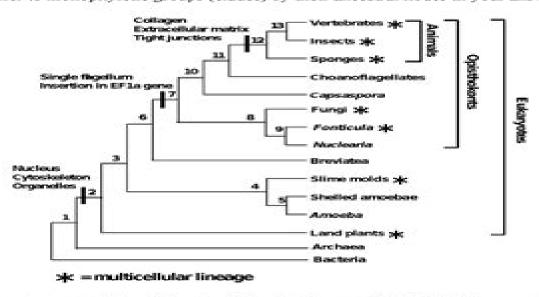
Consider the phylogenetic tree shown below. Nodes are labeled with numbers (1 – 13) above each node. You can refer to monophyletic groups (clades) by their ancestral nodes in your answers.



a) Is the group composed only of Fungi and Fonticula monophyletic? Justify your answer.

No. A monophyletic group is defined as a group of the groups/species, their most recent common ancestors, and the descendants of this ancestors. A monophyletic group would be Fungi, Fonticula, their recent ancestor (8), and Nuclearia (the ancestor's descendant).

b) Identify a monophyletic group of your choice that includes choanoflagellates. List all of the members of this group and the node of the common ancestor.

The node of the common ancestor of choanoflagellates would be 11 and would include the descendant vertebrates, insects, sponges, and choanoflagellates in the monophyletic group.

c) Which node represents the most recent common ancestor of land plants and animals?

Node 2.

d) Assume that multicellularity can be gained but not lost.

## **Biology Phylogenetic Tree Exercise Answers**

Ligia R. Rodrigues, George Sakellaris, Giovanni Sannia

#### **Biology Phylogenetic Tree Exercise Answers:**

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and government The complexity of the global system has reached proportions that seriously challenge our abilities to understand the consequences of our use of technology modification of natural ecosystems or even how to govern ourselves For this reason complex mathematics is eschewed when simpler structures will suffice allowing the widest possible audience to apply and benefit from the available tools and concepts of systems science in their own work The book shows in detail how to functionally and structurally deconstruct complex systems using a fundamental language of systems It shows how to capture the discovered details in a structured knowledge base from which abstract models can be derived for simulation The knowledge base is also shown to be a basis for generating system design specifications for human built artifacts or policy recommendations policy mechanisms for socio economic ecological systems management The book builds on principles and methods found in the authors textbook Principles of Systems Science co authored with Michael Kalton but without prerequisites It will appeal to a broad audience that deals with complex systems every day from design engineers to economic and ecological systems managers and policymakers Bioinformatics Challenges at the Interface of Biology and Computer Science Teresa K. Attwood, Stephen R. Pettifer, David Thorne, 2016-09-26 This innovative book provides a completely fresh exploration of bioinformatics investigating its complex interrelationship with biology and computer science It approaches bioinformatics from a unique perspective highlighting interdisciplinary gaps that often trap the unwary The book considers how the need for biological databases drove the evolution of bioinformatics it reviews bioinformatics basics including database formats data types and current analysis methods and examines key topics in computer science including data structures identifiers and algorithms reflecting on their use and abuse in bioinformatics Bringing these disciplines together this book is an essential read for those who wish to better understand the challenges for bioinformatics at the interface of biology and computer science and how to bridge the gaps It will be an invaluable resource for advanced undergraduate and postgraduate students and for lecturers researchers and professionals with an interest in this fascinating fast moving discipline and the knotty problems that surround it Bio-Inspired Computing for Information Retrieval Applications Acharjya, D.P., Mitra, Anirban, 2017-02-14 The growing presence of biologically inspired processing has caused significant changes in data retrieval With the ubiquity of these technologies more effective and streamlined data processing techniques are available Bio Inspired Computing for Information Retrieval Applications is a key resource on the latest advances and research regarding current techniques that have evolved from biologically inspired processes and its application to a variety of problems Highlighting multidisciplinary studies on data processing swarm based clustering and evolutionary computation this publication is an ideal reference source for researchers academics professionals students and practitioners Bio-Inspired Artificial Intelligence Dario Floreano, Claudio Mattiussi, 2023-04-04 A comprehensive introduction to new approaches in artificial intelligence and robotics that are inspired by self organizing biological processes and structures New approaches to artificial intelligence spring from the idea that intelligence emerges as much from cells

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intended as a textbook for engineering and computer science students attending advanced undergraduate and graduate courses in bioinformatics and computational biology The book develops bioinformatics concepts from the ground up starting with an introductory chapter on molecular biology and genetics to enable physical science students to appreciate the challenges in biological data management sequence analysis and systems biology The book is divided into five parts The first one includes a survey of existing biological databases and tools that have become essential in today s biotechnology research The second part covers methodologies for retrieving biological information including fundamental algorithms for sequence comparison scoring and determining evolutionary distance The third part of the book focuses on modeling biological sequences and patterns as Markov chains covering core principles for analyzing and searching for sequences of significant motifs and biomarkers and developing stochastic ergodic hidden Markov models for biological sequence families The fourth one is dedicated to systems biology and covers phylogenetic analysis and evolutionary tree computations as well as gene expression analysis with microarrays In turn the last part of the book includes an introduction to machine learning algorithms for bioinformatics and outlines strategies for developing intelligent diagnostic machine learning applications RNA sequence data and deep learning systems for mass spectrometry data All in all this book offers a unique hands on reference guide to bioinformatics and computational biology This second edition has been updated to cover additional and most recent databases and machine learning and deep learning applications in RNA sequence and mass spectrometry data analysis Moreover it presents significant enhancements to the chapter dedicated to microarray analysis and more practical examples Bio-Based Compound Production and Their Innovative Industrial Applications with additional end of chapter problems Ligia R. Rodrigues, George Sakellaris, Giovanni Sannia, 2022-01-17

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