

1058

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

I. Attempt the following:-

- a) Define empirical energy.
- b) Write formula for estimation of energy term for bonded atoms?
- c) Explain genetic algorithm.
- d) Define pharmacophore keys.
- e) Define QSAR.
- f) Discuss 3D Fingerprints.
- g) What is Tanimoto coefficient?
- h) Discuss phase I clinical trial stage.

(8x1½)

**UNIT – I**

- II. a) Define molecular modeling and what can we learn from it.
  - b) Write a note on conformational analysis.
- III. a) Discuss role of molecular mechanism & its application.
  - b) What is ideal minimization technique and why.

(2x6)

**UNIT –II**

- IV. a) What are the different approaches for drug designing?
  - b) Differentiate between high throughputs vs. rational drug design methods.
- V. a) What is the importance of molecular descriptor in QSAR?
  - b) Why indicator variable is used in QSAR models?

(2x6)

P.T.O.



(2)

**UNIT – III**

- VI. a) Discuss Pharmacogenomics & its applications.  
 b) What a note on using genomics for target identification. (2x6)
- VII. a) What are molecular descriptor & name three descriptors?  
 b) Write a note on molecular similarity & its applications. (2x6)

x-x-x