

Graph Theory for Biologists

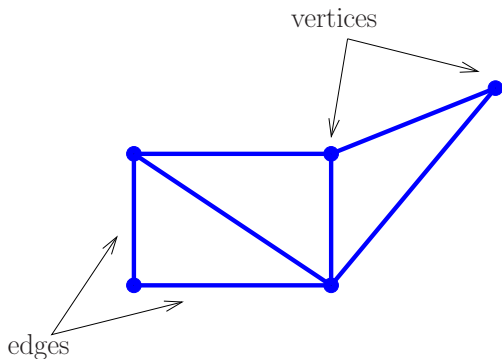
Andrew Treglown

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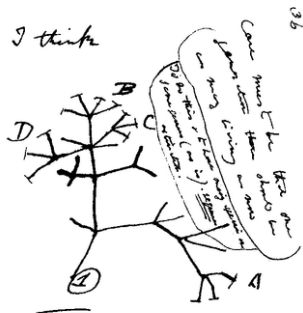
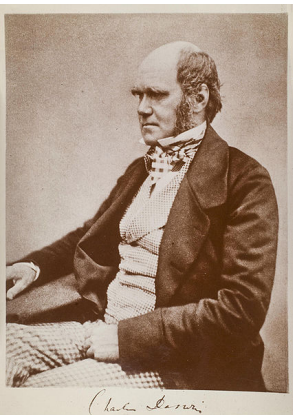
24th June 2010

What is a graph?

Graph = collection of points (**vertices**) joined together by lines (**edges**)

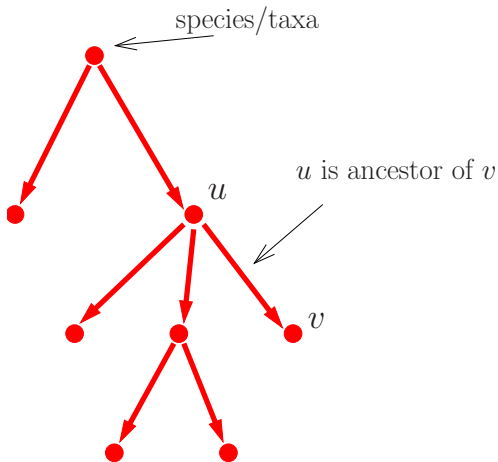


Darwin's tree of life



Then between A & B. various
sort of relation. C & B. The
first predation, B & D
rather greater distinction
than genera would be
formed. - binary relation

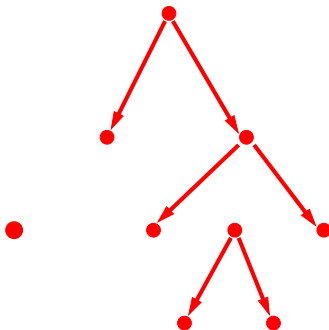
Phylogenetic trees



What kind of problems do mathematical biologists look at?

Graph reconstruction/Hereditary properties

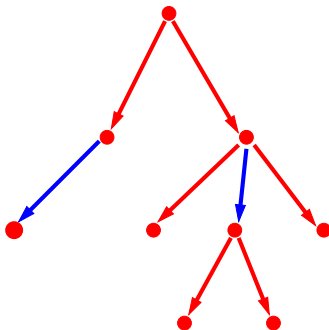
- Given **some** information about our tree can we obtain **more** information (or even work out what the entire tree looks like)



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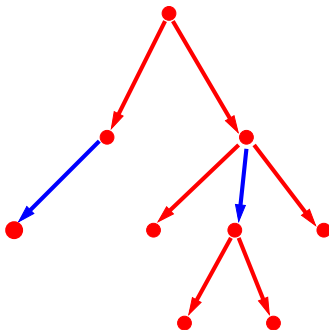
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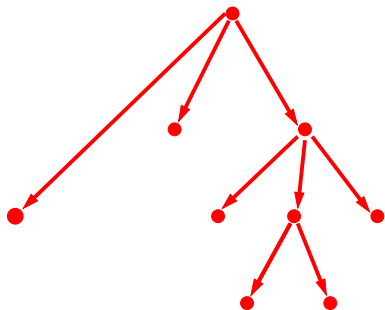
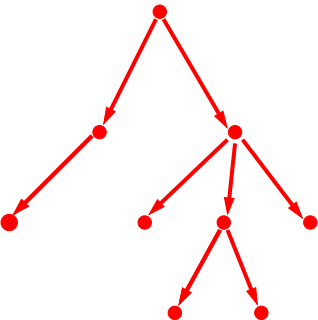
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- Stephen Willson (Iowa State) works on such problems

The tree space

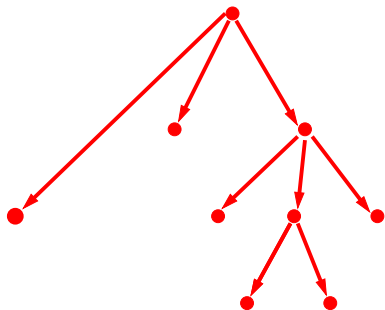
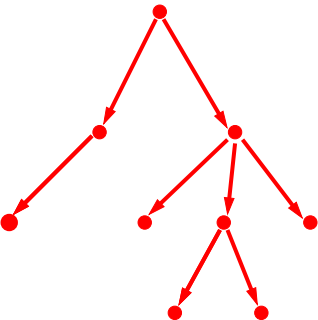
Given different pieces of information we may obtain different 'candidates' for our phylogenetic tree



- Can we compute the tree space efficiently?

The tree space

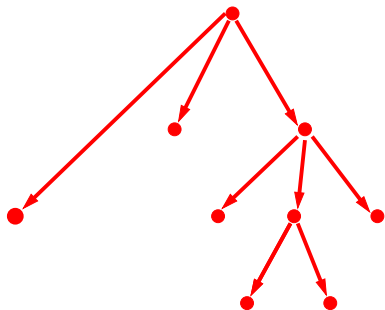
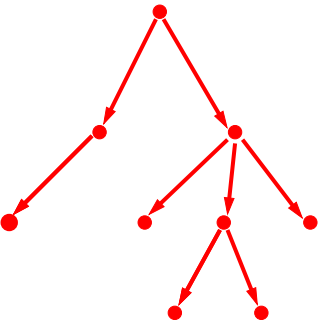
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- Can we find the optimal tree (“tree of best fit”)?

The tree space

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- Can we find the optimal tree (“tree of best fit”)?
- Katherine St. John (New York) works on such problems