

Roll No: _____

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M.Sc. Biochemistry Semester II Examination, May 2024

Paper Title: Protein & Proteomics
Paper Code: MBC 202

Time: 2½ h

Max Marks: 60

Note: This paper has 3 Sections, A, B, and C. Attempt all three sections. Section A: Answer all the questions; Section B: Answer any two; Section C: Answer all the questions, each question in this section has an internal choice.

Abbreviations Used: CLO- Course Learning Outcome, PLO- Program Learning Outcome, BTL- Bloom Taxonomy Level; BT level 1- Remember; BT level 2-Understand; BT level 3-Apply; BT level 4-Analyse; BT level 5-Evaluate, BT level 6-Create.

Section A: Attempt all questions.

(1x12=12 Marks)

Q No.1	Questions (Fill in the blanks/True and False/short answers)	BTL	CLO	PLO
i.	Proteins are synthesized from _____ templates.	1	1	3
ii.	The folding of a protein into its functional shape is guided by its _____ structure.	1	3	3
iii.	X-ray crystallography technique is used to determine the _____ structure of proteins.	3	4	2
iv.	_____ proposed a hypothesis that states that a protein's native structure is determined solely by its amino acid sequence.	4	2	3
v.	The function of peptidyl prolyl isomerase is _____.	2	1	2
vi.	_____ refers to the specific three-dimensional arrangement of amino acids in a protein molecule.	5	5	3
vii.	Protein denaturation disrupts the _____ interactions that maintain a protein's native structure.	2	2	3
viii.	Mass spectrometry is a technique used to analyze the _____ of proteins.	3	3	3
ix.	The presence of alpha-helices, beta-sheets, and loops in a protein constitutes its tertiary structure. (True/False)	4	1	3
x.	The secondary structure of a protein is determined by the sequence of amino acids. (True/False)	1	2	3
xi.	Name one amino acid with sulfhydryl group.	3	3	2
xii.	Define protein sequencing.	2	1	3

Section B: LONG ANSWER QUESTIONS
Attempt any TWO out of four questions

(2x12=24 Marks)

Q No. 2	Questions	BTL	CLO	PLO
a)	Explain the significance of protein sequencing in understanding protein structure and function. Discuss any two techniques used for protein sequencing and their advantages and limitations.	1	1	
b)	Describe the structure and function of proteins, highlighting the relationship between protein structure and its biological activity. Include examples of different protein classes and their roles in cellular processes.	2	3	2
c)	Compare and contrast different techniques used in proteomics research, such as mass spectrometry, two-dimensional gel electrophoresis, and protein microarrays. Discuss the strengths, limitations, and applications of each technique.	3	2	3
d)	Explain how protein misfolding can lead to the formation of amyloid fibrils and the development of neurodegenerative diseases.	4	4	2

Section C: SHORT ANSWER QUESTIONS

Attempt ALL questions, each question has an internal choice

(6x4=24 Marks)

Q. No 3	Questions	BTL	CLO	PLO
a)	Explain the concept of protein denaturation and describe the various factors that can induce protein denaturation. OR Discuss the role of molecular chaperones in protein folding and quality control.	4	3	2
b)	What are the advantages of using 2D-PAGE in proteomics research? OR Describe the steps involved in protein purification and characterization.	1	1	3
c)	Discuss the structural features of the peptide bond and its implications for protein conformation. OR Explain how the linear sequence of amino acids dictates the folding pathway and conformational stability of a protein.	4	5	2
d)	Name and write the principle of the technique used to identify proteins and protein expression pattern in biological samples. OR Compare and Contrast the structure of hemoglobin with myoglobin.	3	2	3
e)	Define secondary structure in proteins and describe the common elements of secondary structure, such as alpha helices and beta sheets. OR Discuss the structural motifs found in secondary structures and their role in stabilizing protein conformation.	5	2	3
f)	Discuss the structural features of globular proteins and fibrous proteins, highlighting their differences in secondary and tertiary structures, as well as their biological functions and properties. OR Discuss the technique of MALDI TOF MS indicating its applications. How is this technique different from APIMS or ESI MS?	3	2	2

END OF THE QUESTION PAPER