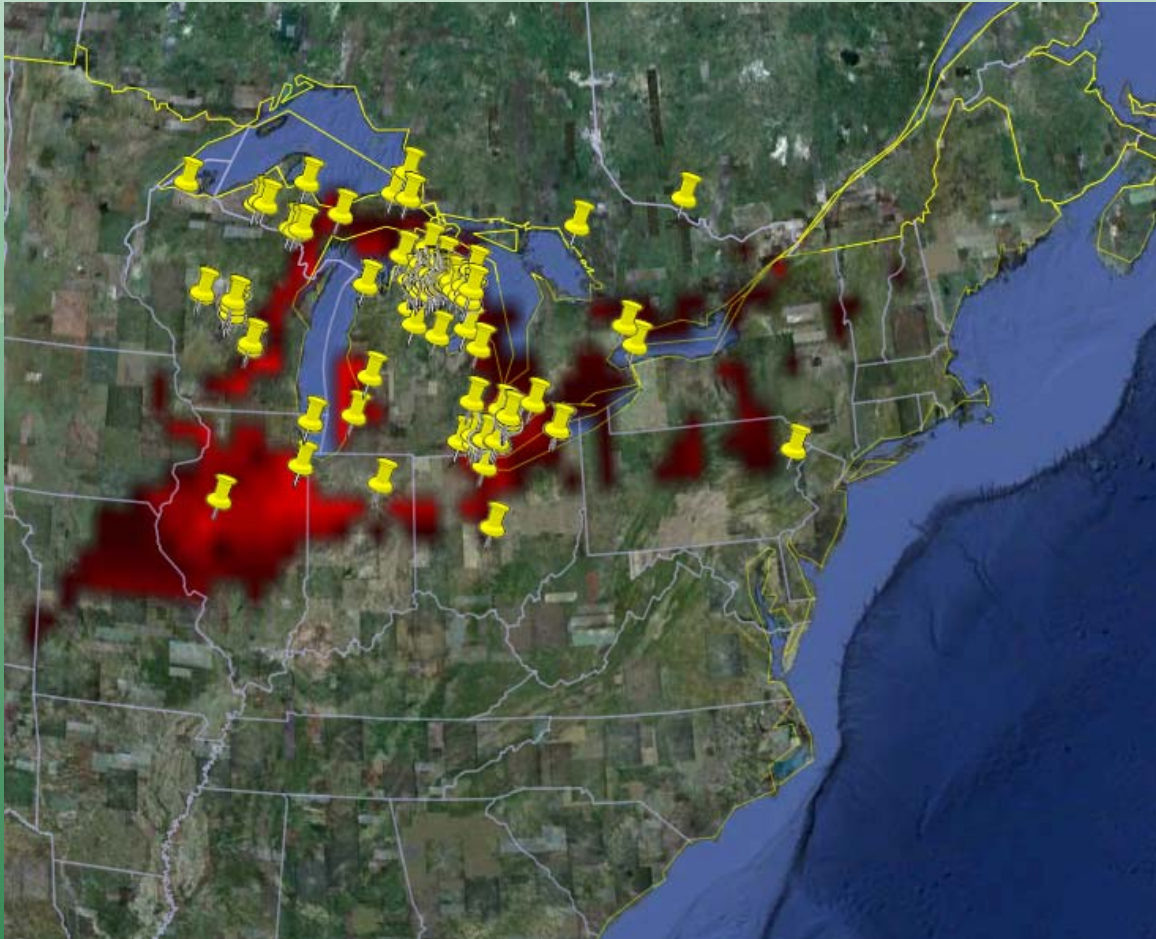
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<http://lifemapper.org>



What is Lifemapper?



Current museum (GBIF) vouchered occurrences for *Dendroica kirtlandii*



What is Lifemapper?

Integration

Web Services Standards

Interoperability Issues, Business Process Specifications, Management Specifications, Metadata Specifications, Reliability Specifications, Security Specifications, Transaction Specifications

Work Flow Tools

VisTrails

Archiving

DataONE

```

<dataset id="ds.1">
<title>Sample Dataset Description</title>
<creator id="23445" scope="document">
<individualName>
<surName>Smith</surName>
</individualName>
</creator>
</dataset>
</eml:eml>
    
```

EEML

Repeatable Transparent Science

Ellison, A. 2010. J. Ecology

Analytical method(s) used	Analytical tool(s) used	Comments
linear and quadratic regressions	none specified →	not repeatable
ordinary least-squares regression	SYSTAT 8.0	possibly repeatable; current available version is 12.0
Poisson regression	NAG statistical add-in for Excel →	not repeatable; software discontinued
"Mitchell-Olds and Shaw test" (Mitchell-Olds and Shaw 1987)	none specified →	not repeatable; software unavailable (but algorithm available); which of three tests proposed by Mitchell-Olds and Shaw was also not specified
chi-square exact test	StatXact	possibly repeatable; no version given
meta-analysis using mixed-effects model	MetaWin 2.0	repeatable; commercial software version still available
Poisson regression	not specified →	not repeatable
ordinary least-squares regression on "some" data sets of Mittelbach	software not specified →	not repeatable



Background

- Workflows
 - A series of connected steps describing a process
- Metadata
 - Information that describes a data set or process.
 - Data about data
- Open Source Software
 - Software that includes access to the source code used to create it. This is provided to encourage study, contribution and sharing.



So What's the Problem?

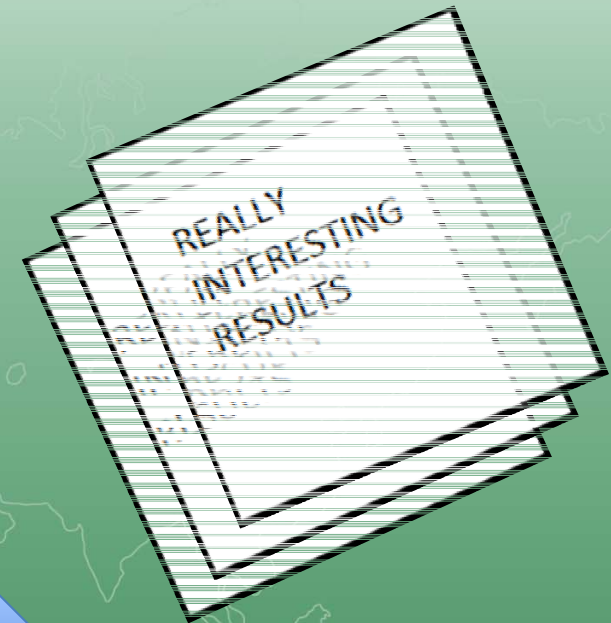
- Difficult and time consuming to assemble biodiversity experiments by hand
- Scientists often don't have adequate computing resources
- Experiments can be difficult or impossible to reproduce



Study of Experiment Reproducibility



???



METHODS

1. -----
2. -----
3. -----



Ellison, Aaron. 2010. Repeatability and transparency in ecological research. Ecology 90.

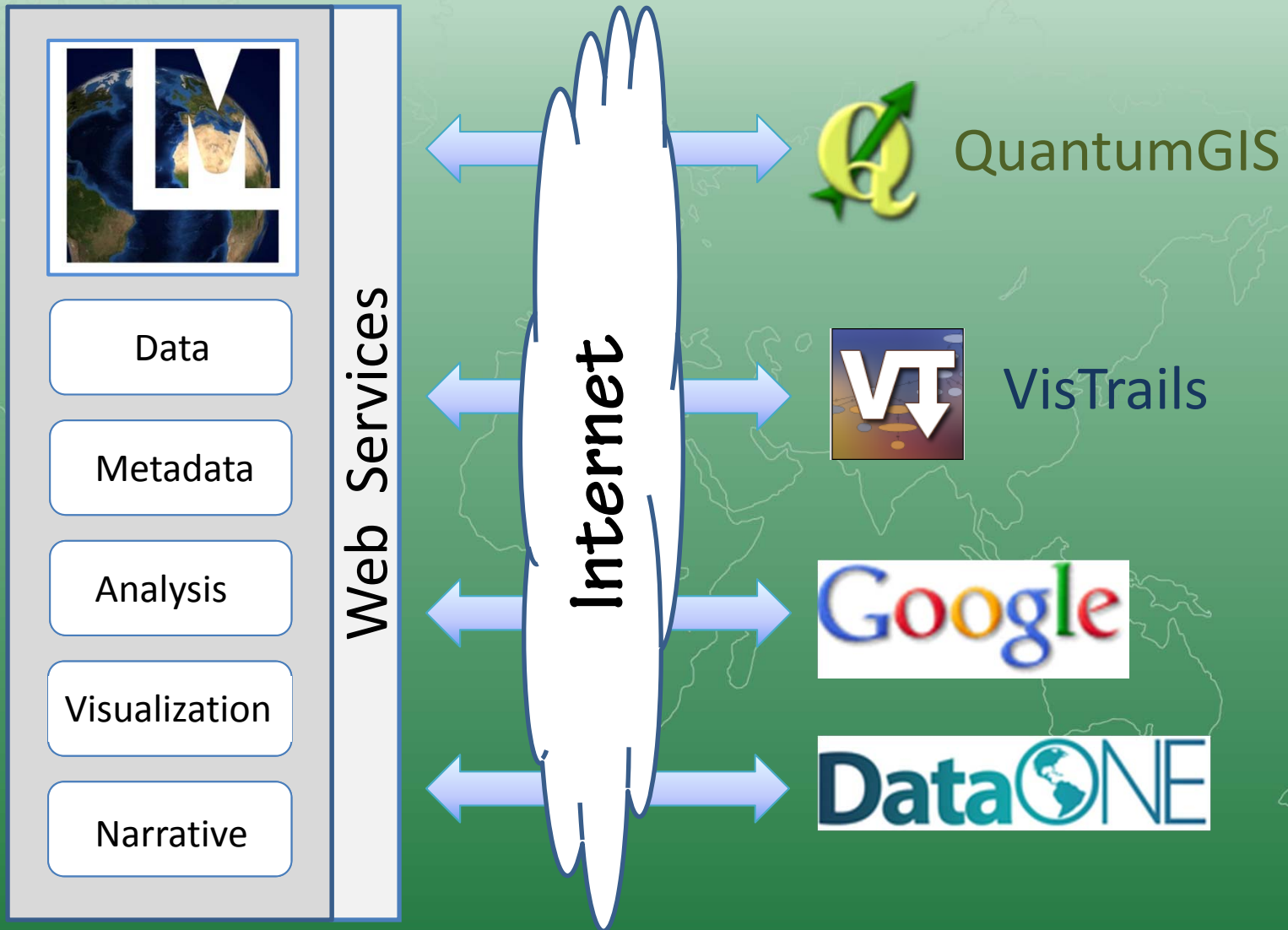


What we have done

- Metadata for all of our Species Distribution Modeling services
- Simple process metadata
 - Documents how an experiment is ran through our cluster including what versions of software
 - Also describes what web services would be called to execute the experiment again



Web Services



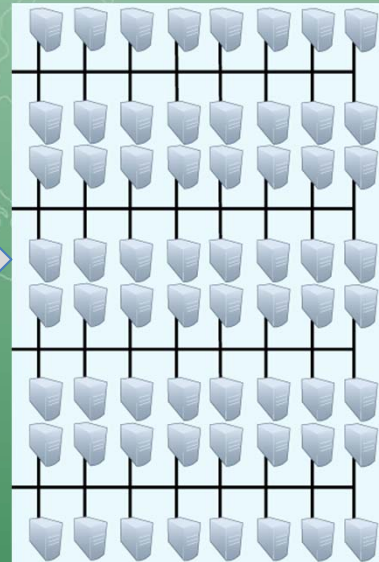
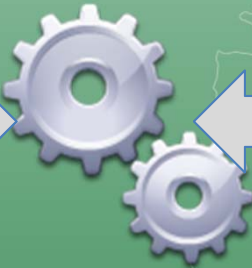


Lifemapper Backend

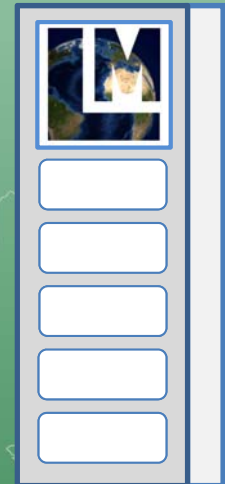


Data Archive

Pipeline



Compute Resources

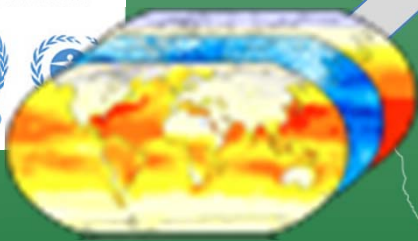
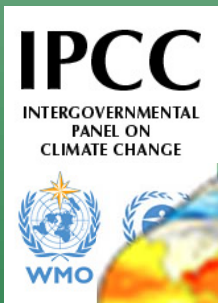




Species Distribution Modeling



Species Occurrence Data



Environmental Data

open
Modeller

SDM Modeling Algorithm



Predicted Habitat



What we are doing

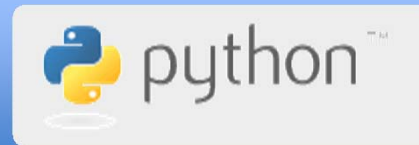
- Publishing metadata to a public repository
- Client extensions
- Lifemapper Range and Diversity



Clients



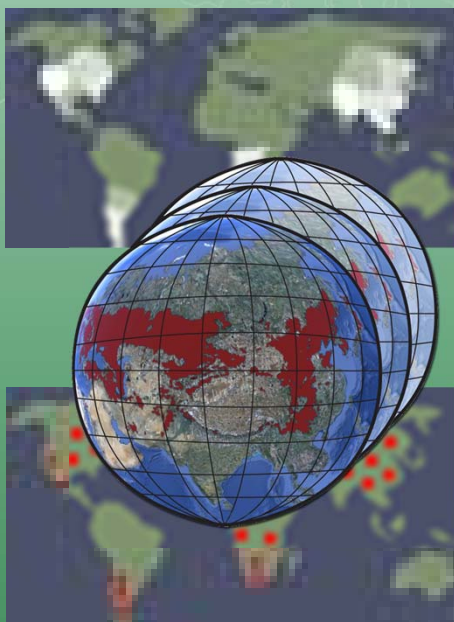
WWW



Lifemapper Web Services



LmRAD: Range and Diversity

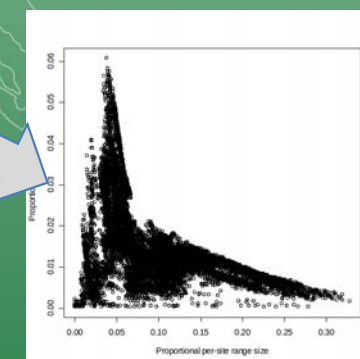
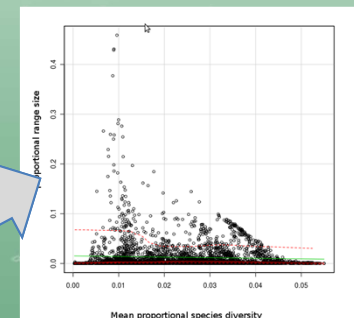


Species Habitat Data

M =

0	0	0	0	1	0	0	0	0	0	0	1	0
0	0	0	0	0	0	0	0	0	1	0	0	0
0	0	0	0	0	0	1	0	0	0	1	0	1
1	0	1	0	1	0	0	0	0	1	0	1	0
1	0	0	1	0	0	0	0	0	0	0	0	0
0	0	1	0	0	0	1	0	0	0	0	0	0
0	0	1	0	0	0	1	0	0	0	1	0	1
0	0	1	0	0	1	0	0	0	0	0	0	0
0	0	1	0	0	0	0	1	0	0	0	0	0
0	0	1	0	0	0	1	0	0	0	0	0	0
0	0	1	0	0	0	0	0	0	0	0	0	0
1	0	0	1	0	0	0	0	0	0	0	0	0
0	0	1	0	0	1	0	0	0	0	0	0	0

Presence Absence Matrix (PAM)



Range and Diversity
Quantifications



Species Distribution and Diversity

- Biodiversity patterns
 - Species abundance, distribution and diversity
 - Multiple scales and extents
 - Used for conservation and management decisions
- Challenges
 - Large extents ($> 10,000 \text{ km}^2$)
 - Fine resolution ($< 1000 \text{ m}^2 \approx 30\text{m} \times 30\text{m}$)
 - Many species (10,000 +)



QGIS with Lifermapper MacroEcology plug-in


File Edit View Layer Settings Plugins Raster Vector MacroEcology Help

- Sign In
- New User
- New Experiment**
- Resume Experiment
- Populate Grid

Layers

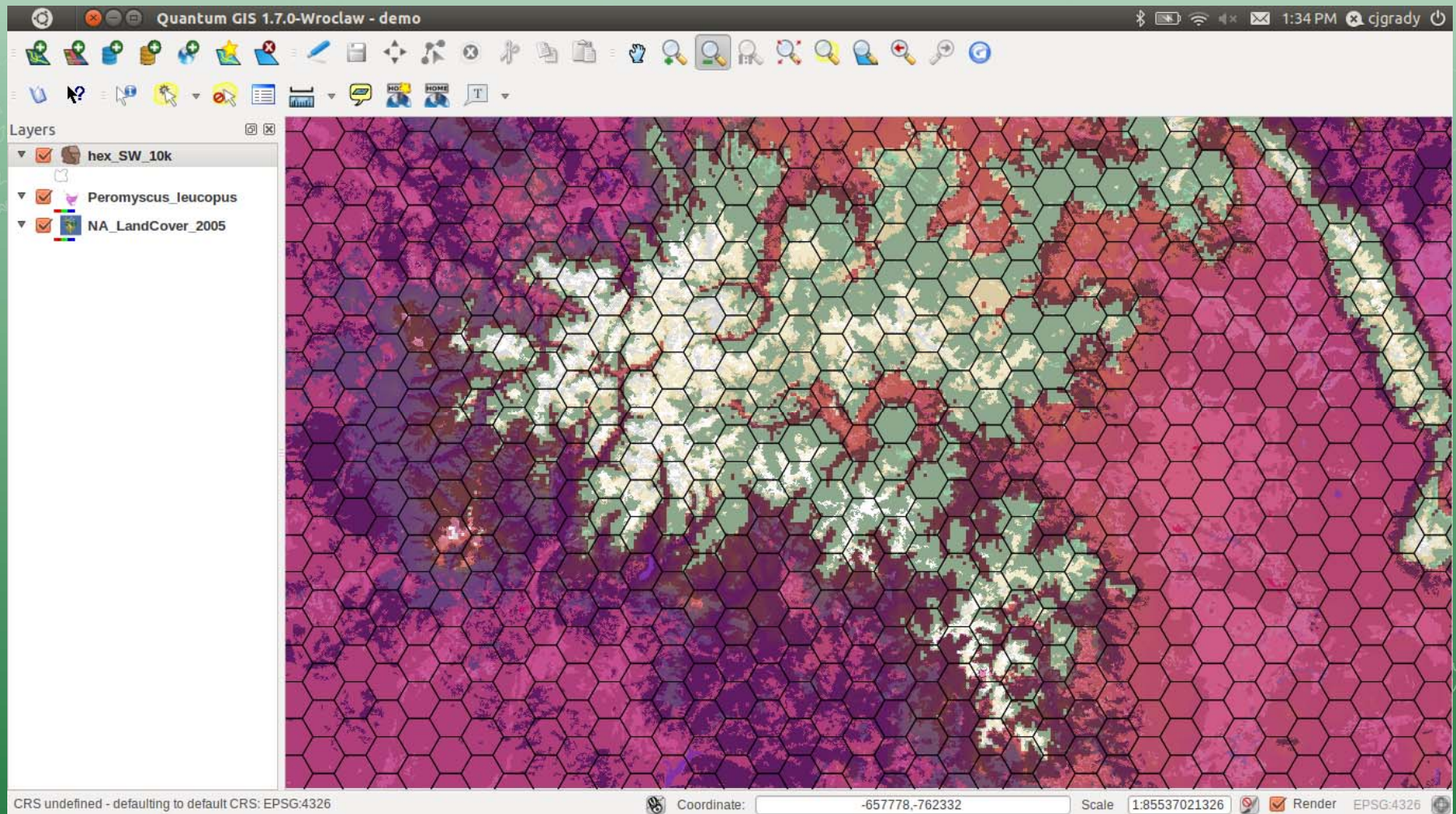
- Peromyscus_leucopus
- NA_LandCover_2005

Coordinate: -4083532.5375726 Scale: 4934194123563 Render EPSG:4326



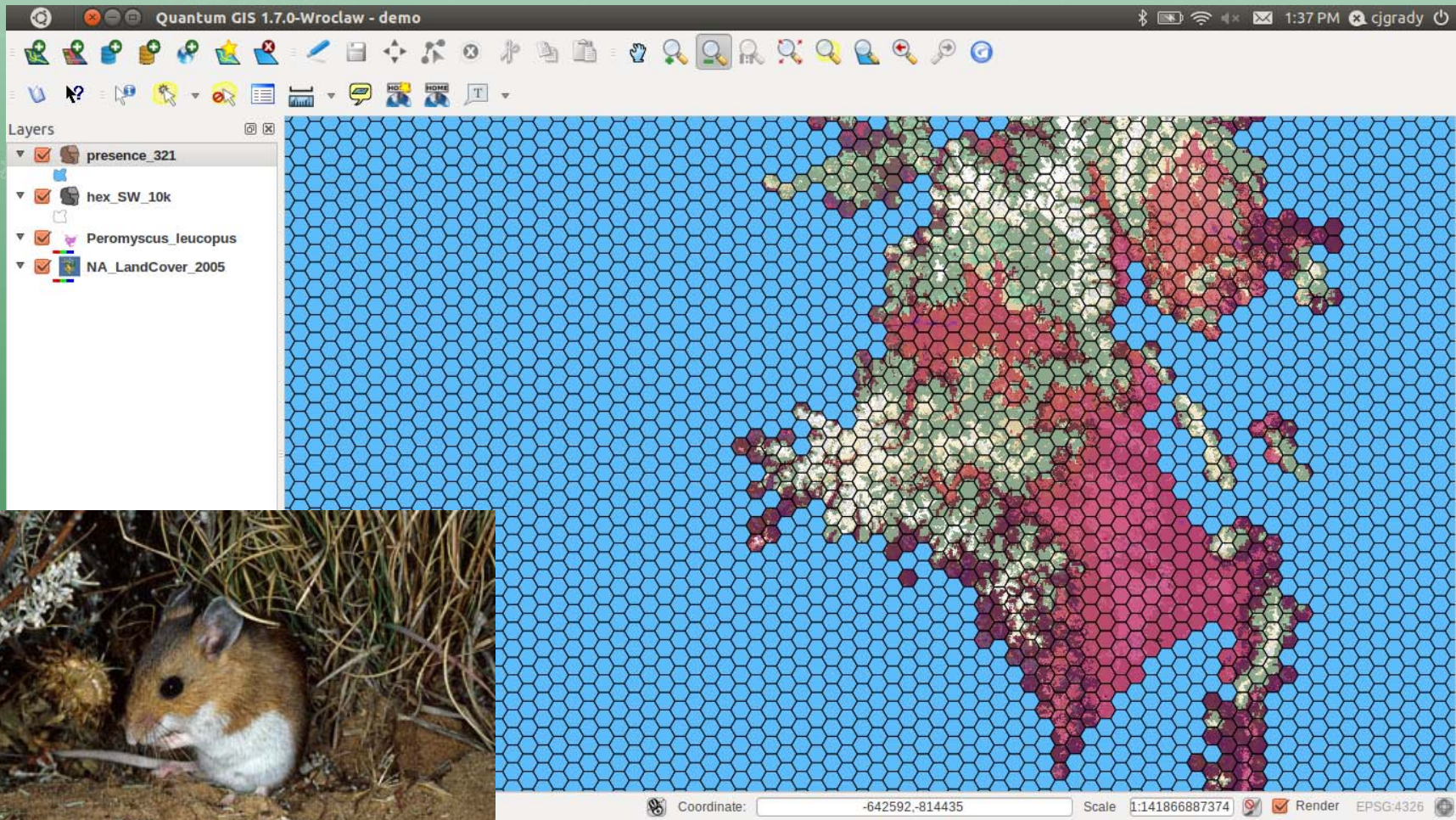


Hexagonal Grid



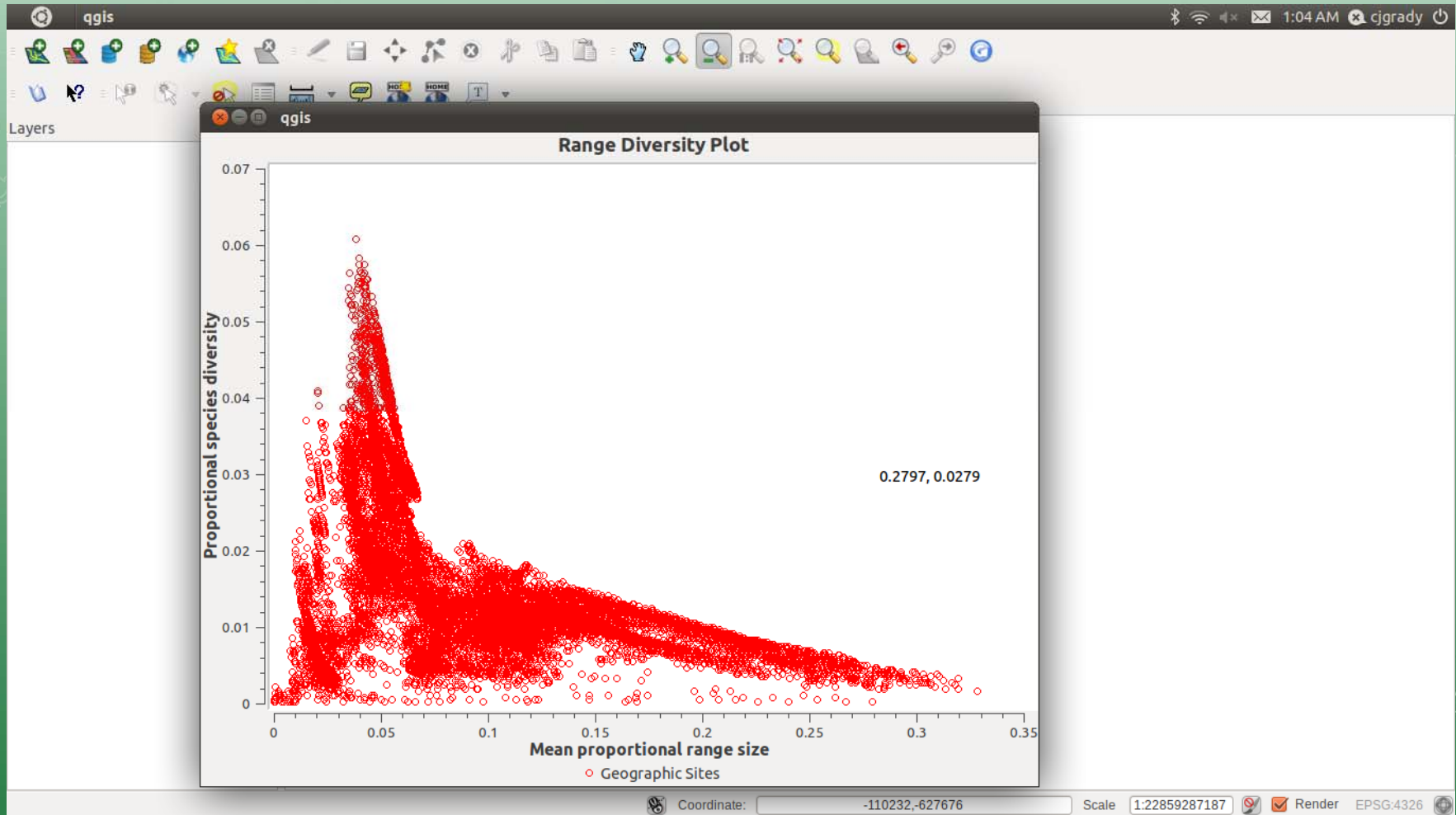


Mouse 'Presence'



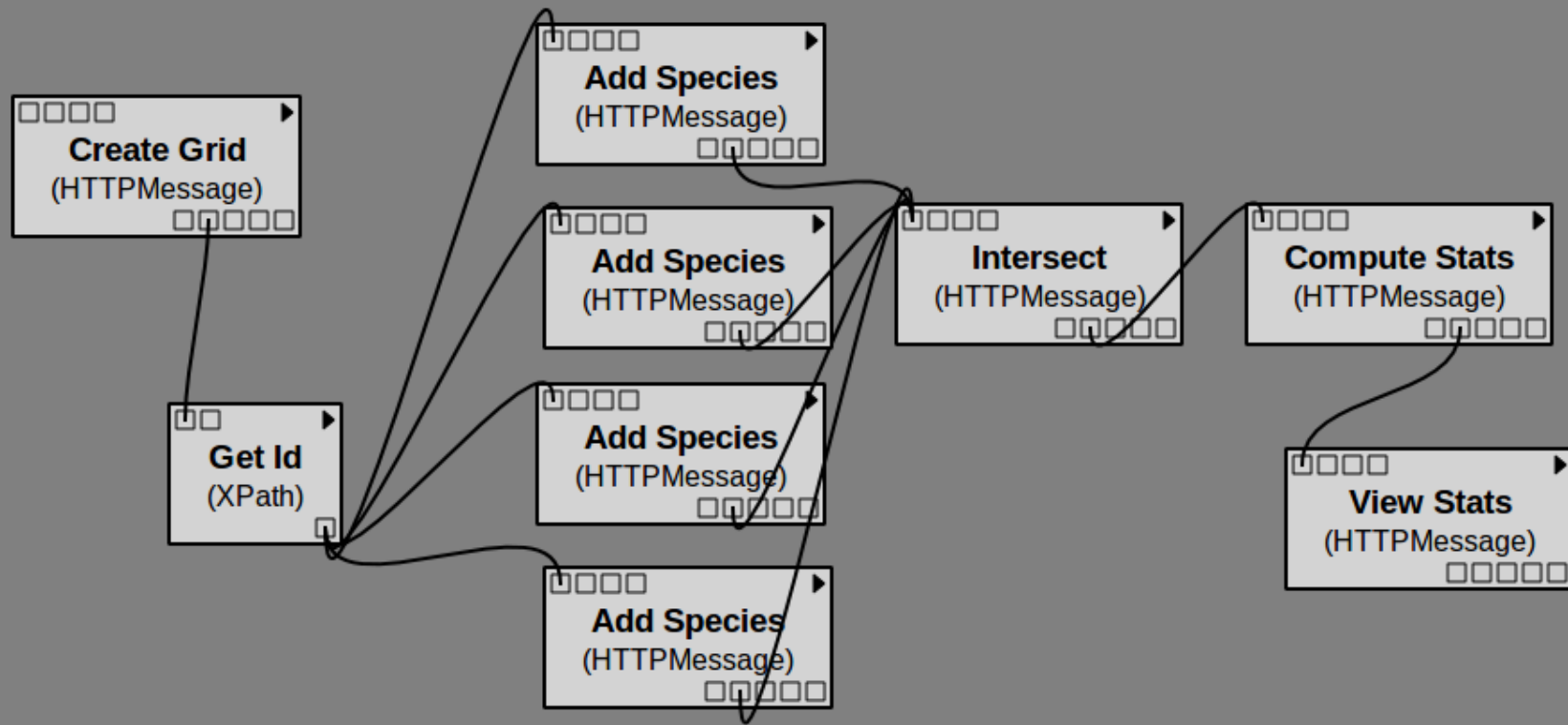


Sites / Species Plot





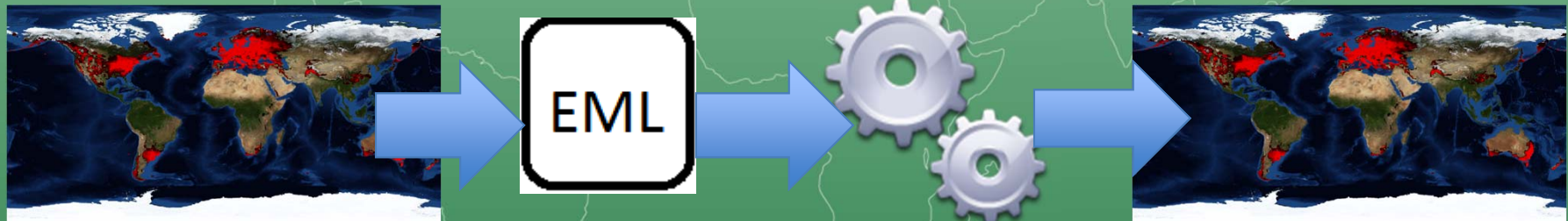
VisTrails





Reproducibility

- Simple process metadata
- Process metadata extensions
- Lifemapper client metadata reader





Collaborations

- KU Biodiversity Institute



- NSF
Cyber-Commons



- Change Thinking



- CI Team





<http://www.youtube.com/watch?v=VCFixtqlimg>



Future Directions

- Publish metadata through standard APIs
- Contribute process metadata extensions to community
- Gesture based interface
- Explore extensions into cloud and other grid computing environments



Parallel Processing





Summary

- Provide end-users with clients to assemble and manage biodiversity modeling experiments
- Allow users to harness the computing power available through our cluster to perform computationally intensive tasks
- Include process metadata to document how an experiment was performed



Funding

U.S. National Science Foundation



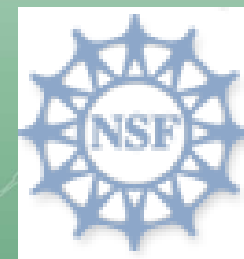
NSF EPSCoR 0553722

NSF EPSCoR 0919443

EHR/DRL 0918590

BIO/DBI 0851290

OCI/CI-TEAM 0753336





Questions

- cjgrady@ku.edu
- <http://lifemapper.org>