[Time: 3 Hours] [Max. Marks: 75]

Computer Aided Drug Design Q.P. CODE: 5159

Your answers should be specific to the questions asked. Draw neat, labeled diagrams wherever necessary.

LONG ESSAY (Answer any Three)

3 X 10 = 30 Marks

- 1. Explain briefly different physiochemical used to determine the Quantitative structure activity relationship of drug molecules.
- 2. Explain different energy minimization methods and compare global minimum conformation and bioactive conformation.
- 3. Discuss the fragment based drug design and homology modeling.
- Explain the pharmacophore based and structure based In-silico virtual screening protocols. 4.

SHORT ESSAY (Answer any Nine)

9 X 5 = 45 Marks

- 5. Explain in brief Hammett equation and sigma parameters.
- 6. Enumerate the different methods in the calculation of partition co-efficient.
- 7. Discuss the Free Wilson analysis and give its applications.
- 8. What are the various statistical methods adopted for the QSAR analysis.
- 9. Discuss the different approaches of 3D QSAR.
- 10. Discuss the agents used in the inhibition of HMG Co-A reductase enzyme.
- 11. Discuss the different types of drug receptor interactions.
- 12. Explain how ADMET properties are predicted for new molecules.
- Write a note on receptor/enzyme interaction and its analysis. 13.
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 of pharmacophore? 14. How do you identify features of pharmacophore?

