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Interactive Visualization of Evolutionary Trees and Gene Sequences

February 2, 2006
UBC CS Discovery Forum

Computer Graphics

- create or manipulate images with computer
 - movies, games, photorealistic simulation



77 K polygons
24 aera lights
solution render time : around 7200 sec

Computer Graphics

- create or manipulate images with computer
 - movies, games, photorealistic simulation
 - but wait, there's more!

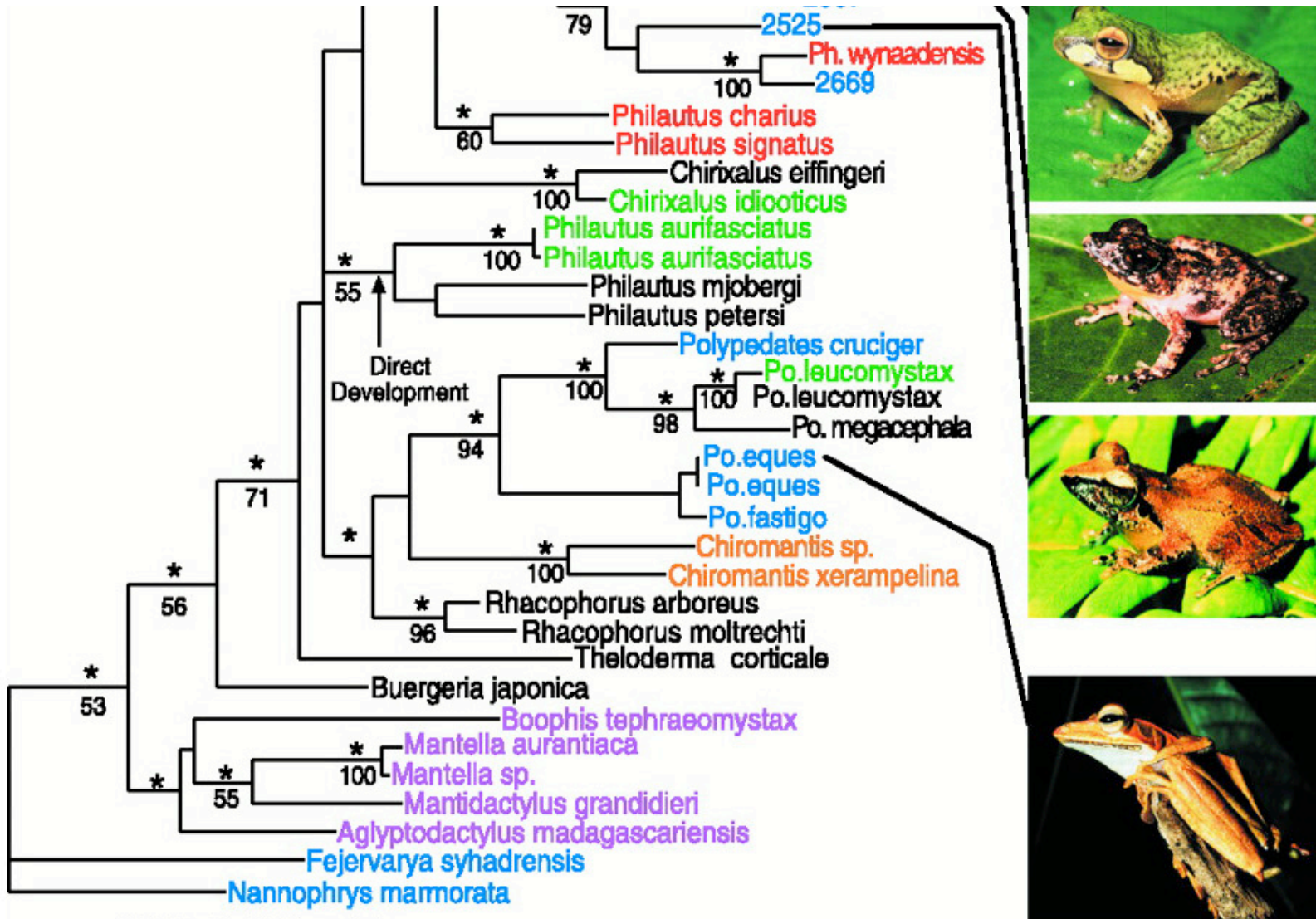


77 K polygons
24 aera lights
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Visualization

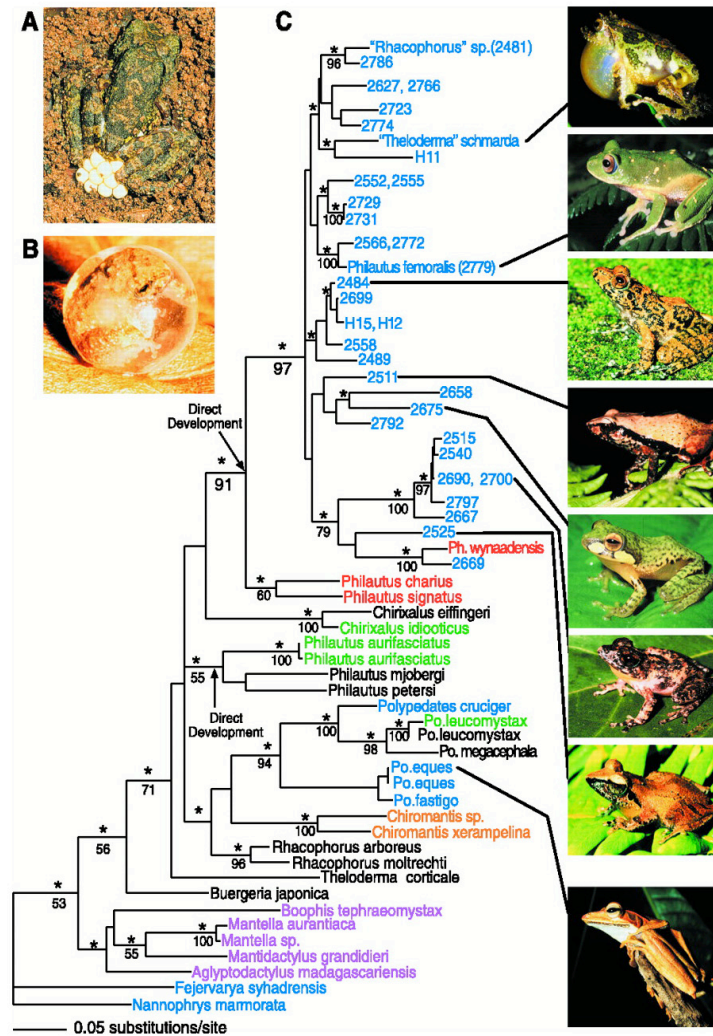
- using interactive computer graphics to help people understand information better
- biological data:
evolutionary trees and gene sequences

Evolutionary Tree



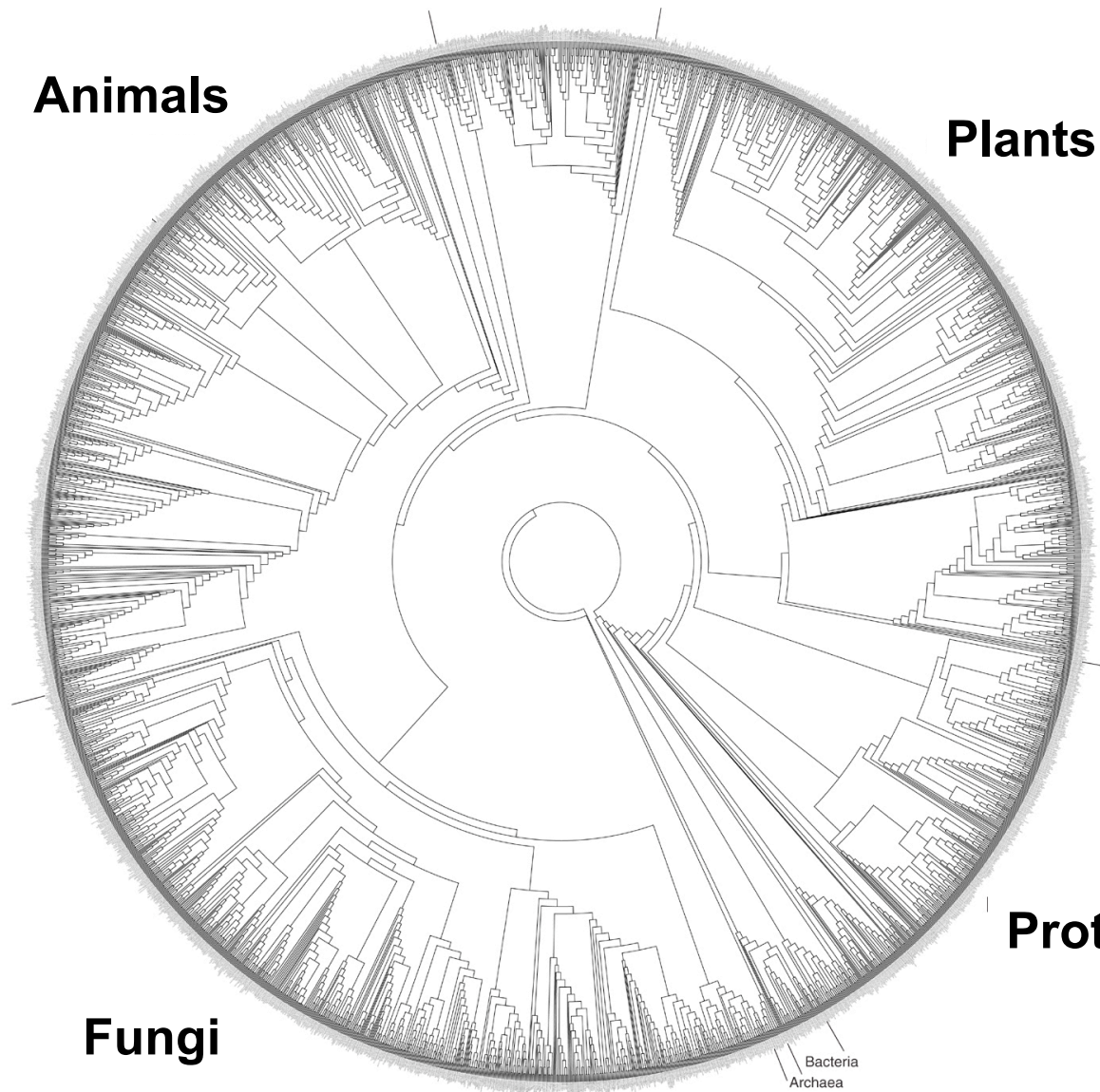
M Meegaskumbura et al., Science 298:379 (2002)

Common Dataset Size Today



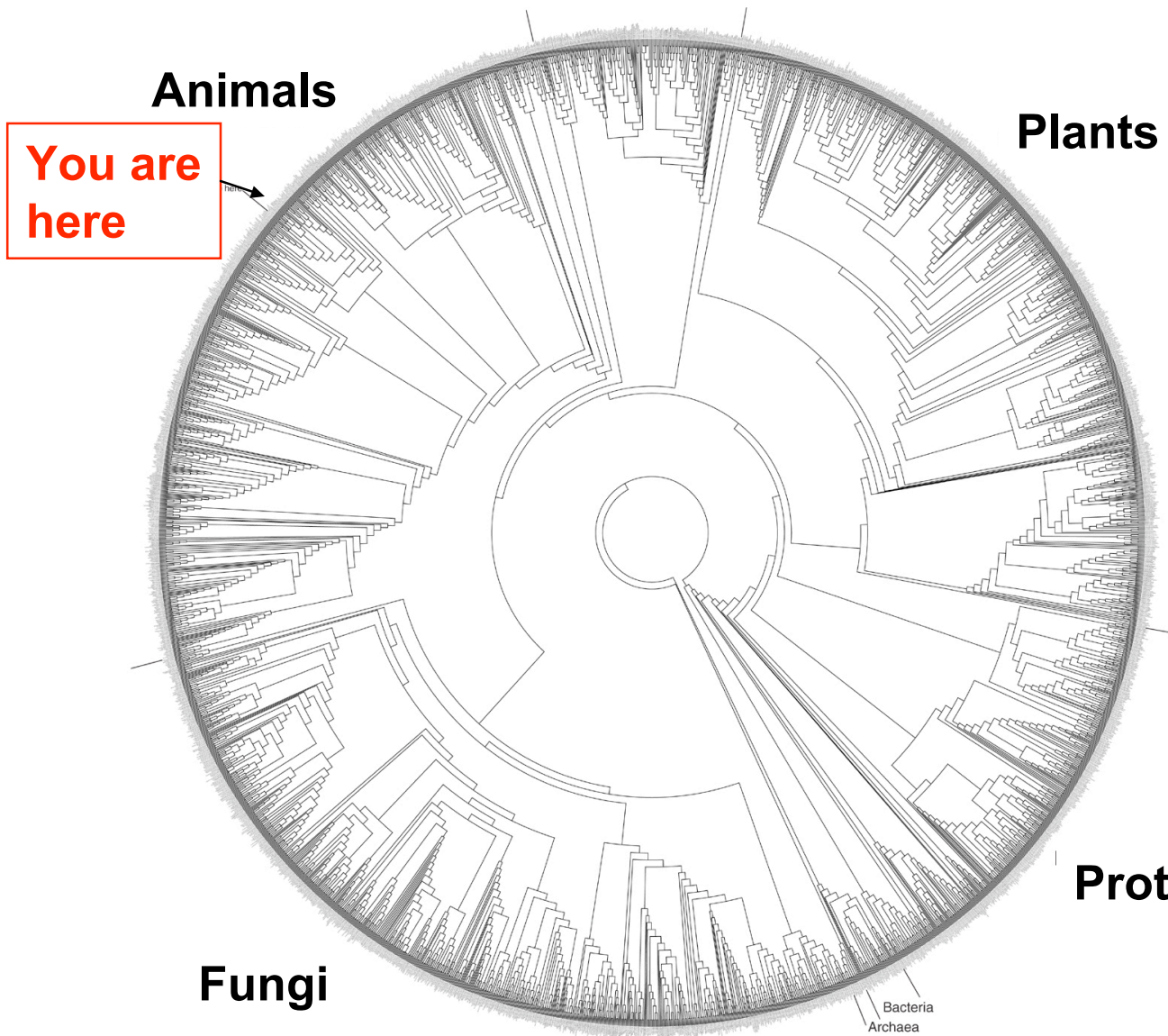
M Meegaskumbura et al., Science 298:379 (2002)

Future Goal: 10M Node Tree of Life



David Hillis, *Science* 300:1687 (2003)

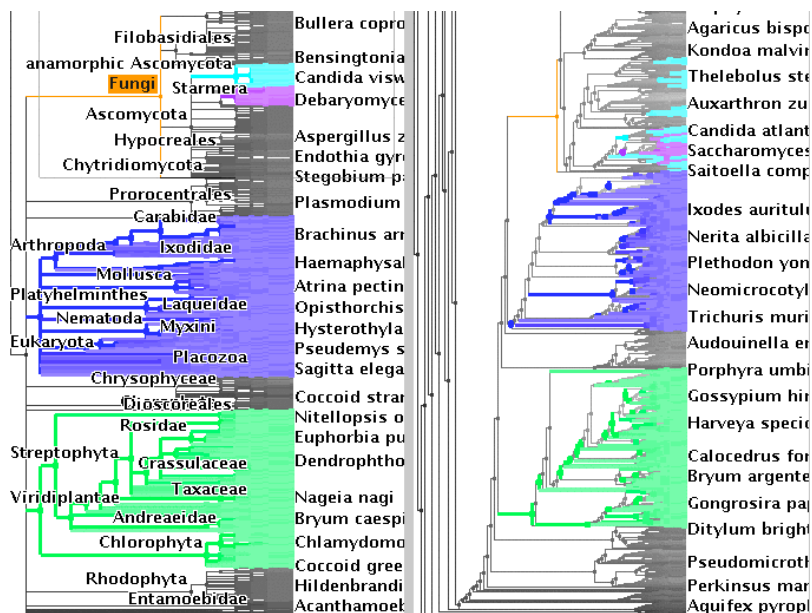
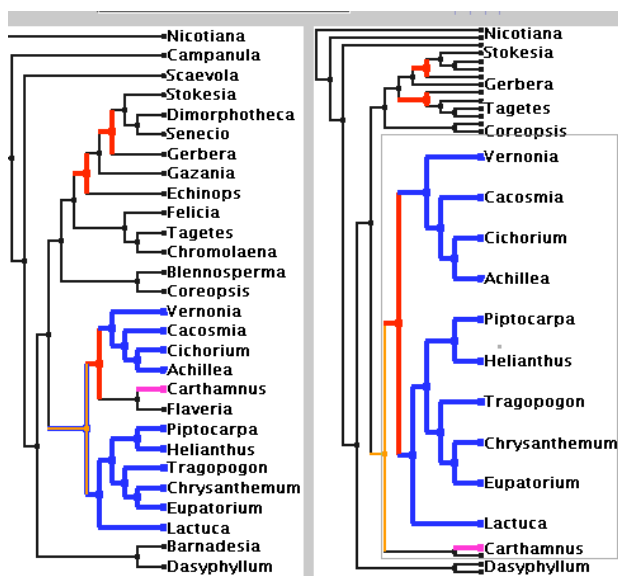
Future Goal: 10M Node Tree of Life



David Hillis, *Science* 300:1687 (2003)

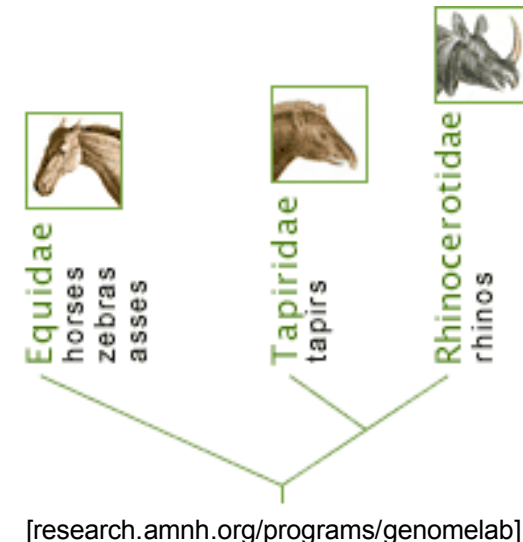
TreeJuxtaposer

- side by side comparison of evolutionary trees
 - both focus and context with stretchable surface
- demo - downloadable from <http://olduvai.sf.net/tj>

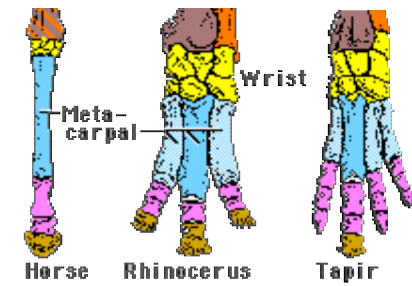


Reconstructing Trees from Genes

- know leaves, infer interior nodes
 - similarity: parallel evolution or common ancestor?



- old: morphology
 - observable similarities



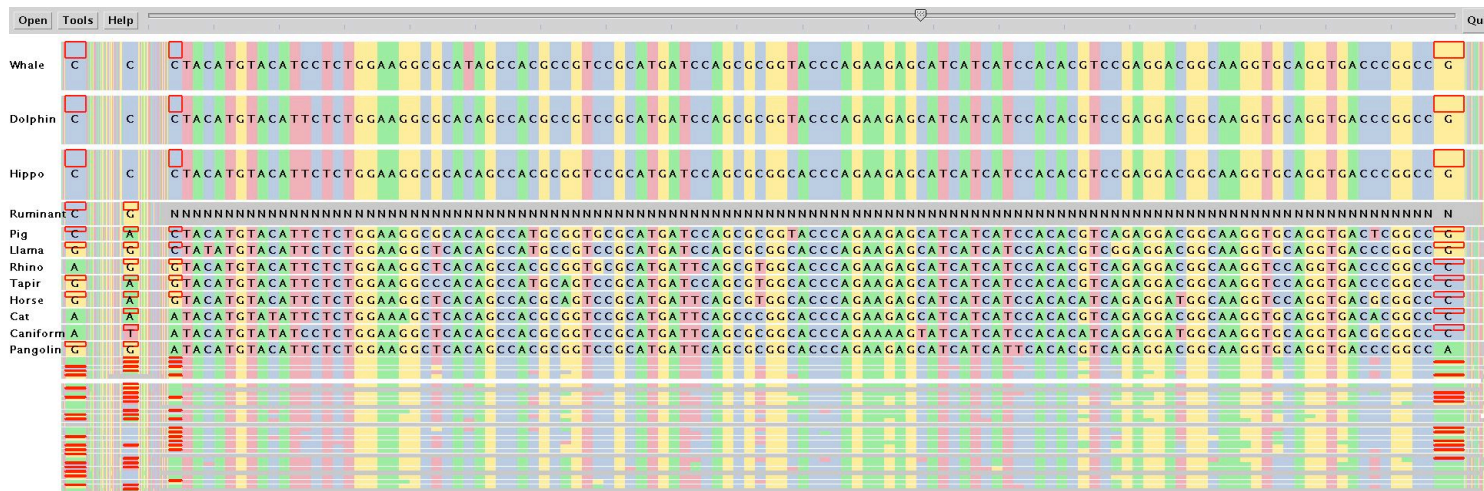
- new: molecular
 - DNA sequences - nucleotides
 - protein sequences - amino acids

horse: ...CCTGAACCG...
tapir: ...ACTCTACCG...
rhino: ...GCTCTACCG...

[gwis2.circ.gwu.edu/~atkins]

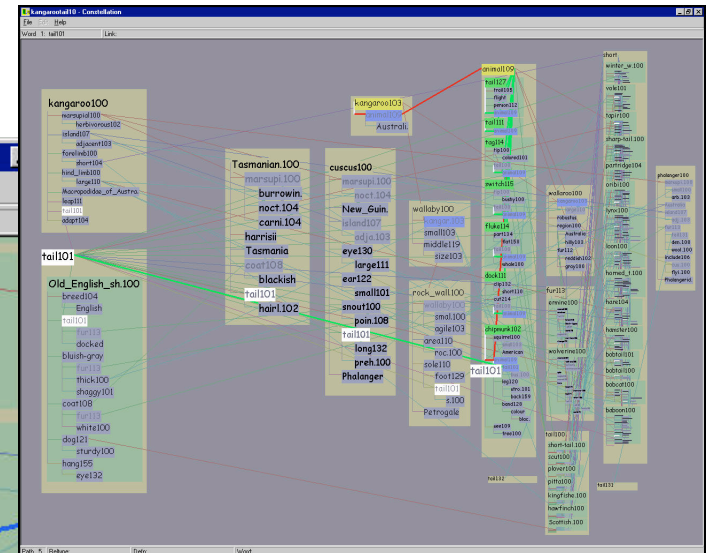
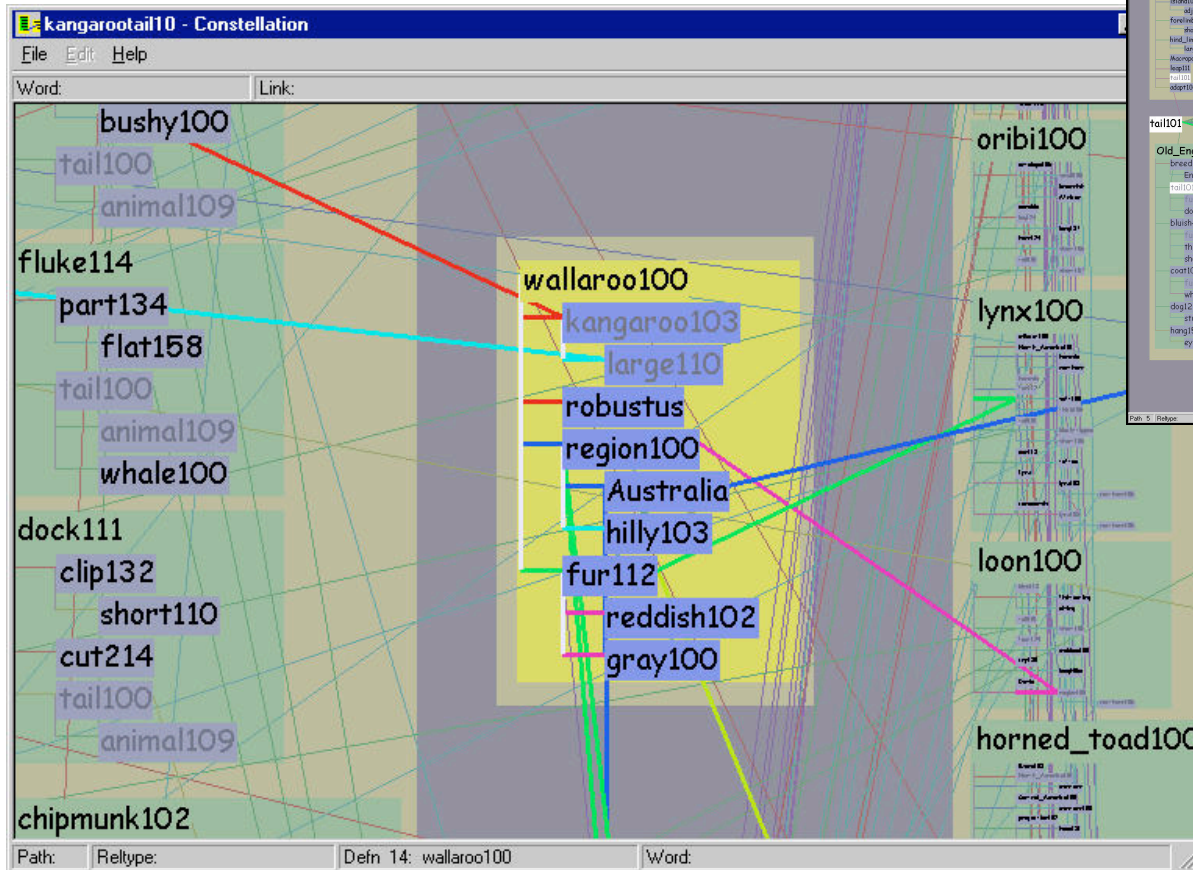
SequenceJuxtaposer

- comparison of aligned gene sequences
 - focus and context with stretchable surface
- demo - downloadable from <http://olduvai.sf.net/sj>



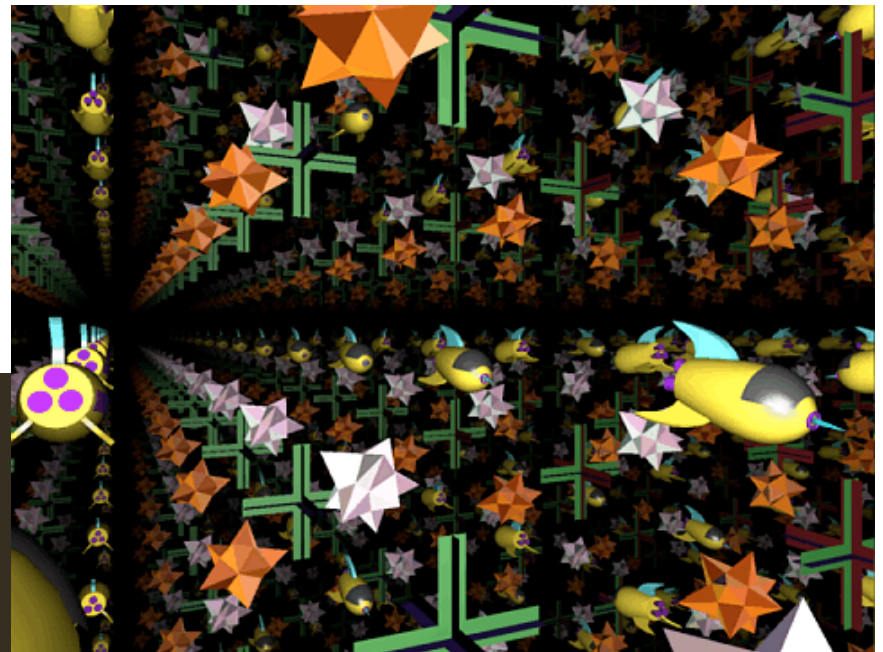
CS: Collaboration with Many Fields

- computational linguistics



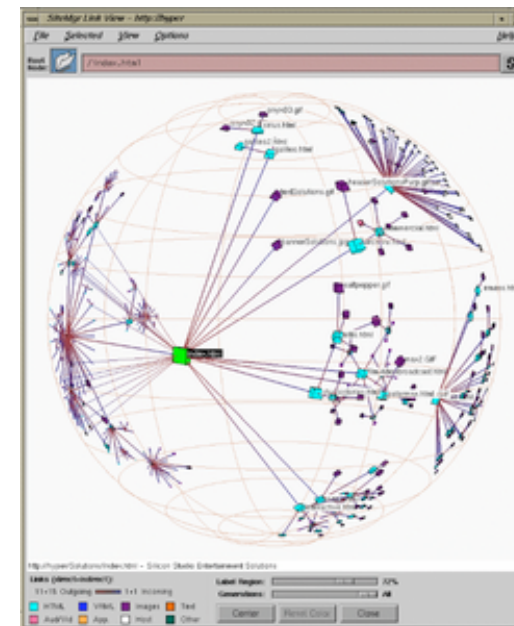
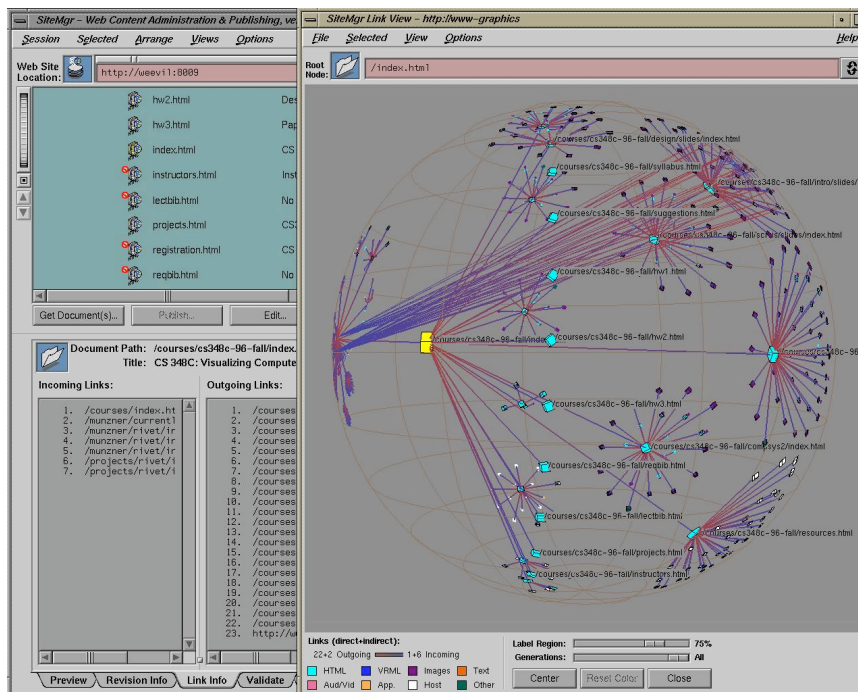
CS: Collaboration with Many Fields

- mathematics: topology and geometry



CS: Collaboration with Many Fields

- software for web site designers
 - now also useful for biologists!
 - downloadable from graphics.stanford.edu/~munzner/h3



More Information

- <http://www.cs.ubc.ca/~tmm>