

(i) Printed Pages : 2

Roll No.

(ii) Questions : 7

Sub. Code :

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Exam. Code :

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B.Sc. (Hons.) Bioinformatics 5th Semester
(2122)

**MOLECULAR MODELING AND COMPUTER
AIDED DRUG DESIGN**

Paper : BIN-5005

Time Allowed : Three Hours]

[Maximum Marks : 60

Note :— Attempt **five** questions in all by selecting **two** questions from each Unit & the first compulsory question.

1. Compulsory Question

- (i) Define Chemo-informatics and give its scope.
- (ii) Briefly explain the concept of potential energy surface.
- (iii) State Lipinski Rule of Five.
- (iv) Differentiate between structure based and ligand based drug design.

4×3=12

UNIT—I

2. (a) What is energy minimization and its importance ? Give any 1 energy minimization method.

6

- (b) Write a note on molecular docking & scoring. 6
3. (a) Explain the concept of force fields. 8
- (b) Differentiate between quantum mechanics & molecular mechanics. 4
4. (a) What is molecular dynamics simulations and its applications. 8
- (b) Briefly explain coordinate system in molecular modeling. 4

UNIT—II

5. (a) Discuss basic concept of ADME and its importance. 6
- (b) Discuss protein-ligand interactions in terms of computer aided drug design. 6
6. (a) Discuss virtual drug screening process and its importance. 8
- (b) Write a note on the time-lines of drug discovery process from bench to bedside. 4
7. (a) Explain concept of 1D, 2D & 3D molecular descriptors. 8
- (b) Give the basic principle and applications of QSAR. 4