## M. Sc. Bioinformatics Sem.-II (C.B.C.S.) (2013 Course) / Advanced Diploma in Bioinformatics Sem.-II (C.B.C.S.) (2013 Course) : WINTER - 2018

## SUBJECT: GENOMICS AND PROTEOMICS

Day:

Friday

Time: 02.00 PM TO 05.00 PM

W-2018-1258 Date: 26/10/2018 Max Marks: 60 **N.B.:** Q. No. 1 and Q. No. 5 are COMPULSORY. Out of the remaining, attempt ANY 1) TWO from each Section. Each section should be solved in **SEPARATE** answer books. 2) 3) Figures to the right indicate FULL marks. Draw neat labeled diagrams WHENEVER necessary. 4) **SECTION I Q.1** Explain the principal of: (10)a) Sanger sequencing b) PCR DNA fingerprinting c) d) BLAS Pipmaker e) Answer the following: **Q.2** (10)What is third generation sequencing? Explain any one method in brief. a) What was the basic idea behind HGP? Explain its outcomes. b) Q.3 Write short notes on: (10)Structural Genomics a) Genomic Data Browsers **Q.4** Give an account of comparative genomics of organisms. (10)OR Explain the concept of pharmacokinetics in detail. Give emphasis on ADMET properties. **SECTION-II** Define: **Q.5** (10)**IEF** a) **b)** PAGE Protein chip d) InterPreTS c) **GRID** e) **Q.6** Answer the following: (10)Write a note on protein separation techniques. a) How do you analyze images in 2D gels? Explain. b) **Q.7** Write short notes on: (10)a) Mass Spectrometry **b)** Protein Sequencing What are protein-protein interactions? How do you analyze it? Explain with (10) **Q.8** example. OR Write in brief on DIP and MINT.