

**M. PHARM. (PHARMACEUTICAL CHEMISTRY) SEM –II**  
**(CBCS- 2019 COURSE): Winter - 2021.**  
**SUBJECT: COMPUTER AIDED DRUG DESIGN**

**Day:** Wednesday  
**Date:** 01-12-2021

**Time.** 02:00 PM-05:00 PM  
**Max. Marks:** 75

**W-20784-2021**

**N.B.:**

- 1) **Q. No.1 and Q. No.5 are COMPULSORY.** Out of remaining questions attempt **ANY TWO** questions from each section.
- 2) Answer to both the sections should be written in **SEPARATE** answer book.
- 3) Figures to the right indicate **FULL** marks.

**SECTION-I**

- Q.1** What are physiochemical properties? List out the different physiochemical properties and describe the importance of any two of them. **(08)**
- Q.2** Explain in detail Hansch and Free Wilson model with examples. **(15)**
- Q.3** Describe detailed methodical process of molecular docking. Write different mathematical components of scoring functions. **(15)**
- Q.4** Write short notes on **ANY TWO** of the following: **(15)**
- a) Energy minimization
  - b) HIV protease structure and binding interactions with any one HIV protease inhibitors
  - c) Topliss approach of substitution

**SECTION-II**

- Q.5** What is pharmacophore model and explain in detail. **(07)**
- Q.6** Explain importance of ADMET properties of new molecules in view with drug design. **(15)**
- Q.7** What are the methods to predict binding cavity size of an enzyme or receptor? **(15)**
- Q.8** Write short notes on **ANY TWO** of the following: **(15)**
- a) Fragment based screening
  - b) Importance of drug design
  - c) Strategies for virtual screening

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