## M.Sc. (Part-II) Semester—IV (CBCS Pattern) Examination COMPUTER SOFTWARE

## Paper-XVI

## (Computing Techniques in Bio-Informatics)

11m	e : 1	hree Hours] [Maximum	Marks: 80
N.B	. :	(1) Due credit will be given to neatness and adequate dimensions.	
		(2) Illustrate your answers necessary with the help of neat sketches.	
1.	(A)	What are the applications of Bio-informatics? Explain.	8
	(B)	State and explain the categories of databases.	8
		OR	
2.	(A)	Explain retrieval system in detail.	8
	(B)	State and explain prediction of protein structure.	8
3.	(A)	Explain sequence database with example.	8
	(B)	Explain:	
		(i) DALI	
		(ii) FASTA.	8
		OR	
4.	(A)	Explain following:	
	,	(i) CAZY	
		(ii) BLAST.	8
	(B)	Explain enzyme database in detail.	8
5.	(A)	Explain data submission in brief.	8
		Describe different data analysis tools.	8
		OR	
6.	(A)	State and explain protein submission.	8
	(B)	Explain prediction tools in detail.	8
7.	(A)	State and explain sequence comparison with example.	8
		Describe UPGMA in detail.	8
	,	OR	
8	(A)	Explain substitution matrices in brief.	8
	(B)	State and explain structure algorithm.	8
9.	(A)	State and explain Monte Carlo methods.	8
	(B)	Explain PERL for bio-informatics.	8
		OR	
10.	(A)	Explain molecular dynamics simulation package.	8
		State and explain commercial and academic bio-informatics software.	8

