Total No. of Questions : 10]	SEAT No. :
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[4959]-1141

B.E. (Information Technology) **BIOINFORMATICS**

(2012 Course) (Elective-IV) (414464 A) (Semester-II) *Time* : 2½ *Hours*] [Max. Marks: 70 Instructions to the candidates: Answer Q1 or Q2, Q3 or Q4, Q5 or Q6, Q7 or Q8, Q9 or Q10. 2) Figures to the right side indicate full marks. Assume suitable data if necessary. 3) What is scope of Bioinformatics? Explain protein structure databases. **Q1)** a) [6] Explain any two rendering tools used in visualization. b) [4] OR Define Bioinformatics. Explain Bioinformatics application related to the **Q2)** a) following areas. [6] i) Phylogenetic Analysis. Genome Annotation. ii) b) Explain sources of veriability in microarray preparation & reading. [4] Explain major steps in pattern recognition & discovery process with **Q3**) a) diagram. [8] Difference between clustering & classification. [2] b) OR What is meant by sensitivity & specificity of statistical analysis tool. [2] **Q4)** a) Explain knowledge discovery process or datamining methods with neat b) diagram. [8]

Q5)	a)	Explain similarities & differences between BLAST & FASTA tools for sequence elignment. [8]		
	b)	Wri	te short note on:	[8]
		i)	Heuristic methods for sequence alignment.	
		ii)	Prediction algorithm for phylogenetic.	
			OR	
Q6)	a)	-	plain FASTA algorithm with recommended steps for similarity search etail.	ning [8]
	b)	Exp	plain different prediction algorithm for genes.	[8]
Q7)	a)	Exp	plain comparitive modeling process with neat diagram.	[8]
	b)	Exp	plain process of drug discovery in detail.	[8]
			OR	
Q8)	a)	Discuss the components of modeling & simulation system alongwit process.		vith [8]
	b)	Write short note on:		[8]
		i)	Methods for protein modeling.	
		ii)	Model refinement & evaluation.	
Q9)	a)	Wri	te short note on:	[12]
		i)	Tools for modeling & simulation.	
		ii)	Hidden Markov model.	
	b)	Exp	plain recent trends in Bioinformatics.	[6]
			OR	
<i>Q10)</i> a)		Wh	at are Future trends in Bioinformatics?	[6]
b)		Wri	te short note on:	12]
		i)	Structural Bioinformatics in drug discovered.	
		ii)	System Biology in human health & disease.	

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