# STELLA MARIS COLLEGE (AUTONOMOUS) CHENNAI –600 086 (For candidates admitted from the academic year 2011 – 12 & thereafter)

**SUBJECT CODE: 11BI/PC/RB44** 

### M. Sc. DEGREE EXAMINATION, APRIL 2016 BIOINFORMATICS FOURTH SEMESTER

**COURSE** 

: CORE

PAPER TIME	: RECENT ADV : 3 HOURS	ANCES IN B	IOINFORMA	ATICS MAX. M	ARKS:	100
		SECTI	ON – A			
ANSWER A	ALL QUESTIONS				(20 X 1:	=20)
1. File name asr	extension for SMILE me b. sr		c. sml	d. sm	18	
	he chemical structure i N=C=O b. C=N=C=					
3. The chem a. Ga	informatics software ates b. Craig			Del Mar		
a. se	pharmacogenetics help lection of right drug lection of right dose	doctors to trea	-	ferent to one for o	each patio	ent
a. ch	tific field Cheminform emistry sysical synthesis	natics is the stu b. Biology d. All the a	•	of		
containing a. SF	R model is based on a g two1 P4-hybridized P-hybridized	nitrogen atom. b. S	isting of a six SP <sub>3</sub> -hybridized SP <sub>2</sub> -hybridized		atic ring	
7. R is an ir a. S	nplementation of the b. C++	programming i	language of d. Java			
a. ge c. no	the black spots mean in one was strongly expre- one DNA has bonded to one was neither strongle	ssed o the DNA in t	b. gene w the gene locat		sed	
	etural fragments, finge D descriptors b. 1D o			tors d. 3E	) descript	ors
	ety margin of pre clinic erapeutic dose b. le	cal trial is thal dose c. o	ptimal dose	d. both a and	l b correc	t

11. Gene expression Omnibus segregates data into three principle components. There are, a. sequence, protein, amino acid b. header, extensive, empirically c. platform, sample, series d. database, sequence, dataset
12. The treatment of Alzhemer's disease depends on variation of the following gene a. APOE b. CYP c. ACE d. All are correct
13. In R programming when the same or similar tasks need to be performed multiple times done by  a. data shift b. data frame c. database d. loop
<ul><li>14. Which of the following is the amount of a drug absorbed per the amount administered?</li><li>a) Bioavailability b) Bioequivalence c) Drug absorption d) Bioinequivalence</li></ul>
15. The mapping for T cell epitopes by web based tools a. EpiMatrix b. ClustiMer c. JanusMatrix d. All the above
16. Toxicogenomics combines toxicology with other high throughput molecular profiling technologies such as a. transcriptomics b. proteomics c. metabolomics d. All the above
17. DNA microarrays allow detection of Gene mutations using? a. Polymerase Chain Reaction b. Cloning c. Southern Blotting d. Hybridization
<ul> <li>18. How can scientists determine the function of or differences between cell types? They can examine the:</li> <li>a. Number of nucleotide bases in genes versus intergenetic sequences.</li> <li>b. Amount of mRNA expressed for each gene in a cell type, and then compare that information between cell types.</li> <li>c. Amount of mutations between genes in the intergenetic spaces.</li> <li>d. Number of tRNA copies for a particular cell type.</li> </ul>
<ul> <li>19. How is a microarray constructed? In each spot, there are:</li> <li>a. Copies of all the genes for an organism.</li> <li>b. Multiple copies of one gene; each spot has copies for a different gene.</li> <li>c. Multiple copies of intergenetic sequences, which bind to genes in the samples.</li> <li>d. Copies of intergenetic sequences, which promote the replication of DNA in a sample.</li> </ul>
<ul><li>20. How do the beads in the column separate mRNA from all other RNA? The beads contain:</li><li>a. Sequences that magnetically separate the mRNA.</li><li>b. A glue-like substance derived from spider webs.</li></ul>

c. Poly-T's.

d. A sequence of uracil's that bind to the Poly-A tail.

#### **SECTION - B**

## ANSWER ANY FOUR QUESTIONS. EACH ANSWER SHOULD NOT EXCEED 500 WORDS. $(4 \times 10 = 40)$

- 21. What is Pharmacokinetics? Explain briefly about the safety metabolisms
- 22. Describe the different tools used for Vaccine development
- 23. Define PubChem. Explain the role of PubChem on searching the molecules
- 24. Illustrate the salient feature of R programming on database management.
- 25. Write short notes on molecular descriptor and Finger prints
- 26. Explain in details about application of DNA microarray on gene sequencing.
- 27. Give an account of role of Immunoinformatics in personalized medicine

#### **SECTION - C**

## ANSWER ANY TWO QUESTIONS. EACH ANSWER SHOULD NOT EXCEED 1200 WORDS. $(2 \times 20 = 40)$

- 28. Describe the pharmacogenomics and toxicogenomics of Alzheimer disease
- 29. Explain in details about the 3-D QSAR and its applications.
- 30. Write the features of any one chemical drawing packages used for searching & construction of 3 D structures of chemical molecule.
- 31. Explain the process involved in designing the Microarray experiments and data analysis

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