



# **“Tree of Life” Teaching Tool**

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University of Pennsylvania  
DMD Senior Design  
May 1, 2009  
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# Presentation Outline

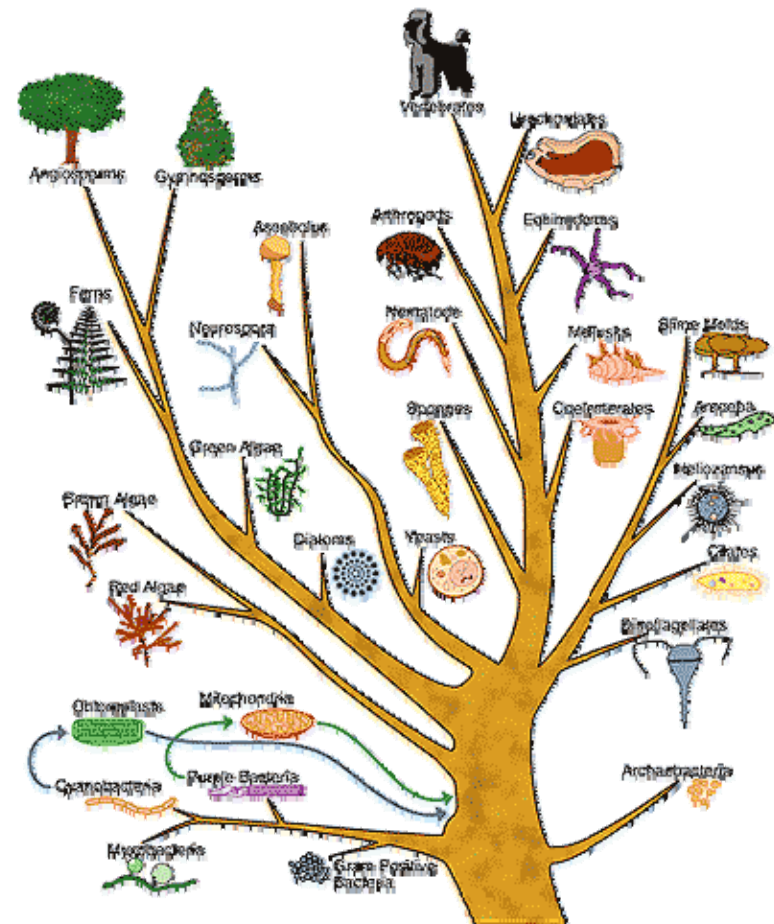
- Describing My Project
- Expanding PhyloWidget
  - Algorithms
  - Results and Achievements
  - Demo
- Wrap-Up
  - Future Directions
  - Tools/Languages
  - Thanks
- Questions



# The Problem

## *Phylogenetics*

the study of how organisms are related by evolution





# The Problem

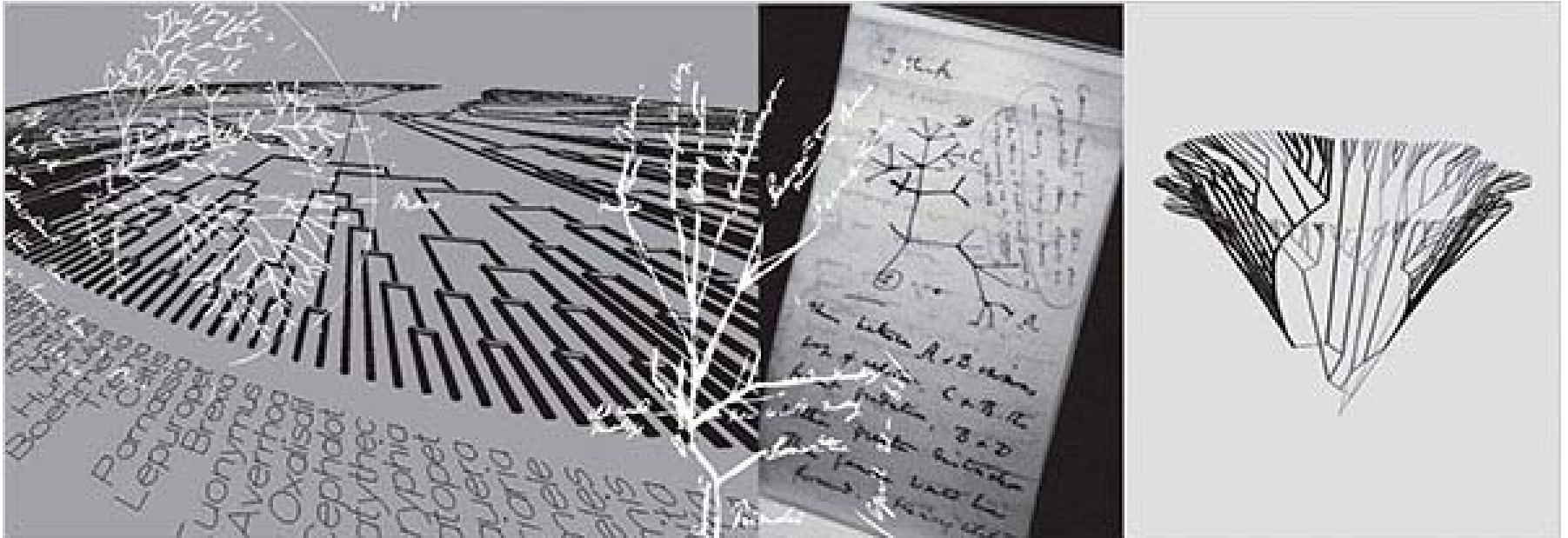
Data is only as valuable as a user's ability to interact with it and learn from it.

My program allows students to use this great body of knowledge in a way that is better suited to their needs.



# The Problem

## Tree Visualization in The New York Times





# Prior Work

*Tree of Life Web Project*

- Catalogue and classify data about *biodiversity* and *phylogenetics*
- Aid *learning* about and *appreciation* of biological diversity and the evolutionary Tree of Life





# Prior Work

*TreeBASE II*

published phylogenetic research  
and the trees generated by those  
studies



# Prior Work

*Tamara Munzner's TreeJuxtaposer*

- Scalability in Tree and Display Size
- Guaranteed Visibility of Landmark Nodes
- Limited Label Overlap



# Prior Work

*Tamara Munzner's TreeJuxtaposer*



Figure 1: Left, Right Top: Biologists faced with inadequate tools for comparing large trees have fallen back on paper, tape, and highlighter pens. Right Bottom: TreeJuxtaposer is a scalable tool for interactive exploration and comparison of trees.

# Prior Work

*Tamara Munzner's TreeJuxtaposer*

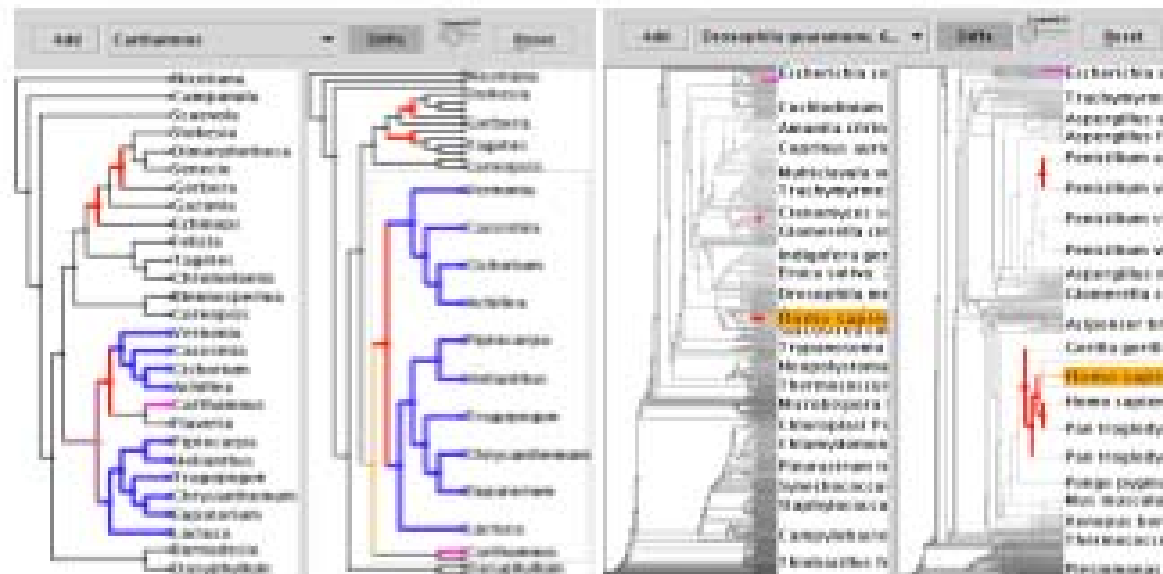
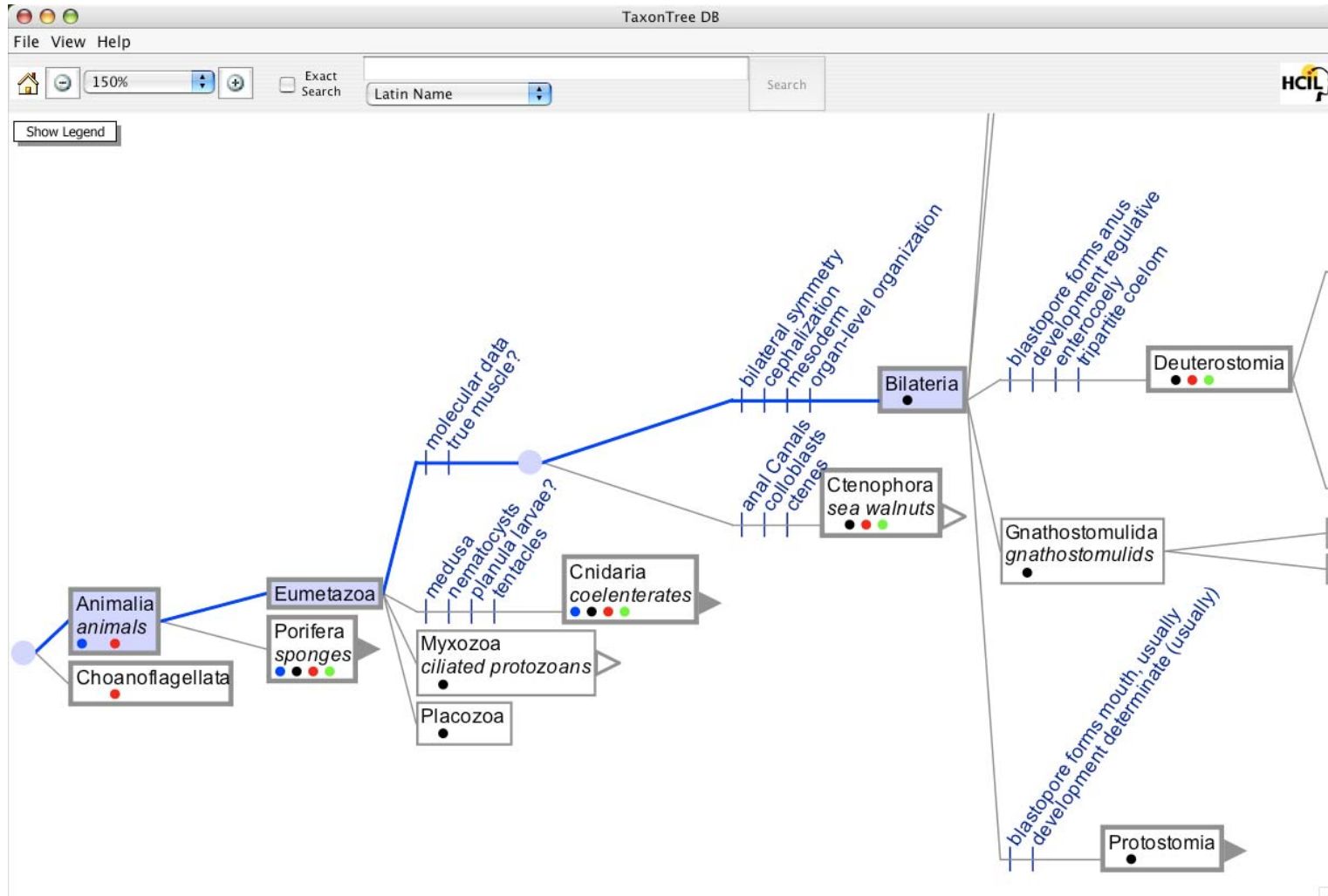


Figure 5: Tree comparison between two variants of a single phylogenetic reconstruction run, with the exact location of structural differences (in biological terms, nonmonophyletic clades) marked in red. The left tree is in the undistorted overview position, while parts of the right side have been expanded. **Left:** A small 55-node tree. **Right:** A larger 1600 node tree.



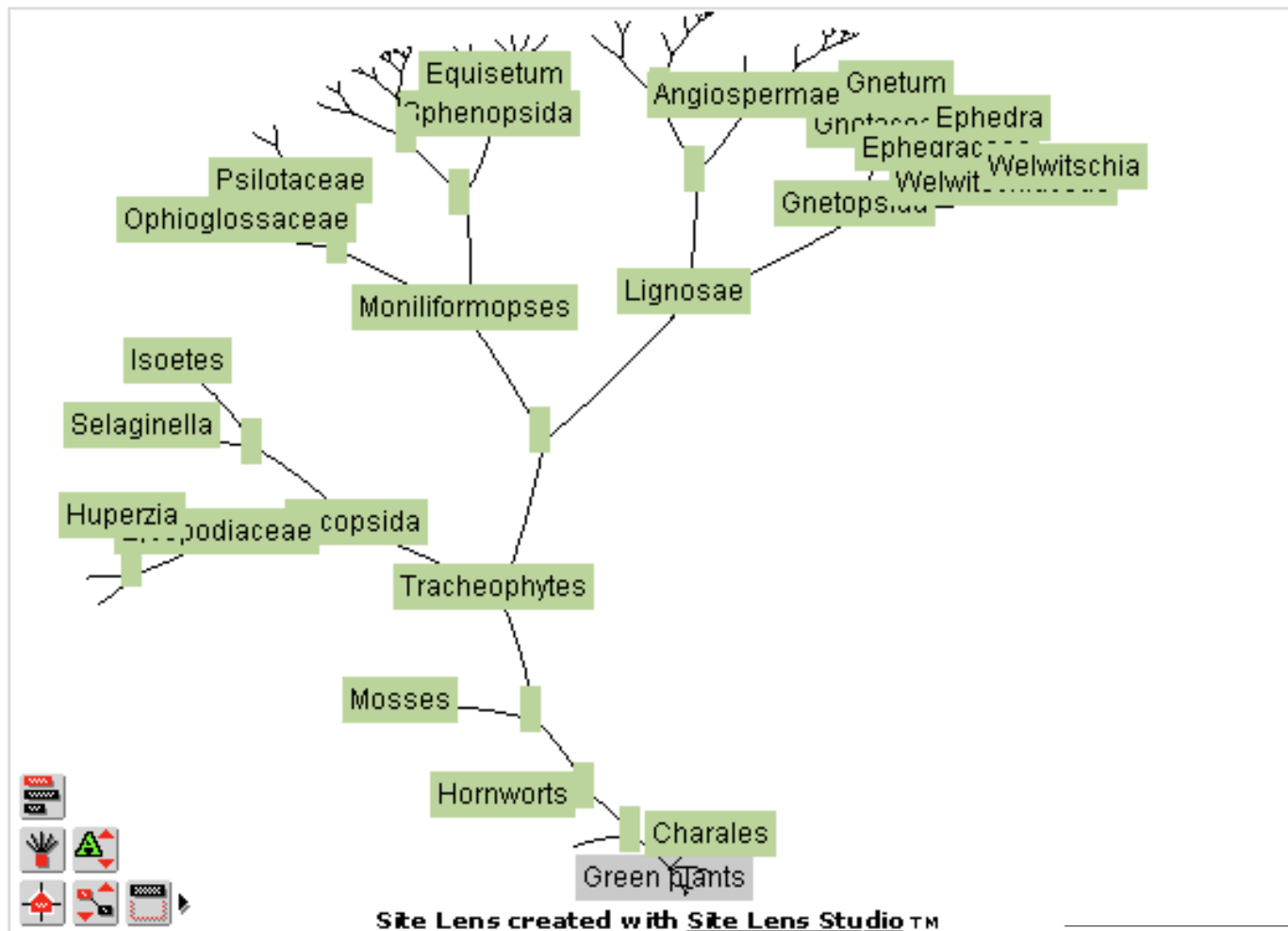
# Prior Work

*Denise Green and Rebecca Shapley*

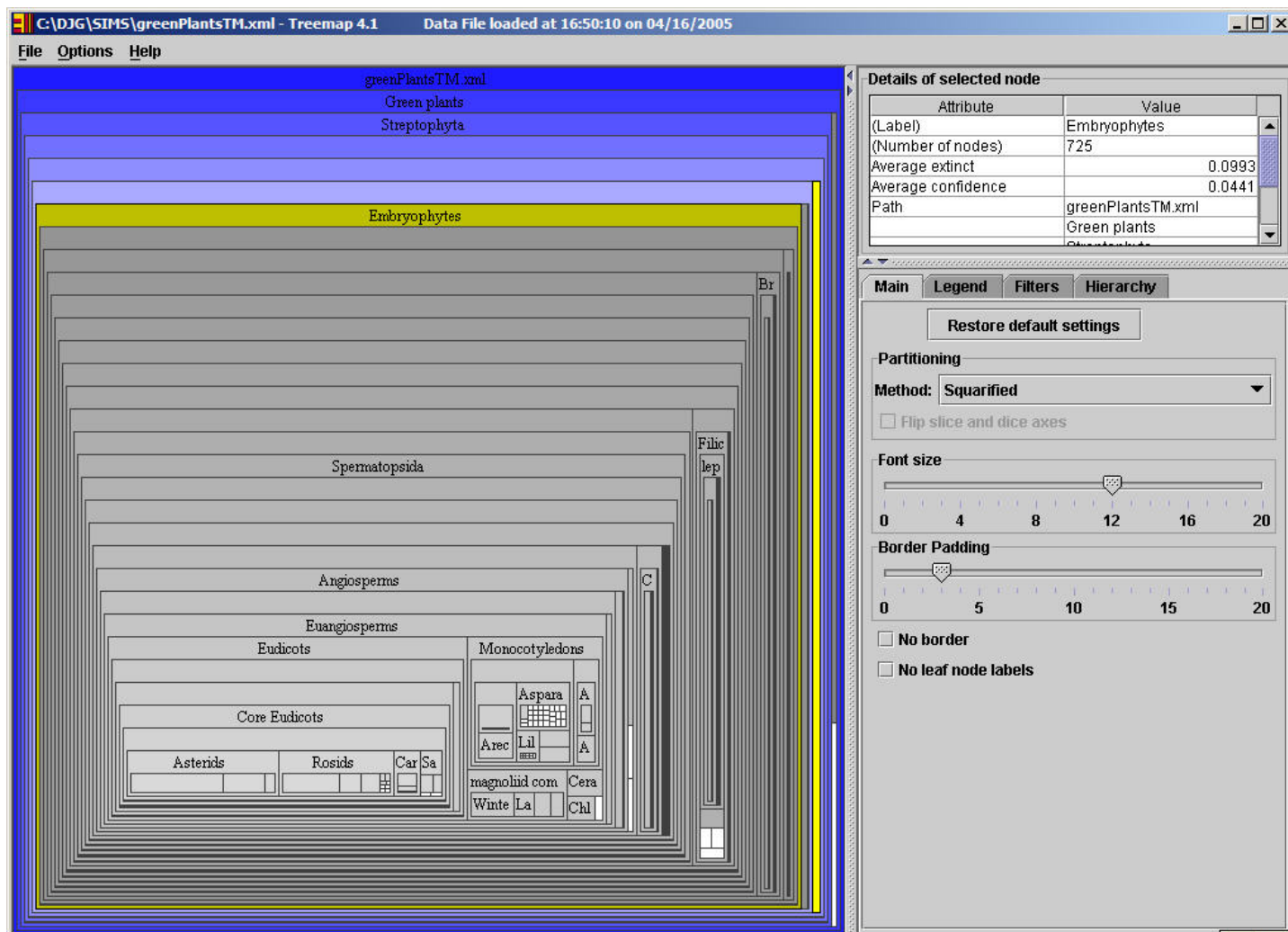


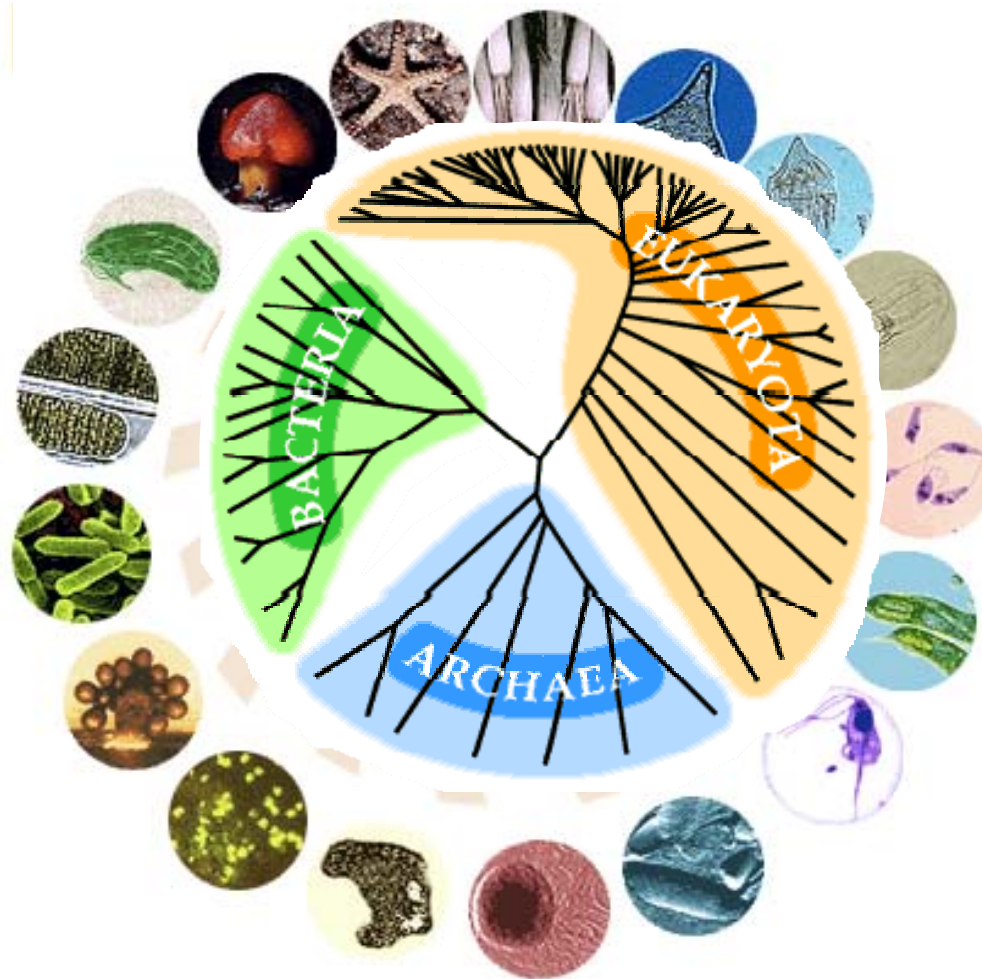
# Prior Work

*Denise Green and Rebecca Shapley*



*Denise Green and Rebecca Shapley*







# Prior Work

*Denise Green and Rebecca Shapley*



General Recommendations:

- Solve the “Overwhelm Factor”
- Simple, Teachable Tree Views
- Connect with the Familiar:  
History, Common Names, and  
Pictures



# Prior Work

*Denise Green and Rebecca Shapley*

## Relevant Recommendations

- *Images of organisms*
- *Common name search*
- Variable branch lengths
- Rank-free circular layout
- Animated changes
- Non-overlapping labels
- Resizeable window and text







# Prior Work

*Denise Green and Rebecca Shapley*

A *fully integrated system*\* ... would be fantastic. Having a website that seamlessly relates [data from independent sources] would *save an unimaginable amount of teaching and preparation time.*

\* Emphasis added



# Prior Work

*PhyloWidget*

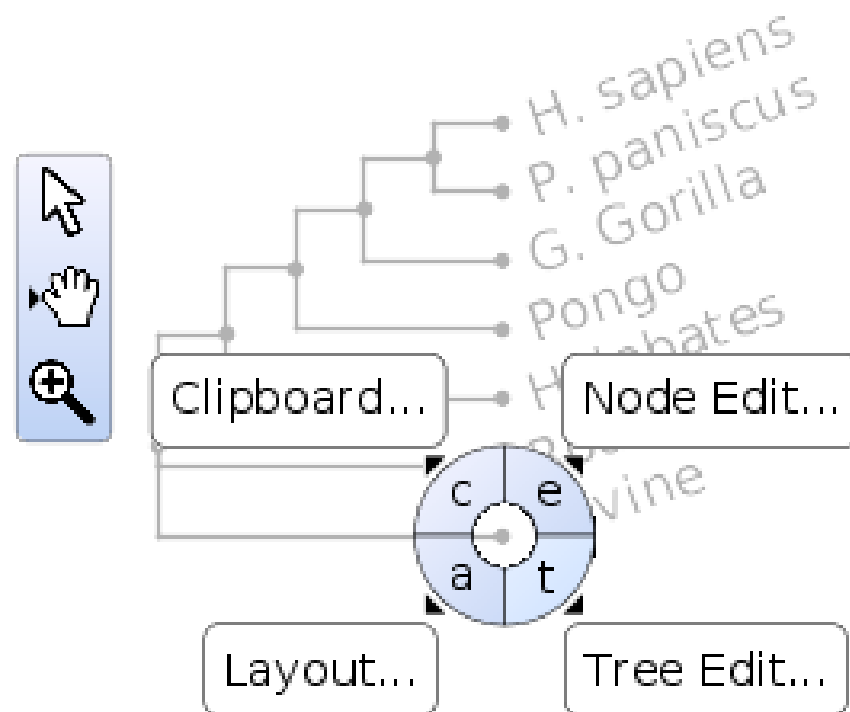


*“viewing, editing, and publishing  
phylogenetic trees online”*

# Prior Work

*PhyloWidget*

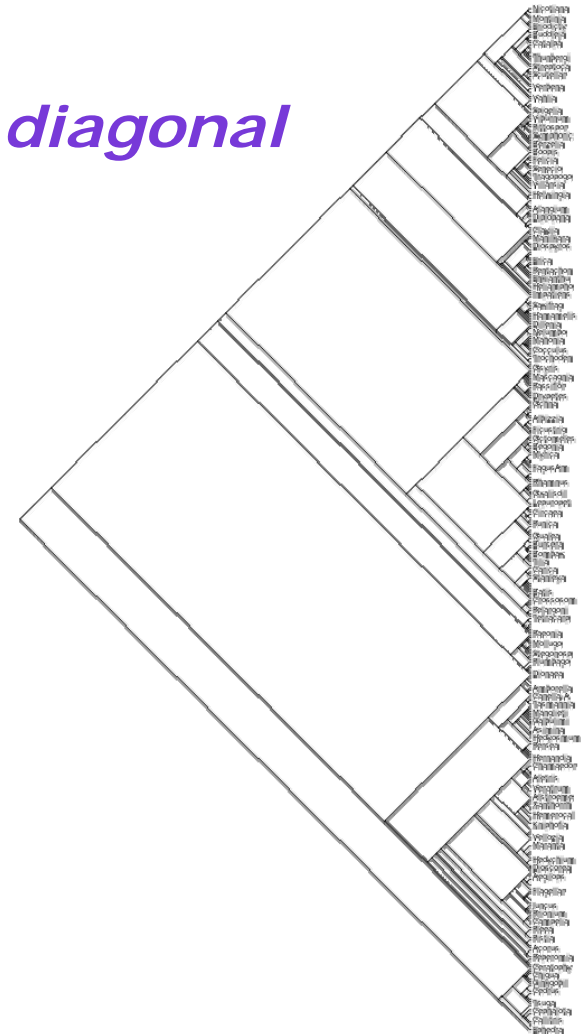
File View Tree Search:



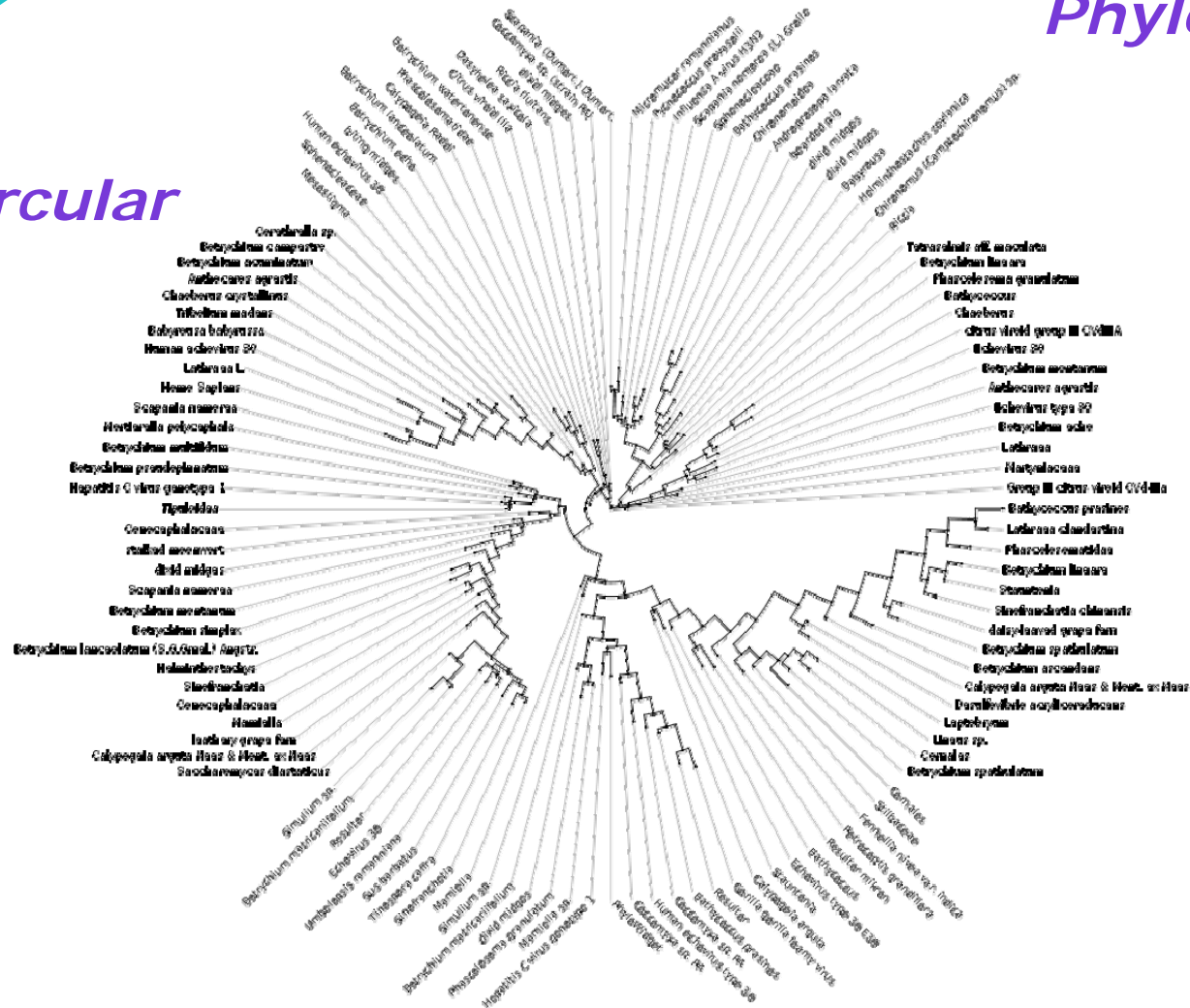


# Prior Work

## *PhyloWidget*



*circular*





# Prior Work

*PhyloWidget*

I chose to use PhyloWidget as the basis of my project. Why?

- Take advantage of the existing interface and tree renderer
- Focus on fulfilling the need for integrated images and common names in educational environments





# Prior Work

*More Introductions*

Newick Tree Format

NHX New Hampshire eXtended

NBII National Biological Information Infrastructure

Morphbank

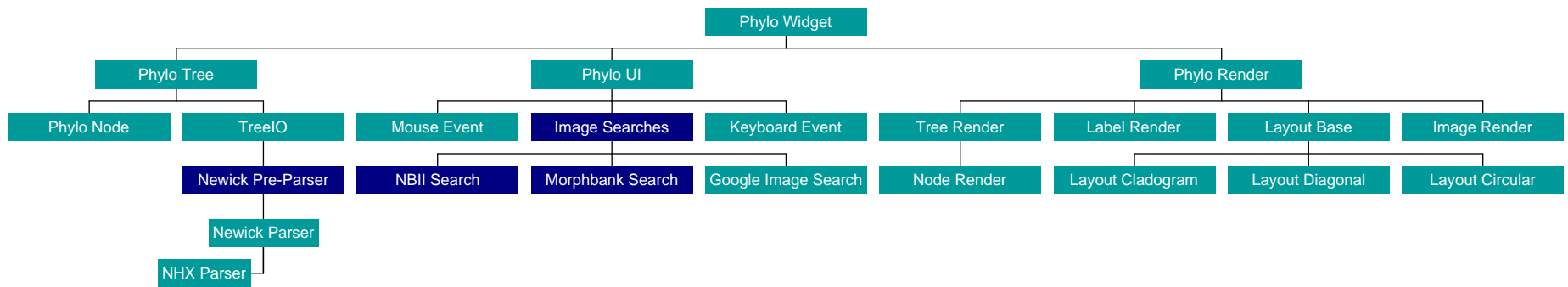
UBIO





# Algorithms

## *Overall Flowchart*







# Algorithms

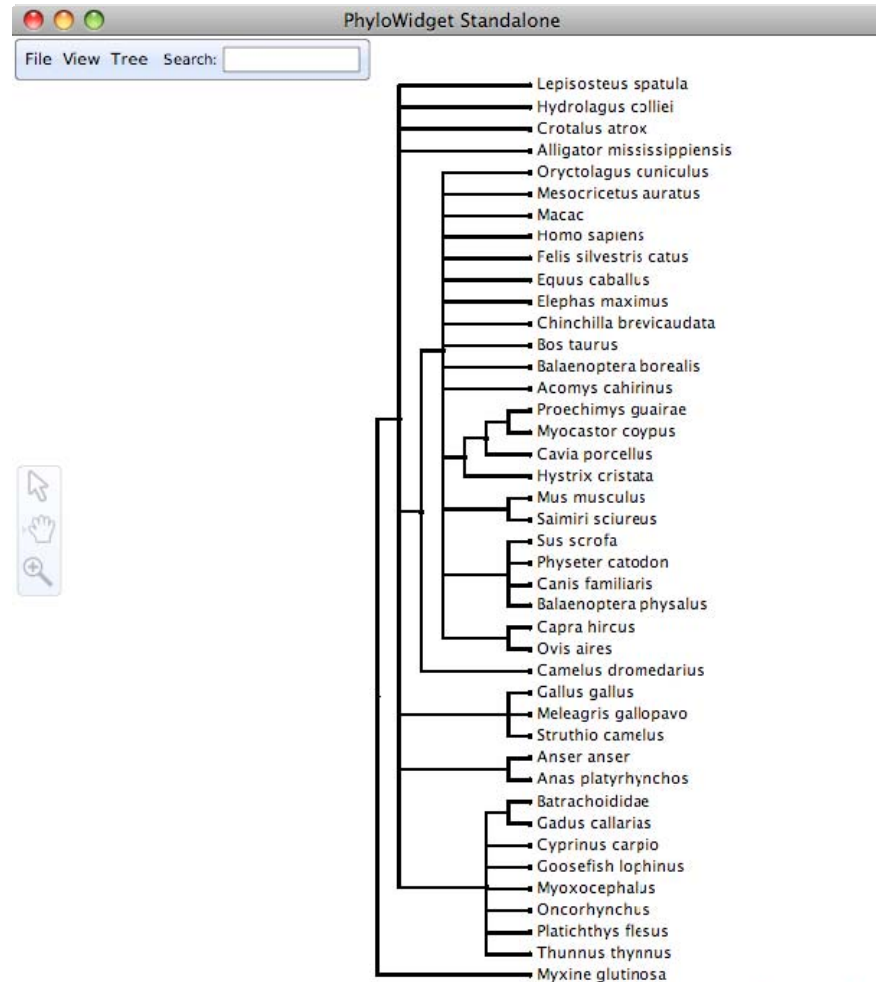
## *Pre-Parsing a Newick File*

```
#NEXUS
BEGIN TREES;

TRANSLATE
1 Alligator_mississippiensis,
2 Chinchilla_brevicaudata,
3 Felis_silvestris_catus,
4 Balaenoptera_borealis,
...
40 Sus_scrofa,
41 Bos_taurus,
42 Mammalia,
43 Macaca
;
TREE tree_0 = [&R]
  (((((((((11,20),24),19),2,7,18,(36,23),37,43,5,(16,17,6,40),29,25,41,3,4,(34,39)),
  10),(33,8,21),30,1,15,(38,12),9,((31,26),32,14,28,13,27,35)),22);
ENDBLOCK;
```

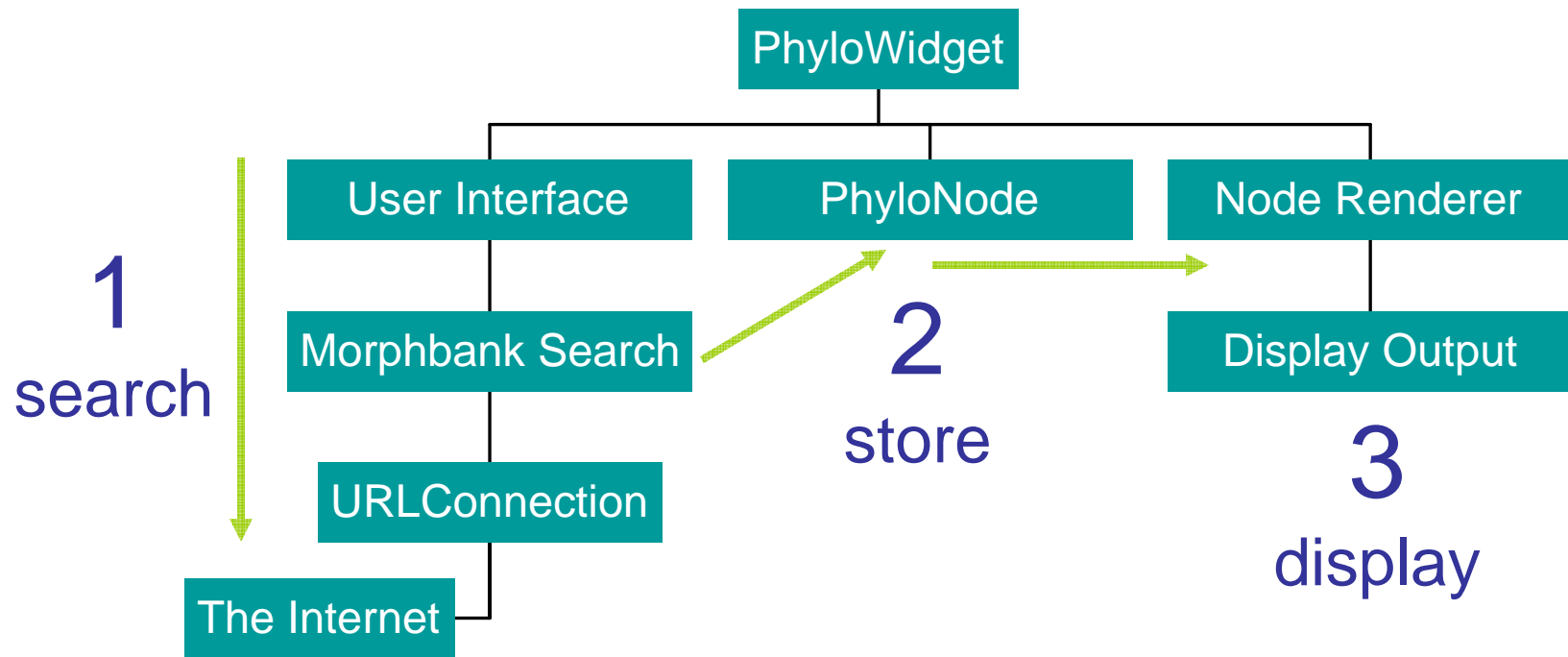
# Algorithms

## *Pre-Parsing a Newick File*



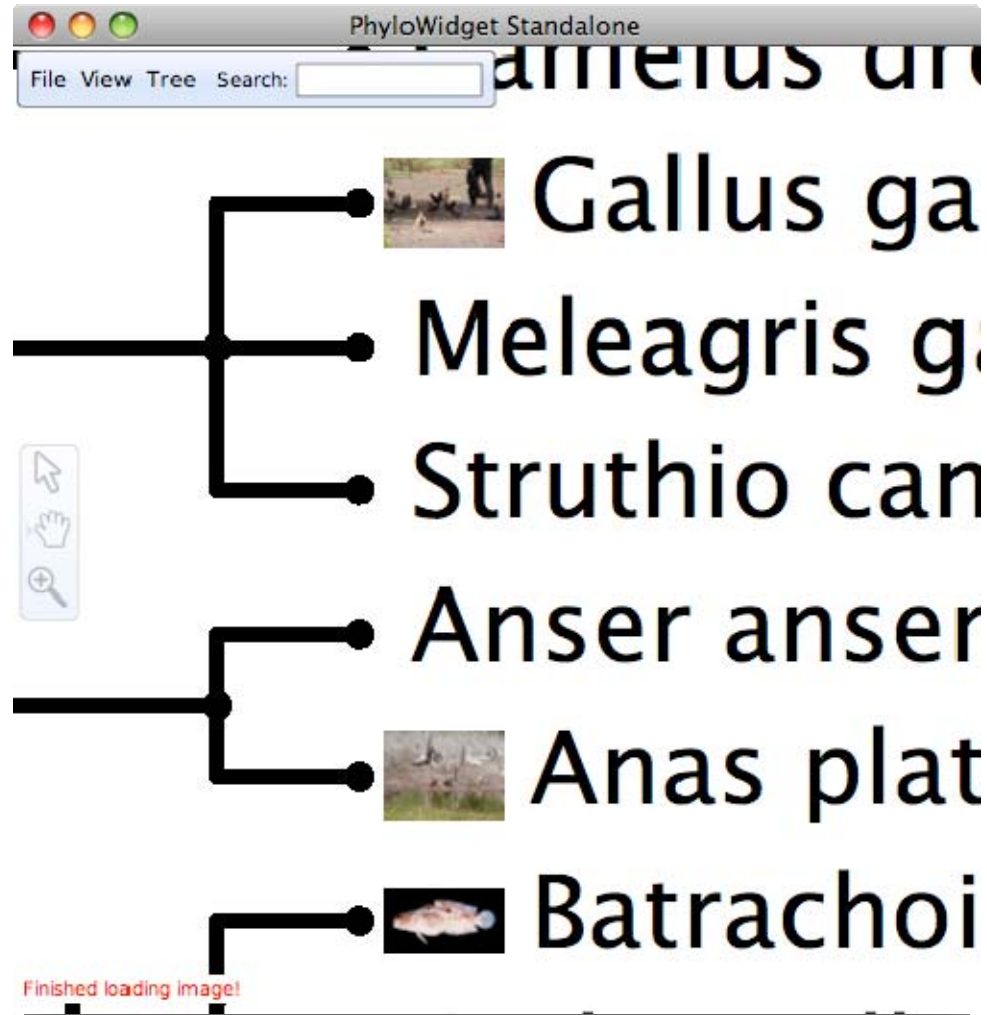
# Algorithms

*Image Search and Display*



# Algorithms

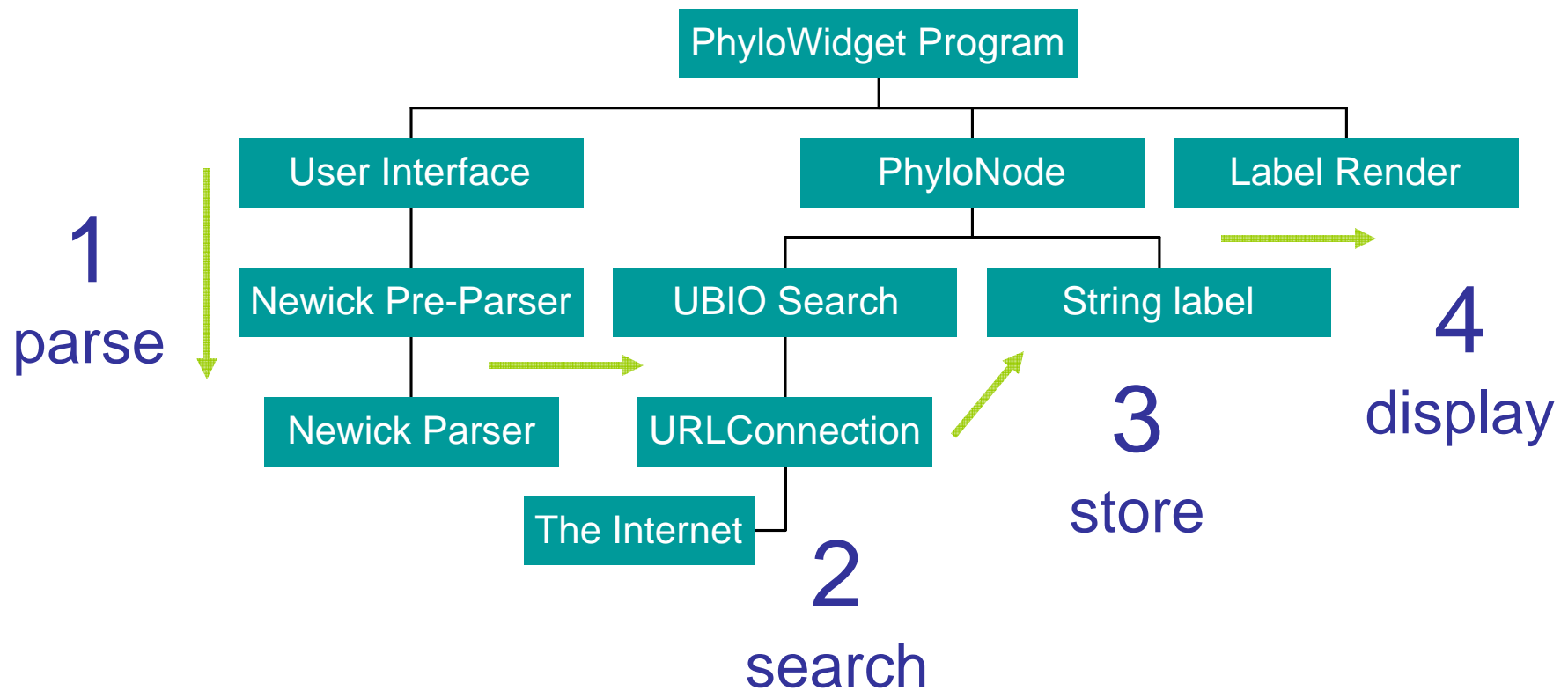
*Image Search and Display*





# Algorithms

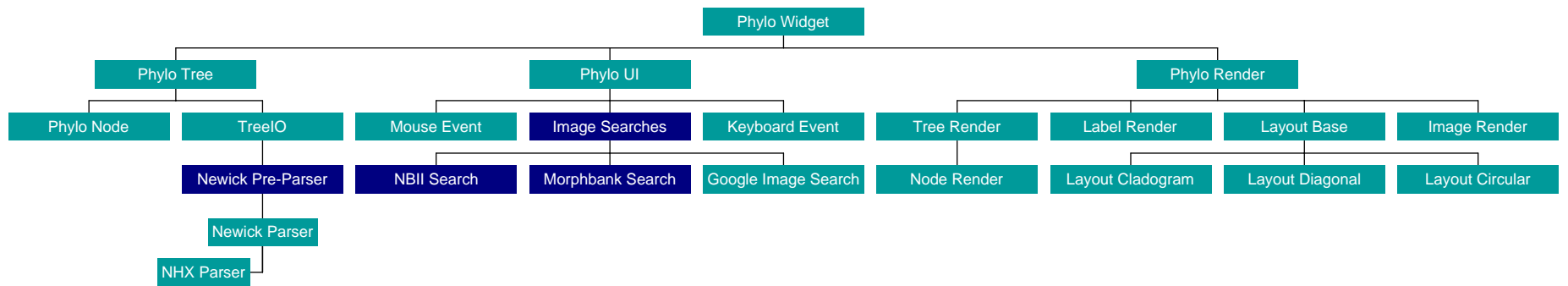
*Common Name Search and Display*





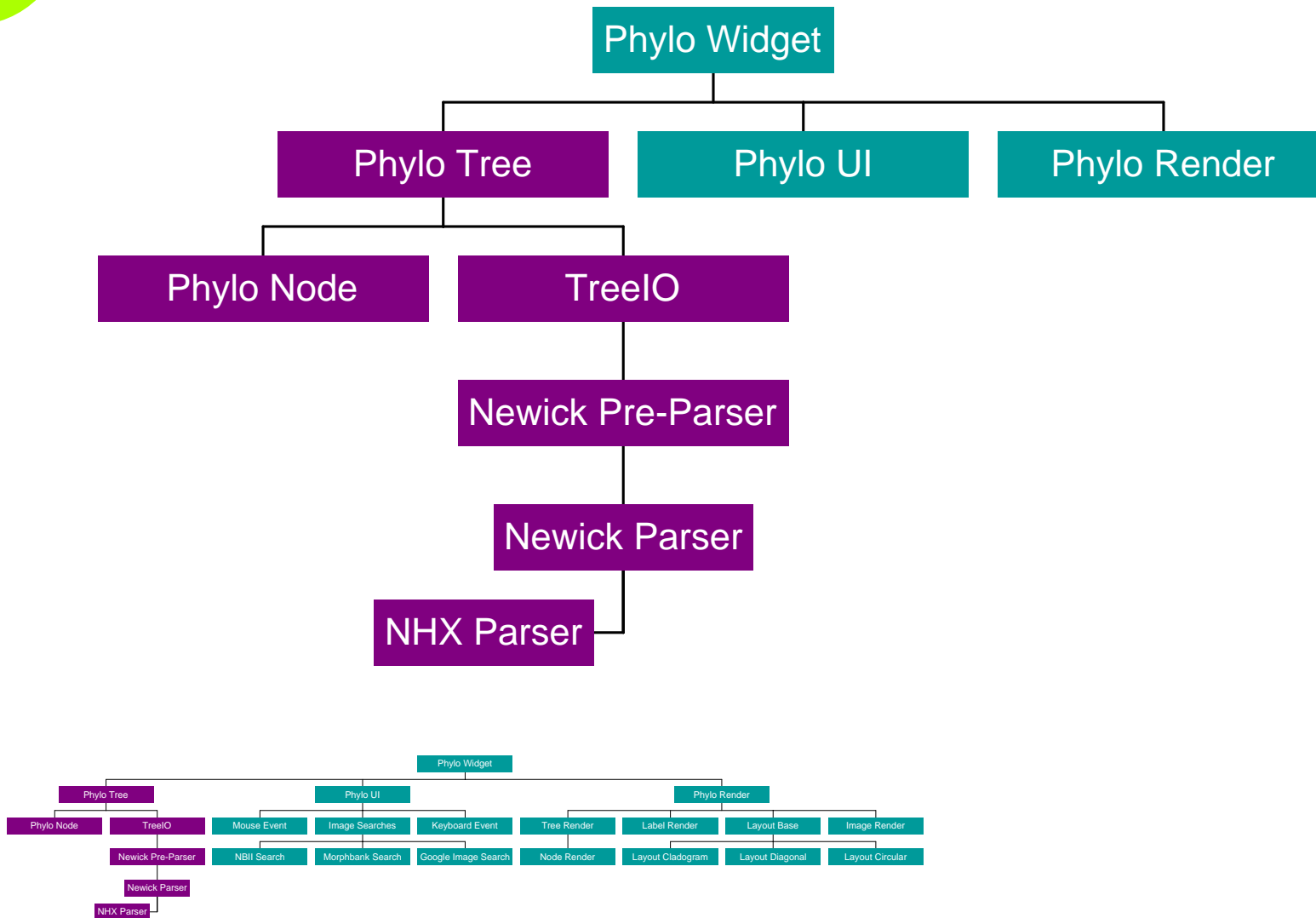
# Algorithms

## *Overall Flowchart*



# Algorithms

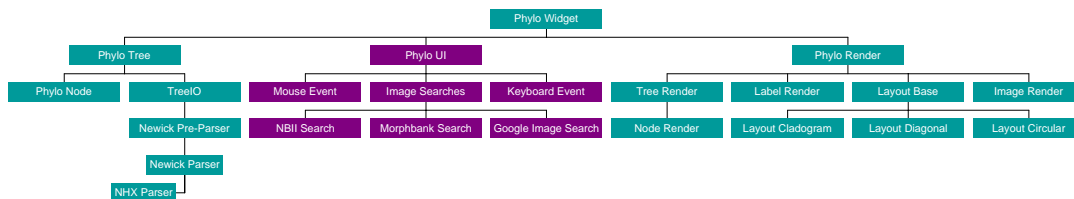
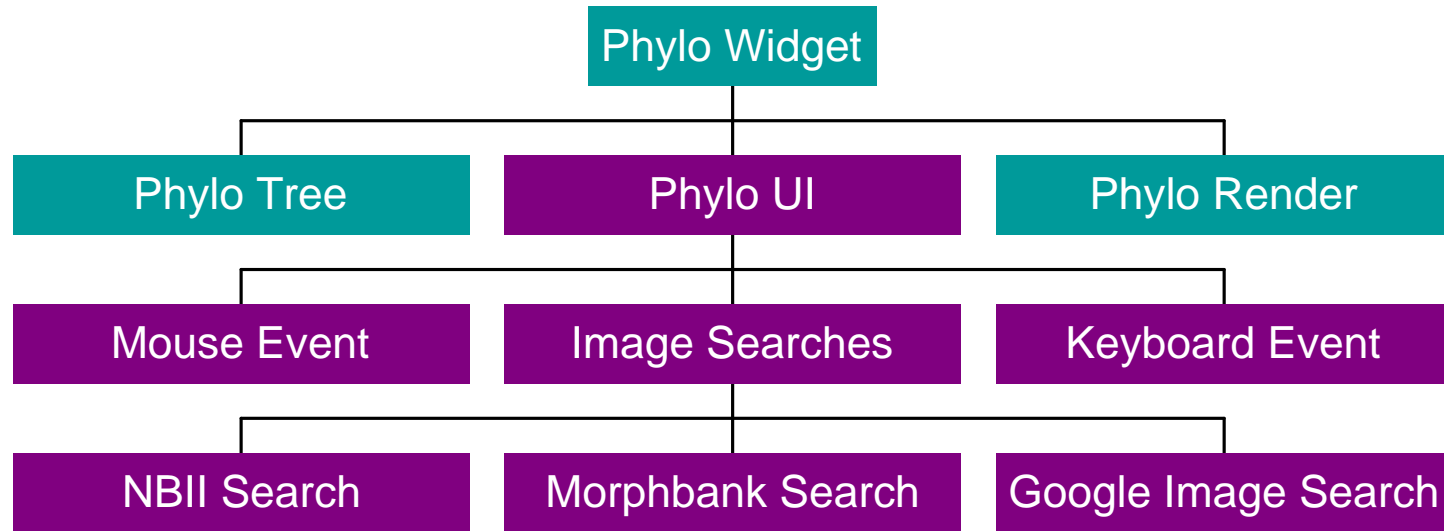
## *Overall Flowchart*





# Algorithms

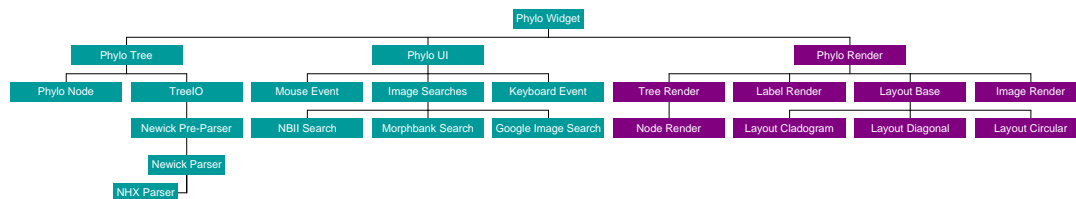
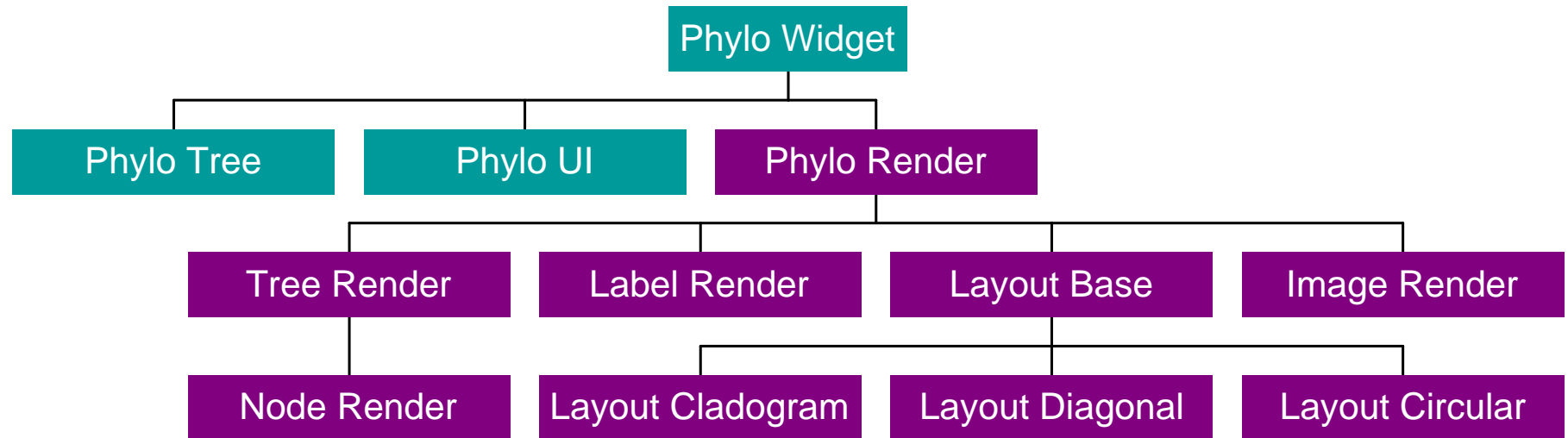
*Overall Flowchart*





# Algorithms

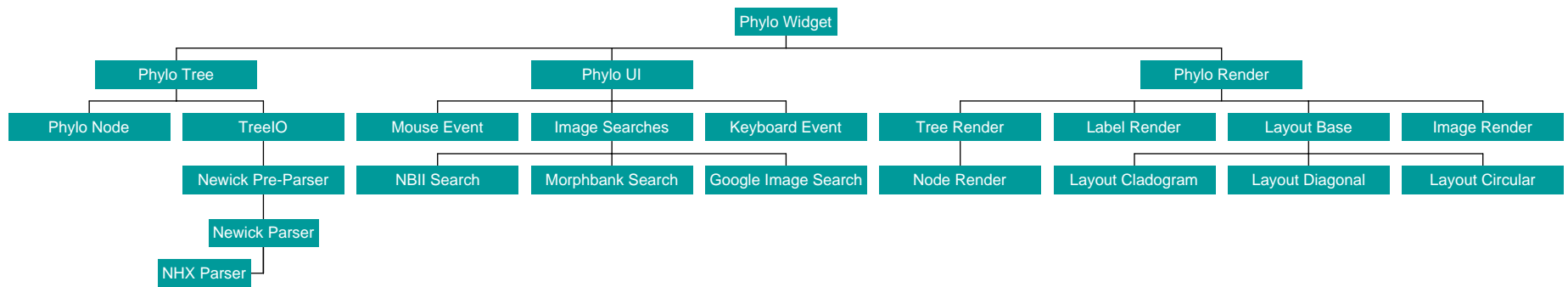
## Overall Flowchart





# Algorithms

## *Overall Flowchart*

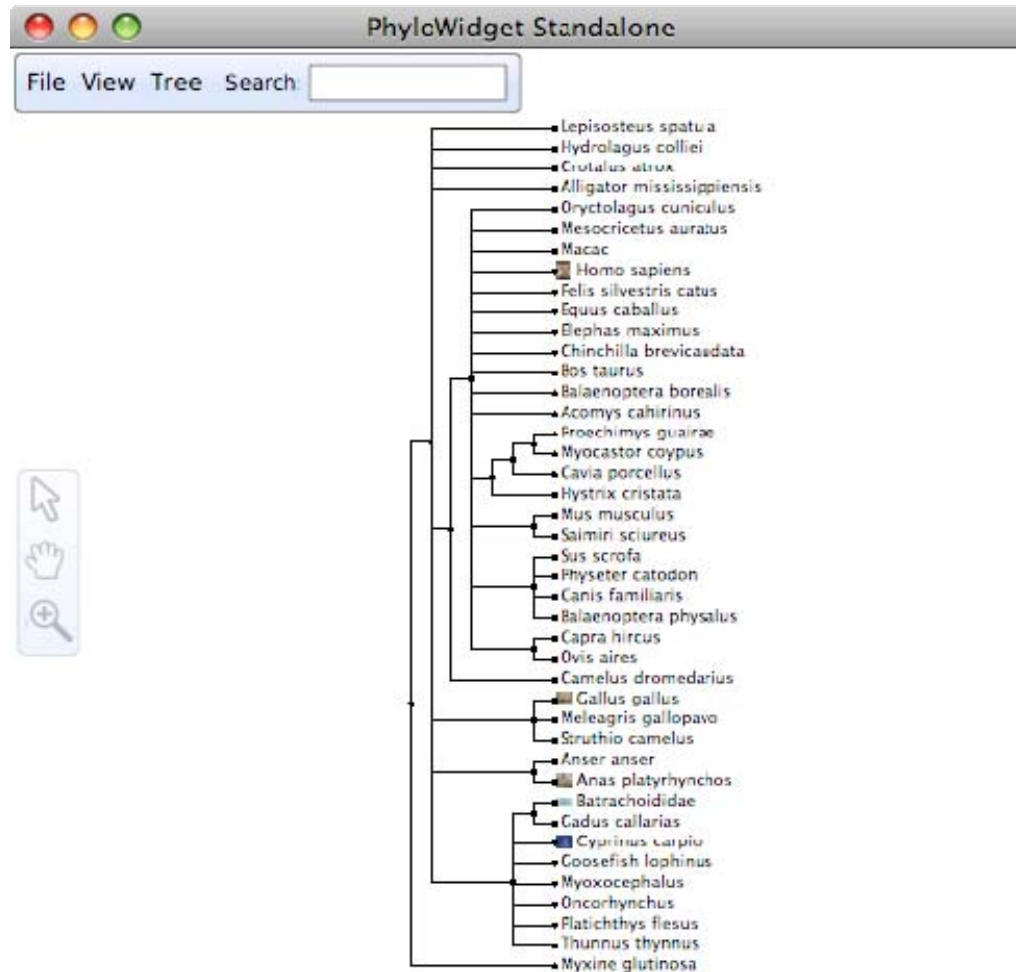




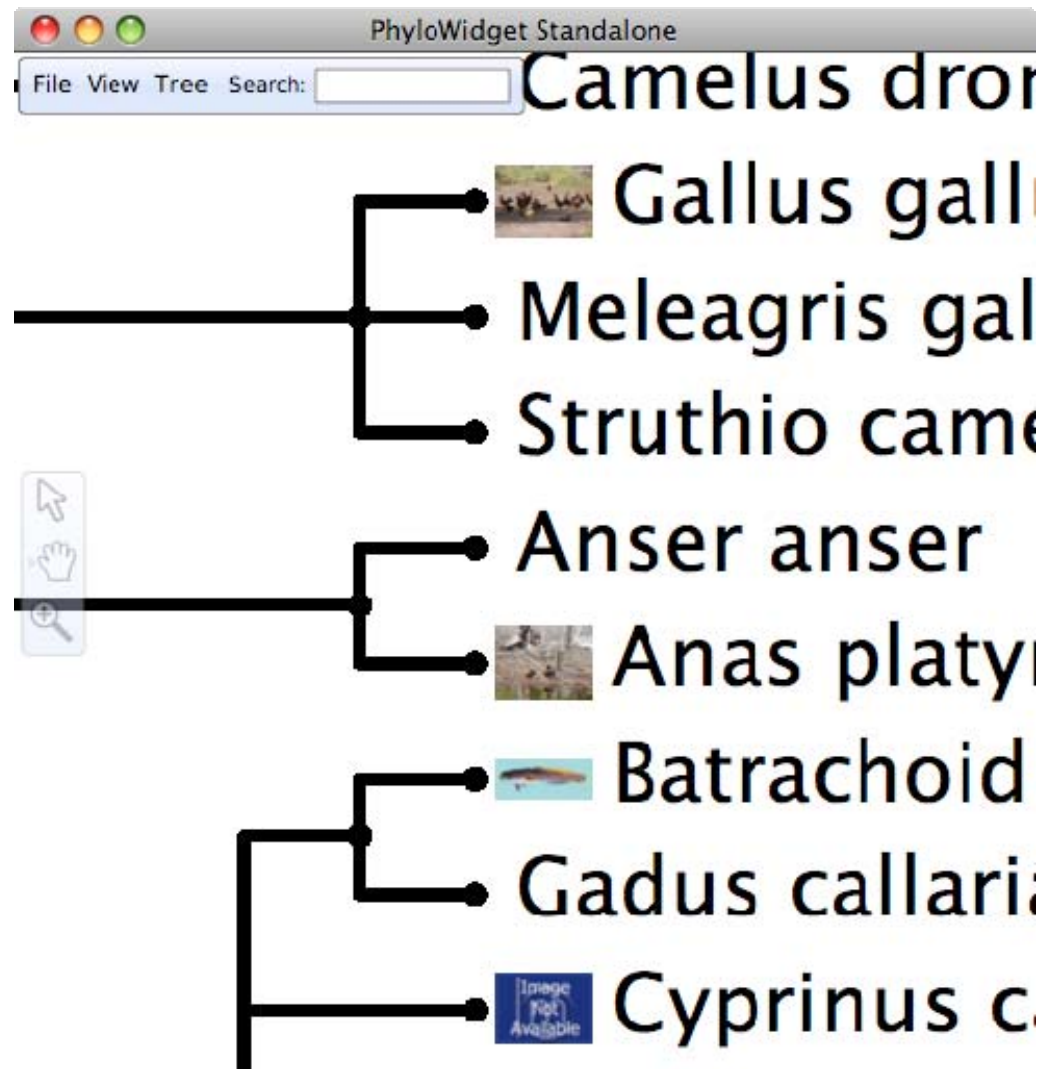
# Demo

- Still images
- Screen Capture
- Live Demo

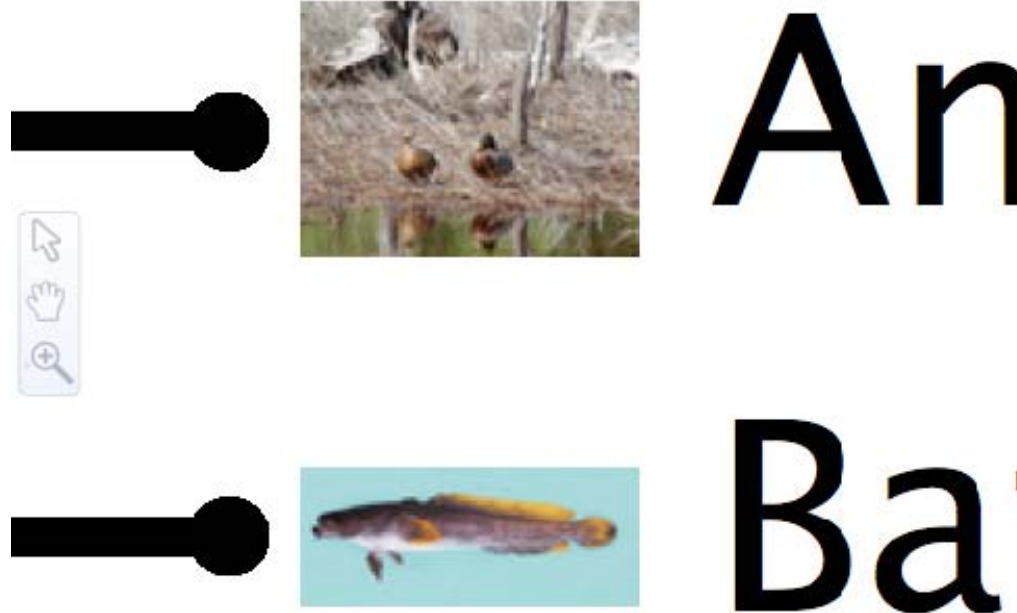
# Demo



# Demo



# Demo





# Demo

- Still images
- Screen Capture
- Live Demo



# Achievements

*Tangible Products*

- Newick pre-parser
- Morphbank search
- NBII search
- UBIO search and display





# Achievements

*Intangible Outcomes*

- Integrate existing tools in a form useful to students and researchers
- Understand and build upon another's work
- Participate in large-scale international project
- Lay the ground work for software with high potential in education
- Structure a long-term project into a manageable series of goals



# Future Work

*Looking for a Senior Project?*

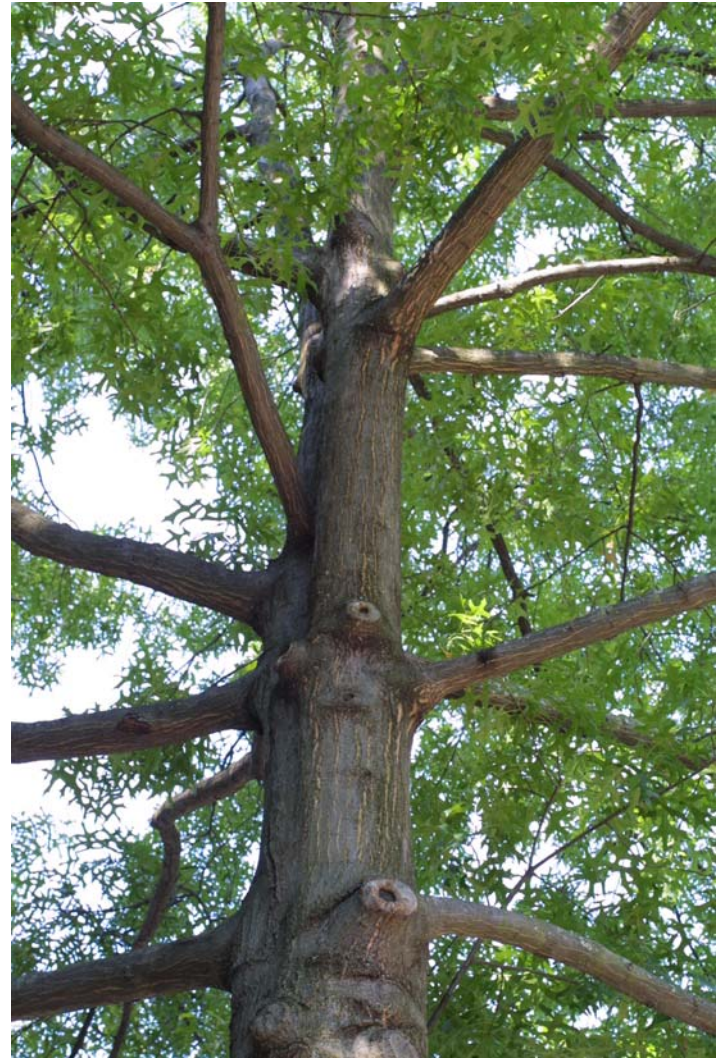
- Design a student-friendly interface
- Implement a TreeBASE II search
- Provide a sidebar with supplementary data and full quality images





# Tools & Languages

- Java
- Eclipse
- Subclipse (SVN Client)
- Processing (UI Library)



# Thank You

- Val
- Joe Kider
- Norm
- Amy
- Greg Jordan





Questions?

