# GUJARAT TECHNOLOGICAL UNIVERSITY

BE- VII<sup>th</sup> SEMESTER-EXAMINATION – MAY/JUNE- 2012 Subject code: 170401 Date: 24/05/2012

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.
- Q.1 (a) What is protein folding? Comment on the protein modular structure and 07 protein evolution.
  - **(b)** Describe different levels of protein organization.
- Q.2 (a) How base sequence in the DNA determines the amino acid sequence in 07 protein?
  - (b) What is protein structure? How native state of protein structure is stabilized? 07

### OR

- **(b)** How can you determine the structures of different proteins?
- Q.3 (a) Compare and contrast the physical and chemical techniques used for the 07 immobilization of enzymes.
  - (b) How protein functions are integrated? Can the functions of different proteins 07 be controlled?

## OR

- Q.3 (a) Explain the principle, procedure and applications of thin layer 07 chromatography.
  - (b) Explain the principle, procedure and applications of affinity 07 chromatography.
- Q.4 (a) Discuss the use of computer in the field of protein science. Explain the 07 expression pattern of proteins in space and time.
  - (b) Describe the interaction of protein with ligand and comment on the catalysis 07 by enzymes.

#### OR

- **Q.4** (a) What are homologous proteins? Describe the evolution of globins and NAD 07 binding domains of dehydrogenases.
  - (b) Discuss competitive, non-competitive and uncompetitive methods of enzyme 07 inhibition.
- Q.5 (a) Explain the "Lock and Key" hypothesis of enzyme action. Enlist different 07 classes of enzymes with suitable example.
  - **(b)** Comment on the specificity of enzyme with suitable example.

#### OR

- Q.5 (a) Explain the principle and procedure of electrophoresis technique used for the purification of proteins.
  - (b) What are molecular chaperons? Give classification of molecular chaperons and describe their roles in protein folding.

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