Classification

Section 18-2

 Key Idea: Scientists have traditionally used similarities in appearance and structure to group organisms. However, this approach has proven problematic.

Traditional Systemics

- Some groups look similar but turn out to be distantly related.
- Other groups look different but turn out to be closely related.

Phylogenetics

- Key Idea: Grouping organisms by similarity is often assumed to reflect phylogeny, but inferring phylogeny is complex in practice.
 Phylogeny is the ancestral
 - relationships between species.

Phylogenetics

Not all similar characteristics are inherited from a common ancestor. Through the process of convergent evolution, similarities may evolve in groups that are not closely related. Similar features may evolve because the groups have adopted similar habitats or lifestyles.

Similarities that arise through convergent evolution are called *analogous characters*.



Cladistics

Key Idea: Cladistic analysis is used to select the most likely phylogeny among a given set of organisms. Cladistics is a method of analysis that infers phylogenies by careful comparisons of shared characteristics.

Cladistics

- Cladistics focuses on finding characters that are shared between different groups because of shared ancestry.
- A derived character is one that evolved in one group but not the other.

Cladogram

Data Table for Cladogram

| Group of organisms | Characters | | |
|-------------------------------|-----------------|-------|---------|
| | Vascular tissue | Seeds | Flowers |
| Mosses (out-group) | 0 | 0 | 0 |
| Ferns | 1 | 0 | 0 |
| Pine trees and other conifers | 1 | 1 | 0 |
| Flowering plants | 1 | 1 | 1 |



Inferring Evolutionary Relatedness

- Key Idea: Biologists compare many kinds of evidence and apply logic carefully in order to infer phylogenies.
- Morphology refers to the physical structure or anatomy of organisms.
 Molecular evidence includes genetic information to infer phylogenies.

 The principle of parsimony holds that the simplest explanation for something is the most reasonable, unless strong evidence exists against that explanation.

Evolutionary Relatedness

Given two possible cladograms, the one that implies the fewest character changes between points is preferred.