

2056

M.Sc. (Bio-Informatics) Fourth Semester
MBIN-8018: Molecular Modeling and Pharmacoinformatics

Time allowed: 3 Hours

Max. Marks: 60

NOTE: Attempt five questions in all, including Question No. 1 which is compulsory and selecting atleast one question from each Unit.

x-x-x

1. Answer briefly:

- a) QSAR
- b) Lead optimization
- c) PubChem
- d) Similarity
- e) Individualized therapy
- f) Fingerprints

(6x2)

UNIT - I

- 2. a) Explain empirical potential energy functions used in molecular mechanics.
- b) Describe molecular surfaces and types such as van der Waals and solvent-accessible surfaces. (2x6)
- 3. a) Discuss different force field methods used in molecular mechanics.
- b) Explain energy minimization methods and their role in conformational analysis. (2x6)

UNIT - II

- 4. a) Explain the principles of rational drug design with suitable examples.
- b) Discuss high-throughput screening and its advantages and limitations. (2x6)
- 5. a) Explain the concepts of similarity and complementarity in drug design.
- b) Elaborate on the development of HANSCH QSAR equation. (2x6)

UNIT - III

- 6. a) Discuss molecular descriptors and fingerprints in cheminformatics.
- b) Describe ADME in drug development. (2x6)
- 7. a) Discuss gene-drug interactions and their importance in personalized medicine.
- b) Explain the role of genomics in drug target identification and disease diagnosis. (2x6)

x-x-x