Exam. Code : 206603

Subject Code: 4620

# M.Sc. (Bio-Informatics) Semester—III ADVANCED ALGORITHMS FOR COMPUTATIONAL BIOLOGY

Paper — BI-632

Time Allowed—3 Hours [Maximum Marks—75

## SECTION-A

Section A is compulsory.

- Explain the following in not more than 4 lines:
  - (a) Complexity theory
  - (b) Data types
  - (c) Double Digest Problem
  - (d) Border block groups
  - (e) HMM
  - (f) Clone libraries
- (g) SVM
  - (h) Genetic algorithm
  - (i) Back propagation
  - (i) Rosetta stone.

 $10 \times 1.5 = 15$ 

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(Contd.)

## SECTION—B

Answer one question from each unit.

#### UNIT—I

- What do you understand by abstract data types? Explain their significance in algorithm designing.
- Illuminate the concepts and principles of algorithm designing.
  Describe various techniques of algorithm designing. 12

## UNIT-II

- 4. What do you understand by integer programming? Write a note on circular maps an map algorithm.
- 5. Illuminate various optimization techniques. 12

## UNIT—III

- 6. Describe the role played by HMM in identification and recognition of motifs in protein.
- Outline the process of cloning and generation of clone libraries. Compare and contrast the libraries of complete and partial digestion.

#### UNIT-IV

- 8. Outline significance and applications of support vector machines in computational biology.
- 9. What are various machine learning techniques? Explain how machine learning is useful in computational biological sciences.

## UNIT-V

- 10. Explain genetic algorithms and their benefits in bioinformatics.
- 11. Write a note on:
  - (a) Docking simulation
  - (b) Microarray data analysis.

6+6=12