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## Sixth Semester B.E. Degree Examination, Dec.2017/Jan.2018 Genomics and Proteomics

Time: 3 hrs.

Max. Marks:100

Note: Answer FIVE full questions, selecting at least TWO questions from each part.

## PART - A

- 1 a. Define polymorphism. Explain different types of polymorphisms seeking human genome.
  (10 Marks)
  - b. Give a detailed historical note on sequencing methods. Explain how Sanger dideoxy method can be automated. (10 Marks)
- 2 a. Shot gun approach is the best method of DNA sequencing for whole genome sequencing.

  Justify the statement. (10 Marks)
  - b. Write a short note on followings
    - i) Human genome project
    - ii) Rice genome.

(10 Marks)

- 3 a. What are EST's? Give an explanatory note on their importance genome informatics.
  - b. Describe the functional genomics studies with model system Drosophila.

(10 Marks) (10 Marks)

4 a. Define Mutations and add an explanatory note on Regulating Transcription.

(08 Marks)

b. Write a short note on following:

i) SiRNA

- on following of
- ii) Post translational modification.

(12 Marks)

## PART - B

- 5 a. What is E values of genome? Explain the general architecture of prokaryotic genome.
  - (12 Marks)

b. Discuss the salient features of mitochondrial genome.

- (08 Marks)
- 6 a. List out the various molecular markers used in genome mapping. Explain how RFLP can be used as molecular marker in mapping.

  (08 Marks)
  - b. Write a short note on Transposon tagging.

(06 Marks)

c. Explain the role of micro array in functional genomics.

(06 Marks)

- 7 a. Write a short notes on:
  - i) Protein purification

(06 Marks)

ii) Merrifield synthesis of peptides

(06 Marks)

b. Write about Mass-Spec analysis of protein expression.

- (08 Marks)
- 8 a. Compare and construct the differences between DNA arrays and non-DNA arrays. (08 Marks)
  - b. Discuss on:
    - i) Stable isotype labeling

(06 Marks)

ii) Proteomics as tool for disease diagnostics

(06 Marks)

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