M. SC. BIOINFORMATICS SEM.-II (C.B.C.S.) (2013 COURSE) / ADVANCED DIPLOMA IN BIOINFORMATICS SEM.-II

(C.B.C.S.) (2013 COURSE) : SUMMER - 2018 SUBJECT : GENOMICS & PROTEOMICS

Time: 2.00 P.M. TO 5.00 P.M. :Thursday Day 5.2018 - 1129 :12-04 2018 Max. Marks: 60 Date **N.B.**: Q.No.1 and Q.No.5 are COMPULSORY. Out of the remaining questions 1) attempt ANY TWO questions from each section. 2) Answers to both the sections should be written in **SEPARATE** answer books. Figures to the right indicate FULL marks. 3) SECTION - I [10] Define **ANY FIVE** of the following: **Q.1** a) SNP b) Pharmacokinetics **Promoters** c) d) Drug Target **EST** e) Paralogues genes f) [10] Q.2 Write short notes on **ANY TWO** of the following: Pyrosequencing a) **OMIM** b) Genome Assembly Q.3 Answer the following: What is gene expression analysis? Explain in detail. [03] Differentiate between metabolic diseases and pathogenic diseases with [03] example. Explain with example different genomic browsers. Why they are not under one [04] roof? OR Write in detail ORF prediction algorithm. **Q.4** Explain in detail MUMmer and VISTA algorithms. [10] Write a note on Gene Synteny and GeneOrder tools. **SECTION - II Q.5** Define ANY FIVE of the following: [10]Proteome a) **GRID** b) Microarray c) d) 2D Gels **BIND** e) **MINT** f) Write short note on ANY TWO of the following: **Q.6** [10] Size exclusion chromatography a) **IEF b**) Scope of proteomics **Q.7** Answer **ANY TWO** of the following: [10] a) Explain in detail mass spectroscopy and its applications. Explain protein sequencing steps in detail. Explain protein engineering in detail and its application. **Q.8** Enlist Clinical and biomedical applications of proteomics. [10] OR Explain in detail different databases used for protein-protein interactions.