B.Tech. - VIEP - COMPUTER SCIENCE AND **ENGINEERING (BTCSVI)**

Term-End Examination

00040 December, 2017

| DICSE-006: DIO-INTURIVIATICS | | | |
|------------------------------|---|--------------------|--|
| Time: 3 h | ours Maximum Marks | Maximum Marks : 70 | |
| | nswer any seven questions. All questions c ual marks. | arry | |
| is b | ly explain about Biological databases. Why io-informatics important? Explain its cations in the field of biology. | 10 | |
| | e the concept of ID Motifs. What are the cent representations in ID motifs? | 10 | |
| 3. (a) | Describe the dynamic programming for sequence alignment. | 5 | |
| (b) | Explain integrated genomic maps in detail. | 5 | |
| 4. (a) | What are the current approaches for bio-informatics and the problems | ,- | |
| | associated with them? | 5 | |
| (b) | Discuss genetic networks in detail. | 5 | |
| BICSE-008 | 8 1 P. | T.O. | |

| 5. | (a) What are the challenges in Moore's law? | | |
|-----------|---|---|---|
| | (b) | Define the structural classification of proteins. | 5 |
| 6. | Expla | ain classical NMR spectroscopy in detail. 1 | 0 |
| 7. | Defin | 8 | |
| | | rentiate between two terminologies and two ogies. | 0 |
| 8. | Diffe | rentiate between the following: 5+ | 5 |
| | (a) | Protein and Amino Acid | |
| | (b) | DNA and RNA | |
| 9. | Desc | ribe the following in brief: $4 \times 2 \frac{1}{2} = 1$ | 0 |
| | (a) | Genome Alignment | |
| | (b) | Gen Bank | |
| | (c) | PDB | |
| | (d) | Microarray Clustering and Classification | |
| 10. | Write follow | e short notes on any two of the wing: $2 \times 5 = 1$ | 0 |
| | (a) | Hidden Markov Model | |
| | (b) | NLP | |
| | (c) | CAPRI | |
| | | | |
| BIC | SE-00 | 8 2 1,00 | 0 |