**Q.P. Code: 20304** 

[Time: 2:30 Hours] [Marks:60]

Please check whether you have got the right question paper.

N.B: 1) **All** questions are **compulsory**.

- 2) **Figures** to the **right** indicates **full** marks.
- 3) Draw neat and labelled diagrams wherever necessary.

## Q.1 Any 3 Out of 5 (5M Each)

- 1. Briefly write about INSDC.
- 2. Write a note on Genbank and the submission tool in Genbank, EMBE and DDBJ.
- 3. Explain the significance of the online biological database.
- 4. Write a brief note on Protein Sequence Database.
- 5. Comment on the various literature databases available online.

## Q.2 Any 3 Out of 5 (5M Each)

- 1. Differentiate between PAM and BLOSUM matrix.
- 2. Demonstrate with example: Homologues, Orthologues and Paralogues.
- 3. What is the difference between the Sequence homology and sequence similarity.
- 4. Write a note on BLOSUM Matrix.
- 5. Explain Dot plot matrix.

## Q.3 Any 3 Out of 5 (5M Each)

- 1. Explain Needleman and Wunsch algorithm.
- 2. Why there is a need for Sequence comparison and Sequence alignment.
- 3. Write a note on Pairwise alignment and its types.
- 4. Describe Smith-Waterman algorithm.
- 5. Briefly explain Interpretation of the result of the analysis of the alignment.

## Q.4 Any 3 Out of 5 (5M Each)

- 1. Explain the process of translation in brief.
- 2. Explain the process of DNA replication in brief.
- 3. Briefly write about the Dayhoff Point Accepted Mutation matrices.
- 4. How sequence alignment plays an important role in bioinformatics.
- 5. Write a note on Domain Database.

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