P3145

## [5154]-710-A

**B.E.** (Information Technology)

**BIOINFORMATICS** 

# (2012 Course) (Semester -II) (Elecitve-IV) (414464A) (End Sem.)

*Time : 2½Hours] Instructions to the candidates:* 

- 1) Answers Q.1 or Q.2, Q.3 or Q4, Q.5 or Q.6, Q.7 or Q.8, Q.9 or Q.10.
- 2) Neat diagrams must be drawn wherever necessary.
- 3) Figures to the right indicate full marks.
- 2) Assume suitable data, if necessary.
- **Q1)** a) Explain central dogma of moleculer biology with neat diagram. [5]
  - b) What is structure visualization? Explain any two rendering tools in structure visualization. [5]

#### OR

- Q2) a) Define Bioinformatics. Explain Bioinformatics application related to following areas[5]
  - i) Proteomics.
  - ii) Sequence Assembly.
  - b) Explain microarray process spotting flow with neat diagram. [5]
- **Q3)** a) Explain knowledge discovery process with neat diagram. [5]
  - b) Write short note on:
    - i) PAM
    - ii) BLOSUM

### OR

| <b>Q4)</b> a) | Explain text mining with NLP process.                           | [5] |
|---------------|---|-----|
| b)            | Explain major steps in pattern recognition & discovery process. | [5] |

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[5]

SEAT No. :

[Total No. of Pages : 2

[Max. Marks : 70

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| <b>Q5)</b> a)  | Explain similarities & differences between BLAST & FASTA tools for sequence Alignment.            |  |
|----------------|---|--|
| b)             | Write short note on: [8]  |  |
|                | i) Heuristic methods for sequence Alignment.  |  |
|                | ii) Phlylogenetic.  |  |
|                | OR  |  |
| <b>Q6)</b> a)  | Explain FASTA algorithm with recommended steps for similarity searching. [8]                      |  |
| b)             | Explain working with BLAST. What are different services available from NCBT related to BLAST? [8] |  |
| <b>Q7)</b> a)  | Explain methods for protein modeling. [8]   |  |
| b)             | Explain comparative modeling process with neat diagram. [8]                                       |  |
|                | OR  |  |
| <b>Q8)</b> a)  | Write short note on: [8]  |  |
|                | i) Structural Bioinformatic in drug discovery.  |  |
|                | ii) Tools for modeling & simulation.  |  |
| b)             | Discuss components of modeling & simulation system along with the process. [8                     |  |
| <b>Q9)</b> a)  | Define genetic Engineering. Discuss current developments in genetic engineering. [9]              |  |
| b)             | What is Biotechnology? What is role of Bioinformatics in biotechnology? [9]                       |  |
|                | OR  |  |
| <b>Q10)</b> a) | How system biology play role in human health disease and future of medicine? [9]                  |  |
| b)             | Explain any two techniques of genetic engineering in detail. [9]                                  |  |
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