

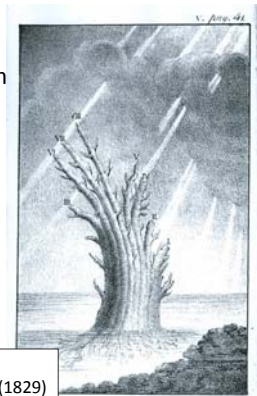
03-327/727 Phylogenetics

Dannie Durand

- Office hours: Tuesday 5pm – 6:30pm
- MI 646

Pieter Spealman

- TBA



Eichwald,
Zoologia specialis (1829)

From "Trees and networks before and after Darwin", Mark A Ragan Biology Direct, 2009

03-327/727 Phylogenetics

Course website:

<http://www.cs.cmu.edu/~durand/Phylogenetics>

Password required to access some material
when logging in from off campus

id: compbio

password: genomics

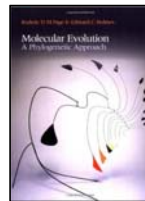
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Textbook

Textbook: *Molecular Evolution: A phylogenetic approach*, Page and Holmes

This book is available, free, in pdf
format on the course website.

Additional readings from journal
articles and scanned textbooks
will be provided on the
syllabus page.



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03-327/727 Phylogenetics: Course work, homework, grading

- 6-8 homework assignments
- 4 data analysis practica
- Two in-class exams
- Final exam
- Project for 727

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03-327/727 Phylogenetics: Course work, homework, grading

- 03-327:
 - Homework: 25 pts
 - Practica: 30 pts
 - In class exams: 15 pts
 - Final: 30 pts
 - Total: 100 pts**
- 03-727:
 - Homework: 25 pts
 - Practica: 30 pts
 - In class exams: 15 pts
 - Final: 30 pts
 - Project: 25 pts
 - Total: 125 pts**

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03-727 Project: Phylogenetic analysis of a molecular data set

Milestones

- Formulate a phylogenetic question involving a sequence family of interest
- Collect appropriate sequences
- Multiple sequence alignment
- Phylogeny estimation
- Assess the reliability of your tree
- Interpret results

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03-727 Project: Phylogenetic analysis of a molecular data set

Deliverables

- Project plan
- Final paper
- In-class presentation
- 3-5 pages (plus appendices)
 - Introduction
 - Methods
 - Results/Discussion
 - Appendix including alignments, trees, etc.

Project can be carried out by a 2-person team

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Topics

- Introduction:
 - a phylogeny is an evolutionary tree that represents a hypothesis concerning evolutionary history
- The underlying evolutionary processes
 - Species evolution
 - Gene and genome evolution
 - Gene trees versus species trees
- How to interpret an evolutionary tree
 - Tree terminology
 - Evolution of traits on a tree
- Molecular phylogenetics: Building evolutionary trees using sequences
 - Finding and aligning sequences
 - Models of sequence evolution
 - Tree reconstruction methods
 - Sources of error, assessing accuracy

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What is a phylogeny?

Phylogeny:

An evolutionary tree that represents a hypothesis concerning the evolutionary history of a group of taxa and their ancestors.

Taxon:

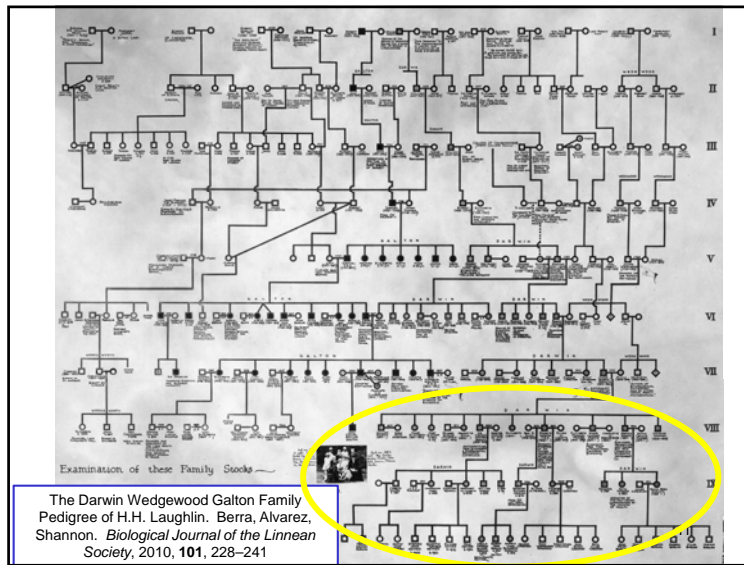
- a unit of classification
- strain, species, individual, gene

A phylogeny is (sort of) like a pedigree or family tree....

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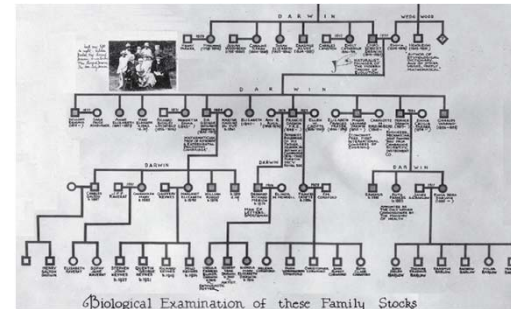
Two motivating examples

- Darwin's family and Darwin's finches
 - Similarities and differences between family trees and evolutionary trees
- Color vision in Primates
 - Why is it useful to think about the evolutionary history of genes independent of the evolutionary history of species?



A pedigree...

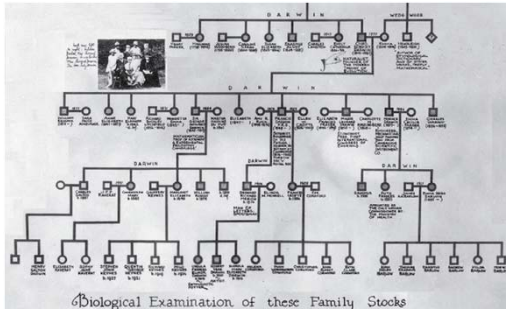
- represents a branching process where
- an individual gives rise to zero, one, or more than one offspring



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Family relationships

- Parent versus child, ancestor versus descendant
- Siblings versus cousins

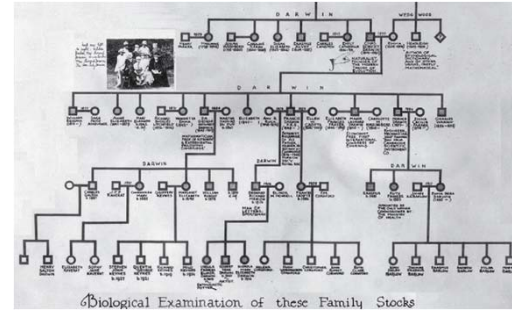


- You are more closely related to your siblings than to your cousins. Why?

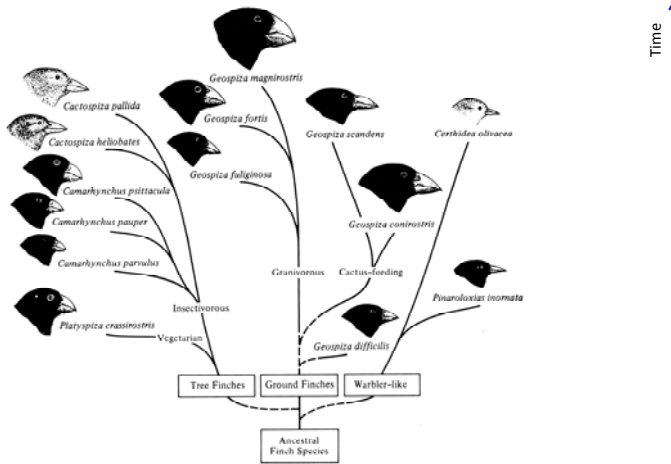
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Some temporal relationships

- A pedigree tells us that your parent was born before you were born, but not who was born first, you or your cousin

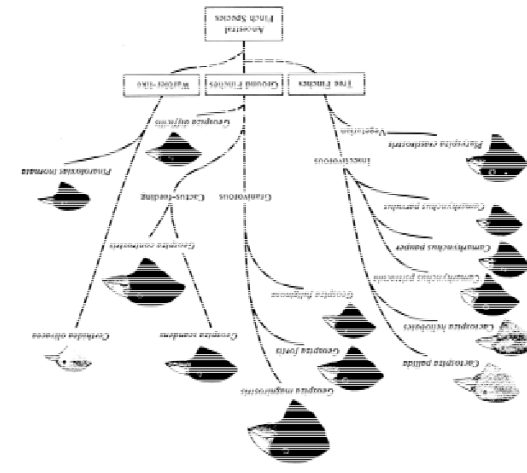


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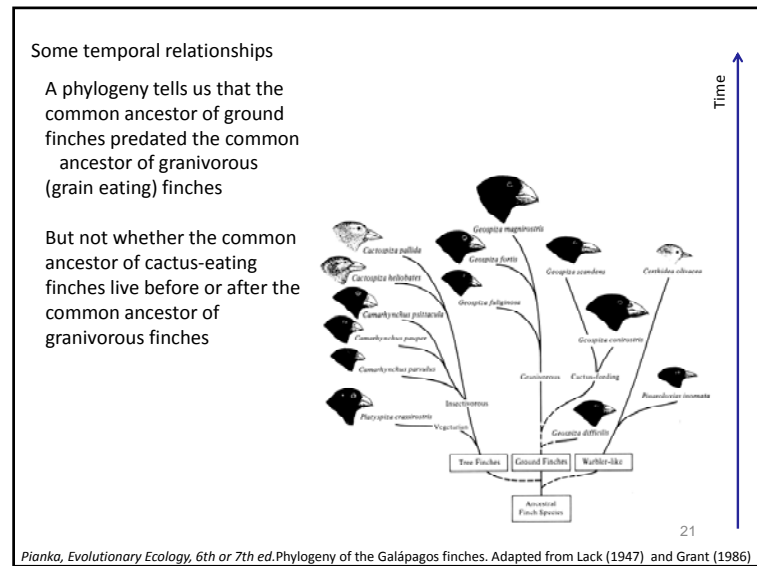
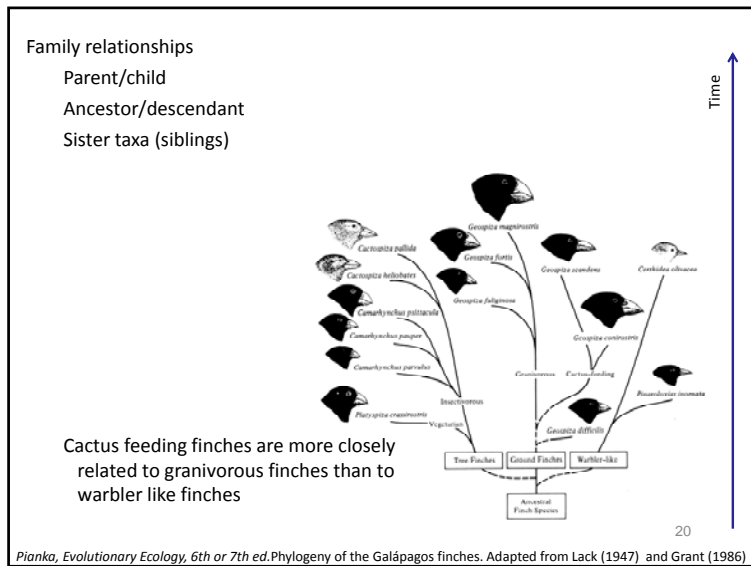
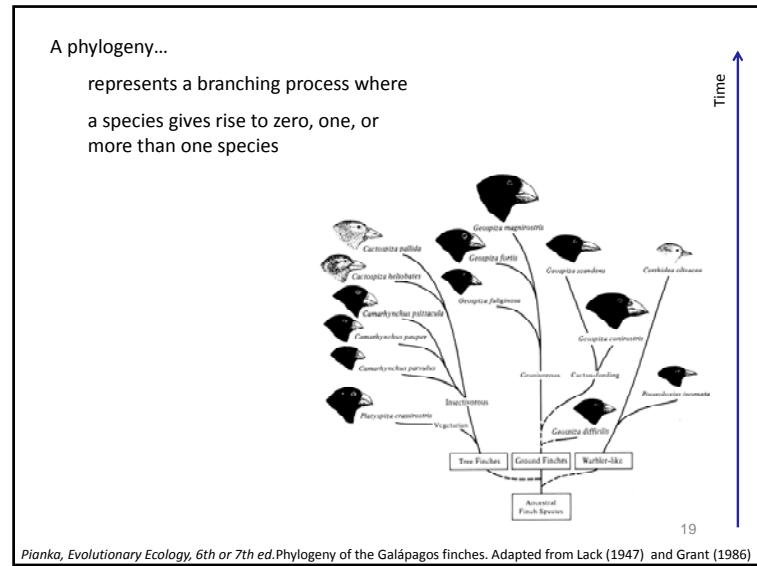
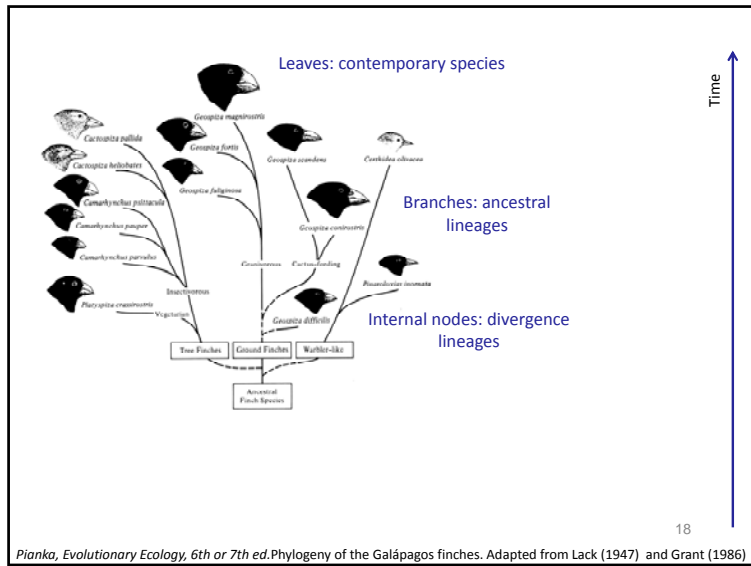
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Pianka, Evolutionary Ecology, 6th or 7th ed. Phylogeny of the Galápagos finches. Adapted from Lack (1947) and Grant (1986)



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Differences between family trees and phylogenies

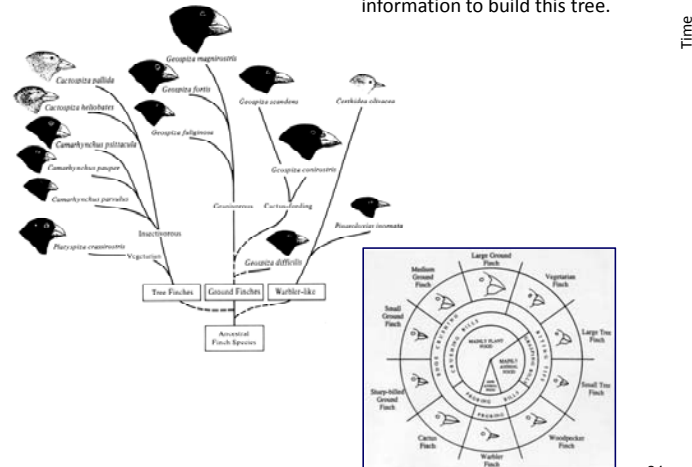
Family trees

- Each node corresponds to a single individual
- The history is derived from the historical record
- Every node has two parents
- Not binary: A couple can have many children

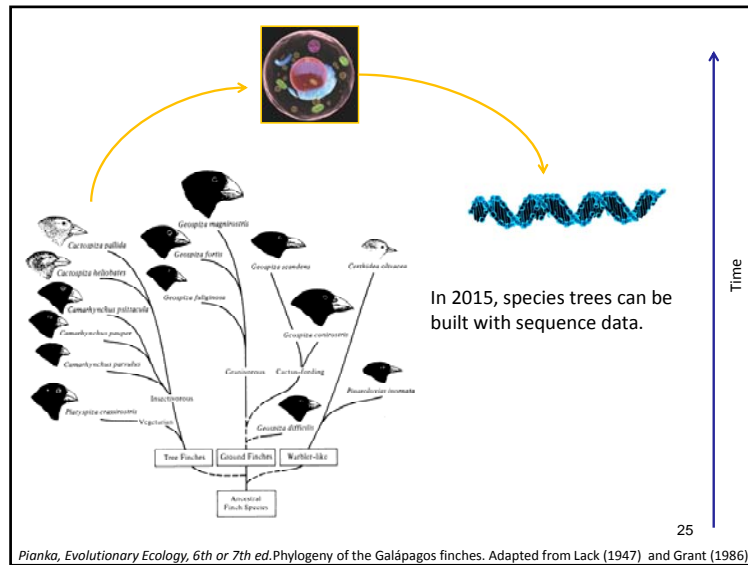
Species trees

- Each node corresponds to a species
- The history is inferred from information about present-day species
- Each node has one “parent”
- Binary: every node has two children, except for leaf nodes

In 1945, Lack used morphological information to build this tree.



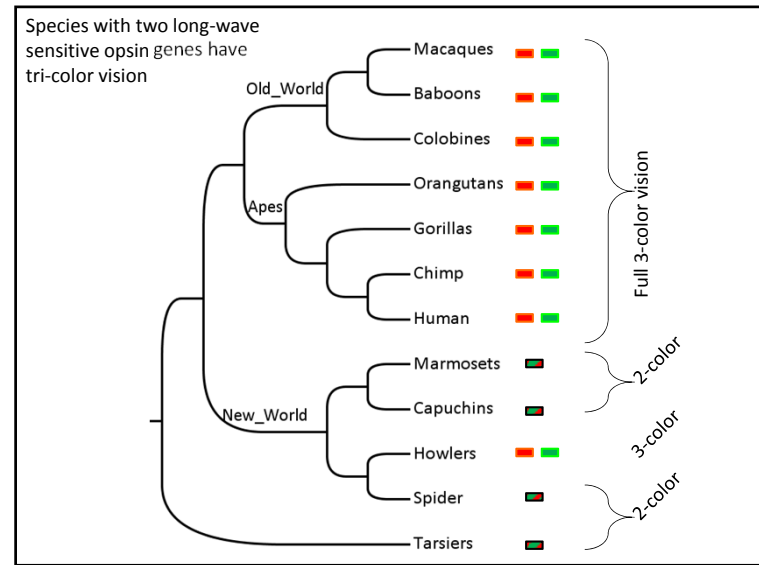
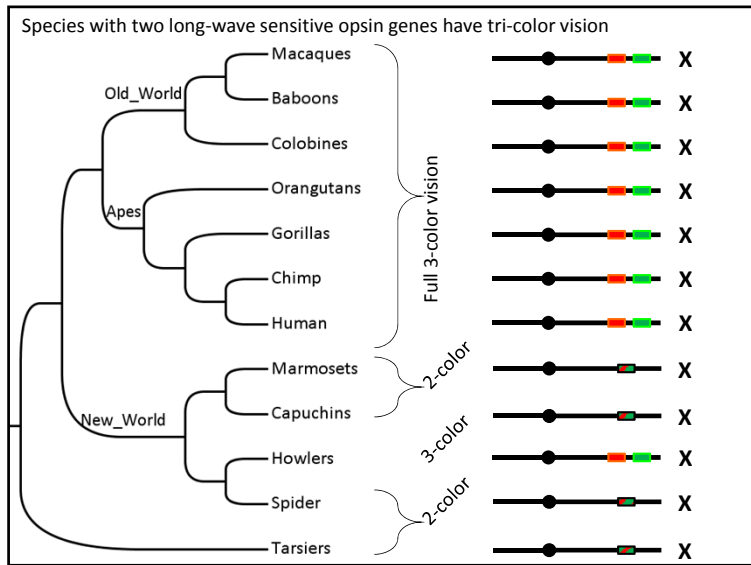
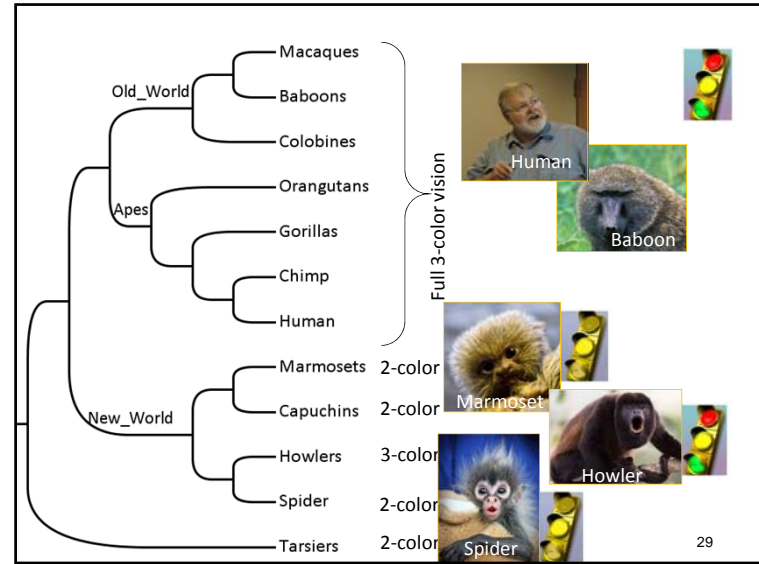
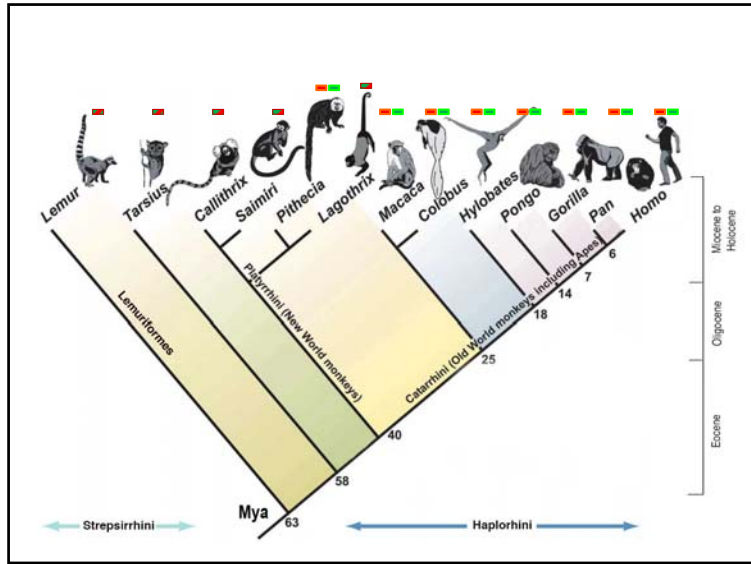
Pianka, *Evolutionary Ecology*, 6th or 7th ed. Phylogeny of the Galápagos finches. Adapted from Lack (1947) and Grant (1986)

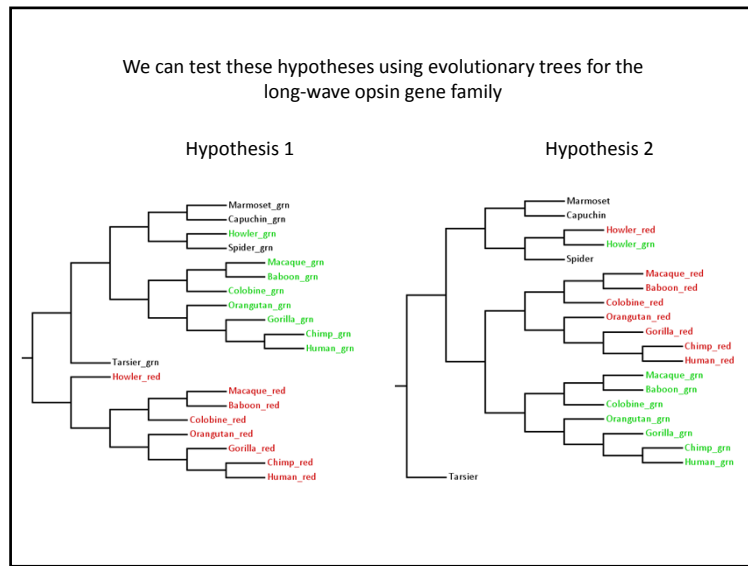
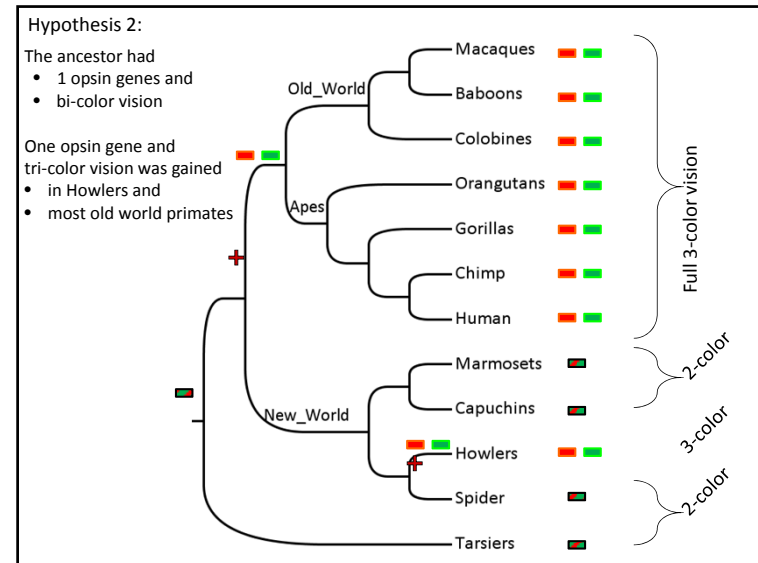
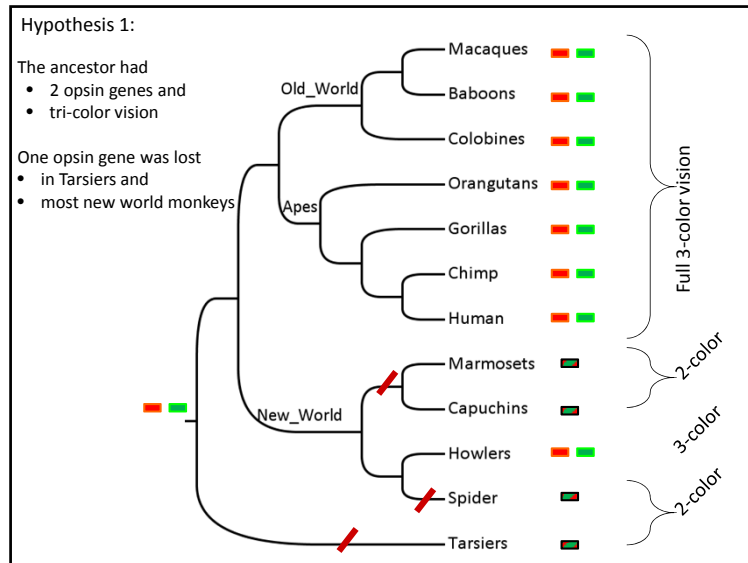


Pianka, *Evolutionary Ecology*, 6th or 7th ed. Phylogeny of the Galápagos finches. Adapted from Lack (1947) and Grant (1986)

Two motivating examples

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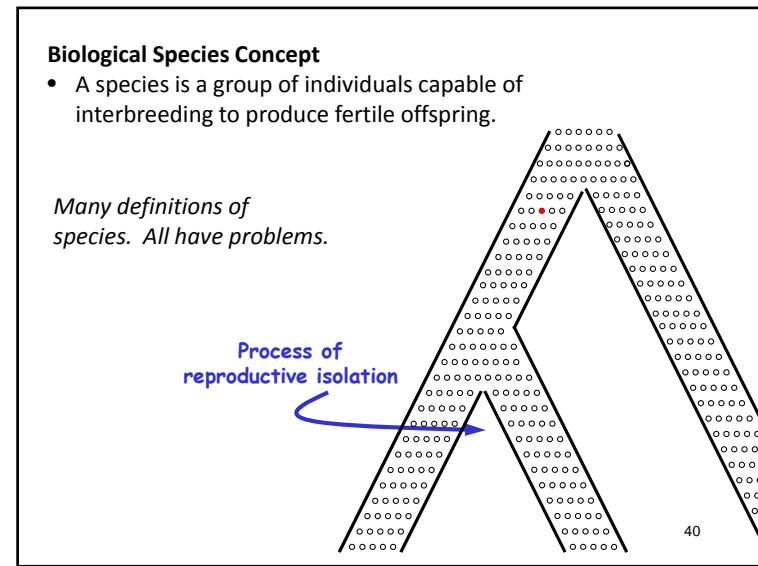
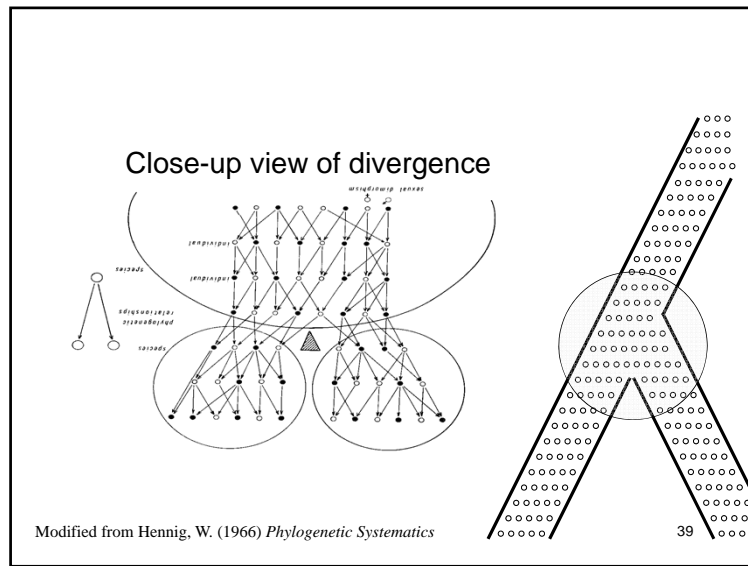
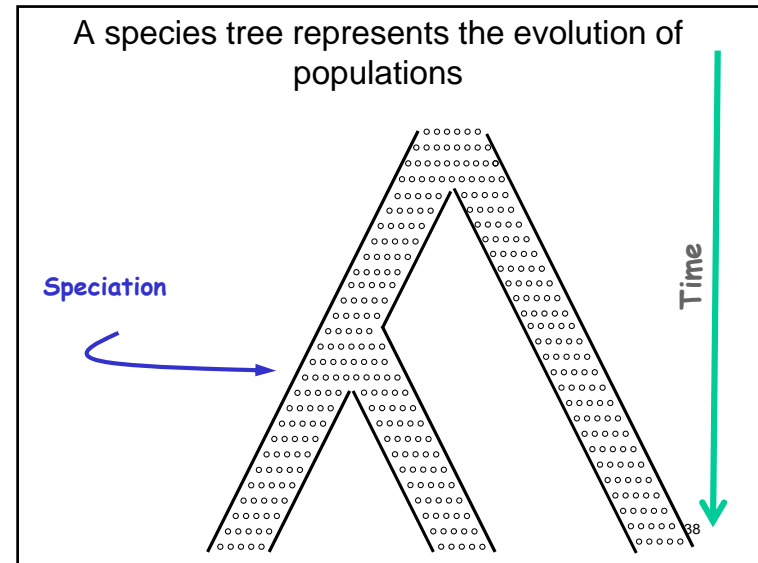
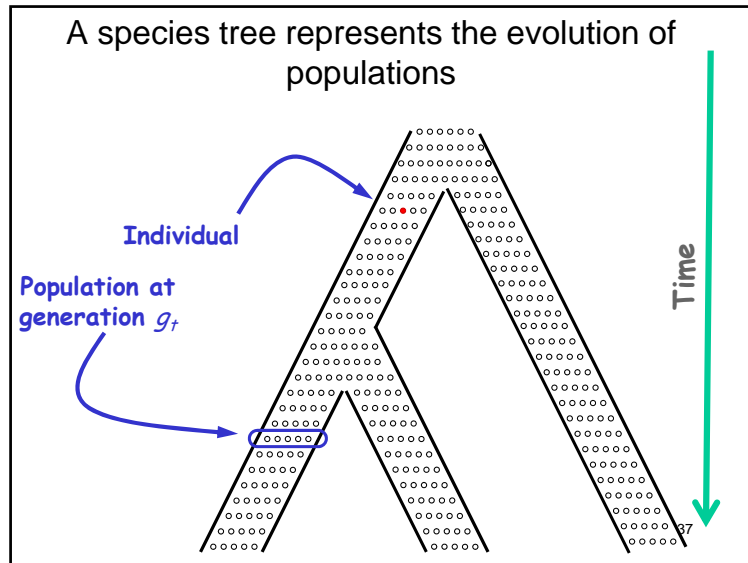




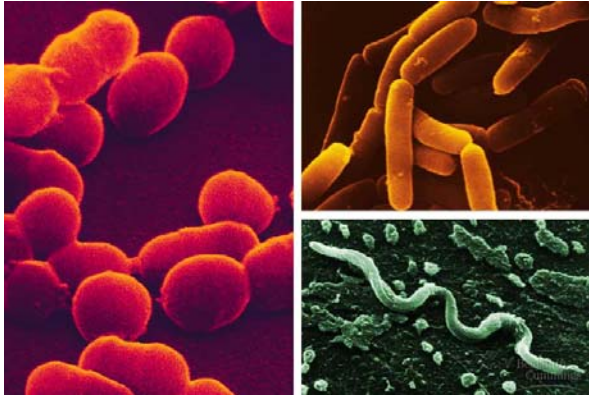
Evolutionary tree reconstruction

What are the processes we are trying to reconstruct?

- Species evolution
- Sequence evolution
- Gene family evolution



One Problem in the Biological Species Concept



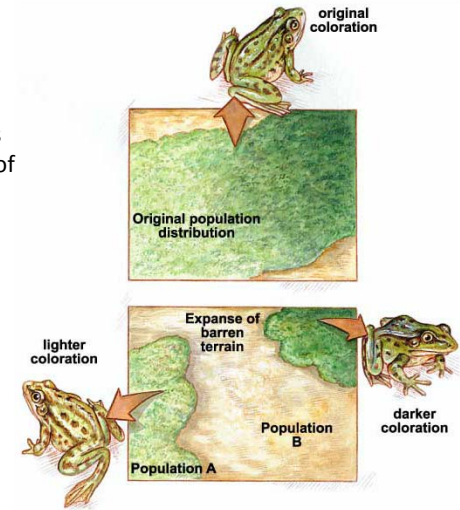
For asexually-reproducing organisms, like these bacteria, what constitutes a species?

How Do Species Arise?

The key to speciation is reproductive isolation of populations.

There are extrinsic and intrinsic reproductive isolating mechanisms.

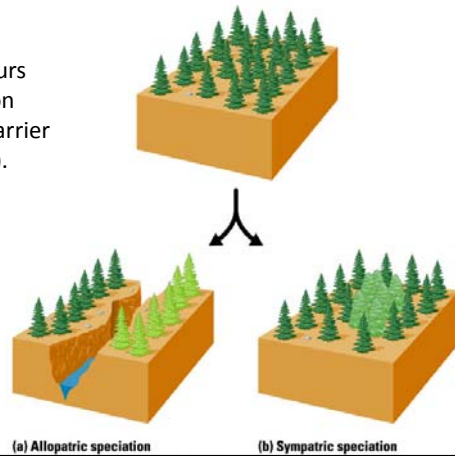
Geographic isolation is the primary extrinsic reproductive isolating mechanism.



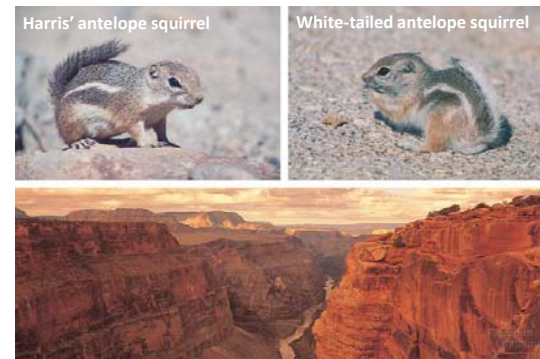
Reproductive Isolation May Occur With or Without Geographic Isolation

Allopatric speciation occurs when geographic isolation creates a reproductive barrier (an extrinsic mechanism).

Sympatric speciation occurs when a reproductive barrier is created by something other than geographic isolation (intrinsic mechanisms).



Allopatric Speciation



Two species of ground squirrel are postulated to have descended from a common ancestral population that was separated by formation of the Grand Canyon.

Many Intrinsic Reproductive Isolating Mechanisms Drive Speciation

Ecological Isolation:



Individuals from separate species do not mate because they do not come into contact (e.g., different habitats within an overlapping range).

Temporal isolation:



Individuals from separate species do not mate because they breed at different times.

Many Intrinsic Reproductive Isolating Mechanisms Drive Speciation

Behavioral isolation:



Individuals from separate species do not mate because they are not attracted to one another.

Courtship rituals are

- critical for mating within a species
- ineffective for attracting members of other species.



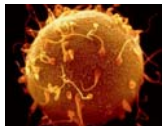
Many Intrinsic Reproductive Isolating Mechanisms Drive Speciation

Mechanical isolation:



Individuals from separate species do not mate because they are not physically compatible.

Gametic isolation:



Even if they are physically compatible, an embryo will not form if the egg and sperm do not fuse properly.

Many Intrinsic Reproductive Isolating Mechanisms Drive Speciation

Hybrid inviability or infertility:

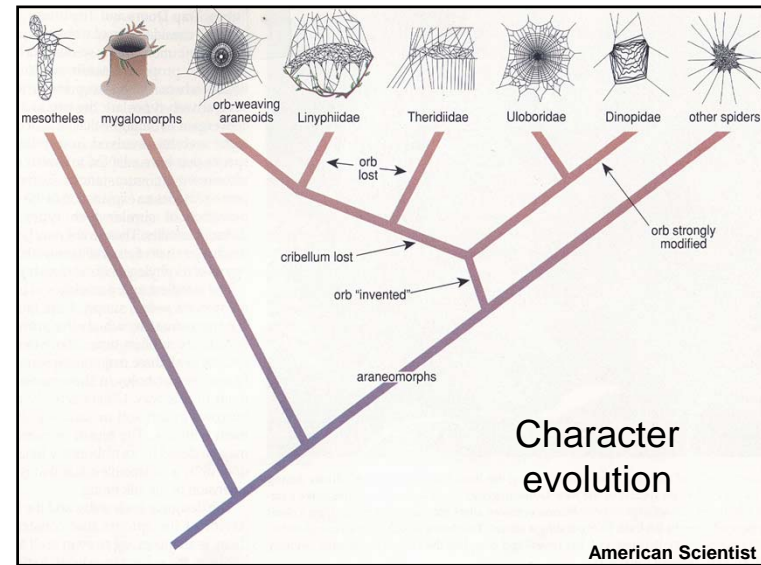
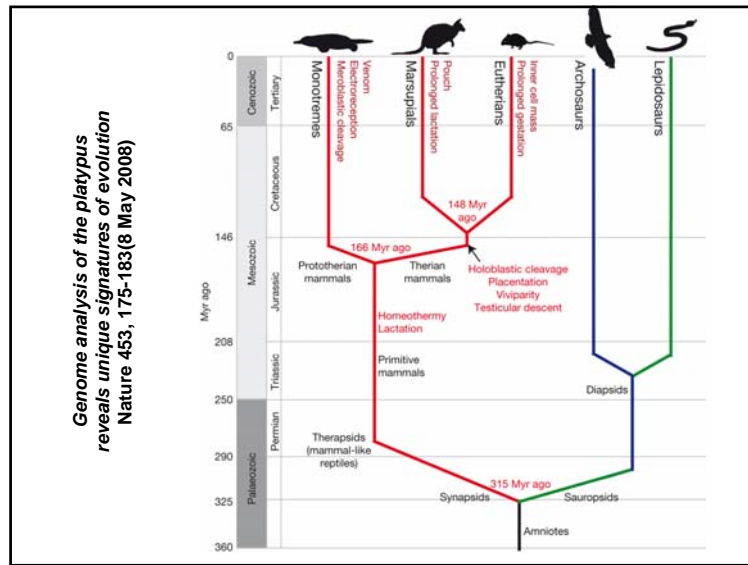
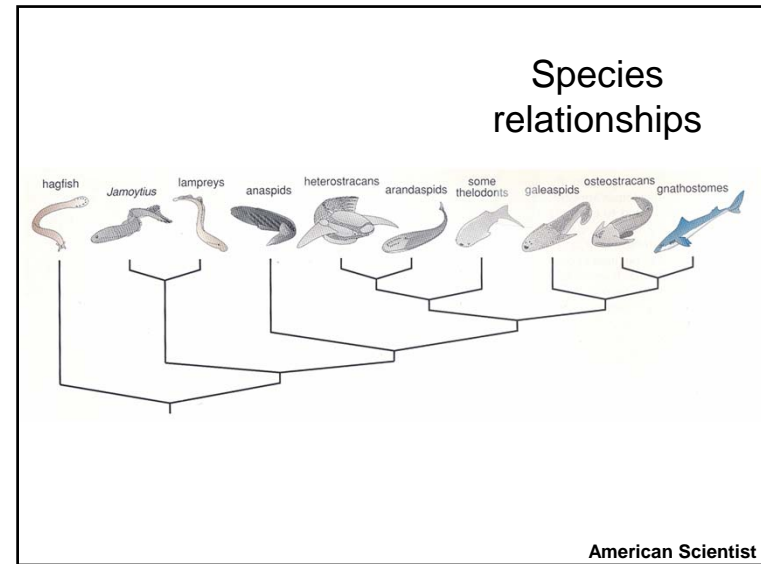


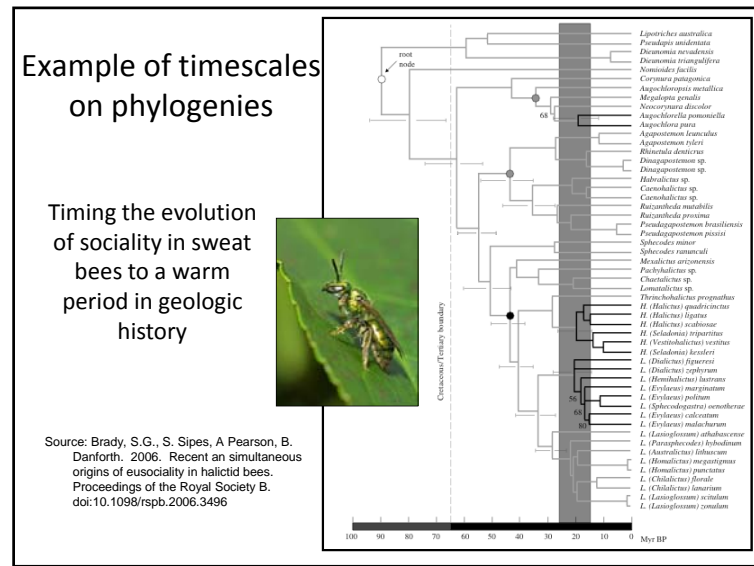
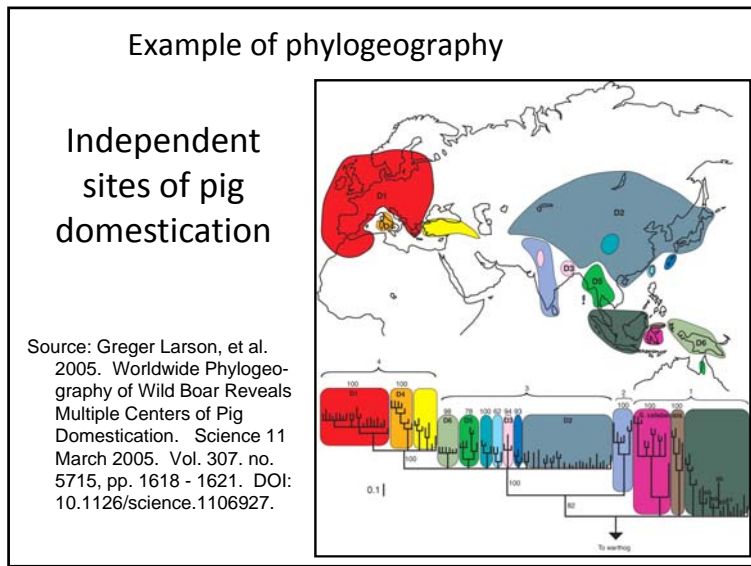
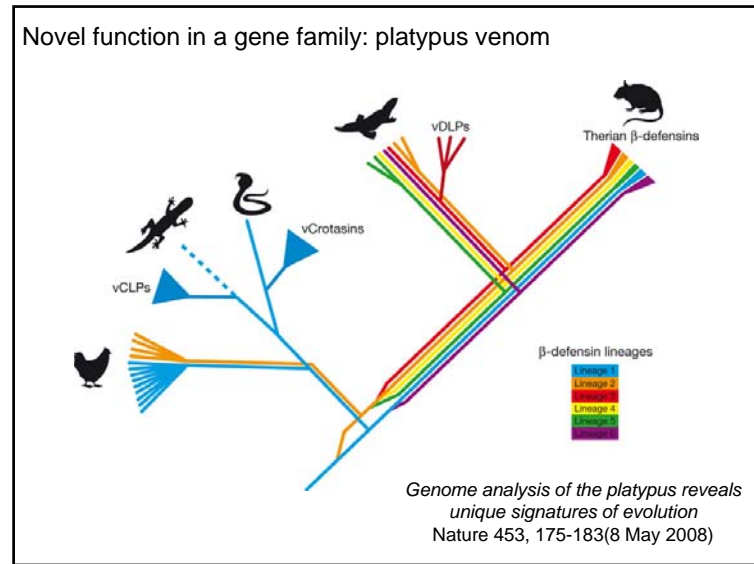
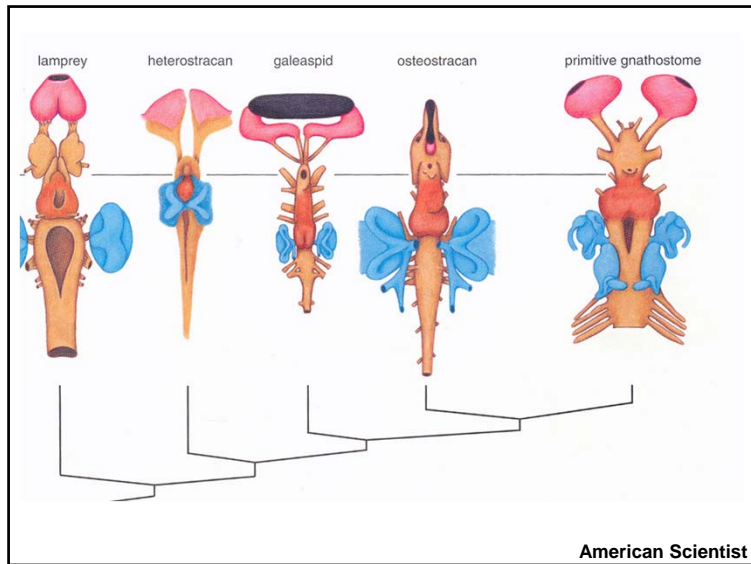
Even if fertilization occurs successfully, the offspring may not survive, or be infertile (e.g., mule).

In this course,

- we will consider speciation as a separation of populations with reproductive isolation;
- species concepts and process of speciation are not the focus.

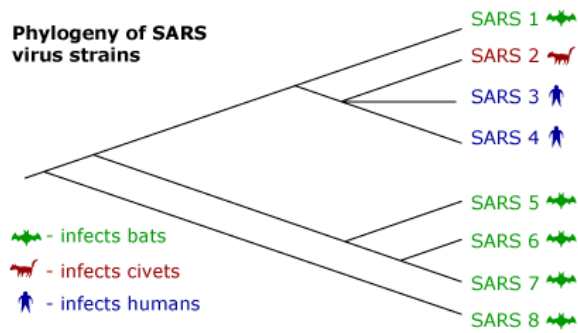
Why phylogeny reconstruction? Some applications





Example of disease phylogeny

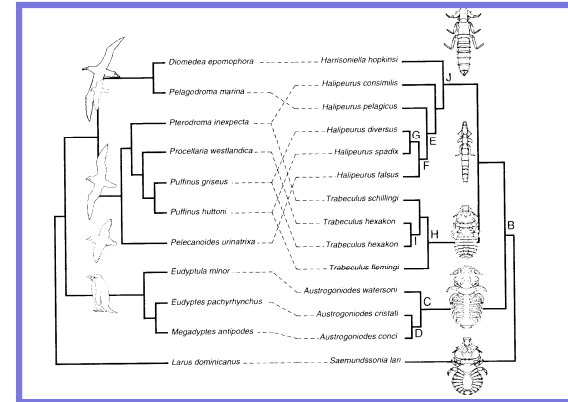
Phylogeny of SARS virus strains



- infects bats
- infects civets
- infects humans

Source: Understanding Evolution: Tracking SARS back to its source.
http://evolution.berkeley.edu/evolibrary/news/060101_batsars
 After: Wendong Li, et al. 2005. Bats Are Natural Reservoirs of SARS-Like Coronaviruses. *Science* 28 October 2005: Vol. 310, no. 5748, pp. 676 - 679. DOI: 10.1126/science.1118391

Example of host-parasite phylogeny



Source: Page, R.D.M., Cruickshank, R.H., Dickens, M., Furness, R.W., Kennedy, M., Palma, R.L., Smith, V.S. 2004. Phylogeny of "Philoceanus complex" seabird lice (Phthiraptera: Ischnocera) inferred from mitochondrial DNA sequences. *Molecular Phylogenetics and Evolution*, 30: 633-652.