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(ii) Questions :9 Sub. Code : 0 9 9 0 Exam. Code : 0 0 3 8

B.Sc. (Hons.) Biotechnology 6th Semester (2054)

GENOMICS AND PROTEOMICS

Paper: BIOT-604-T

Time Allowed: Three Hours] [Maximum Marks: 67

Note:—FIVE questions are to be attempted. Question No. 1 is compulsory. Attempt ONE question from each Unit.

1. Answer briefly:

- (1) UCSC Genome Browser
- (2) Hydrophobicity
- (3) Sedimentation Analysis
- (4) Resolution of 2D PAGE
- (5) β-mercaptoethanol
- (6) Automated sequencing
- (7) Mass spectrometry
- (8) SGD
- (9) Old NTPs
- (10) Clone Contig Method.

 $1.5 \times 10 = 15$

UNIT—I

2.	(a)	Write a note on Maxam and Gilbert methosequencing.	od o:
	(b)	Discuss the principle of pyrosequencing.	6
3.	07-73	te a note on the Genome sequence asse wares.	mbly 13
		UNIT—II	
4.	Wri	te notes on :	
	(a)	ENSEMBL	
	(b)	NCBI genome.	13
5.	(a)	Discuss any tool for genome analysis.	6
	(b)	Explain the Mouse Genome Database.	7
		UNIT—III	
6.	(a)	Explain Edman degradation for protein sequencing	;. 6
	(b)	Discuss any technique for determination of prisizes.	otein 7
7.		ite a note on the physical interactions that determin perty of proteins.	e the
		UNIT—IV	
8.	(a)	Discuss any method for analysis of proteome.	6
	(b)	Explain de novo sequencing using mass spectrom data.	etric 7
9.	Wr	ite notes on :	
	(i)	Sample preparation of 2D PAGE	
	(ii)	Reproducibility of 2D PAGE.	13