

Seat No.: _____

Enrolment No. _____

GUJARAT TECHNOLOGICAL UNIVERSITY

BE - SEMESTER-VI(NEW) – EXAMINATION – SUMMER 2019

Subject Code:2160410

Date:21/05/2019

Subject Name: Bioinformatics

Time:10:30 AM TO 01:00 PM

Total Marks: 70

Instructions:

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

- Q.1** (a) Give the historical overview of Bioinformatics. **03**
(b) What is consensus sequence? **04**
(c) Briefly explain the role of NCBI. Further, discuss MMDB. **07**

- Q.2** (a) Enlist the tools of Entrez. **03**
(b) Write with an example: global alignment. **04**
(c) Discuss the importance of molecular biology in understanding bioinformatics. **07**

OR

- (c) For what attributes, do the phylogenetic analysis help? **07**
- Q.3** (a) What is cladistics and systemics? **03**
(b) How is the score calculated for BLAST and transition-transversion matrices? **04**
(c) Explain the terms: target identification and validation, lead molecule identification and optimization. **07**

OR

- Q.3** (a) Describe Hidden Markov Model. **03**
(b) Narrate the major features of PDB. **04**
(c) Give the importance of multiple sequence alignment. **07**
- Q.4** (a) Write the steps of *Chau-Fasman* method for secondary structure prediction. **03**
(b) What is DNA microarray? **04**
(c) Make a clear list of applications of Bioinformatics and briefly important of all. **07**

OR

- Q.4** (a) Make a list of all the tools for gene prediction. **03**
(b) Discuss maximum parsimony method representing phylogeny. **04**
(c) What are the major tools or techniques in proteomics? Explain any two of them. **07**

- Q.5** (a) “Protein structure is related with its function”. Discuss briefly. **03**
(b) Write the steps for GOR IV method. **04**
(c) What is Single nucleotide polymorphism? Explain. **07**

OR

- Q.5** (a) Define: Progressive alignment. **03**
(b) What is clinical trial? What are its phases? **04**
(c) Explain the types of proteomics studies, based on their characteristics. **07**
