

Species 2 is closer related to species 1 because they have the fewest differences in amino acids.

2. What caused the differences in amino acids?

A mutation caused the differences in amino acids. It probably occurred millions of years ago.

3. If the amino acid sequences are similar in two species, will the nucleotide sequence of their DNA also be similar? Why or why not?

Yes, if the amino acids are similar, the DNA will be similar because DNA codes for proteins and proteins are made of amino acids.

4. List some other proteins that could be used for biochemical comparison.

Insulin, cytochrome, amylase, actin, myosin

5. Write the DNA sequence for the portion of species 1 Hemoglobin protein shown above. Write the *RNA and DNA codons* in the boxes below:

GLY	LYS	VAL	ASP	VAL	ASP	GLU	VAL	GLY	GLY
GGU	AAA	GUU	GAU	GUU	GAU	GAA	GUU	GGU	GGU
CCA	TTT	CAA	CTA	CAA	CTA	CTT	CAA	CCA	CCA

GLU	LYS	LEU	HIS	VAL	ASP	PRO	GLU	ASP	PHE
GAA	AAA	UUA	CAU	GUU	GAU	CCU	GAA	GAU	UUU
CTT	TTT	AAT	GTA	CAA	CTA	GGA	CTT	CTA	AAA

ARG	LEU
CGU	UUA
GCA	AAT

6. How is biochemical comparison different from studying the fossil record?

The fossil record deals with traces of organisms that lived long ago; biochemical comparison looks at organisms that have intact DNA. It can be used to determine evolutionary relationships among species living today.

Analyzing Amino Acid Sequences Answer Key

**Molly Fitzgerald-Hayes, Frieda
Reichsman**



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Chapter Resource 14 Class of Organisms Biology Holt Rinehart & Winston, Holt, Rinehart and Winston Staff, 2004

Basics of Bioinformatics Rui Jiang, Xuegong Zhang, Michael Q. Zhang, 2013-11-26 This book outlines 11 courses and 15 research topics in bioinformatics based on curriculums and talks in a graduate summer school on bioinformatics that was held in Tsinghua University. The courses include Basics for Bioinformatics, Basic Statistics for Bioinformatics, Topics in Computational Genomics, Statistical Methods in Bioinformatics, Algorithms in Computational Biology, Multivariate Statistical Methods in Bioinformatics, Research Association Analysis for Human Diseases, Methods and Examples Data Mining and Knowledge Discovery, Methods with Case Examples Applied Bioinformatics Tools, Foundations for the Study of Structure and Function of Proteins, Computational Systems Biology, Approaches for Deciphering Traditional Chinese Medicine, and Advanced Topics in Bioinformatics and Computational Biology. This book can serve as not only a primer for beginners in bioinformatics but also a highly summarized yet systematic reference book for researchers in this field. Rui Jiang and Xuegong Zhang are both professors at the Department of Automation, Tsinghua University, China. Professor Michael Q. Zhang works at the Cold Spring Harbor Laboratory, Cold Spring Harbor, NY, USA. **Bioinformatics** Mr. Rohit Manglik, 2024-06-15 EduGorilla Publication is a trusted name in the education sector committed to empowering learners with high quality study materials and resources. Specializing in competitive exams and academic support, EduGorilla provides comprehensive and well structured content tailored to meet the needs of students across various streams and levels. **DNA and**

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Voet's Principles of Biochemistry Donald Voet, Charlotte W. Pratt, Judith G. Voet, 2018 Voet's Principles of Biochemistry Global Edition addresses the enormous advances in biochemistry particularly in the areas of structural biology and bioinformatics. It provides a solid biochemical foundation that is rooted in chemistry to prepare students for the scientific challenges of the future. New information related to advances in biochemistry and experimental approaches for studying complex systems are introduced. Notes on a variety of human diseases and pharmacological effectors have been expanded to reflect recent research findings. While continuing in its tradition of presenting complete and balanced coverage, this Global Edition includes new pedagogy and enhanced visuals that provide a clear pathway for student learning. 4e de couverture

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