(i)	Printed Pages: 2	Roll No

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

B.Sc. (Hons.) Biotechnology 6th Semester (2053)

GENOMICS AND PROTEOMICS Paper: BIOT-604-T

Time Allowed: Three Hours] [Maximum Marks: 67

Note: — Attempt five questions in all, including question no. 1 which is compulsory and taking one from each Unit. All questions carry marks as indicated.

- (a) Define DNA.
 - (b) Discuss SNP.
 - (c) What is ENSEMBL?
 - (d) Write two advantages of 2 D PAGES.
 - (e) Write two Model organisms for human.
 - (f) What is "NCBI Browser"?
 - (g) What is genetic map?
 - (h) Define Genome.
 - (i) Discuss non covalent bonds,
 - (j) What are covalent linkages?

 $1.5 \times 10 = 15$

UNIT—J

2.	(a)	Discuss the Automated Genome sequencing and its importance	
	(b)	What is a Shotgun method for sequencing?	6,7
3.	(a)	What is a Maxam & Gilbert sequencing method?	
	(b)	Write a note on limitations of human genome.	6,7
		UNIT—II	
4.	(a)	Write a note on manage and distribution genome.	
	(b)	Discuss UCSC database and its applications.	6,7
5.	(a)	Write a note on web server for genome analysis.	
	(b)	Write a note on ENSEMBL and its significance.	6,7
		UNIT—III	
6	(a)	Write a note on chemical interaction define in protein stru	cture.
	(b)	Write a note on short range interaction.	6,7
7.	(a)	Discuss about Gel filtration and its applications.	
	(b)	Discuss SDS — PAGE and its uses.	6,7
		UNIT—IV	
8.	(a)	Write a note on Proteome and its significance.	
	(b)	Write a note on 2D-PAGE and its applications.	6,7
9.	(a)	Discuss about the mass spectroscopy based metho- protein identifications.	ds for
	(b)	Discuss De novo sequencing using mass spectrometric	e data.
			6,7