

Seat No.: \_\_\_\_\_

Enrolment No. \_\_\_\_\_

**GUJARAT TECHNOLOGICAL UNIVERSITY**  
**MCA - SEMESTER-V • EXAMINATION – SUMMER 2013**

**Subject Code: 650015**

**Date: 20-05-2013**

**Subject Name: Bioinformatics**

**Time: 02.30 pm - 05.00 pm**

**Total Marks: 70**

**Instructions:**

1. Attempt all questions.
2. Make suitable assumptions wherever necessary.
3. Figures to the right indicate full marks.

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|-------------|---|------------|
| <b>Q.1</b>  | (a) Explain central dogma of molecular biology with diagram                       | <b>07</b>  |
|             | (b) Explain biologist model with example  | <b>07</b>  |
| <b>Q.2</b>  | Explain the following in detail   |            |
|             | (a) What is public biological database with types of databases                    | <b>07</b>  |
|             | (b) What is data annotation and data format                                       | <b>07</b>  |
|             | <b>OR</b>   |            |
|             | (b) Explain motifs and profiles with examples                                     | <b>07</b>  |
| <b>Q.3</b>  | Attempt all questions   |            |
|             | (a) Write a short note on   | <b>04+</b> |
|             | 1) Genbank  | <b>03</b>  |
|             | 2) PDB  |            |
|             | (b) Explain pairwise sequence comparison in detail                                | <b>07</b>  |
|             | <b>OR</b>   |            |
| <b>Q.3</b>  | (a) What is phylogenetic tree ? and Explain types of phylogenetic tree in detail. | <b>07</b>  |
|             | (b) What is mutation and explain point and segmental mutation in detail           | <b>07</b>  |
| <b>Q.4</b>  | Attempt following questions   |            |
|             | (a) Explain chemical composition of Protein structure in detail                   | <b>07</b>  |
|             | (b) What is Data mining and explain data mining techniques                        | <b>07</b>  |
|             | <b>OR</b>   |            |
| <b>Q. 4</b> | (a) Explain SCOP, CATH and UNIPROT in detail.                                     | <b>07</b>  |
|             | (b) Explain Blast algorithm and its options                                       | <b>07</b>  |
| <b>Q.5</b>  | Explain in brief  |            |
|             | (a) Write a short note on   | <b>03+</b> |
|             | 1) Global alignment   | <b>04</b>  |
|             | 2) Gap penalty  |            |
|             | (b) What is Genomes and Genomics  | <b>07</b>  |
|             | <b>OR</b>   |            |
| <b>Q.5</b>  | (a) Explain CGI and XML in detail   | <b>07</b>  |
|             | (b) Explain SwissPort, Ghostview, GV, Xgobi with example.                         | <b>07</b>  |

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