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MBOI-004

M.S. IN BIOTECHNOLOGY (MSBOMM/MSBOCC)

Term-End Examination

00214

December, 2014

MBOI-004 : APPLICATIONS OF GENOMICS AND PROTEOMICS

Time: 3 hours

Maximum Marks: 100

Note:

- (i) Section I is compulsory.
- (ii) In Section II, answer any five questions.
- (iii) Assume suitable data wherever required.
- (iv) Draw suitable sketches wherever required.
- (v) Italicized figures to the right indicate maximum marks.

SECTION I

- 1. What are proteins? Define the physico-chemical nature of proteins. Give the different protein purification methods.

 2+4+4
- 2. Explain the following terms:

 5×2

- (a) Proteomics
- (b) Genomics

- (c) Stationary phase and mobile phase in Chromatography technique
- (d) Probe and Target Sequence in a Microarray experiment
- (e) STS
- **3.** Explain the whole genome shotgun sequencing method with the help of relevant schematic diagrams.

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SECTION II

- **4.** What is EST? Explain its importance in biomarker discovery research. Explain the SAGE method of measuring gene expression. 2+3+9
- 5. Describe Mass spectrometry and 2D gel Electrophoresis techniques in detail. Explain briefly the role of these two techniques in advancing proteomics research.

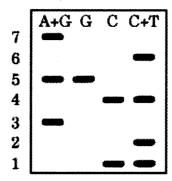
 7×2

6. Consider you are the principal investigator (PI) of a project concerned with the developmental biology of an insect which has four stages in its life cycle. Design a microarray experiment to identify the differentially expressed genes (DEGs) unique to each stage of the life cycle.

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7. Explain Maxam Gilbert DNA sequencing method. From the given auto-radiogram (Figure below), deduce the DNA sequence fragment. Also give the 5' and 3' ends of the fragment.

5+9



8. What are the protein 3D structure determination techniques? Write one limitation of each technique. Explain one of the techniques in detail.

- **9.** Describe the following:
 - (a) Size Exclusion Chromatography
 - (b) Affinity Chromatography
 - (c) Reverse Phase Chromatography

Tabulate the advantages and disadvantages of each technique. 4+4+4+2

10. Describe the process of sequence assembly.Explain any two methods of physical mapping used for the validation of sequence assembly.7+7