

HAF-003-001623 Seat No. _____

B. Sc. (Sem. VI) (CBCS) Examination

June / July - 2017

BT-603: Advance Molecular Tech. & Bioinformatics

Faculty Code: 003

Subject Code : 001623							
Fime : $2\frac{1}{2}$ Hours] [Total Marks : 70							
1	Ansv	wer the following question in one word:	0				
	(1)	The BLAST program was designed by					
	(2)	Southern blotting is used to identify					
	(3)	The technique used to locate specific genes in the chromosome is					
	(4)	Name the scientist who developed the chain termination method of sequencing technique					
	(5)	Pyrosequecning involves sequencing by					
	(6)	The information retrieval tools of NCBI GenBank is					
	(7)	Name the probe used in RT-PCR					
	(8)	Which microorganism genome was first sequenced?					
	(9)	KEGG is used for the study of					
	(10)	Metagenomics is the study of cell.					
	(11)	Name the method used for the study of DNA-Protein interaction					
	(12)	A comprehensive database for the study of human genetics is					
	(13)	NCBl was established in the year					
	(14)	is non PCR based marker.					
	(15)	What are cDNAs spots on a glass slide used for expression studies called?					
	(16)	EST stands for					
	(17)	Field of genomics which deal with DNA structure is called					
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(1	8) BLA	ASTP is protein query against data bank	
(1	•	detriplation in phosphoramidite methodseleased to remove the DMT group.	
(2	·	me one technique used for multiple sequence	
(a)) Wri	te any three out of six:	6
	(1)	Which are the core databases of Uniprot?	
	(2)	What is autosequecning?	
	(3)	How to calculate the annealing temperature in PCR ?	
	(4)	What is inverse PCR ?	
	(5)	Which are the databases of protein?	
	(6)	What is the major functions of NCBI?	
a .	. 33 7 •		0
(b)		te any three out of six :	9
	(1)	Explain the different levels of SCOP.	
	(2)	Criteria used for designing of primer.	
	(3)	Explain the mechanism of chromosome walking.	
	(4)	Explain the mechanism of microarray technique.	
	(5)	Application of autoradiography in molecular biology	
	(6)	Write a note on PDB.	
(c)	Wri	te any two out of five :	10
	(1)	Explain the process of DNA synthesis of phosphoramidite method.	
	(2)	Explain dideoxy method of DNA sequencing.	
	(3)	Explain the mechanism and applications of southern blotting technique.	
	(4)	Write a note on molecular markers used in molecular biology.	
	(5)	Explain the scope and importance of bioinformatics.	

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3	(a)	Write any three out of six:	6
		(1) What is primer?	
		(2) What is micro and mini satellite?	
		(3) Define bioinformatics.	
		(4) Write the use of Taq polymerase.	
		(5) What is restriction mapping?	
		(6) What is pairwise alignment?	
	(b)	Write any three out of six	9
		(1) Explain the mechanism of RT-PCR.	
		(2) Explain the process of Pyrosequecning.	
		(3) Explain the mechanism of chromosome jumping.	
		(4) Write a note on FASTA.	
		(5) How to construct the phylogenetic tree with bioinformatics tools.	
		(6) What is orthologous and paralogous ?	
	(c)	Write any two out of five:	10
		(1) Explain the process of PCR technique.	
		(2) Explain biological database in detail.	
		(3) Explain the mechanism and applications of western blotting technique.	
		(4) Write a note on BLAST and its uses.	

(5)

Explain the mechanism of microarray technique.