## M. SC. (MEDICAL BIOTECHNOLOGY) SEM-III (CHOICE BASED CREDIT SYSTEM): WINTER - 2017

**SUBJECT: GENOMICS & PROTEOMICS** 

Day: Friday Time: 10.00 AM TO 01.00 PM W-2017-1057 Date: 10/11/2017 Max. Marks: 60 N.B: 1) Q.No1 and Q.No 5 are COMPULSORY. Out of the remaining questions, attempt Any **TWO** from each sections. 2) Answer to both the sections should be written in **SEPARATE** answer book. 3) Figures to the right indicate FULL marks. 4) Draw neat labeled diagram WHEREVER necessary. **SECTION-I Q.1** Answer the following: [10] What are pseudo genes? What is their importance in genome? a) b) Define RFLP & SNP. Briefly explain the functions of map repositories. c) What are P elements? d) Define gene order & synteny. **Q.2** Answer the following: (ANY TWO) [10] What are linkage maps? Give practical uses of them. a) Give the steps used in construction of genetic maps. Differentiate between **b**) cytogenetic & physical maps. Write a note on HGP. Explain its outcomes. c) Q.3 Write short notes on: (ANY TWO) [10] Prokaryotic transposons a) OMIM **b**) Gene clustering c) Give a comparative statement on genome size, features and characteristics of [10] **Q.4** different organisms. Explain with suitable examples. Explain in brief the concept of chromosome rearrangement, compositional analysis & composite genes. **SECTION-II** [10] Q.5 Define: d) FSSP **CATH** a) **SCOP** e) MMDB b) **PSSM** c) Answer the following: (ANY TWO) [10] **Q.6** Explain in brief transcriptomics & its applications in data analysis. a) Describe the steps involved in microarray data analysis. b) Write a note on SARF. c) [10] Write short notes on: (ANY TWO) **Q.7** Comparative genomics a) **BLOCKS b**) Motif, pattern & prosite c) Explain automation of proteomic analysis with respect to organ comparison, **Q.8** spatio-temporal comparison, intra-species comparison. Explain the following: a) Phage antibody as a tool b) Protein expression analysis

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