

1/18 - 1/19

LAB 4

Amino Acid Sequences and Evolutionary Relationships

Pre-Lab Discussion

Homologous structures – those structures believed to have a common origin but not necessarily a common function – provide some of the most significant evidence supporting the theory of evolution. For example, the foreilmbs of vertebrates often have different functions and outward appearances, yet the underlying similarity of the bones indicates a common origin. Although homologous structures can be used to demonstrate relationships between similar organisms, they are of little value in determining evolutionary relationships among those structures that are dissimilar.

Another technique used to determine evolutionary relationships is to study the biochemical similarity of organisms. Though molds, aardvarks and humans appear to have little in common physically, a study of their proteins reveals certain similarities. Biologists have perfected techniques for determining the sequence of amino acids in proteins. By comparing the amino acid sequences in homologous proteins of similar organisms and of diverse organisms, evolutionary relationships that might otherwise go undetected can be determined. Biologists believe that the greater the similarity between the amino acid sequences of two organisms, the closer their relationship. Conversely, the greater the differences, the more distant the relationship. Further, biologists have found that such biochemical evidence compares favorably with other lines of evidence for evolutionary relationships.

In this investigation, you will compare amino acid sequences in proteins of several vertebrates. You will also study amino acid differences and infer evolutionary relationships among some diverse organisms.

Problem

How do amino acid sequences provide evidence for evolution?

Materials (per student)

No special materials are needed.

Procedure

Part A. Comparing Amino Acid Sequences

1. Examine Figure 1, which compares corresponding portions of hemoglobin molecules in humans and five other vertebrate animals. Hemoglobin, a protein composed of several long chains of amino acids, is the oxygen-carrying molecule in red blood cells. The sequence shown is only a portion of a chain made up of 146 amino acids. The numbers in Figure 1 indicate the position of a particular amino acid in the chain.

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Rhesus monkey ASN PHE LYS LEU LEU GLY ASN VAL LEU VAL CYS VAL LEU ALA Horse ASN PHE ARG LEU LEU GLY ASN VAL LEU ALA LEU VAL VAL ALA	
Kangaroo ASN PHE LYS LEU LEU GLY ASN RE ILE VAL RE CYS LEU ALA	

- In Data Table 1, notice that the abbreviated names of the amino acids in human hemoglobin are printed.
- 3. In the appropriate spaces in Data Table 1, write the abbreviated name of each amino acid in chimpanzee hemoglobin that is different from that in human hemoglobin. If there are no differences, leave the spaces blank.
- 4. For the remaining organisms, write the abbreviated names of the amino acids that do not correspond to those in human hemoglobin. Note: Always be sure that you compare the amino acid sequence of each organism with that of the human and not the organism on the line above.
- 5. Use Figure 1 to complete Data Table 2.

Part B. Inferring Evolutionary Relationships from Differences in Amino Acid Sequences

Another commonly studied protein is cytochrome c. This protein, consisting of 104
amino acids, is located in the mitochondria of cells. There it functions as a
respiratory enzyme. Examine Figure 2. Using human cytochrome c as a standard,
the amino acid differences between humans and a number of other organisms are
shown.

Species Pairings	# of Differences	Species Pairings	# of Differences
Human-chimpanzee	0	Human-rhesus monkey	1
Human-fruit fly	29	Human-screwworm fly	27
Human-horse	12	Human-snapping turtle	15
Human-pigeon	12	Human-tuna	21
Human-rattlesnake	14	Human-wheat	43
Human-red bread mole	1 48		

- Using Figure 2, construct a bar graph on Graph 1 to show the amino acid differences between humans and other organisms.
 Now examine Figure 3. In this figure the cytochrome c of a fruit fly is used as a standard in comparing amino acid differences among several organisms. Construct a bar graph on Graph 2 to show these differences.

Species Pairing	Number of Differences
Fruit fly-dogfish shark	26
Fruit fly-pigeon	25
Fruit fly-screwworm fly	2
Fruit fly-silkworm moth	15
Fruit fly-tobacco hornworm moti	14
Fruit fly-wheat	47

Figure 3

Observations Data Table 1

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Data Table 2

Organisms	Number of Amino Acid Differences	Positions in Which They Vary
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Human and Gorilla		
Human and Rhesus Monkey		
Human and Horse		
Human and kangaroo		

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aı	alysis and Conclusions A. Use Figure 1 to answer questions 1 and 2.
	On the basis of the hemoglobln similarity, what organisms appear to be most closely related to humans? Explain your answer.
	Among the organisms that you compared, which one appears to be least closely related to humans? Explain your answer.
	on the basis of differences in their cytochrome c, which organisms appear to be most closely related to humans?
	Which organisms appear to be least closely related to humans?
·-	Check the pair of organisms that appears to be most closely related to each other snapping turtle – tuna snapping turtle – rattlesnake snapping turtle – pigeon
	Give a reason for your answer.
9	
5.	Agree or disagree with the following statement and give reasons to support your answer. "Fruit files appear to be more closely related to silkworm moths than they are to screwworm flies."
- 7.	Name the pair of organisms that appears to be equally related to humans on the basis of cytochrome c similarity.