

BIRLA INSTITUTE OF TECHNOLOGY, MESRA, RANCHI
(END SEMESTER EXAMINATION)

CLASS: MSC
BRANCH: BIOINFORMATICS AND COMPUTATIONAL BIOLOGY

SEMESTER : III
SESSION : MO/2025

SUBJECT: BI301 PROTEOMICS, METABOLOMICS AND BIOMARKER DESIGN

TIME: 3 Hours

FULL MARKS: 50

INSTRUCTIONS:

1. The question paper contains 5 questions each of 10 marks and total 50 marks.
 2. Attempt all questions.
 3. The missing data, if any, may be assumed suitably.
 4. Before attempting the question paper, be sure that you have got the correct question paper.
 5. Tables/Data hand book/Graph paper etc. to be supplied to the candidates in the examination hall.
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		CO	BL
Q.1(a)	Give a brief note on the physicochemical properties of the proteins.	[5] 1	1
Q.1(b)	Write the principle and working of SDS-PAGE technique, and explain how this technique helps in the characterization of proteins?	[5] 1	2
Q.2(a)	Explain how Fourier-Transform Infrared (FTIR) spectroscopy technique helps in protein structure determination.	[5] 2	3
Q.2(b)	What do you understand by NMR spectroscopy. Give the significance of isotopic labeling in protein nuclear magnetic resonance (NMR).	[5] 2	3
Q.3(a)	Describe methods and applications of protein sequencing.	[5] 2	2
Q.3(b)	What do you understand by peptide mass fingerprinting (PMF). Give different steps of this technique.	[5] 2	2
Q.4(a)	What are primary and secondary metabolic pathways. Compare between KEGG and MetaCyc pathway databases.	[5] 4	2
Q.4(b)	Write a short note on precision medicines.	[5] 4	1
Q.5(a)	Define biomarkers. Outline the key characteristic features that make a biomarker more reliable and efficient.	[5] 5	2
Q.5(b)	Explain omics-based biomarker designs.	[5] 5	2

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