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phylogenetic trees how do the changes in gene sequences allow us to reconstruct the evolutionary relationships between related species the saying don t judge a book by its cover could be applied to the topic of evolution for example humans share 75 of their dna with chickens bi web japanese maples lace leaf flowering plants dogwood oaks ginkgos conifers bonsai we invite you to visit our specialty nursery in sonoma county we are located at 10300 sonoma highway kenwood california telephone 707 833 1161 email sales at wildwoodmaples com directions to wildwood nursery web key concepts phylogeny is based on evolutionary relationships a phylogenetic tree can be built by repeatedly identifying both the most dissimilar members and the most similar members in a group of organisms dna sequences can be used in the study of phylogenetics student learning targets web the most important thing about pogil activities is that students learn to construct their own content knowledge and develop important 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trees whose branches link together the

relatedness of different organisms

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