Roll No. Total No. of Pages: 02

Total No. of Questions: 09

B.Sc - Honours (Microbiology) (Sem.-5) COMPUTER SCIENCE AND BIOINFORMATICS

Subject Code: BSMB507-20 M.Code: 92513

Date of Examination: 19-12-22

Time: 3 Hrs. Max. Marks: 60

INSTRUCTIONS TO CANDIDATES:

- SECTION-A is COMPULSORY consisting of TEN questions carrying TWO marks each.
- 2. SECTION-B contains FIVE questions carrying FIVE marks each and students have to attempt any FOUR questions.
- 3. SECTION-C contains THREE questions carrying TEN marks each and students have to attempt any TWO questions.

SECTION-A

1. Write briefly:

- a) Discuss the CPU functions.
- b) Define DNA.
- c) Define BLAST
- d) What do you mean by data submission tools?
- e) What is FASTA?
- f) Define file formats.
- g) Discuss the input/output devices.
- h) What are the various storages devices?
- i) Enlist the application of computer sciences in bioinformatics.
- j) Highlight the functional units and their interrelation.

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SECTION-B

- 2. Mention the basic computer organization.
- 3. Elaborate on the phylogenetic trees constitution for microbial diversity.
- 4. Describe about database similarity search.
- 5. Mention the multiple sequence alignment for studying microbial diversity.
- 6. Discuss the various fundamentals of computers.

SECTION-C

- 7. Discuss in detail about stored programs and applications of database similarity search.
- 8. Mention in detail about limitations of bio-information's and protein sequence database.
- 9. What do you know about pair wise sequence alignments and structure databases.

NOTE: Disclosure of Identity by writing Mobile No. or Making of passing request on any page of Answer Sheet will lead to UMC against the Student.

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