

GIET UNIVERSITY, GUNUPUR – 765022 RD19MSC038 Roll No: Total Number of Pages: 1 AR-19 M.SC M.Sc 1ST SEMESTER REGULAR EXAMINATIONS, NOV/DEC 2019-20 BTPC104-BIOINFORMATICS & BIOSTATISTICS Time: 3 Hours Max Marks: 80 The figures in the right hand margin indicate marks. **SECTION A** Q.1 Answer any four of the following: [4 X4 = 16]What is Sequence Alignment? Differentiate between Local and Global alignment with example? What is protein structure? How to visualize a protein structure? What is dot matrix? How to do dot matrix analysis? c What is homology modeling? Differentiate between threading and homology modeling. d What is probability distribution? Write down various probability distribution functions used in statistically analysis? f Find out the mean and variance of first ten natural numbers? 2. Answer all questions from the following $[2 \times 8 = 16]$ What is NCBI? Name the nucleotide database of NCBI. What is PDB? Why it is important? b What is BLAST? Where it is available to use? c What are the significance of Sequence Alignment? d Write down two important softwares used for homology modeling? e What is Multiple Sequence Alignment? Name a tool use in producing MSA? f What is variance? How it is related to standard deviation? g Differentiate between paired t-Test and un-paired t-Test? h **SECTION-B** Answer all Questions: [16 x4 = 64]What is FASTA? How to carry out FASTA and explain with its algorithm? b What is Similarity Search? Explain the BLAST algorithm to carry out the similarity search. 4. What is global alignment? How to do it using Needleman Wunsch Algorithm? a b What is scoring matrices? Explain detail about PAM matrix? 5. What is molecular modeling? Write down steps to carry out the homology modeling? a What is to generate Hidden Markov Model (HMM)? Explain its application in biological b research 6. What is Biostatistics? Write down the application of statistics in biological search? a

Calculate the mode and variance of the following

14

30-40 40-50 50-60

22

32

60-70 70-80

17

18

80-90

11

20-30

11

b

CI: 10-20

5

f: