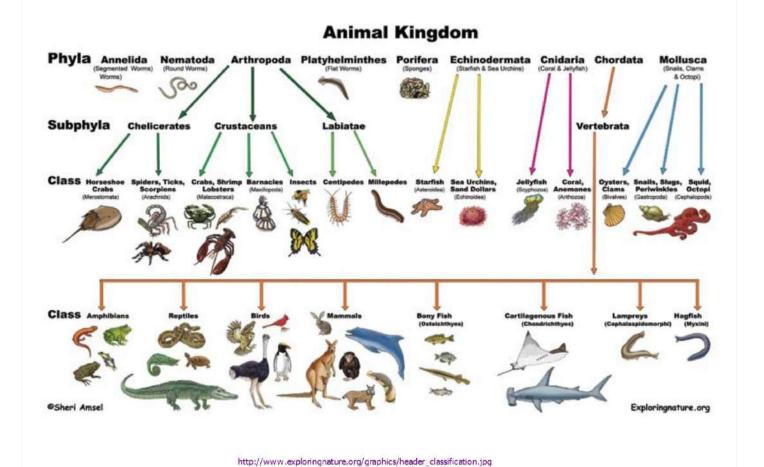
Chapter 19: Taxonomy, Systematics and Phylogeny



19.1 Systematic Biology study of the history of biodiversity uses the characteristicsof living and fossil organisms to infer the relationships of organisms over time

<u>Taxonomy</u>

- branch of systematic biology that identifies, names and organizes biodiversity into related categories (<u>taxa</u> - pl; <u>taxon</u> - sing)
- o in the past, used physical traits
- o now, try to classify into <u>natural groups</u> with a shared evol hist
- use DNA sequences
- phylogeny is contructed from natural groups to show evolutionary history of taxa (evol family tree)

Carolus Linnaeus

- oclassification hierarchy
- o binomial nomenclature (specific epithet)
- eliminates confusion

Names of Taxa

- o domain
- kingdom
- o phylum
- o class
- order
- ু family
- genus
- species

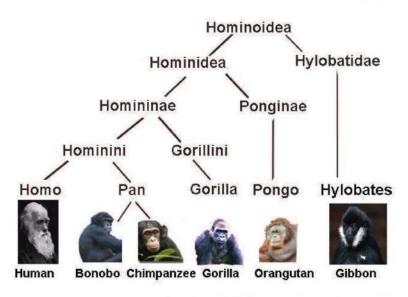
Kingdom	Animalia	Different than a plant or fungus					
Phylum	Chordata	Having a backbone					
Class	Mammalia	Having hair, producing milk and giving birth to live young					
Order	Primates	Having binocular vision, a generalized body plan and an increased reliance on vision					
Family	Hominidae	Being a member of all of the family of higher apes (hum chimpanzees, gorillas, orang-utans, gibbons and siamang					
Sub-family	Homininae	A primate separate from the remainder of the family of apes					
Genus	Homo	In the human line, distinct from other human precursors					
Species	sapiens	Modern human					

http://biologos.org/uploads/static-content/kidder_figure_5.jpg



Order: Primates
Family: Hominidae
Genus: Pongo
Species: Pongo pygmaeus

Pongo abelii



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19-2 The Three-Domain System



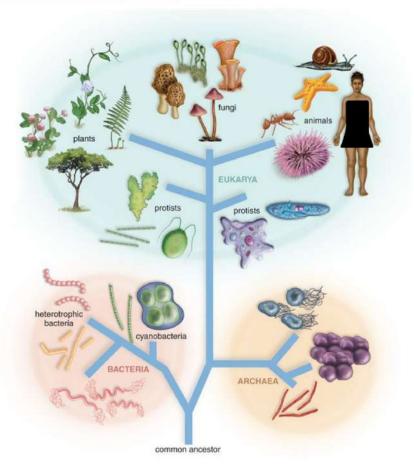


Table 19.1	Major Distinctions Amo	ng the Three Domains of Life
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	Bacteria	Archaea (Control of the Control of t	Eukarya Swall	
Unicellularity	Yes	Yes	Some, many multicellular	
Membrane lipids	Phospholipids, unbranched	Varied branched lipids	Phospholipids, unbranched	
Cell wall	Yes (contains peptidoglycan)	Yes (no peptidoglycan)	Some yes, some no	
Nuclear envelope	No	No	Yes	
Membrane-bounded organelles	No	No	Yes	
Ribosomes	Yes	Yes	Yes	
Introns	No	Some	Yes	

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19.3 Phylogeny

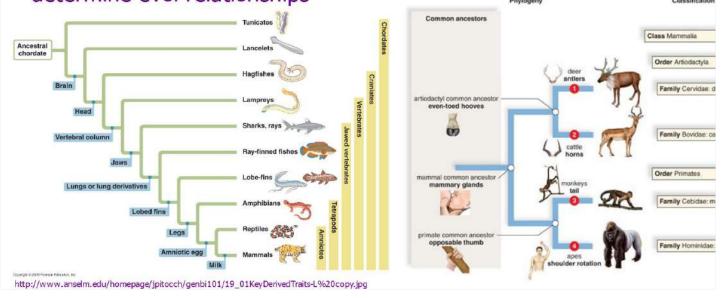
- visual representation of history of biodiversity
- odeveloped using:
 - fossil record
 - comparative anatomy and development
 - sequence, structure and function of DNA and RNA



Interpreting a Phylogeny

- constructed from traits that are shared by and unique to a taxon and their common ancestor
- when a new trait arises, a new path diverges from the old and a new branch is formed
- ancestral traits are traits found in the lineage from common ancestor

 derived traits - are not found in common ancestor and help determine evol relationships



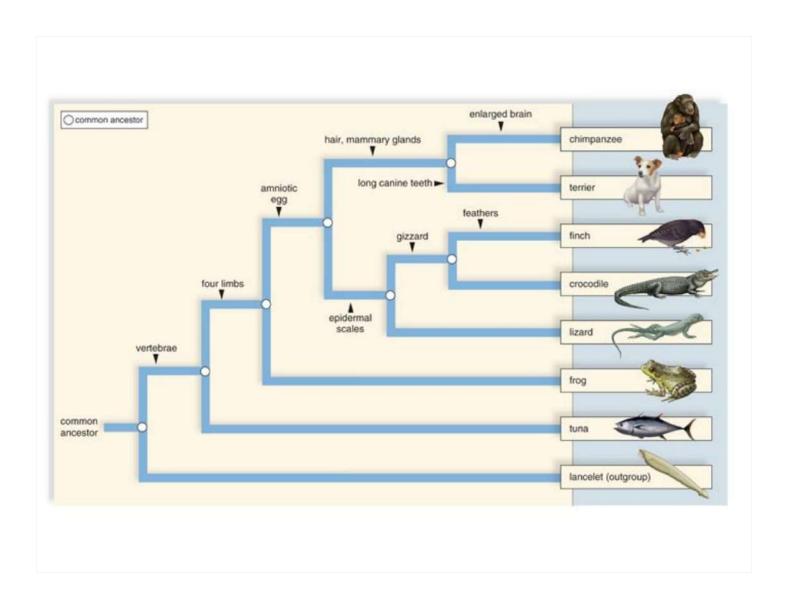
Cladistics

- ouses shared, derived traits to develop a hypothesis of evol hist
- oused to build <u>cladogram</u> type of phylogeny
- oclade common ancestor and its decendants

 parsimony - using simplest solution (smallest number of evol changes)

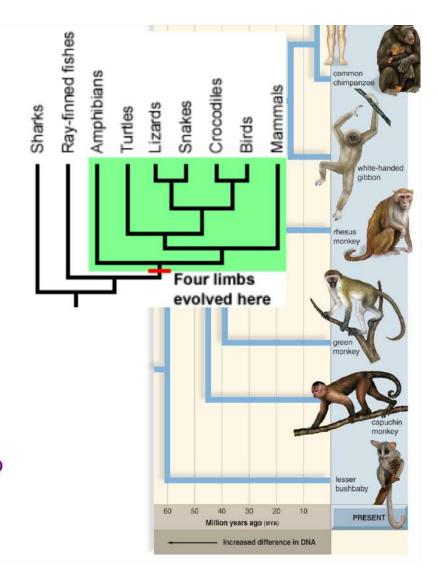
- often change with new info
- outgroup v. ingroup
- each clade contains a uniquely derived trait

		ingroup						(dr	
		chimpanzee	gob	finch	crocodile	lizard	frog	tuna	lancelet (outgroup)
Traits	mammary glands	X	X						
	hair	X	X						
	gizzard			X	X				
	epidermal scales			X	X	X			
	amniotic egg	X	X	X	X	X			
	four limbs	X	X	X	X	X	X		
	vertebrae	X	X	X	X	X	X	X	
	notochord in embryo	X	X	X	X	X	X	X	X



Tracing Phylogeny

- o morphology is misleading
- fossil record helpful, but incomplete
- morphology homologous/analagous traits
- behavioral traits can be used to separate species (mating calls)
- molecular traits mutations in DNA accumulate over time
 - more diff means more separation
 - more closely related also means fewer amino acid diff in proteins



cont'd...

- mtDNA mutates faster and can be used to differentiate closely related species
- o molecular data also useful for agriculture, medicine and forensics

Molecular Clock

- o neutral mutations can be used to provide a timeline of evol hist
- o accumulate but do not affect fitness