

# 7.012 Quiz 1 Answers

<b>A ≥ 87</b>	~25% of test takers
<b>B ≥ 73</b>	~34% of test takers
<b>C ≥ 60</b>	~24.3% of test takers
<b>D ≥ 50</b>	~10.1% of test takers
<b>F ≥ 49</b>	~6.5% of test takers

Regrade requests (with a note attached indicating the problem and part you want looked at)

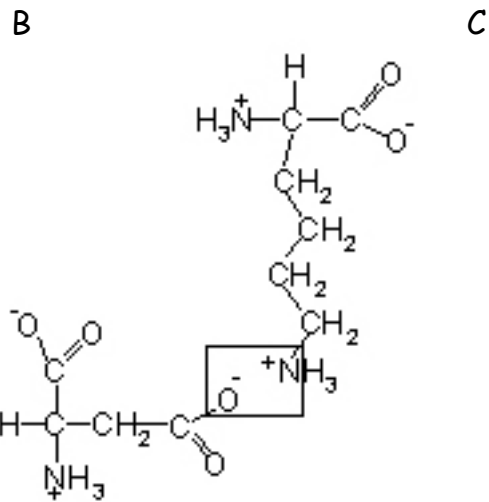
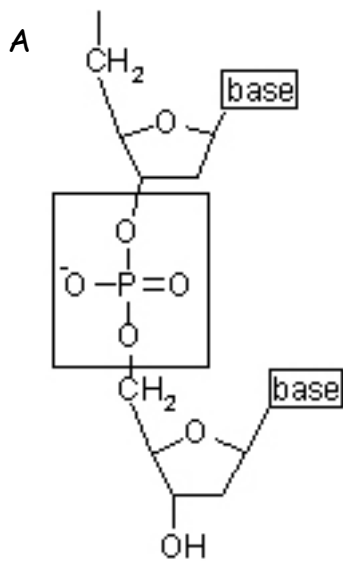
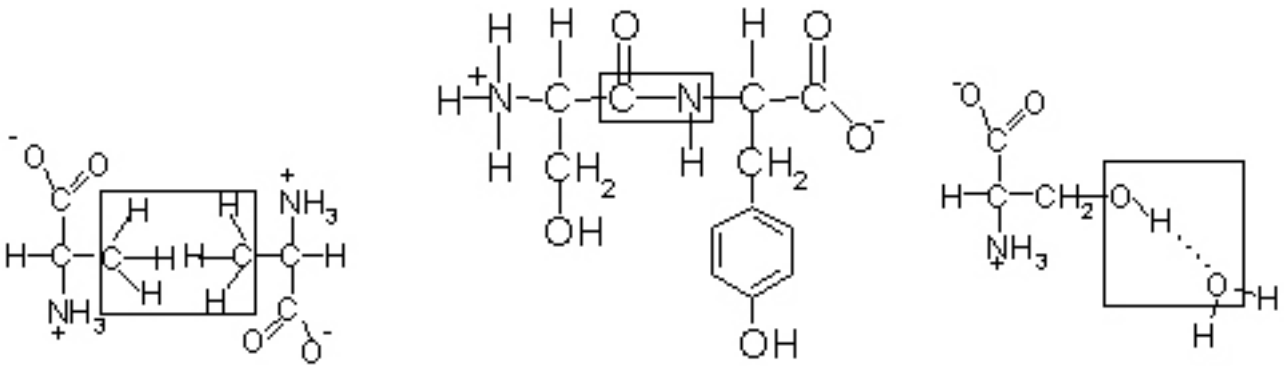
Monday 10/04/04

<b>Question</b>	<b>Value</b>	<b>Score</b>
<b>1</b>	<b>16</b>	
<b>2</b>	<b>20</b>	
<b>3</b>	<b>16</b>	
<b>4</b>	<b>16</b>	
<b>5</b>	<b>15</b>	
<b>6</b>	<b>17</b>	
	<hr/> <b>100</b>	

**Question 1**

a) Match each boxed chemical interaction in the structures below with the appropriate description. 10 pts

- \_\_\_\_\_ B \_\_\_\_\_ peptide bond
- \_\_\_\_\_ E \_\_\_\_\_ ionic bond
- \_\_\_\_\_ A \_\_\_\_\_ van der Waal's interaction
- \_\_\_\_\_ D \_\_\_\_\_ phosphodiester bond
- \_\_\_\_\_ C \_\_\_\_\_ hydrogen bond



**D**

**E**

Name \_\_\_\_\_

b) Which one of the following structural features is NOT based on hydrogen bonds?  
Circle all that apply. 3 pts

i) complementary base pairing in DNA

ii) protein primary structure

iii) protein secondary structure

iv) complementary base pairing in RNA

c) Your friend tried to remove some writing on a plastic box. He used a napkin dampened with water, which did not work. Then as you advised, he used ethanol (instead of water), and successfully removed the writing. Circle why ethanol was **better** than water in this case. 3 pts

**[formula of ethanol:  $\text{CH}_3\text{-CH}_2\text{-OH}$ ]**

i) Ethanol has a greater surface tension, so it extracts the ink molecules better from the plastic.

ii) Ethanol can form hydrogen bonds with the ink molecules.

iii) One end of ethanol is non-polar. This makes ethanol a better solvent for the ink molecules.

iv) Ethanol is more hydrophilic, so it washes off the writing more efficiently.

**Question 2**

**a) Fill in the percentages. If the value can't be determined write "ND". 6 points**

- i) According to Chargaff's rule, if the DNA content of a cell was composed of 15% A, then C would make up 35%, G would make up 35%, and T would make up 15% % of the cell's DNA.
- ii) If the RNA content of a cell was composed of 15% A, then C would make up ND%, G would make up ND%, and T would make up 0% % of the cell's DNA.

**b) Match the following with all that apply. 6 points**

- |            |                              |  |
|------------|------------------------------|--|
| <u>ace</u> | Lactase                      | A. Lowers the activation energy of a reaction      |
| <u>ag</u>  | ribozyme                     | B. Raises the activation energy of a reaction      |
| <u>f</u>   | "The Transforming Principle" | C. Breaks down milk sugar to galactose and glucose |
|            |                              | D. Lipid   |
|            |                              | E. Protein   |
|            |                              | F. DNA   |
|            |                              | G. RNA   |

**c) Fill in the blanks. 2 pts each**

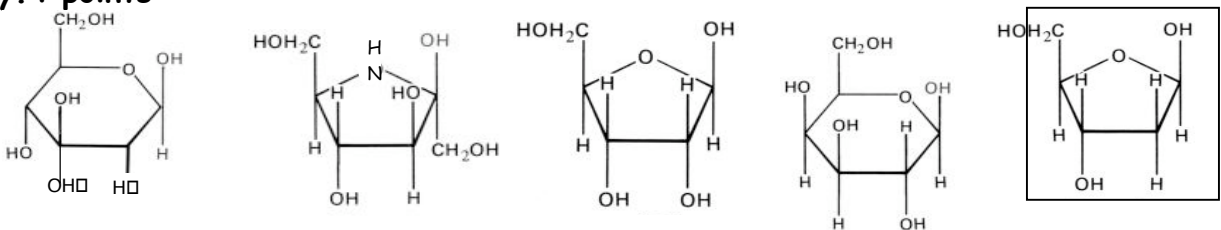
i) DNA is synthesized in the \_\_\_\_\_ direction. Circle one.

- C→N      3'→5'      O→P      N→C      5'→3'      P→O

ii) Proteins are synthesized in the \_\_\_\_\_ direction. Circle one.

- C→N      3'→5'      O→P      N→C      5'→3'      P→O

**d) Which of the following is used to make deoxyribonucleic acid? Circle all that apply. 4 points**



**Question 3**

Eukaryotes have proteins called histones that are involved in tightly packing DNA. Below is a schematic of the interaction between a molecule of DNA and the surface of a histone protein. "X" represents an amino acid side chain sticking out from the surface of the protein towards the DNA backbone.

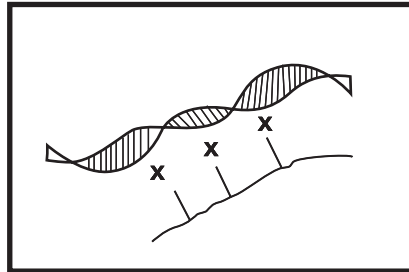


Figure by MIT OCW.

a) Circle the type(s) of amino acids you would expect to find with side chains in the positions marked "X"?

negatively charged

non-polar

positively charged

hydrophobic

2 pts.

b) Circle the **strongest** interaction between the DNA backbone and the histone protein.

hydrogen

ionic

van der Waals

X-linked

2 pts

Eight histone proteins function as subunits in a multi-protein complex called a nucleosome. Portions of two subunits (HA and HB) interact in the core of the nucleosome. The figure below shows the  $\alpha$ - helices where these two proteins interact.

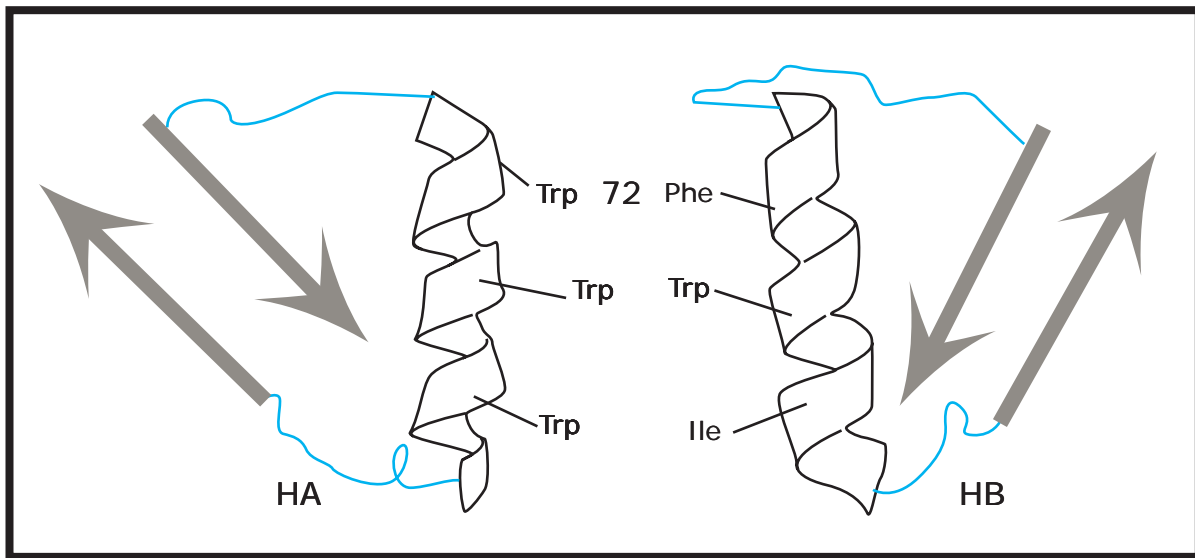


Figure by MIT OCW.

c) Based on the amino acids labeled in the diagram, what interactions keeps HA and HB together? Circle all that apply. 3 pts

covalent

hydrogen

hydrophobic

ionic

van der Waals

d) If tryptophan 72 mutates to become an arginine residue, indicate how the interaction between HA and HB would change. 2 pts.

becomes stronger

becomes weaker

remains the same

can not be determined

e) Explain your answer in d) in twelve (12) words or less. 3 pts

**Positive charged doesn't interact with hydrophobic side chain.**

f) Based on the information given, circle **all** levels of structure possessed by histones within a nucleosome. 4 pts

primary

secondary

tertiary

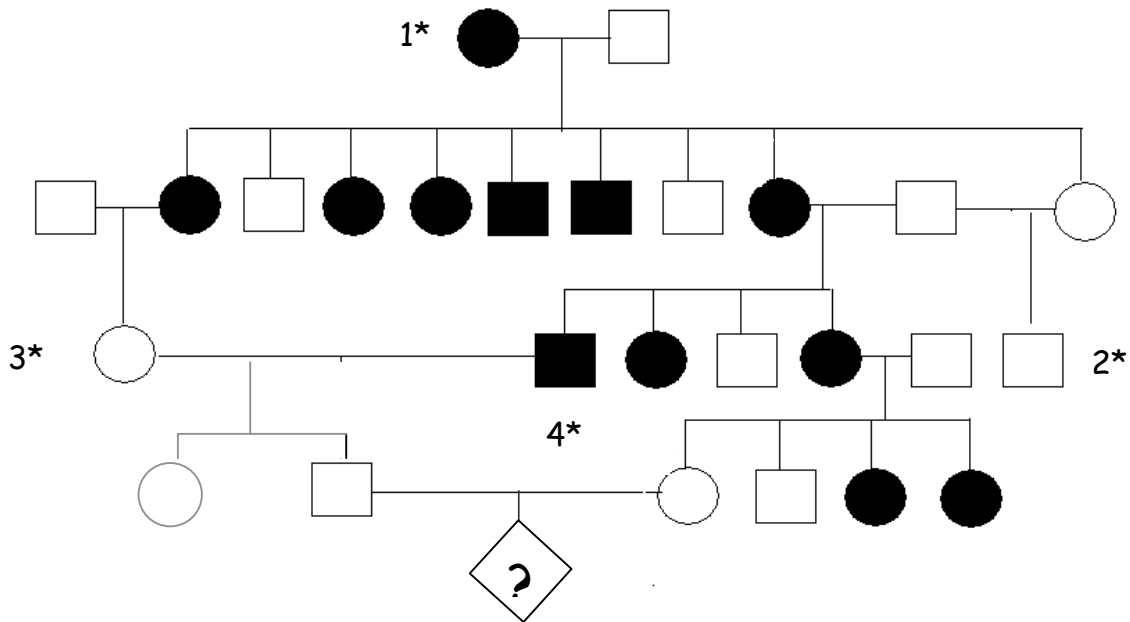
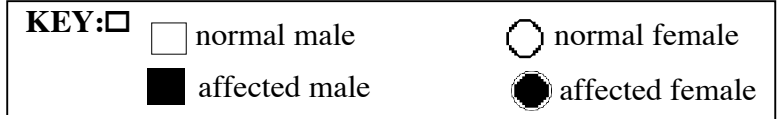
quaternary

quinternary



**Question 4**

Polydactyly is a trait where an individual develops more than five fingers per hand or toes per foot. The pattern of inheritance of polydactyly is shown in the following pedigree. Assume that all individuals marrying in from outside the family do not carry a disease allele. Assume complete penetrance.



a) What is the mode of inheritance of this trait? Circle all that apply. 4 pts

- Autosomal       Dominant       Mitochondrial       Recessive  
 Sex-limited       X-linked       Y-linked

b) What are the genotypes of the numbered individuals? If more than one genotype is possible, write all. Only show the genotype relating to the disease phenotype. Use the letters D, d, X<sup>D</sup>, X<sup>d</sup>, Y<sup>D</sup>, Y<sup>d</sup>, M<sup>D</sup> or M<sup>d</sup> to indicate the genotype corresponding to the polydactyly alleles. 8 pts

1\*     Dd          2\*     dd          3\*     dd          4\*     Dd    

c) What is the probability that the unborn individual marked with (?) will have polydactyly, 4 pts  
 if it were male?     0%          if it were female?     0%



**Question 5**

You are studying flies with different eye colors, so you cross them to determine the mode of inheritance. **Using standard nomenclature**, fill in the blanks with all possible genotypes of the flies described. Use the letters B, b, G, g, H, h, A, a, R and/or r to designate the alleles. If sex chromosomes are involved, use the format  $\rightarrow X^B, X^b, X^G, X^g, Y^B, Y^b$ , etc. 7 pts

a)  $F_0$ : Black-eyed flies \_\_\_\_\_  $gg$  \_\_\_\_\_ x Gray-eyed flies \_\_\_\_\_  $Gg$  \_\_\_\_\_



$F_1$  253 Black-eyed flies \_\_\_\_\_  $gg$  \_\_\_\_\_

228 Gray-eyed flies \_\_\_\_\_  $Gg$  \_\_\_\_\_

Gray-eyed flies x Gray-eyed flies ( $F_1$  flies)



$F_2$  138 Black-eyed flies \_\_\_\_\_  $gg$  \_\_\_\_\_

284 Gray-eyed flies \_\_\_\_\_  $Gg$  \_\_\_\_\_

124 Hazel-eyed flies \_\_\_\_\_  $GG$  \_\_\_\_\_

b)  $F_0$ : Amber-eyed male flies \_\_\_\_\_  $X^rY$  \_\_\_\_\_ x Red-eyed female flies \_\_\_\_\_  $X^RX^R$  \_\_\_\_\_ 8 pts



$F_1$ : All red-eyed \_\_\_\_\_  $X^RX^r, X^RY$  \_\_\_\_\_

Red-eyed male  $F_1$  x Red-eyed female  $F_1$



$F_2$  354 Red eyed males \_\_\_\_\_  $X^RY$  \_\_\_\_\_

355 Amber-eyed males \_\_\_\_\_  $X^rY$  \_\_\_\_\_

706 Red-eyed females \_\_\_\_\_  $X^RX^R, X^RX^r$  \_\_\_\_\_

### Question 6

You have a strain of yeast that glows purple when compound A is added to the medium. You suspect that compound A is a precursor of the "glowing" pathway. You mutagenize the original haploid strain and isolate six mutants that do not glow purple in the presence of compound A.

Since haploid yeast can be mated to make diploids, you mate each mutant strain with your original WT purple yeast to determine whether the mutant phenotypes are dominant or recessive. You also mate each mutant strain with each other to see which strains can complement each others' phenotypes. The results are summarized below.

("+" means colonies glow purple, "-" means they do not glow purple)

Strains Mated	m1	m2	m3	m4	m5	m6	WT
m1	-	-	+	+	+	-	+
m2		-	+	+	+	-	+
m3			-	+	-	+	+
m4				-	+	+	+
m5					-	+	+
m6						-	+

a) Are all the mutant phenotypes recessive?  Yes.  No. 2 pts.

If no, indicate the mutants having dominant phenotypes.

b) Place the mutants into complementation groups. 6 pts

m1 m2 m6

m3 m5

m4

c) How many genes have you identified that are involved in this pathway? 2 pts

**3**

In the presence of compound A, the different mutants glow different colors (other than purple). You suspect that these colors may be due to a buildup of different compounds in a biochemical pathway. To figure out this novel biochemical pathway, you make the following observations.

- °When you add compound A to mutant 1, the strain accumulates compound R and glows red.
- °When you add compound A to mutant 4, the strain accumulates compound B and glows blue.
- °When you add either compound B or R to Mutant 5, the strain glows purple.
- °When you add compound A to a double-mutant yeast strain having both Mutation 2 and Mutation 4, it glows blue.

Based on these results solve the biochemical pathway.

Circle the model that is consistent with your epistasis data. Note that "Enz 1" denotes the enzyme encoded by the gene mutated in the m1 strain.

<b>Model 1</b>	Compound A	Enz 5	Compound B	Enz 4	Compound R	Enz 6	Purple
Model 2	Compound R	Enz 2	Compound B	Enz 5	Compound A	Enz 4	Purple
Model 3	Compound A	Enz 4	Compound R	Enz 2	Compound B	Enz 5	Purple
Model 4	Compound A	Enz 1	Compound R	Enz 5	Compound B	Enz 4	Purple
Model 5	Compound A	Enz 5	Compound R	Enz 4	Compound B	Enz 3	Purple
Model 6	Compound A	Enz 3	Compound R	Enz 6	Compound B	Enz 4	Purple

7 pts