

Seat No.

HB-003-1194003 M. Sc. (Sem. IV) (CBCS) Examination April - 2023 Microbiology : MICRO-421 (Biomolecular Engineering)

Faculty Code : 003 Subject Code : 1194003

Time : $2\frac{1}{2}$ Hours / Total Marks : 70

- 1 Answer the following (Any Seven, Each of 02 Marks)
 - (1) Enlist forces involved in the tertiary structure of a protein.
 - (2) Which functional groups of amino acids form a peptide bond in protein ?
 - (3) Define Domain.
 - (4) What are molecular chaperones ?
 - (5) What is codon bias ?
 - (6) Enlist the steps involved in PCR.
 - (7) Define gene shuffling.
 - (8) Give the principle of pyro sequencing.
 - (9) Define recombinant clones.
 - (10) Define gene library.
- 2 Answer the following (Any Two, Each of 07 Marks) 14
 - (a) Write a note on the role of secondary structure of a protein in protein function.
 - (b) Describe peptide geometry in detail.
 - (c) Explain domain and topology with reference to the catalytic action.

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(b) What are molecular chaperones ? Explain its cellular functions. OR Answer the following (each of 07 Marks) 14 3 Describe in vitro protein folding and its biotechnological (a) significance. (b) Explain molecular chaperone-assisted protein folding. Answer the following (each of 07 Marks) 14 Describe different methods of protein engineering. (a) Explain evolution and mutators strain in detail. (b) Answer the following (Any Two, each of 07 Marks) 14 5 What is Real-Time PCR ? Explain. (a) Provide an account on strategies for primer designing for (b) known and unknown sequences. Write a short note on next-generation sequencing. (c)

Answer the following (each of 07 Marks)

What is protein folding ? Describe.

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(a)

(d) Write a short note on molecular tagging of the expressed protein.

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