PARUL UNIVERSITY

PARUL INSTITUTE OF APPLIED SCIENCES

MID SEMESTER INTERNAL EXAMINATION, SEPTEMBER 2018

B. Sc. Semester V

Title of the paper: Omics Technologies

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Subject: Microbiology/Biotechnology/Biochemistry

Paper Code: 11102305

Date:	10/ 09/2018		Time: 11.30AM to	01:00PM				
Maximum Marks: 40								
Instructions:								
1.	All questions are compulsory and opt	tions are g	given in first and second	question				
	only.							
2.	Numbers to the right of question indi	icate the n	narks of respective quest	ion.				
	· / /							
Q. 1	Attempt any one question of the foll	owing.		(08)				
	(i) Describe different types of Omics fields and respective technologies							
	used to study them.							
	(ii) Write an essay on 'Nucleic acid	Sequencin	g' also explaining NGS					
	technology.							
Q. 2	Attempt any three questions of the f	ollowing.		(12)				
	(i)Write a note on Genome reduction							
	(ii) Describe '1000 Plant Genome Project'.							
	(iii) Write a brief note on NCBI.							
	(iv) Name a few Bioinformatics cen							
	(v) What is Complete Genome Project? Name a few projects.							
Q.3 Do as directed. Attempt all five questions. (05)								
	(i) Mention the contribution of Allan	n Coulson	in relation to DNA					
	sequencing.							
	(ii) What is GRC? Explain in one or two lines.							
		(iii) Explain the purpose of a Database.						
	(iv) What does NCBI stand for?	1 37 - 1	411					
Q. 4	(v) Explain the terms Transcriptome and Volatilome. Write correct option in your answer sheet for following 15 multiple (15)							
Ų. 1	choice questions.	sheet for i	ollowing 15 multiple	(15)				
	enoice questions.							
MCQ 1 The comprehensive study of the sugars and carbohydrates is called								
	(A) Glycomics	(B)	Glucomics					
	(C) Glucosics	(D)	Carbidomics					
MCC	The second concerned with	the structu	are, function, comparison,	and				
	evolution of genomes?(A) Genomics		n					
	(C) Epigenomics	(B)	Functional genomics Comparative genomics					
MCC		(D)	nsulin in					
	1955?	defice of f						
	(A) Frederick Sanger	(B)	Adam Gilbert					
MCC	(C) James Watson	(D)	Frederick Crick	11-1-1-11-1				
MCQ 4 Who published a sequencing procedure using DNA polymerase with radiolabelled nucleotides that was called the Plus and Minus technique?								
	(A) Alan Coulson and Fredrick	s and Mini (B)	Frederick Sanger					
	Sanger							
	(C) Alan Coulson	(D)	Walter Gilbert					

100 5	Camer	olete genome of human published	i in	- 1384		
1CQ 5		2001	(B)	2009		
	(A)	2001 2003	(D)	2007		
1016 8	(C)		_	. Shotoun Sequencer		
MCQ 6		stands for Epigenomics Simple	(B)	Environmental Shotgun Sequencer		
	(A)	Sequencing		S. S		
	. ~	Environmental Shotgun	(D)	Easy System Sequencing		
	(C)	Environmental 2				
		Sequencing is the study of genetic	materia	1 recovered directly from		
MCQ 7		onmental samples.	- 1			
		Metagenomics	(B)	Ecogenomics		
	(A)	Environmental genomics	(D)	None of A,B or C		
Note that a bottom with a standard	(C)	Environmental general				
MCQ 8		stands for	(B)	Proteomics Information Reserve		
	(A)	Protein Information Resource	(D)	Proteomics Information Reserve		
•	(C)	length of <i>Homo sapiens</i> genome		Account of		
MCQ9			(B)	$9.5 \times 10^{5} \text{ bp}$		
	(A)	10121	(D)	$3.2 \times 10^9 \text{ bp}$		
	(C)	2.1 x 10 ¹² bp mical method of Nucleic acids se	equencin	g was proposed by		
MCQ 10		Maxam and Gilbert	(B)	Frederick Sanger		
	(A)		(D)	Steve Jobs		
	(C)	. L'Eratione are	e reversi	ble modifications on a cell's DNA or		
MCQ 11	histones that affect gene expression without altering the DNA sequence.					
		Genetics	(B)	Lpigenene		
	(A)		(D)	None of A,B or C		
	(C)	Metagenetics man species, whose genome incl				
MCQ 12			(B)	22 pairs of autosomes and 2 sex		
	(A)	23 pairs of autosome	(-)	chromosomes		
	(0)	48 pairs of chromosome	(D)	23 pair of autosome and 2 sex		
	(C)	48 pairs of emoniosome	(-)	chromosome		
	. г.	ample of Secondary database is				
MCQ 13			(B)	Swissprot		
	(A)		(D)	Genbank		
1 (00 1	(C)	PIR e term 'Genomics' was coined b				
MCQ 14			(B)	Tom Roderick		
	(A)	I I amer: Ingleson	(D)	Walter Gilbert		
MCO 1	(C)	a number of Cyanobacteria genu	s whose	complete genome is sequences is		
MCQ 1			(B)	76		
	(A (C		(D)	82		
	11.	1 40	(~)			

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