B.Sc. MICROBIOLOGY FIFTH SEMESTER (SPECIAL REPEAT) BIOINFORMATICS

BMB-503 (Use Separate Answer Scripts for Objective & Descriptive) Full Marks: 70 Duration: 3 hrs. [PART-A: Objective] Time: 20 min. Marks: 20 1x20 = 20Choose the correct answer from the following: 1. Which of the following is an example of tertiary structure of protein? a. Greek key motif b. Helix turn helix c. EF hand d. All 2. Beta pleated sheet is found mostly in b. Globular protein a. Fibrous protein d. All c. Transmembrane protein 3. Which of the peptides will be least deflected by magnetic field in mass spectrometry? a. 120 KDA b. 350 KDA c. 1256 KDA d. 100 KDA 4. Which of the following is a E. Coli metabolic pathway and transcriptional regulation database? a. WIT b. Regulon DB d. Metacyc c. Ecocyc 5. Which of the following database is of short sequence pattern and profile of protein? b. iProclass a. PROSITE d. Pfam c. PDB 6. Which database classifies proteins based on class and homology? b. SCOP a. CATH d. TrEMBL c. swissprot 7. Which of the following is the best measure of central tendency? a. Median b. Mode d. None of the above c. Mean 8. The _ ignores the negative sign.

b. Mean Deviation

d. None of the above

b. Mean = n(1-p), Variance = np(1-p)

d. Range

a. Ouartile Deviation

c. Standard Deviation

9. In a binomial variable X with parameters n and p

a. Mean = np(1-p), Variance = np

c. Mean = np, Variance = np(1-p)

10. The number of persons died due to a rare diseaa. Binomial distributionc. Normal distribution	nse is an example of b. Poisson distribution d. None of the above
11. Degree of freedom is associated with a. t test c. F test	b. Chi-square test d. All of the above
12. If the calculated value of the test statistic is less a, the null hypothesis is not rejected c. no conclusion	than its critical value, then b. the null hypothesis is rejected d. None of the above
13. The two variables X and Y have linear relation, between X and Y is a. +1 c. ±1	if the correlation coefficient b1 d. 0
 14. If one of the regression coefficients is greater that a. less than or equal to 1 c. equal to 1 15. Which of the following transposons is found in a. Ty elements c. Retroviral elements 	b. greater than or equal to 1d. None of the above.
16. Highly variable repeats of almost 15 base pairsa. microsatellitec. VNTR	are called b. minisatellites d. Both b and c
17. Which is the algorithm for using encrypted filea. DESc. Both a and b	transfer? b. Triple DES d. DFS
18. Which of the entity in RDBMS is called an attrib a. row c. Table	bute? b. column d. All
19. Which of the following file transfer protocol is a. FTPc. TCP	used over remote network? b. SFTP d. SSH
20. What is the size of yeast genome? a. 24 mb c. 2400 mb	b. 16 mb d. 28 mb

(PART-B : Descriptive)

Time: 2 hrs. 40 min. Marks: 50

[Answer question no.1 & any four (4) from the rest]

1.	over pairwise alignment? Maximum parsimony is considered better than maximum likelihood. Comment	=6
	b. Differentiate between motif and domain AND alpha helix and beta sheet.	2+2=4
2.	a. Describe the process of 2D gel electrophoresis with suitable diagram	6
	b. Write a note on lead identification and lead optimisation in drug discovery.	4
3.	a. The average height of 10 males is 66 inches and standard deviation is 9 inches. Test the hypothesis that the average height is greater than 64 inches. [Given $t_{0.05,9} = 1.833$]	6
	b. Distinguish between correlation and regression.	4
4.	a. Write the characteristics of an ideal measure of central tendency	4
	b. Find the coefficient of variation of the following distribution Class limit: 59 – 61 61 – 63 63 – 65 65 – 67, 67 – 69 Frequency: 4 30 45 15 6	6
5.	a. Wrte the properties of Poisson distribution	4
	b. If X is normally distributed with mean μ = 20 and standard deviation σ = 2, Find P(18 < X < 23). [Given, Φ (1) = 0.3413 and Φ (1.5) = 0.4322]	6
6.	Which is the most suitable database to retrieve nucleotide sequences? Explain the features of that database and also the file format in which it can be retrieved in detail.	10
7.	a. Differentiate between local and global alignment. Which type of alignment is used in homology finding?	5
	b. Discuss about the data or sequence submission tools highlighting	5

- 8. a. Eukaryotic genome is superior to prokaryotic genome. Subtantiate you answer with atleast 5 points.

 b. Evplain this poi and among angles along with Ramachandran plot

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 - b. Explain phi, psi and omega angles along with Ramachandran plot and primary structure of protein.

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