

Bonine I

# Phylogenetics (Ch 27)

Thursday  
29 January 2009  
ECOL 182R UofA  
K. E. Bonine

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Tucson ~native

University of Arizona (undergrad)

- Ecology & Evolutionary Biology
- Economics

University of Wisconsin, Madison (graduate)

Zoology, Evolutionary Physiology  
Reptiles & Amphibians

Teaching at UA since 2002

- Herpetology
- Vertebrate Physiology
- Conservation Biology
- Environmental Biology
- Introductory Biology

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M 1:10-2pm  
W 11:10-11am  
in BSE113

Mostly Middle-Third of this course (plants etc.)

Text readings are highly encouraged

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**Website:** For my lecture material:

<http://www.eebweb.arizona.edu/courses/182-spring2009-Bonine/182-BONINE-sp2009.htm>

There is a [link from Dr. Schaffer's 182R website](#).

**Text:** The text is Freeman, Scott. 2008. *Biological Science* (Third Edition). Pearson Benjamin Cummings, San Francisco, CA. Available at UA Bookstore.

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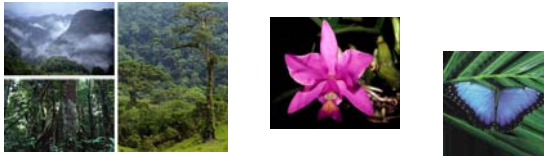
## Posted Lecture Notes

Items in orange will not be available on-line (except for today), but they will be presented during lecture.

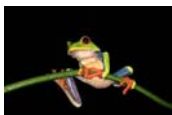
Note that slides are numbered for easy reference.

I will strive to post the lecture PDF file on your D2L site [before](#) lecture.

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How do we organize/categorize biodiversity? Why would we want to?



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## Linnaean Taxonomy (1700s)

- **Taxonomy** is the effort to name and classify organisms.
- In Linnaeus' taxonomic system for classifying organisms, each organism is given a unique two-part scientific name consisting of the **genus** and the **species**.
  - (1) A **genus** is made up of a closely related group of species.
  - (2) A **species** is made up of individuals that regularly breed together and/or have characteristics that are distinct from those of other species.

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## Taxonomic Levels

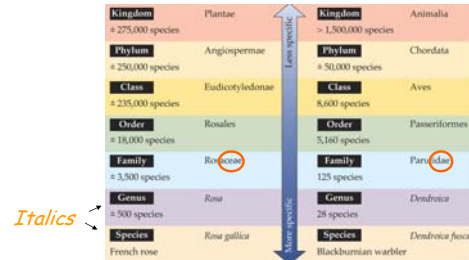
- Linnaeus' system is **hierarchical** with nested **taxa**. The taxonomic levels from least to most specific are as follows:

domain  
kingdom  
phylum  
class  
order  
family  
genus  
species

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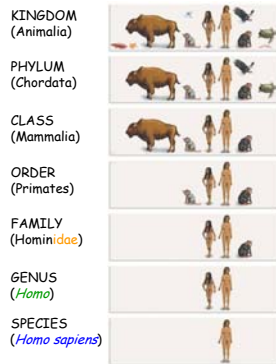
## Linnaean hierarchy

- Group of organisms treated as unit is a **taxon** (plural **taxa**)
- Hierarchy of taxonomic categories



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## Linnaeus' Taxonomic Levels



9

## Theory of common descent

- Any two organisms can trace back to a common ancestor
- We all belong to a big family tree, some more closely related than others
- On the right is a history of **individuals**; can also draw up the history of **species**



## Phylogeny

- Phylogeny** = history of exactly how a group of organisms are descended from their common ancestor
- Phylogenetic tree** = representation of that history

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## Lamarck



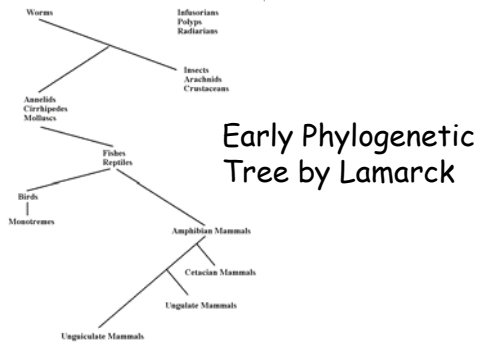
1821. From Desmond 1989, p. 43.

<http://www.nceas.ucsb.edu/~alroy/lefa/Lamarck.jpg>

<http://www.tulane.edu/~darwin/Herbarium/Herbarium/Koch%20images/Lamarck2.jpg>

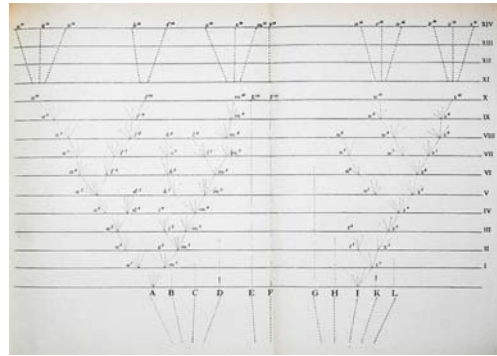


"Cineraria"  
Jean Baptiste Pierre Antoine de Monet de Lamarck, (1744-1829). *Tableau encyclopedique et methodique des trois regnes de la nature... botanique*. Paris, 1791-1823



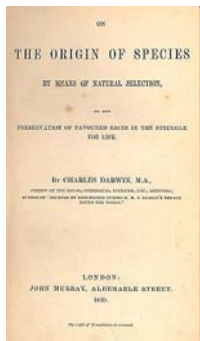
<http://bill.snr.arizona.edu/classes/182/Zool%20Phil%20Tree%20crop-700.gif>

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Tree diagram used to show the divergence of species. It is the only illustration in *The Origin of Species* – Darwin 1859.

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Title page of the 1859 edition of *On the Origin of Species*

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200 years old!

HAPPY BIRTHDAY MR. DARWIN!!!

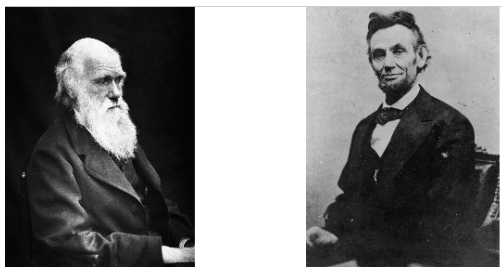
CHEER, BALLOONS AND FUN ACTIVITIES! POETRY, SCIENCE AND MUSIC!

12 Feb 2009

Celebration 3-5pm Thursday Feb 12 Student Union Memorial Ballroom

sponsored by EEB, Poetry Center, the UGA Bookstore & others at UGA

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They Still Free Minds and Peoples  
Darwin & Lincoln's Birthday, February 12th, 1809

[http://media.richarddawkins.net/images/2008/darwinday/016\\_FrederickBrenion.jpg](http://media.richarddawkins.net/images/2008/darwinday/016_FrederickBrenion.jpg)

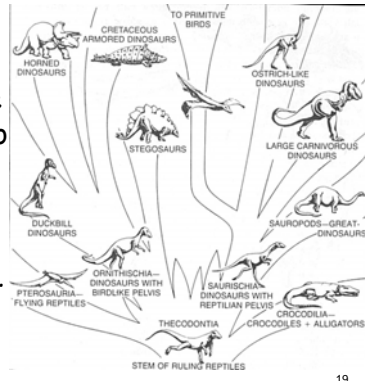
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- A **phylogenetic tree** is a graphical representation of the evolutionary relationships among species. Phylogenies can be established by analyzing similarities and differences in **traits**.



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Often we are attempting to identify relationships from the distant past...



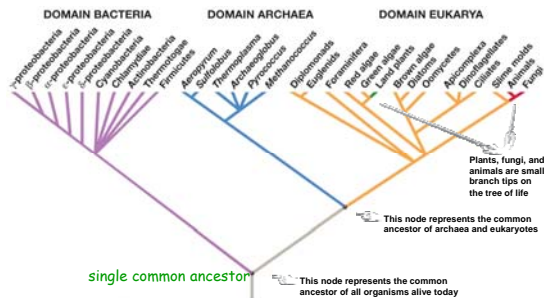
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## Phylogenetic Trees

- **Systematics**, the study of organismal diversity with respect to evolutionary (or not) relationships between organisms (patterns of descent).
  - **Taxonomy** - a subdiscipline that relates to classification
- What evolutionary relationships could be useful/helpful to understand?

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## Phylogenetic Tree of Life



Remember that >99% of all species are EXTINCT!

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## Old species often become new species...

Darwin's eventual conclusions stemming from his first question about the birds and plants of the Galapagos were to feature in one of the most important passages in *Origin of species* (pp. 397-406). The passage ended with one of his key points about evolution by natural selection:

The relations just discussed ... [including] the very close relation of the **distinct species** which inhabit the islets of the same archipelago, and especially the striking relation of the inhabitants of each whole archipelago or island to those of the nearest mainland, are, I think, **utterly inexplicable on the ordinary view of the independent creation of each species**, but are explicable on the view of **colonisation** from the nearest and readiest source, together with the **subsequent modification and better adaptation of the colonists to their new homes**.

[http://darwin-online.org.uk/EditorialIntroductions/Chancellor\\_Keynes\\_Galapagos.html](http://darwin-online.org.uk/EditorialIntroductions/Chancellor_Keynes_Galapagos.html)



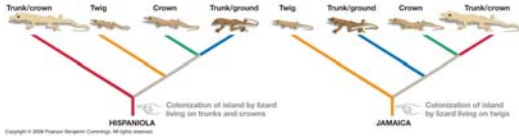
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## Adaptive Radiation

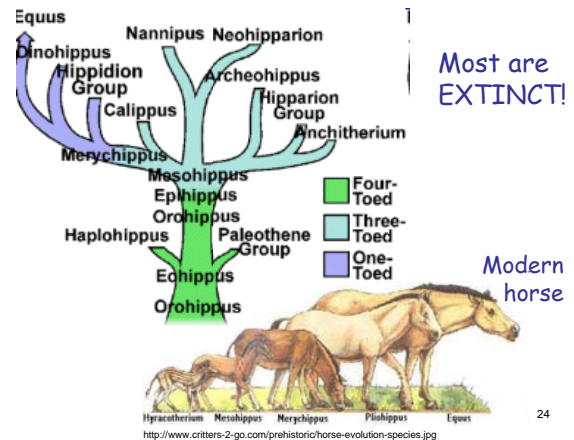
(a) Species of Anolis vary in leg length and tail length. Some species are ground dwelling; others live in distinct regions of shrubs or trees.



(b) The same adaptive radiation of Anolis has occurred on different islands, starting from different types of colonists.

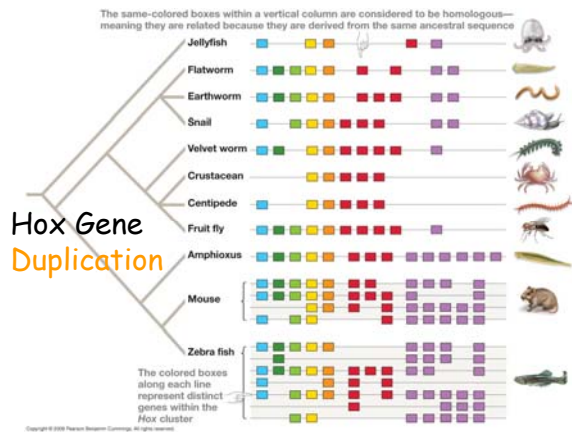


Expand and **diversify** into areas **without** competition (Area either **1) new** or **2) former residents extinct**.)

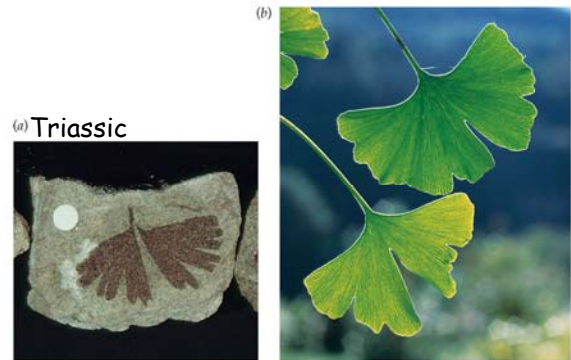


<http://www.critters-2-go.com/prehistoric/horse-evolution-species.jpg>

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## Living fossils: Gingko



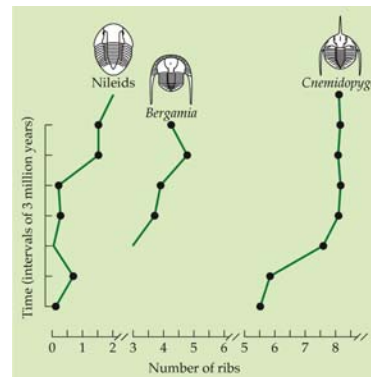
## Living fossils: Horseshoe crabs



(b) *Limulus polyphemus*

~identical for 300 mya  
sandy coastlines are harsh environments that changed little, so they didn't need to evolve anything different

## Gradual change: trilobite rib number



fossils show number of ribs

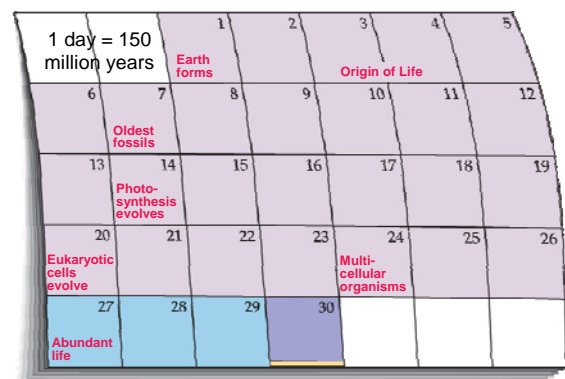
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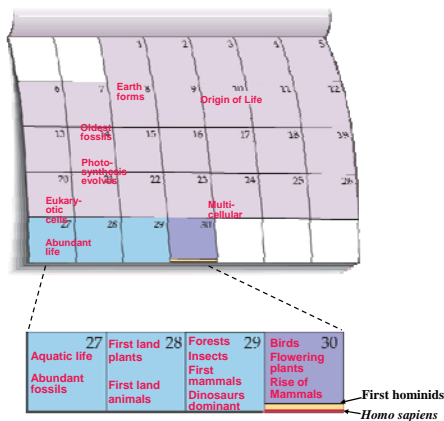
## Biological history

22.1 Earth's Geological History (Part 2)	
RELATIVE TIME SPAN	MAJOR EVENTS IN THE HISTORY OF LIFE
Cenozoic	Quaternary 1.8 mya <sup>a</sup> Humans evolve; many large mammals become extinct
	Tertiary 65 mya Diversification of birds, mammals, flowering plants, and insects
Mesozoic	Cretaceous 144 mya Dinosaurs continue to diversify; flowering plants and mammals diversify. Mass Extinction at end of period (~75% of species disappear)
	Jurassic 206 mya Diverse dinosaurs; radiation of ray-finned fishes
	Triassic 248 mya Early dinosaurs; first mammals; marine invertebrates diversify; first flowering plants; Mass Extinction at end of period (~60% of species disappear)
Paleozoic	Permian 290 mya Reptiles diversify; amphibians decline; Mass Extinction at end of period (~90% of species disappear)
	Carboniferous 354 mya Extensive "fern" forests; first reptiles; insects diversify
	Devonian 417 mya Fishes diversify; first insects and amphibians. Mass Extinction at end of period (~75% of species disappear)
Precambrian	Silurian 443 mya Jawless fishes diversify; first ray-finned fishes; plants and animals colonize land
	Ordovician 490 mya Mass Extinction at end of period (~75% of species disappear)
	Cambrian 543 mya Most animal phyla present; diverse algae
Precambrian	600 mya Ediacaran fauna
	1.5 bya <sup>b</sup> Eukaryotes evolve; several animal phyla appear
	3.8 bya Origin of life; prokaryotes flourish
4.5 bya	

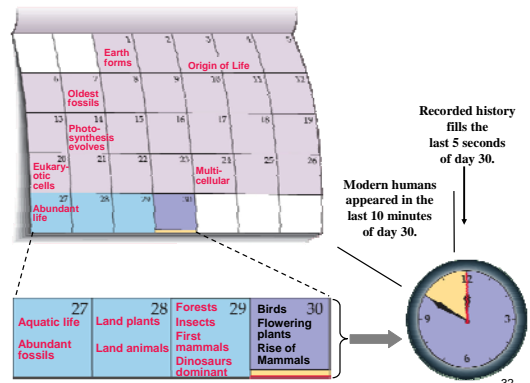
<sup>a</sup>mya, million years ago; bya, billion years ago.

## Biological history

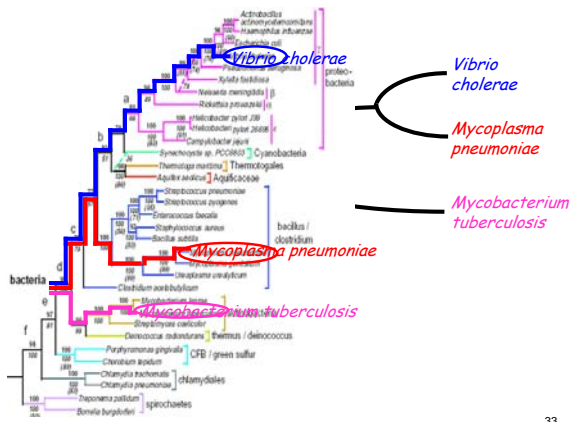




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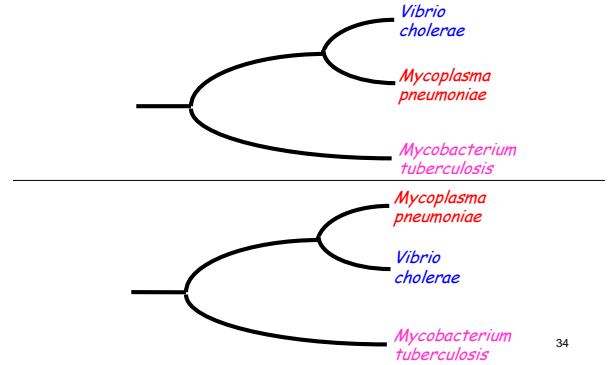


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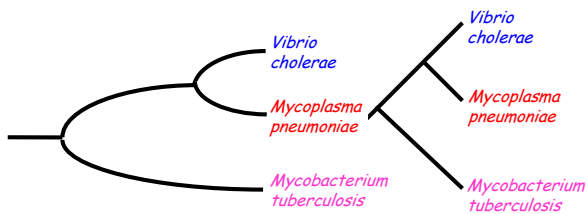
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These trees are the same



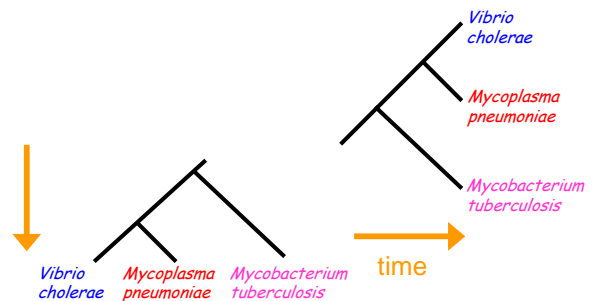
34

These trees are the same



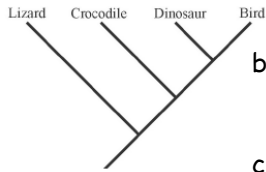
35

These trees are the same



36

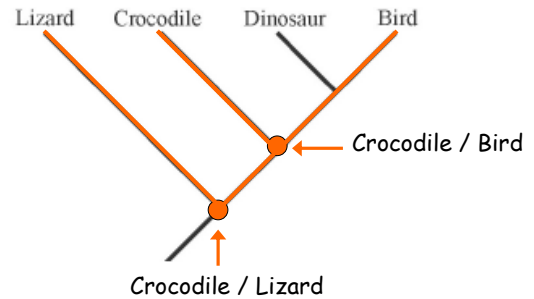
Which of these statements follows from the tree?



- Crocodiles are more closely related to lizards than to birds
- Crocodiles are more closely related to birds than to lizards
- Crocodiles are equally related to lizards and birds

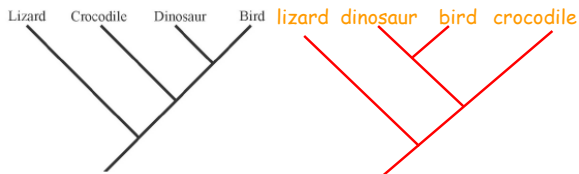
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Focus on nodes = common ancestors



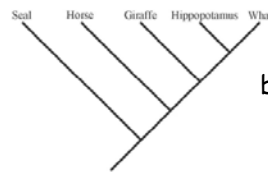
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Another method: redraw tree with focus on crocodile



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Which of these statements follows from the tree?



- A seal is more closely related to a horse than to a whale.
- A seal is more closely related to a whale than to a horse.
- A seal is equally related to a whale and a horse.

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## Ancestral traits

- Traits inherited from ancestor in distant past should be shared by large number of species
- Traits that first appeared in more recent ancestor should be shared by fewer species
- Ancestral trait** = shared traits inherited from a common ancestor
- Derived trait** = different from ancestral form

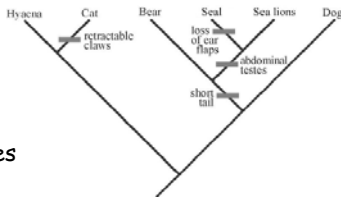
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## Nucleic acids are an ancestral trait

- Nucleic acids (DNA/RNA) as genetic material is ancestral to all life on Earth
- Specific genetic code is ancestral to most

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The ancestor to this tree had a long tail, ear flaps, external testes, and fixed claws. Based on the tree, what traits does a **sea lion** have?



- a. long tail, ear flaps, external testes, and fixed claws
- b. short tail, no ear flaps, external testes, and fixed claws
- c. short tail, no ear flaps, abdominal testes, and fixed claws
- d. short tail, ear flaps, abdominal testes, and fixed claws
- e. long tail, ear flaps, abdominal testes, and retractable claws

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### Ancestral vs derived may depend on scale of taxa

- In rodents: continuously growing incisors are **ancestral**: all rodents have them
- In mammals they are **derived**, unique to rodents

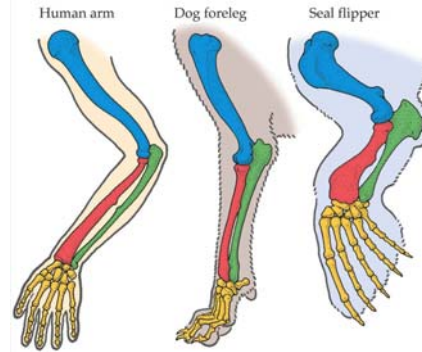
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### Homologous traits

- Features (DNA sequence, behavior, morphology) shared by species **descended from common ancestor** are called **homologous**
- e.g., vertebral column is homologous in vertebrates

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### Mammalian limbs are homologous, even with different functions



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### Homologous structures derived from leaves

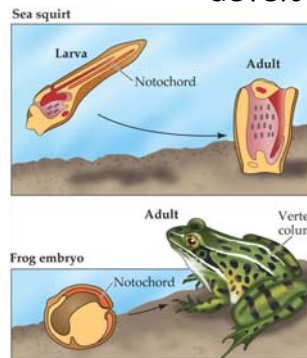


*Cheiridopsis tuberculata*

*Heliconia sp.*

*Sarracenia purpurea*

### Some homologies are hidden by development



Notochord homology seen only in early development, invisible in adult

Both in Phylum Chordata 48



## Deducing phylogeny

### Look at shared traits

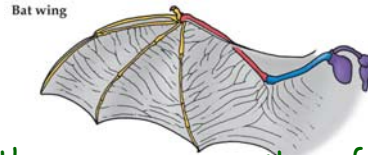
- What is ancestral state?
- How has it been modified?

Two processes make this difficult

- 1. **Convergent evolution** = similar selective pressures make **independently evolved traits look superficially similar**
- 2. **Evolutionary reversal** = character reverts from derived state back to ancestral state

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Bird and bat bones are homologous, but not wings (convergent evolution)



Did the common ancestor of birds and bats have wings?

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## Homoplastic traits

Convergent evolution and evolutionary reversal generate **homoplasies**: traits that are similar for some reason other than inheritance from a common ancestor

Bird and bat wings are **homoplastic**

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*Phrynosoma asio*,  
Long-spined  
Horned Lizard  
MEXICO



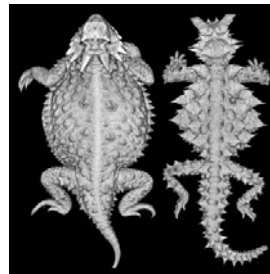
[http://digimorph.org/specimens/Phrynosoma\\_asio/](http://digimorph.org/specimens/Phrynosoma_asio/)

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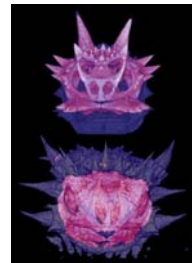
*Moloch horridus*, Thorny Devil, AUSTRALIA



Spiny head: **Homoplasy** or Homology?



*Phrynosoma cornutum* (left) vs. *Moloch horridus* (right)



*P. cornutum* (top) vs. *M. horridus* (bottom); semi-transparent flesh in blue, bone in red

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## All of the following can result in homoplasy except

- a. similar selection pressures.
- b. reverse evolution.
- c. parallel evolution.
- d. descent from a recent common ancestor.
- e. convergent evolution.

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## Creating a phylogeny

1. Choose the characters and identify the possible forms (traits) of the characters
2. Determine ancestral and derived traits
3. Distinguish homologous from homoplastic traits

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## Traditional character choice

- Most often **morphology**
  - describes size and shape of body parts
  - can be seen directly in fossil record
- Also physiological, behavioral, molecular and structural traits as available
- The more characters measured, the more the inferred phylogenies should converge on each other and on the real evolutionary pattern

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## Simple phylogeny example

- Choose 8 vertebrate species and hypothesize their phylogeny
- Traits either present (+) or absent (-)
- Assume that each derived trait evolved only once and that no derived traits were lost

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TAXON	JAWS	LUNGS	CLAWS OR NAILS	GIZZARD
Lamprey	-	-	-	-
Perch	+	-	-	-
Salamander	+	+	-	-
Lizard	+	+	+	-
Crocodile	+	+	+	+
Pigeon	+	+	+	+
Mouse	+	+	+	-
Chimpanzee	+	+	+	-

TAXON	FEATHERS	FUR	MAMMARY GLANDS	KERATINOUS SCALES
Lamprey	-	-	-	-
Perch	-	-	-	-
Salamander	-	-	-	-
Lizard	-	-	-	+
Crocodile	-	-	-	+
Pigeon	+	-	-	+
Mouse	-	+	+	-
Chimpanzee	-	+	+	-

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## Determine ancestral and derived traits

- Chimp and mouse share mammary glands and fur, other animals lack them
- Ancestral or derived? **derived**

TAXON	FEATHERS	FUR	MAMMARY GLANDS	KERATINOUS SCALES
Lamprey	-	-	-	-
Perch	-	-	-	-
Salamander	-	-	-	-
Lizard	-	-	-	+
Crocodile	-	-	-	+
Pigeon	+	-	-	+
Mouse	-	+	+	-
Chimpanzee	-	+	+	-

## Determine ancestral and derived traits

- Use similar reasoning on other traits
- Lamprey has no derived traits, so it is an **outgroup**

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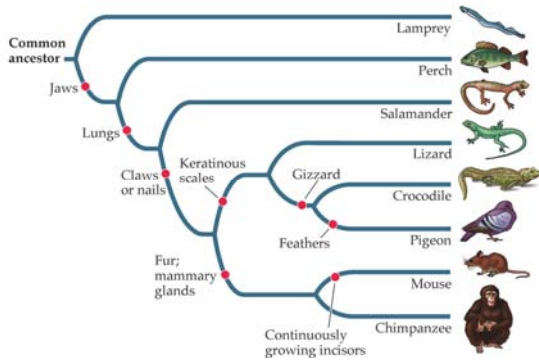
## Your Turn...

TAXON	DERIVED TRAIT*			
	JAWS	LUNGS	CLAWS OR NAILS	GIZZARD
Lamprey	-	-	-	-
Perch	+	-	-	-
Salamander	+	+	-	-
Lizard	+	+	+	-
Crocodile	+	+	+	+
Pigeon	+	+	+	+
Mouse	+	+	+	-
Chimpanzee	+	+	+	-

TAXON	DERIVED TRAIT*			
	FEATHERS	FUR	MAMMARY GLANDS	KERATINOUS SCALES
Lamprey	-	-	-	-
Perch	-	-	-	-
Salamander	-	-	-	-
Lizard	-	-	-	+
Crocodile	-	-	-	+
Pigeon	+	-	-	+
Mouse	-	+	+	-
Chimpanzee	-	+	+	-

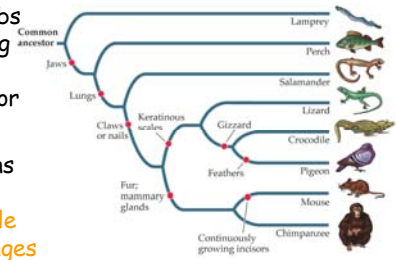
## Simple phylogeny example



- **Not always so simple**
- Assumed derived traits appeared once and were never lost

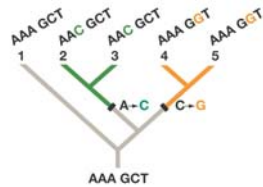
- If we had included snakes, this wouldn't have worked: they have scales but no claws
- Snakes lost limbs and claws during descent from a common ancestor with lizards

- **Parsimony** means we assume the **smallest possible number of changes**

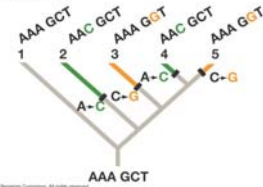


## Parsimony

(a) Two changes



(b) Four changes



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## Modern character choice

Today, most phylogenies come from **gene sequences**

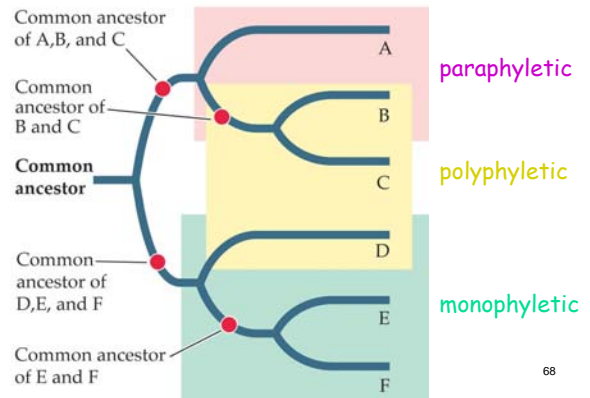
- DNA/RNA/protein

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## Classification should reflect evolutionary relationships

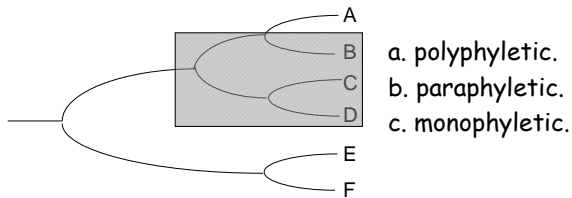
- A **monophyletic** group (or **clade**) **best** contains all descendents of a particular ancestor and no other organisms
- A **polyphyletic** taxon contains members with more than one recent common ancestor
- A **paraphyletic** group contains some, but not all of the descendents of a particular ancestor

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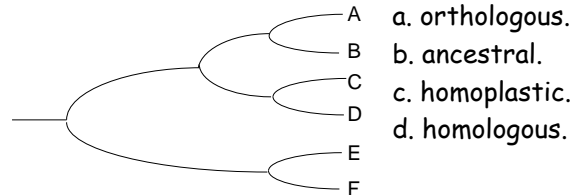
A group consisting of the shaded species is best described as



- polyphyletic.
- paraphyletic.
- monophyletic.

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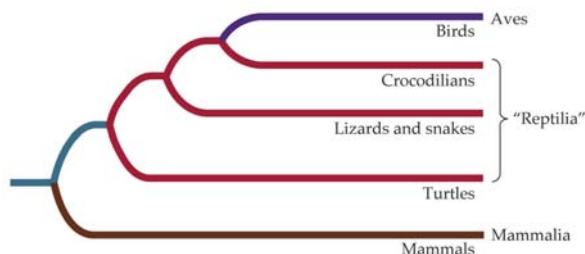
A particular trait is found only in species A and E. Assuming that the phylogenetic tree is correct, the simplest model describes this trait as



- orthologous.
- ancestral.
- homoplastic.
- homologous.

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Should reptiles be a group?



Molecular evidence shows birds and crocodilians closer than crocodilians and other reptiles

Reptiles are **paraphyletic**, since birds excluded

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Birds look very different

- Birds have rapidly evolved unique derived traits since they separated from reptiles
- Groups called **grades** have changed rapidly. May be an appropriate group even if paraphyletic
- General tendency to eliminate paraphyletic groups as we learn more, but some familiar categories, such as reptiles, won't disappear in a hurry

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## Uses of phylogenetic trees

1. How are different species related?
2. What traits do you expect a newly discovered species to have?
3. How many times has a trait evolved?
4. Which molecular change is responsible for adaptation?
5. When did lineages split?  
use DNA sequence and the "molecular clock"

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## The Evolution of Macromolecules

- The **neutral theory of molecular evolution**:
  - postulates that, at the molecular level, the **majority of mutations are selectively neutral**.
- Thus, macromolecule evolution, and much of the genetic variation within species, does not result from positive selection of advantageous alleles nor stabilizing selection.
- Mutation fixation rate is theoretically constant and equal to the neutral mutation rate - a molecular clock.
  - The concept of the **molecular clock** states that **macromolecules should diverge from one another at a constant rate**.

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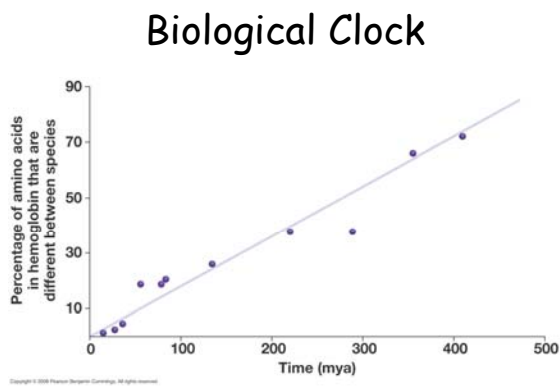


Figure 26.5 *Cytochrome c Has Evolved at a Constant Rate*

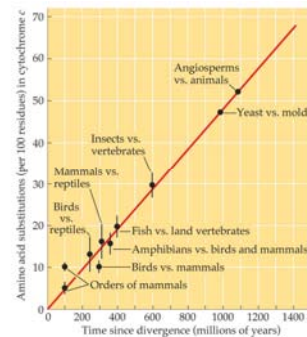
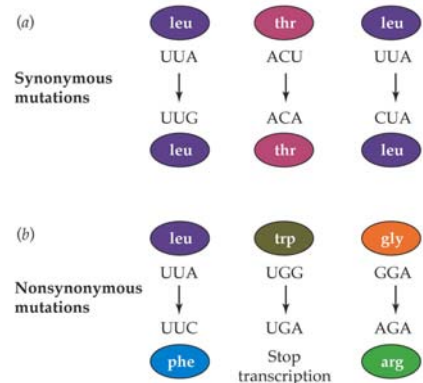


Figure 26.1 *When One Base Does or Doesn't Make a Difference*

## The Evolution of Macromolecules

- Molecular evolution differs from phenotypic evolution in one important way:
  - In addition to **natural selection**, **random genetic drift** and **mutation** exert important influences on the rates and directions of molecular evolution.
  - What does this say about our study of "Evolutionary Theory"?
- Many mutations, called **silent or synonymous** mutations, do not alter the proteins they encode
- A **nonsynonymous mutation** does change the amino acid sequence

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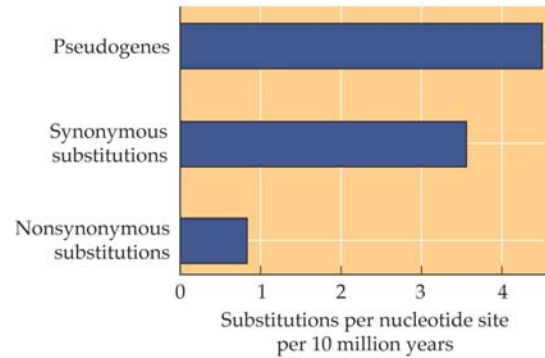
## Genetic Code (DNA → amino acid)

		Second Letter				
		T	C	A	G	
First Letter	T	TTT } Phe TTC } TTA } Leu TTG }	TCT } Ser TCC } TCA } TCG }	TAT } Tyr TAC } TAA } Stop TAG }	TGT } Cys TGC } TGA } Stop TGG } Trp	Third Letter T C A G
	C	CTT } Leu CTC } CTA } CTG }	CCT } Pro CCC } CCA } CCG }	CAT } His CAC } Gln CAA } CAG }	CGT } Arg CGC } CGA } CGG }	
	A	ATT } Ile ATC } ATA } Met ATG }	ACT } Thr ACC } ACA } ACG }	AAT } Asn AAC } Lys AAA } AAG }	AGT } Ser AGC } Arg AGA } AGG }	
G	GTT } Val GTC } GTA } GTG }	GCT } Ala GCC } GCA } GCG }	GAT } Asp GAC } Glu GAA } GAG }	GGT } Gly GGC } GGA } GGG }	T C A G	

<http://plato.stanford.edu/entries/information-biological/GeneticCode.png>

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Figure 26.3 Rates of Base Substitution Differ



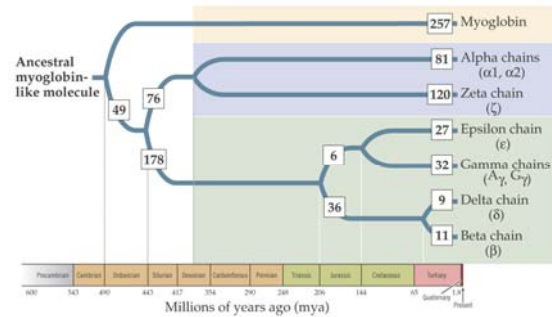
LIFE: THE SCIENCE OF BIOLOGY, Seventh Edition, Figure 26.3 Rates of Base Substitution Differ  
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## The Evolution of Genome Size

- Several rounds of duplication and mutation may lead to formation of a **gene family**, a group of **homologous genes with related functions**.
- There is evidence that the globin gene family arose by gene duplication.
  - = Neofunctionalization
- To estimate the time of the first globin gene duplication, a **gene tree** can be created.
- Based on the gene tree, the two globin gene clusters are estimated to have split about 450 mya.

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Figure 26.9 A Globin Family Gene Tree



LIFE: THE SCIENCE OF BIOLOGY, Seventh Edition, Figure 26.9 A Globin Family Gene Tree  
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