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B.M.S. College of Engineering, Bengaluru-560019

Autonomous Institute Affiliated to VTU

July 2023 Semester End Main Examinations

Program: B.E.

Branch: Biotechnology

Course Code: 19BT6DE4GIN

Course: Genomic Informatics

Semester: VI

Duration: 3 hrs.

Max Marks: 100

Date: 19.07.2023

Instructions: 1. Answer any FIVE full questions, choosing one full question from each unit.
2. Missing data, if any, may be suitably assumed.

Important Note: Completing your answers, compulsorily draw diagonal cross lines on the remaining blank pages. Revealing of identification, appeal to evaluator will be treated as malpractice.			UNIT – I	CO	PO	Marks
	1	a)	Compare all three generations of sequencing technologies.	CO1	PO	10
		b)	List various NGS file formats. Discuss any one briefly.	CO1	PO	05
		c)	How do you carry out DNA sequencing by exploiting torrent of H ⁺ ions? Illustrate with explanation.	CO1	PO	05
			OR			
	2	a)	Identify a method where light emission is used for genome sequencing. Write its principle and steps involved with figures.	CO1	PO	10
		b)	“DNA sequencing can be done for single large molecule of DNA in real time”. Justify this statement by identifying the method and steps involved with figures.	CO2	PO5	10
			UNIT – II			
	3	a)	Using velvet how do you assemble genome? Identify the algorithm behind it and write the steps involved with diagrammatic representations.	CO2	PO5	10
		b)	Develop a pipeline for denovo genome assembly. Explain it with neat figures.	CO2	PO5	10
			UNIT – III			
	4	a)	How do you assemble genome with greedy approach? Explain it with at least two examples.	CO3	PO2 PO12	10
		b)	Without genome browsers it is difficult to understand, view and analyze the genomic data. Justify this by stating the importance of genome browser, provide examples of genome browsers and explain one in detail.	CO3	PO2 PO12	10

		UNIT – IV			
5	a)	lncRNA play an important role in normal and disease process. Justify this statement by explaining role in diagnosis and treatment of cancer with suitable examples.	CO4	PO1	10
	b)	How can you employ NGS in cancer research? Substantiate by explaining various methods of NGS.	CO4	PO1	10
		OR			
6	a)	RNA sequencing plays a crucial role in identifying biomarkers. Justify this by outlining the procedure and any four applications in cancer research.	CO4	PO1	10
	b)	What is epigenome? Identify different approaches and sequencing strategies for epigenome with neat figures.	CO4	PO5	10
		UNIT – V			
7	a)	Provide reasons to state that NGS is valuable in clinical oncology.	CO4	PO5	10
	b)	Classify polymorphism. Give examples of polymorphic genes in major depression.	CO4	PO5	10
