D S	DL-003-006403 Seat No
ъ. ъ	c. (Bioinformatics) (Sem. IV) (CBCS) Examination April / May - 2015
	B. Sc. BI: 403 Applied Bioinformatics
	Faculty Code : 003 Subject Code : 006403
Time	$:2\frac{1}{2}$ Hours] [Total Marks :70]
	PART - A
	(Max. 30 Minutes)
	QUANTA is developed by company and used
(	(a) Schrodinger & Docking
(	(b) Accelrys & Modelling and Simulation
(	(c) ChemAxon & Virtual Screening
(	(d) OpenEye Scientific Software & Ligand optimization
2	Parallel sequencing was used in which technique
(	(a) Illumina
(	(b) Roche method
(	(c) Maxim and Gilbert method
(	(d) Both (a) and (b)
3	The product for molecular modeling is which is developed

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(c)

 $Modeller \ \& \ OpenEye$ 

(d)

Yasara & ChemAxon

(a) Maestro & Schrodinger (b) Maestro & Accelrys

4	Whi	nich is the enzyme used for removal of T+C base				
	(a)	Hydrazine	(b)	Acid catalysed depurinase		
	(c)	Piperdine	(d)	Dimethyl sulphate		
5	Mar	vinSketch & MarvinView	v are	the product of		
	(a)	PerkinElmer	(b)	Cambridge Soft		
	(c)	ChemAxon	(d)	OpenEye		
6	Mod	ified sequencing method	used	sequencing gel having		
	(a)	Urea	(b)	PAGE +urea		
	(c)	Agarose	(d)	Dye		
7	Flur	ophore began to be used	l in :			
	(a)	Capillary Array Electro	phore	esis		
	(b)	Automated Method				
	(c)	Modified Sanger's Metho	od			
	(d)	Maxim and Gilbert met	hod			
8	Diffe	Piperdine (d) Dimethyl sulphate  vinSketch & MarvinView are the product of  PerkinElmer (b) Cambridge Soft  ChemAxon (d) OpenEye  ified sequencing method used sequencing gel having  Urea (b) PAGE +urea  Agarose (d) Dye  ophore began to be used in:  Capillary Array Electrophoresis				
	(a)	Removal of oxygen in o	ldNT	P		
	(b)	hemAxon (d) OpenEye ed sequencing method used sequencing gel having frea (b) PAGE +urea  garose (d) Dye hore began to be used in : apillary Array Electrophoresis utomated Method dodified Sanger's Method faxim and Gilbert method fince between dNTP and ddNTP emoval of oxygen in ddNTP NTP extends DNA strand dNTP extends DNA strand				
	(c)	ddNTP extends DNA st	rand			
	(d)	both (a) and (b)				

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[ Contd...

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9	Cha	Challenges in Proteomics with Respect to plant Physiology				
(a) Chromatographic separation						
	(b)	Diabetes research				
	(c)	Neurological behavioral studies				
	(d)	Protein regulation and its relation with regulatory networks				
10		ch is the specialized format used to store the chemical pounds?				
	(a)	XML – based Chemical Markup Language				
	(b)	SMILES				
	(c)	Both (a) and (b)				
	(d)	None of the above				
11 IBM stands for?						
	(a)	International Business Machines Corporation				
	(b)	Indian Bureau of Mines				
	(c) International Business Management					
	(d)	None of the above				
12	Sho	rt overlapping DNA sequences are called				
	(a)	Clones (b) Consensus sequence				
	(c)	Contigs (d) Fragments				
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13	Cur	Current NIH & NCBI Directors are respectively				
	(a)	Francis S. Collins and David J. Lipman				
	(b)	Eric Green and Francis S. Collins				
	(c)	David J. Lipman and Eric Green				
	(d)	None				
14	Reco	ombinant clones are used for				
	(a)	Automated method (b) Assembly				
	(c)	Shot gun sequencing (d) HTS				
<b>15</b>	Deo	Deoxy-adenosine 5' ( $\alpha$ thio) - triphosphate :				
	(a)	Used for incorporating nucleotide				
	(b)	Used as dye for high resolution				
	(c) Used for base incorporation					
	(d) Filter sequence					
16	Physical gap is filled using					
	(a)	A libraries				
	(b)	Primer directed walking				
	(c)	Contigs				
	(d)	Gene libraries				
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<b>17</b>	SDI	F stands for							
	(a)	) Structure dynamic format							
	(b)	Structure data format							
	(c)	Script data file							
	(d)	None							
18	TEp	predict is tool for							
	(a)	MHC-1	(b)	MHC-2					
	(c)	Both	(d)	None					
19 By using the "reference sequence" developed in the human									
	project, individual differences are now being detected using								
	(a)	NPs	(b)	PSNs					
	(c)	SNPs	(d)	"snaps"					
20	Disc	iscovery Studio is the product of and used for							
	(a) Schrodinger & Docking								
	(b)	b) Accelrys & Docking, and ADMET prediction							
	(c)	ChemAxon, Docking and ADMET prediction							
	(d)	OpenEye Scientific Soft	ware	& Docking and					
		Pharmacophore generat	ion						

## PART-B

1	(a)	Exp	lain any Three	6
		(1)	Base calling	
		(2)	List out any four National Bioinformatics company.	
		(3)	Toxicogenomics	
		(4)	Accelrys	
		(5)	Discovery Studio	
		(6)	Biosafety	
	(b)	Exp	lain any Three	9
		(1)	Proteome	
		(2)	Technology Transfer	
		(3)	IPR	
		(4)	Genome technology	
		(5)	PBR	
		(6)	Neuroinformatics tools and database	
	(c)	Atte	empt any Two	10
		(1)	Write an essay on growth of International Bioinformatics research.	
		(2)	Transcriptome and proteome	
		(3)	Describe in detail about high throughput sequencing method.	
		(4)	Explain Cheminformatics and its tools and databases	
		(5)	What are the agreements and organizations overseeing IPR ?	
		(6)	Biosafety & Bioethics	

2	(a)	Exp	lain any Three	6
		(1)	Therapeutic target	
		(2)	Bioinformatics patents	
		(3)	Bioethics	
		(4)	Clustering	
		(5)	Ligand	
		(6)	Example for assembly tool	
	(b)	Exp	lain any Three	9
		(1)	Traditional knowledge	
		(2)	Cheminformatics tools and database	
		(3)	What are the tools available to assemble the sequenced fragments?	
		(4)	What is the modification in Sanger's method?	
		(5)	What are the factors affecting commercialization of Bioinformatics?	
		(6)	Comparative proteomics	
	(c)	Atte	empt any Two	10
		(1)	NGS	
		(2)	How SNPs are used for designing the personalized dr	ug?
		(3)	Immunoinformatics	
		(4)	IPR impacts on Biotechnology Research in India	
		(5)	Four levels of Biosafety.	