



(Following Paper ID and Roll No. to be filled in your Answer Book)

PAPER ID : 154753

Roll No.

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B. Tech.

(SEM. VII) (ODD SEM.) THEORY
EXAMINATION, 2014-15
**MOLECULAR MODELING AND
DRUG DESIGN**

Time : 3 Hours]

[Total Marks : 100

Note: Attempt all questions.

1. Attempt any two parts. **10x2=20**
 - (a) What do you mean by molecular modeling? Explain in detail.
 - (b) Discuss the postulates of quantum mechanics.
 - (c) Enumerate the drawbacks of mechanical models as compared to graphical models.

2. Attempt any two parts. **10x2=20**
 - (a) What do you understand by protein folding? Discuss the process of protein folding.
 - (b) Discuss the Electrostatic interaction and Van der Waals interactions.
 - (c) Write short notes on the following
 - (i) Conformational searching
 - (ii) Molecular dynamics

3. Attempt any two parts. **10x2=20**
- (a) What do you understand by homology modeling? Discuss the homology modeling of proteins.
 - (b) How Ramachandran plot is used for validation of protein models? Explain.
 - (c) Discuss the ab initio method for computational modeling of proteins.
4. Attempt any four parts **5x4=20**
- (a) What do you mean by linear and non linear QSAR model?
 - (b) What is drug discovery? Explain the various steps of drug design.
 - (c) What are artificial neural networks? Explain its application in QSAR.
 - (d) Write short notes on genetic algorithm.
 - (e) What are descriptors in QSAR? Explain.
5. Attempt any two parts **10x2=20**
- (a) What do you mean by virtual screening? Explain in detail.
 - (b) What is molecular docking? Discuss the protein-ligand docking with suitable example.
 - (c) Write short notes on the following
 - (i) Combinatorial libraries
 - (ii) ADME properties of lead.
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