

What is a species?

- A human-made construct to categorize variation. Species are difficult to define because of the amount of variation in nature.
- Most commonly, a species is an interbreeding group of animals or plants that are reproductively isolated through anatomy, ecology, behavior, or geographic distribution from all other such organisms.

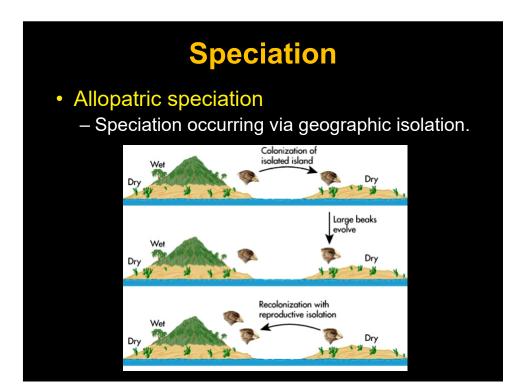
Species concepts

- · Biological species concept
 - Interbreeding populations, reproductively isolated from others
- Phylogenetic species concept
 - Smallest distinguishable cluster of ancestors & descendants
- Evolutionary species concept
 - Unique evolutionary lineages
- · Recognition species concept
 - Unique traits/behaviors that allow identification of mates

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Reproductive isolating mechanisms

- Habitat isolation species A & B occupy different habitats (e.g., arboreal vs. terrestrial)
- Temporal isolation Species A & B breed in different months, or are active by day vs. night
- Behavioral isolation Courtship behavior by males of species A does not elicit a response by females of species B
- Mechanical incompatibility species A & B cannot mate successfully because of anatomical difference

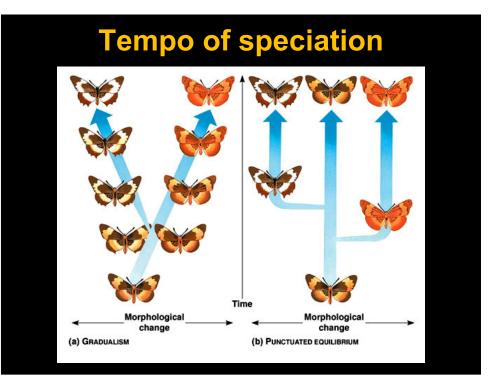


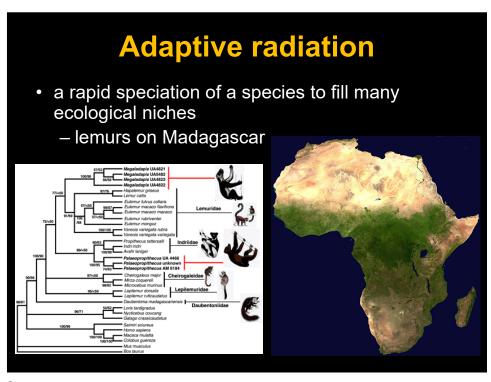
Modes of speciation • Speciation is the formation of one or more species via (reproductive) isolation. species 3 species 3 species 2 species 2 species 1 species 1 Cladogenesis Anagenesis **CLADOGENESIS: ANAGENESIS:** Branching of a species Speciation within a or lineage lineage

Tempo of speciation

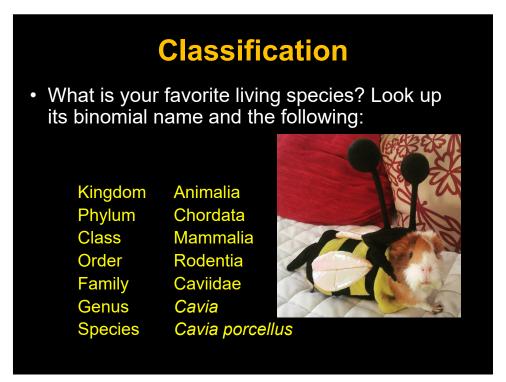
- Phyletic gradualism (slow)
 - Darwinian view of slow, incremental evolutionary change.
- Punctuated equilibrium (fast)
 - Model of evolution characterized by rapid bursts of change, followed by long periods of stasis.

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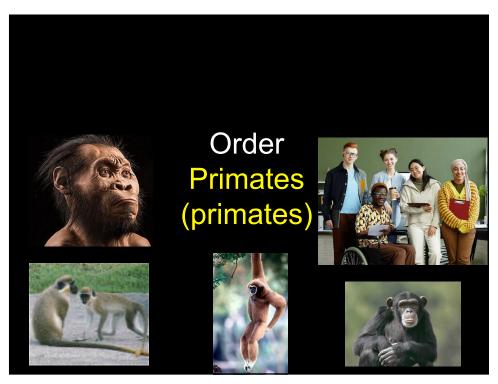






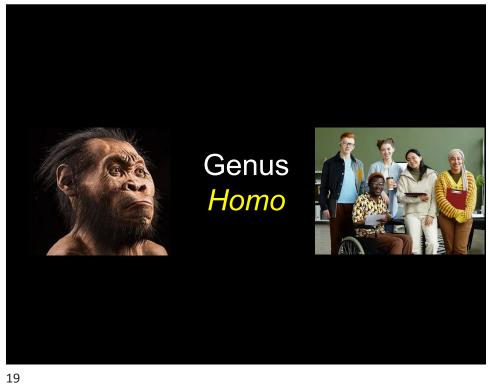














Taxonomy & systematics

- Taxonomy: theory & practice of naming & classifying biological organisms.
- Systematics: branch of biology that describes patterns of organismal variation & relationships.
- · Binomial nomenclature
 - Genus species like Homo sapiens

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Phylogeny & classification

- Classification should reflect the evolutionary history of life (phylogeny).
- Phylogenies are the evolutionary histories of groups of related organisms.
 - trace adaptations from common ancestors
 - show relationships & the time scale of splitting between ancestors & descendants (usually in the form of a *phylogenetic tree*)
- The 'Tree of Life' is very bushy, with different groups evolving new adaptations.

Reconstructing phylogeny

Q: How do we trace ancestry back millions of years?

A: Look at living & fossil groups, try to identify features that related groups share.

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Establishing evolutionary history

- The genotype & phenotype of living animals (generally only phenotype for extinct animals) is the primary evidence for classification.
- A species' genotype/phenotype is viewed as a suite of characters/traits.
 - Characters are discrete, variable features that differ across groups (species, genera, orders, etc.)

Character examples

- Number of teeth.
- Number of vertebrae.
- Alleles at a locus.
- Feathers vs. fur vs. scales.
- Bony morphology of the cranium.
- Type of reproductive system.

Can be morphological, behavioral or genetic.

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Character value depends on context

- The value of a character (whether it is phylogenetically informative or not) cannot be determined without a context.
- A character which is not informative in one context may be pivotal in another.
- Characters often apply at particular levels in classification.

What makes an informative character?

- Variable between groups, but not within groups.
- Shared by more than one group.
 (Provides information about grouping, unlike unique characters)

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Shared character examples

 Among primates, some monkeys of the Americas have prehensile tails.



Among mammals, dogs & cats have carnassial teeth.

Unique character examples

 Among mammals, bats have wings.



 Among primates, humans are bipedal.



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Unique vs. shared is generally easily discernible

 If you have access to all of the information about your study group, it should be relatively easy to determine if characters are unique.

Unique vs. shared examples

 RELATIVE: Among mammals, bats are unique in having wings, but within bats there are many species - all of which have wings.





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Early approaches: phenetics

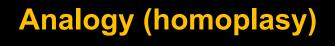
- Method of classification that groups organisms based on overall similarity (can be molecular, phenotypic or anatomical).
- Phenetics assumes that all similarities imply relatedness.
- But, do all similarities imply relatedness?

Different types of similarities

- Homology vs. analogy
- Homology
 - similarities due to common ancestry
 - imply shared descent (common ancestry)
- Analogy
 - similarities due to common function
 - does not imply shared descent or common ancestry

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Homology Traits that are the same in different organisms because they were inherited from a common ancestor e.g., arm bones in amphibians, reptiles, birds & mammals Human Cat Whale Bat





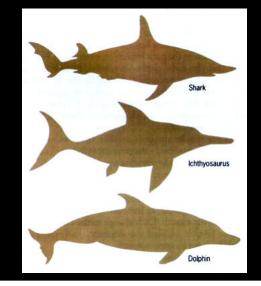




- common function, NOT common ancestry
- convergent (parallel) evolution
- e.g., wing structure of insects, birds, bats

Analogy (homoplasy)

Similarities due to common function, NOT ancestry

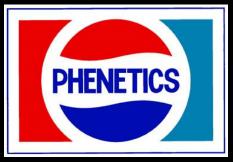


Fish

Reptile

Mammal

 Groups organisms based on overall similarity.
 Does NOT distinguish between similarity due to homology (common ancestry) or analogy (common function).



· Can lead to incorrect classifications.

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 Method of classification using shared, derived characters to link taxa together in nested groups.

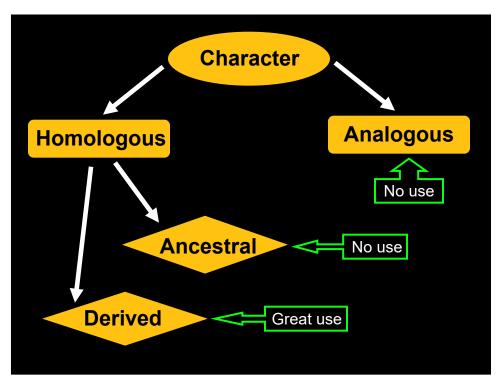


 Shared, derived characters are more informative as they provide grouping information. Ancestral (primitive) characteristics do not.

Different types of similarity

- There are 2 types of homologies:
 - Ancestral or primitive (plesiomorphy)
 - Derived (apomorphy)
- Thus there are 4 possible character types:
 - shared ancestral (symplesiomorphies)
 - shared derived (synapomorphies)
 - uniquely derived (autapomorphies)
 - Independently derived (analogies / homoplasies)

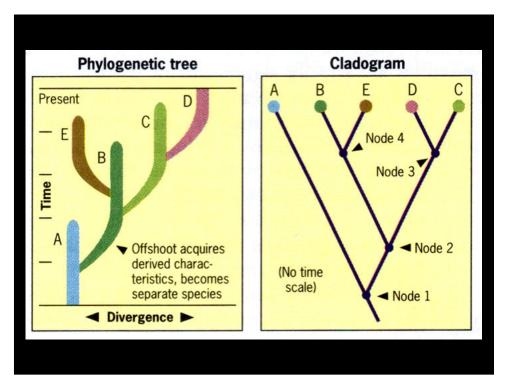
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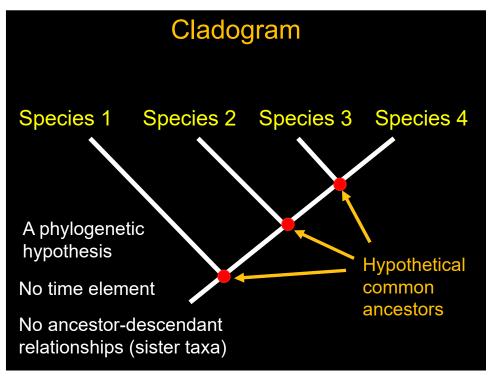


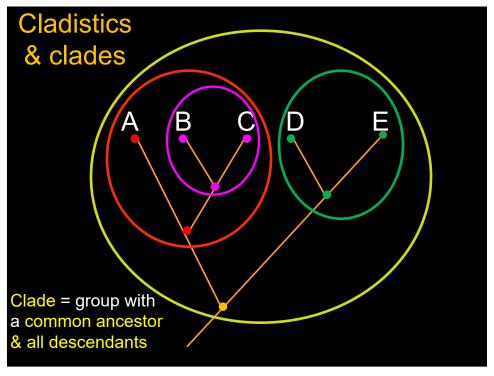
Trees

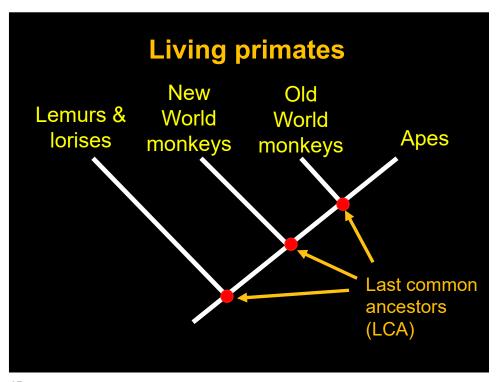
- Cladogram
 - strictly a branching tree
 - shows the sequence of character evolution
 - does not show ancestor-descendant relationships
- Phylogenetic tree
 - branch lengths indicate time & ancestordescendant relationships can be represented

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Constructing relationships

We can examine relationships between primates by exploring their physical traits.

- Shared trait present in more than one taxon.
- Ancestral trait expressed in an ancestor and the descendant; this may be shared across multiple taxa. Also known as a primitive trait.
- Derived trait trait expressed in descendants but not expressed in an ancestor. These features may be shared across taxa or unique to an individual taxon.

Constructing relationships

- · There is no end point of evolution!
 - No species is more evolved than any other species. Taxa CANNOT be primitive, only traits can.
- Traits are ancestral or derived relative to other taxa.
- Traits can develop in a mosaic fashion.
 - Most taxa have derived AND ancestral traits.

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Example

 Some primates have tails, some do not.





