### B.TECH/BT/6<sup>TH</sup> SEM/BIOT 3242(BACKLOG)/2021

## **BIOPHYSICS OF MACROMOLECULES** (BIOT 3242)

#### Time Allotted : 3 hrs

Full Marks: 70

Figures out of the right margin indicate full marks.

## Candidates are required to answer Group A and any 5 (five) from Group B to E, taking at least one from each group.

*Candidates are required to give answer in their own words as far as practicable.* 

## Group – A (Multiple Choice Type Questions)

- 1. Choose the correct alternative for the following:
  - (i) Which of the following atoms is least likely to participate in hydrogen bond? (a) 0 (b) S (d) N (c) F
  - (ii) Which of the following polar and uncharged acid? is a amino (a) Histidine (b) Proline (d) Threonine (c) Alanine
  - (iii) Which interaction takes place when a partially positive end of one molecule interacts with a partially negative end of another molecule? (a) dipole-dipole interaction (b) ion-dipole interaction (d) van der Waals force
    - (c) ion-ion interaction
  - (iv) Which of the following statement is correct? (a) Charged amino acid residues are never found in the core of a protein (b) Charged amino acid residues are seldom found in the core of a protein (c) Hydrophobic amino acid residues are always found in the core of a protein (d) Tyrosine is always found in the core of a protein
  - (v) If no. of amino acid residues is 110, what will be the molecular weight of the protein? (a) 15000 (b) 11000 (c) 12100 (d) 13800

#### Which one is a non-proteinogenic amino acid? (vi) (b) Pyrrolysine (a) Selenocysteine (c) N-formyl methionine (d) GABA.

(vii) In which region the wave length 600 nm belongs to? (a) UV (b) visible (c) near IR (d) far IR

 $10 \times 1 = 10$ 

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- (viii) Under which condition Lambert-Beer law can be applied?(a) when the solution is dilute and transparent
  - (b) when the solution is concentrated and transparent
  - (c) when the solution is in colloid form
  - (d) when solution is dilute and has suspended particles
- (ix) What is the use of Faraday Cup in Mass Spectrometry?
  (a) Ion analyzer
  (b) Ion detector
  (c) Ion production
  (d) None of the analysis
  - (c) Ion production (d) None of the above
- (x) Virtual, erect and magnified image as that of the object is formed in
  (a) Telescope
  (b) Photocopier
  - (c) Camera

(d) Magnifying glass

# Group – B

- 2. (a) Write notes on Reverse Turns.
  - (b) Compare A, B and Z DNA.

5 + 7 = 12

- 3. (a) Draw and explain the titration curve of Aspartic acid.
  - (b) pKa of any functional group is greatly affected by it's chemical environment-Explain with examples

6 + 6 = 12

 $(3 \times 4) = 12$ 

# Group – C

- 4. Justify the following statements.
  - (i) Mixing of benzene and water increases energy of a system
  - (ii) Melting point of an alpha helix depends on its chain length
  - (iii) Tertiary structure of a protein is maintained by weak forces
- 5. (a) Based on the two-state model, derive the equation for kinetic study of denaturation of a protein. Discuss how the kinetic constant can be determined experimentally using the equation.
  - (b) Explain cooperative nature for melting of an alpha helix

(4+4)+4=12

# Group – D

- 6. (a) State the basic principles of fluorescence spectrometry. What is the major difference between fluorescence and UV-visible spectroscopy?
  - (b) Draw a schematic diagram of a fluorescence spectrophotometer. State the parameters that influence the shape of the curve of fluorescence spectrum.

(4+2) + (4+2) = 12

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- 7. (a) What are the different types of vibrations in a molecule? Draw diagrams of different types of vibrations in a three-atomic non-linear molecule.
  - (b) Discuss three applications of IR.

(3+3)+6=12

# Group – E

- 8. (a) Explain with a diagram the working principle of TEM.
  - (b) Write down the different areas where Mass Spectrometry is used.

8 + 4 = 12

- 9. (a) Briefly explain the production of X-rays.
  - (b) What is the principle behind Mass Spectrometry?

8 + 4 = 12

Department & Section	Submission Link
BT	https://classroom.google.com/c/MzAxNjYwNzg0NDEx/a/MzY1MzU3MzQ5NTM5/details