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2011 MOLECULAR MODELLING AND DRUG DESIGN

Time Allotted: 3 Hours Full Marks: 70

The figures in the margin indicate full marks.

Candidates are required to give their answers in their own words as far as practicable.

GROUP - A (Multiple Choice Type Questions)

1. Choose the correct alternatives for any ten of the following:

 $10 \times 1 = 10$

- i) Of the following which is NOT a minimization algorithm?
 - a) Steepest descent
 - b) Conjugate gradient
 - c) Newton-Raphson
 - d) Modified Neglect of Diatomic Overlap.
- ii) The Boltzmann factor is given by
 - a) $\exp(-E/RT)$
 - b) $\exp(-x^2)$
 - c) $\exp(-kT)$
 - d) none of these.

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iii) Which of the following is NOT a force field?

- a) AMBER
- b) CHARMM

c) OPLS

d) GEO.

iv) In Lipinski rule the molecular weight varies

- a) 350 to 500
- b) 100 to 150
- c) 400 to 500
- d) 230 to 250.

v) In drug design IND means

- a) International New Drug
- b) Investigation of New Disease
- c) Investigational New Drug
- d) Indian New Disease.

vi) In drug design MTD means

- a) Maximum Treated Dose
- b) Maximum Tolerated Dose
- c) Maximum Time Does
- d) Measurement to Treated Dose.

vii) IC_{50} means

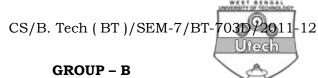
- a) concentration of a drug that is required for50 per cent inhibition in an assay
- b) concentration of a drug that is required for50 per cent activation in an assay
- c) 50 mg of a drug that is required for 50 per cent inhibition in an assay
- d) 50 μg of a drug that is required for 50 per cent activation in an assay.

viii) A pharmacophore is defined as

- a) a molecule that carries essential features responsible for a drug's biological activity
- b) a molecule without biological activity
- c) a molecule that carries non-essential biological information
- d) a kinetically fast reacting molecule.
- ix) A method for protein structure determination from which data is incorporated into the Brookhaven protein data bank is
 - a) Nuclear magnetic resonance spectroscopy
 - b) Fluorescence spectroscopy
 - c) Infrared spectroscopy
 - d) Atomic force microscopy.



- x) Generally in Phase 1 the number of total human subjects is in the range of
 - a) 20 to 80
- b) 50 to 100
- c) 200 to 300
- d) 500 to 1000.
- xi) 'QSAR' stands for
 - a) Quantum Structure Action Relationship
 - b) Quantitative Structure Activity Relationship
 - c) Quantum Similarity Activity Relationship
 - d) None of these.
- xii) Lead is a
 - a) Molecule with no property
 - b) Starting molecule of docking
 - c) Molecule with biological and pharmacological property
 - d) A software tool.



(Short Answer Type Questions)

Answer any three of the following.

 $3 \times 5 = 15$

- 2. What is meant by 'ADME' of a drug? How is it useful in drug-design? 2 + 3
- 3. Define the following:
 - ED_{50} a)
 - LD_{50} b)
 - Pharmacokinetics c)
 - Pharmacodynamics d)
 - QSPR. e)
- 4. Briefly describe the advantages and disadvantages of QSAR $2\frac{1}{2} + 2\frac{1}{2}$ in drug-design.
- 5. What is a Lennard-Jones potential? What are some of the microscopic parameters that are obtained from simulation results that are then compared to experimentally obtained 2 + 3results.
- 6. Briefly describe the 'Steepest descent' method used in energy minimization.



GROUP - C

(Long Answer Type Questions)

Answer any three of the following.

 $3 \times 15 = 45$

- 7. "Protein stability and folding plays an increasingly important role in the search of biopharmaceuticals (i.e. protein-based drugs)." Elaborate on the statement with the use of the following 3 informational categories:
 - (i) Ramachandran-Sasisekharan plot
 - (ii) The phenomenon of protein folding itself
 - (iii) The hydrophobic effect.

5 + 5 + 5

- 8. a) Give a step-wise description of the Metropolis Monte Carlo algorithm. What makes this algorithm unique?
 - b) To what types of biological system processes have Monte Carlo simulations been successfully applied?
 - c) What is simulated annealing and what is it ideally suited for?
 - d) What are the factors that decide choice between Monte Carlo and molecular dynamics for carrying out a molecular simulation?
 5 + 2 + 4 + 4
- 9. a) What is one of the biggest advantages of combinatorial chemistry over classical synthetic chemistry? Elucidate with an example from medicinal chemistry.

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- b) Briefly describe the three methods by which compounds in a combinatorial chemical library are identified. 6 + 9
- 10. Ludi is one of the approaches of De nuovo Drug designing.
 - a) Briefly describe what methodology it follows.
 - b) Briefly enumerate the functional steps of Ludi.
 - c) What are the major differences of Ludi and Ludi/CAP methods? 4+8+3
- 11. a) Name four physico-chemical parameters and their mathematical representation that can be referred to as molecular descriptors specific to drug-design.
 - b) Use a flow-chart to depict the drug discovery process.
 - c) Briefly describe how clinical trial of a drug is conducted.

6 + 4 + 5

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