USN												15BT5
-----	--	--	--	--	--	--	--	--	--	--	--	-------

Fifth Semester B.E. Degree Examination, Dec.2018/Jan.2019 **Bioinformatics**

Tir	ne: í	3 hrs.	Max. Marks: 80									
	Note: Answer any FIVE full questions, choosing ONE full question from each module.											
		Module-1										
1	a. b. c.	ove a short account on: OMIM. PDB flat file. SCOP.										
	d.	BLOSUM.	(16 Marks)									
2	a.	OR Explain about the algorithm behind CLUSTALW.	(00 Mayles)									
	b.	Comment on : i) PSI – BLAST ii) PROSITE.	(08 Marks) (08 Marks)									
			(00 Marks)									
_		Module-2										
3	a.	Explain in detail the distance based tree building methods.	(12 Marks)									
	b.	Calculate the maximum number of rooted and unrooted trees that can be										
		taxa A, B, C, D, E.	(04 Marks)									
		OR OR										
4	a.	Diagramatically explain the functional sites in a eukaryotic DNA and	evoluin a tool for									
		prediction of eukaryotic genes.	(08 Marks)									
	b.	Write about the tools available for predicting secondary structural elemen	nts in proteins.									
			(08 Marks)									
5	a.	Write the method and principle helical Savera's and in the same of										
5	b.	Write the method and principle behind Sanger's sequencing method. Given a raw genome sequence data, write about the different analysis tha	(08 Marks)									
	٠.	with it.	(08 Marks)									
			(00 Marks)									
		OR										
6	a.	Brief on SNPs and their importance.	(08 Marks)									
	b.	Write about comparative genomics and the different themes involved.	(08 Marks)									
		Module-4										
7	a.	Give a account on:										
		i) Force fields in MM ii) Energy minimization of a biological syst	em. (12 Marks)									
	b.	What is the need for structural super position of proteins?	(04 Marks)									
		OR										

Comment on : i) Rotamer libraries ii) Rasmol iii) Canonical DNA form. (12 Marks) b. Brief about the different graphical representation of a protein. (04 Marks) Module-5

9 a. Construct restriction map on a plasmid for restriction enzymes Eco RI and HindIII. The following are the fragments got by their digestion: Eco RI – 7 Kb and 3 Kb, Hind III – 7 Kb and 3 Kb and dual digestion of EcoRI + HindIII – 3 Kb and 2 Kb.

(04 Marks)

b. Comment on i) Vector NTI ii) REBASE.

(06 Marks)

c. What are Primers? Discuss the criteria to be considered while designing the primers.

(06 Marks)

OR

10 a. Brief on bioactive conformation of ligands and method to derive them.

(04 Marks)

b. Comment on QSAR.

(06 Marks)

c. What are Pharmacophoric patterns? Write about its significance and derivation methods.

(06 Marks)