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# M.Sc. (Bio-Informatics) Fourth Semester MBIN-8017: Genomics and Proteomics – II

## Time allowed: 3 Hours

#### Max. Marks: 60

**NOTE:** Attempt <u>five</u> questions in all, including Question No. I which is compulsory and selecting atleast one question from each Unit.

x - x - x

I. Answer the following:-

- a) Name any toe diseases due to SNPs.
- b) What are VNTRs? Give suitable example.
- c) Give full forms of DIP and STRING.
- d) What is gene fusion?
- e) Differentiate between Pfam A and Pfam B.
- f) What is hybridoma technology?
- g) Briefly explain FRET.
- h) Differentiate between polymorphism and mutation.  $(8x1\frac{1}{2})$

### <u>UNIT – I</u>

II.	a) What are SNPs? Discuss any one method for their detection.	
	b) Compare and contrast genetic, physical and cytological maps.	(6,6)
III.	Write notes on the following:-	
	a) RFLP	
	b) SSCP detection of SNPs	(6,6)
	<u>UNIT – II</u>	
IV.	a) How are protein-protein interactions detected using yeast 2- hybrid method?	
	b) Discuss the technique of page display and give its applications.	(6,6)
V.	a) How can Mass spectroscopy be used to characterize protein complexe	s?
	b) Write a note on genome wide protein interaction studies.	(6,6)

### <u>UNIT – III</u>

- VI. Write notes on the following:
  - a) MUMMER
  - b) Hydropathy plot
  - c) SCOP

(3x4) P.T.O.

(6,6)

VII. a) How is fold recognition used to predict protein structure?b) Discuss BLASTZ and how does it differ from BLAST.

### *x*-*x*-*x*