Total No. of printed pages = 2

MPL 203 T

Roll No. of candidate

23/52/22

HIMT & SIPS) \*\*
Hatkin apara,

2022

## M.Pharm. 2nd Semester End-Term Examination

## PRINCIPLES OF DRUG DISCOVERY

(New Regulation)

Full Marks - 75

Time - Three hours

The figures in the margin indicate full marks for the questions.

Answer any five questions. All questions carry equal marks.

- 1. Explain Target and lead with examples. Write about various sources of discovery of new chemical entities (NCEs). (15)
- 2. (a) Explain rational approach for drug design.
  - (b) Explain in brief the role of proteomics in target identification. (8+7 = 15)
- 3. What is QSAR? Give advantages and disadvantages of QSAR. Explain the Model Validation technique mentioning LOO and LNO. Discuss about the CoSASA modelling using steroid backbone template. (1+4+5+5 = 15)
- 4. (a) Write a note on COMFA. (8+7 = 15)
  - (b) How computer aided drug design is useful in new drug discovery and development.
- 5. (a) Describe the methods of lead optimization. (8+7=15)
  - (b) What do you understand about De Novo drug design?
- 6. (a) Write a note on prediction of protein structure. (8+7 = 15)
  - (b) Describe the process of Homogeneous Radioactive Bioassays (HRB) in High-Throughput Screening.

7. Write a note on

(8+7=15)

- (a) Drug likeness screening.
- (b) Pharmacophore (lead for drug structures) based screening.
- 8. (a) Write a principle involved in design of pro-drug.

(8+7=15)

(b) Explain the term molecular docking.

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