



FACULTY OF SCIENCE

DEPARTMENT OF BIOTECHNOLOGY AND FOOD TECHNOLOGY

DIPLOMA IN BIOTECHNOLOGY

MODULE BIC12B1/BIC21B1

Biochemistry 2

CAMPUS DFC

NOVEMBER MAIN EXAMINATION 2019

DATE: 21/11/2019

SESSION: 16:30 – 18:30

ASSESSOR(S):

DR K KONDAH

INTERNAL MODERATOR

MR. L. ALAGIOZOGLOU

DURATION: 2 HOURS

MARKS: 75

NUMBER OF PAGES: 9

INSTRUCTIONS TO INVIGILATORS:

1. CALCULATORS ARE PERMITTED (ONLY ONE PER STUDENT).
2. THIS QUESTION PAPER MUST BE RETURNED WITH SCRIPT.
3. QUESTION 1 TO BE ANSWERED IN THE MCQ SECTION OF ANSWER SCRIPT.

INSTRUCTIONS TO STUDENTS:

1. ANSWER ALL THE QUESTIONS SHOWING ALL CALCULATIONS WHERE NECESSARY.
 2. RETURN EXAMINATION PAPER WITH SCRIPT.
 3. ANSWER QUESTION 1 IN MCQ SECTION OF THE ANSWER SCRIPT.
 4. ANSWER QUESTIONS 2-4 IN THE MAIN SECTION OF ANSWER SCRIPT.
 5. STANDARD GENETIC CODE, CHART OF AMINO ACIDS AND pKa VALUES FOR AMINO ACIDS ARE PROVIDED AT THE BACK OF THE ASSESSMENT.
 6. IT IS IN YOUR BEST INTEREST TO WRITE CLEARLY AND LEGIBLY
 7. GOOD LUCK!
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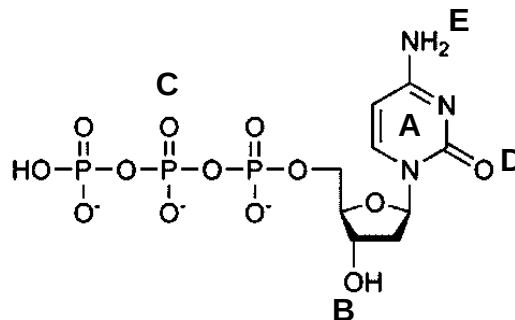
Question 1:

[25x1 = 25 marks]

Answer the multiple choice questions by clearly selecting the best option.

- 1.1. A single H₂O molecule binds to _____ other H₂O molecules in the lattice structure of ice
 - a) 7
 - b) 5
 - c) 4
 - d) 3
 - e) 2
- 1.2. How much of a 0.55 M solution of NaOH is required to make 15 mL of a 0.2 M solution of NaOH?
 - a) 41.25 mL
 - b) 41.25 µL
 - c) 41.25 L
 - d) 5.45 µL
 - e) 5.45 mL

Study the biomolecule below and answer questions 1.3 and 1.4.



- 1.3. This is an example of a
 - a) Nucleic acid
 - b) Deoxyribonucleotide
 - c) Ribonucleotide
 - d) Deoxyribonucleoside
 - e) Ribonucleoside
- 1.4. What is the name of structure A?
 - a) Thymine
 - b) Cytosine
 - c) Uracil
 - d) Adenine
 - e) Guanine
- 1.5. Molecules that carry charged groups of opposite polarity are known as
 - a) Racemic
 - b) Hydropathic
 - c) Enantiomers
 - d) Chiral
 - e) Zwitterions
- 1.6. Which amino acid does not belong in the group?
 - a) G
 - b) E
 - c) R
 - d) D
 - e) K

- 1.7. Which of the following statements best reflects the properties of enzymes?
- a) Enzymes lower the activation energy of a reaction to speed it up.
 - b) Enzymes can change an endothermic reaction to an exothermic one.
 - c) Enzymes alter the equilibrium constant of a reaction.
 - d) Enzymes are non-specific and can bind any molecule.
 - e) Enzymes are globular molecules made of polysaccharides.
- 1.8. If the pH of physiological saline is 7.4, what is the concentration of H^+ ?
- a) 4×10^{-8}
 - b) 4×10^8
 - c) 2.5×10^7
 - d) 2×10^{-9}
 - e) 2×10^9

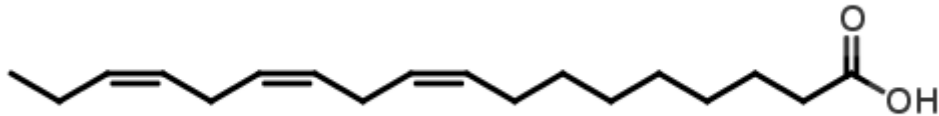
Use the following sequence to answer questions 1.9 and 1.10.

3'-AATGCTTAAGGTCAAGGC-5'

- 1.9. Which restriction enzyme will cut this sequence?
- a) *Sma*I (CCC*GGG)
 - b) *Hind*III (A*AGCTT)
 - c) *Pst*I (CTGCA*G)
 - d) *Bam*HI (G*GATTC)
 - e) *Eco*RI (G*AATTC)
- 1.10. After electrophoresis, _____ DNA fragments will be visible on the gel.
- a) 1
 - b) 2
 - c) 3
 - d) 4
 - e) 5
- 1.11. Which of the following is responsible for the ABO blood groups?
- a) Glycolipid
 - b) Polysaccharide
 - c) Peptide
 - d) Palmitic acid
 - e) Peptidoglycan
- 1.12. Which of the following is NOT a characteristic of lipids?
- a) They can form polymers
 - b) They are hydrophobic
 - c) They are soluble in methanol
 - d) They are sources of energy
 - e) They are composed of fatty acids
- 1.13. Codons are found in
- a) mRNA
 - b) rRNA
 - c) tRNA
 - d) 5'-3' DNA
 - e) 3'-5' DNA
- 1.14. A bond formed between glycine and tyrosine is a
- a) Phosphodiester bond
 - b) Hydrogen bond
 - c) Peptide bond
 - d) Glycosidic bond
 - e) Van der Waal's force

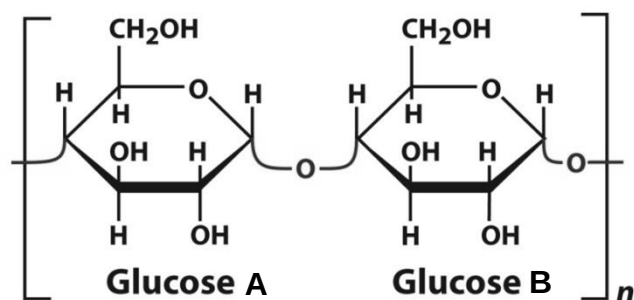
- 1.15. A buffer is a
- solution consisting of a strong acid and a strong base that resists large changes in pH.
 - solution consisting of a strong acid and a strong base that resists small changes in pH.
 - solution consisting of a weak acid and its conjugate base that resists large changes in pH.
 - solution consisting of a weak acid and its conjugate base that resists small changes in pH.
 - neutral solution.
- 1.16. Semi conservative DNA replication results in progeny where
- One strand is from the parent and one strand is from the daughter
 - Both strands are from the parent
 - Both strands are from the daughter
 - One strand is from DNA and one strand is from RNA
 - One strand is the original template and one strand is mutated

Study the biomolecule below and answer questions 1.17 and 1.18.



- 1.17. The biomolecule is
- Saturated
 - Monounsaturated
 - Polyunsaturated
 - Hydrophilic
 - A beta-sheet
- 1.18. The systematic name for the biomolecule is
- 18:3 n-3
 - 18:3 n-9
 - 16:3 n-9
 - 16:3 n-3
 - 17:3 n-3
- 1.19. Glutathione is a _____ that helps inactivate oxidative compounds capable of causing cellular damage.
- Glycoprotein
 - Lipid
 - Nucleic acid
 - Tripeptide
 - Enzyme
- 1.20. The helices in collagen form a _____ coil; the helices in keratin form a _____ coil.
- Left-handed; left-handed
 - Right-handed; left-handed
 - Right-handed; right-handed
 - Left-handed; right-handed
 - Random; structured

Study the polysaccharide provided below and answer questions 1.21 to 1.23.



- 1.21. The glucose is represented as a
 - a) Chair conformation
 - b) Linear conformation
 - c) Furanose ring
 - d) Helix ring
 - e) Pyranose ring
- 1.22. What is the anomeric form of Glucose A?
 - a) α
 - b) β
 - c) D
 - d) L
 - e) N
- 1.23. Which of the following is the correct systematic name for the disaccharide shown above?
 - a) β -D-glucopyranosyl-(1 $\rightarrow$ 4)- β -D-glucopyranoside
 - b) α -D-glucopyranosyl-(1 $\rightarrow$ 4)- α -D-glucopyranoside
 - c) α -D-glucopyranosyl-(1 $\rightarrow$ 4)- α -D-glucopyranoside
 - d) α -D-glucopyranosyl-(1 $\rightarrow$ 4)- β -D-glucopyranoside
 - e) β -D-glucopyranosyl-(1 $\rightarrow$ 4)- α -D-glucopyranoside
- 1.24. Sucrose is a non-reducing sugar because
 - a) The anomeric carbon on fructose is not involved in a glycosidic bond.
 - b) The anomeric carbon on glucose is not involved in a glycosidic bond.
 - c) The anomeric carbons on both glucose and fructose are not involved in a glycosidic bond.
 - d) The anomeric carbons on both glucose and fructose are involved in a glycosidic bond.
 - e) C6 on both glucose and fructose are involved in a glycosidic bond.
- 1.25. The complementarity between enzyme and substrate is known as _____ model.
 - a) Pins and needles
 - b) Soap and water
 - c) Shoes and socks
 - d) Needle and thread
 - e) Lock and key

Question 2:**[12 marks]**

Study the DNA sequence below and answer the questions that follow.

3'- AGG CTG TTT TGC AAA TCT -5'

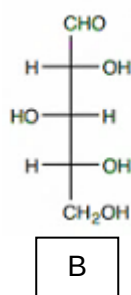
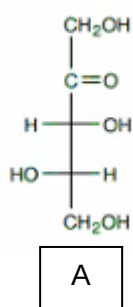
- 2.1. Write out the sequence of the complimentary DNA strand indicating direction. (1)
- 2.2. Write out the sequence of the mRNA strand indicating direction. (1)
- 2.3. What is the G+C content of this DNA molecule? Show your calculations. (2)
- 2.4. List two differences between DNA and RNA. (2)
- 2.5. Translate the mRNA sequence using the one letter amino acid abbreviated form clearly indicating termini. (2)
- 2.6. Group the amino acids in question 2.5 above by polarity. (3)
- 2.7. How many of these amino acids have ionisable side chains? (1)

Question 3:**[22 marks]**

- 3.1. A buffer solution was made by dissolving 75 g $\text{C}_2\text{H}_5\text{COOH}$ (propionic acid) in 600 mL of 0.5 M $\text{C}_2\text{H}_5\text{COONa}$. To this buffer is added 0.025 moles of NaOH. Assume the change in volume when the NaOH is added is negligible. The K_a of $\text{C}_2\text{H}_5\text{COOH}$ is 1.30×10^{-5} . Calculate the pH of the buffer after the addition of the NaOH. M_r Na = 23, C = 12, O = 16, H = 1. Show all calculations to obtain full marks. (7)
- 3.2. Draw a clear ionisation pattern from acid to base for tyrosine and cysteine indicating the pH for each ionisation. (11)
- 3.3. Calculate the pI for both tyrosine and cysteine. Show all your calculations. (2)
- 3.4. Comment on the charge of both tyrosine and cysteine if they were placed in the buffer solution from question 3.1. (2)

Question 4:**[16 marks]**

4.1. Study the molecules below and answer the questions that follow.



- 4.1.1. Classify molecules A and B according to the Fischer convention. (2)
- 4.1.2. Classify molecules A and B according to their ring structure should they cyclise. (2)
- 4.2. Draw a table comparing cellulose and chitin. (8)
- 4.3. Write brief notes on cholesterol in animals and its role in the cell membrane. (4)

STANDARD GENETIC CODE

		2nd base					
1st base	U	U	C	A	G	3rd base	
		UUU Phenylalanine	UCU Serine	UAU Tyrosine	UGU Cysteine		U
		UUC Phenylalanine	UCC Serine	UAC Tyrosine	UGC Cysteine		C
		UUA Leucine	UCA Serine	UAA Stop	UGA Stop		A
		UUG Leucine	UCG Serine	UAG Stop	UGG Tryptophan		G
	C	CUU Leucine	CCU Proline	CAU Histidine	CGU Arginine		U
		CUC Leucine	CCC Proline	CAC Histidine	CGC Arginine		C
		CUA Leucine	CCA Proline	CAA Glutamine	CGA Arginine		A
		CUG Leucine	CCG Proline	CAG Glutamine	CGG Arginine		G
	A	AUU Isoleucine	ACU Threonine	AAU Asparagine	AGU Serine		U
		AUC Isoleucine	ACC Threonine	AAC Asparagine	AGC Serine		C
		AUA Isoleucine	ACA Threonine	AAA Lysine	AGA Arginine		A
		AUG Methionine (Start)	ACG Threonine	AAG Lysine	AGG Arginine		G
	G	GUU Valine	GCU Alanine	GAU Aspartic Acid	GGU Glycine		U
		GUC Valine	GCC Alanine	GAC Aspartic Acid	GGC Glycine		C
		GUA Valine	GCA Alanine	GAA Glutamic Acid	GGA Glycine		A
		GUG Valine	GCG Alanine	GAG Glutamic Acid	GGG Glycine		G

TABLE OF pKa VALUES FOR AMINO ACIDS

Amino Acid	pKa Value		
	Name	Alpha Carboxy	+Alpha Amino Side Chain
Glycine		2.34	9.60
Alanine		2.34	9.69
Valine		2.32	9.62
Leucine		2.36	9.60
Isoleucine		2.36	9.68
Methionine		2.28	9.21
Phenylalanine		1.83	9.13
Tryptophan		2.38	9.39
Proline		1.99	10.60
Serine		2.21	9.15
Threonine		2.63	9.10
Cysteine		1.71	10.78
Tyrosine		2.2	9.11
Asparagine		2.02	8.84
Glutamine		2.17	9.13
Aspartic Acid		2.09	9.82
Glutamic Acid		2.19	9.67
Lysine		2.18	8.95
Arginine		2.17	9.04
Histidine		1.82	9.17

AMINO ACIDS

