

GUJARAT TECHNOLOGICAL UNIVERSITY
M. Pharm. – SEMESTER – II • EXAMINATION – SUMMER • 2015

Subject Code: 2920201**Date: 16-05-2015****Subject Name: Drug Design and Discovery****Time: 10:30 am - 01:30 pm****Total Marks: 80****Instructions:**

1. Attempt any five questions.
2. Make suitable assumptions wherever necessary.
3. Figures to the right indicate full marks.

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|------|-----|--|----|
| Q.1 | (a) | Discuss a path of new drug discovery by rational approach. | 06 |
| | (b) | Why new drug discovery is essential? Which are the constraints in new drug discovery? | 05 |
| | (c) | Discuss in detail about different approaches used in new drug discovery? | 05 |
| Q.2 | (a) | What are the methods used for lead optimization? Give basic principle and requirements of two widely used methods. | 06 |
| | (b) | How stereochemistry influence biological response? Discuss in detail with suitable example. | 05 |
| | (c) | Write a note on molecular dynamics | 05 |
| Q.3 | (a) | Which are the physicochemical parameters used in Hansch LFER model? Give examples of two representative of each parameter and method of determining them experimentally. | 06 |
| | (b) | How can we validate Hansch model? | 05 |
| | (c) | Write a note on importance of bioisosterism in lead modification. | 05 |
| Q.4 | (a) | How drug acts in the body? How activity of drug is represented? Which method is most preferable? Why? | 06 |
| | (b) | Write a note on CoMFA. | 05 |
| | (c) | Interpret following equation obtained through Hansch technique and give your suggestion for further lead modification.
$\text{Log } I_{50} = 1.04(\pm 0.23) \pi - 0.83(\pm 0.18) \sigma - 0.66(\pm 0.20) E_s + 4.32(\pm 0.35)$ $n = 19, \quad r = 0.943 \quad r^2 = 0.889, \quad s = 0.34, \quad F_{3,15} = 22.8$ | 05 |
| Q.5 | (a) | How can we optimize the pharmacokinetics of lead without altering parent structure of compound? Discuss different methods used for it with suitable example. | 06 |
| | (b) | Write in brief about methods used for ligand based drug design. | 05 |
| | (c) | Write a note on virtual screening. | 05 |
| Q. 6 | (a) | Lead molecule active <i>in vitro</i> doesn't guarantee similar potency <i>in vivo</i> . – Justify | 06 |
| | (b) | Give detail account on designing enzyme inhibitors. | 05 |
| | (c) | What are the importance of molecular mechanics in de novo drug designing? Which are the different algorithms used for its calculations? | 05 |
| Q.7 | (a) | Which technique is preferred for designing drug when target structure is known? Which are the steps involve in this technique? What are the requirements for this technique? | 06 |
| | (b) | How target structure is derived? How can we determine binding site in the target? | 05 |
| | (c) | Write in detail about methods used to identify hit molecule. | 05 |