

DR-003-2016043 Seat No. _____

B. Sc. (Sem. VI) Examination

March / April - 2022

Biotechnology: Paper - BT-603

(Advance Molecular Biology & Bioinformatics)

Faculty Code: 003

Subject Code: 2016043

Time: 2	$2\frac{1}{2}$ H	Iours] [Total Marl	ζs :	70
1 (a)	Ans	wer the question : (one mark each)		4
	(1)	Asymmetric PCR is used to generateDNA.		
	(2)	Dideoxy method is used for DNA		
	(3)	Unreacted nucleotide in phosphoramidite method of oligonucleotide synthesis is washed by		
	(4)	To sequence larger molecules, individu chromosomes are purified and broken into		
		or larger random fragments, which are cloned in vectors designed for large molecules.	to	
(b)	Ans	wer the questions: (any one out of two)		2
	(1)	What are the types of hybridization technique	?	
	(2)	Define probe.		
(c)	Ans	wer the questions: (any one out of two)		3
	(1)	Explain the principle of pyrosequencing.		
	(2)	Explain Sangers method of sequencing.		
(d)	Ans	wer the questions: (any one out of two)		5
	(1)	Explain the mechanism of artificial DNA synthes	is.	
	(2)	Enlist and explain the types of PCR.		
DR-003-2	20160	43] 1 [(Cont	d

2	(a)	Answer the questions : (one mark each)	
		(1) Sometimes successive rounds of screening of a genomic library are carried out and an ordered collection of clones is done in a linear fashion, then the, process is called as	
		(2) Restriction enzyme recognize sequence.	
		(3) In 1990 the first gene therapy was given to treat which deficiency?	
		(4) Method used to describe in vivo DNA protein interaction	
	(b)	Answer the questions: (any one out of two)	2
		(1) Enlist PCR and non PCR based molecular markers.	
		(2) What is restriction mapping?	
	(c)	Answer the questions: (any one out of two)	3
		(1) Explain the mechanism of DNA foot printing technique.	
		(2) Difference between, microsatellite and minisatellite.	
	(d)	Answer the questions: (any one out of two)	5
		(1) What is gene therapy? Explain type of gene therapy.	
		(2) Explain the methods of chromosome jumping and walking.	
3	(a)	Answer the questions : (one mark each)	4
•	(00)	(1) Human genome project completed in the year	_
		(2) The identification of drugs through the genomic study is called	
		(3) The first draft of human genome project was first published in 2001 in the journal.	
		(4) Full form of NCBI.	
	(b)	Answer the questions: (any one out of two)	2
		(1) What are the objectives of human genome project?	
		(2) What is biological database?	
	(c)	Answer the questions: (any one out of two)	3
		(1) Application of human genome project.	
		(2) Importance of biological database.	
	(d)	Answer the questions: (any one out of two)	5
		(1) Enlist and explain any two major, bioinformatics resources.	
		(2) Define bioinformatics and explain the branches and role of bioinformatics in biotechnology.	

2

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DR-003-2016043]

4	(a)	Answer the questions : (one mark each)	4
		(1) is a metabolic database.	
		(2) RCSB is a database of protein	
		(3) is the search and retrieval tool of	
		NCBI.	
		(4) InterPro is used to study	
	(b)	Answer the questions: (any one out of two)	2
		(1) Which are the primary and secondary database of protein?	
		(2) What is CATH?	
	(c)	Answer the questions: (any one out of two)	3
		(1) Explain the levels of SCOP.	
		(2) What are different types of models used by InterPro?	
	(d)	Answer the questions: (any one out of two)	5
		(1) Write a note on UniProt.	
		(2) Write a note on importance of PubMed.	
5	(a)	Answer the questions : (one mark each)	4
		(1) Full for of BLOSUM.	
		(2) The initiation of FASTA format has symbol.	
		(3) QSAR method involves properties.	
		(4) Primer3 is tool.	
	(b)	Answer the questions: (any one out of two)	2
		(1) What is comparative genomics?	
		(2) What is global and local alignment?	
	(c)	Answer the questions: (any one out of two)	3
		(1) What are the general rules of primer designing?	
		(2) What are the types of BLAST?	
	(d)	Answer the questions: (any one out of two)	5
		(1) Write a note on computer aided drug discovery.	
		(2) What is multiple sequence alignment? Write the types of alignment and applications of MSA.	