

5/2016

(3 Hours)

[Total Marks : 80]

Note : 1. Question no. 01 is compulsory.

2. Attempt any 03 questions out of remaining 05 questions

- Q1] i) Define docking and explain the process of protein ligand docking. 10M
- ii) Differentiate between covalent and non covalent interactions. 10M
- Q2] i) Describe the various methods of Machine learning and there role in bioinformatics. 10M
- ii) Describe 3D QSAR 10M
- Q3] i) Describe the features and technical challenges of interoperability. 10M
- ii) Explain the process of energy minimization 10M
- Q4] i) Define optimization and explain the process of drug optimization. 10M
- ii) List and describe the role of various types of chaperones 10M
- Q5] Describe : 20M
1. molecular mechanics
 2. UMLS
 3. Simmulated annealing
 4. Molecular dynamics
- Q6] i) Describe the features of ramchandran plot and explain how to detect the quality of protein using ramchandran plot. 10M
- ii) Explain the features of torsion angle and Cartesian coordinates. 10M