



Avinashilingam Institute for Home Science and Higher Education for Women

(Deemed to be University under Category 'A' by MHRD, Estd. u/s 3 of UGC Act 1956)

Re-accredited with 'A+' Grade by NAAC. Recognised by UGC Under Section 12B

Coimbatore - 641 043, Tamil Nadu, India

Master's Degree Examination – June / July 2021

II Semester

Class: I PG
Major: Bioinformatics

Time : 3 Hours
Max Marks : 100

20MBIC10 Introduction to OMICs Technologies

Part A

10 X1 = 10

Choose the Correct Answer

- Identify the size of the human genome from the following. CO1 K1
 - 2 billion base pairs
 - 3 billion base pairs
 - 4 billion base pairs
 - 1 billion base pairs
- Choose the correct definition of metagenomics from the following. CO1K3
 - Study of genetic materials of human
 - Study of genetic materials of bacteria
 - Study of genetic materials recovered from environmental samples
 - Study of genetic materials of model organisms
- Infer the expansion of EST from the following. CO2K2
 - Expressed Structure Tags
 - Expressed Sequence Tags
 - Expressed Sequence Time
 - Expressed Structure Time
- Select the technique used in gene expression analysis. CO2K4
 - Microarrays
 - Electrophoresis
 - Chromatography
 - Alignment
- Identify the mobile metabolomic interface from the following: CO3K1
 - Medwin
 - Metlin
 - Metmob
 - Metome
- Name the metabolic pathway database from the options. CO3K1
 - BRENDA
 - NCBI
 - Uniprot
 - KEGG
- Infer the definition of Proteomics from the given options. CO4K4
 - The Study of proteins in an organism
 - The study of proteome at a given point in time
 - The Study of Protein in a cell
 - The study of genome that encodes proteins
- Choose the protein sequencing method from the following. CO4 K3
 - Sanger's method
 - Mass Spectrometry
 - Next generation Sequencing
 - Pyrosequencing
- Name the term used for the technique of comparison of genomes. CO5K1
 - Comparative genomics
 - Structural Genomics
 - Functional genomics
 - Artificial Genomics
- Identify the term used to refer to the statistical analysis that combine the results of previous multiple scientific studies. CO5K1
 - Metagenomics
 - Meta-analysis
 - Sub-analysis
 - Statistical analysis

Part B

5x6=30

Answer ALL Questions

Each answer should not exceed 400 words or two pages

- 11.a. Illustrate about C-value Paradox with examples. CO1K3
(or)
- 11.b. Demonstrate the techniques involved in metagenomics. CO1K3
- 12.a. Appraise on the applications of SAGE. CO2K4
(or)
- 12.b. Conclude on the salient features of array express with examples. CO2K4
- 13.a. Categorize the resources for metabolic pathway analysis and explain in brief. CO3K4
(or)
- 13.b. Evaluate the principle and applications of Capillary Electrophoresis with an example. CO3K5
- 14.a. Determine the method of separation of proteins using gel Electrophoresis. CO4K3
(or)
- 14.b. Report on the various methods of phylogenetic profiling for predicting protein-protein interactions. CO4K3
- 15.a. Interpret the applications of Comparative genomics with examples. CO5K3
(or)
- 15.b. Summarize the salient features of Evolutionary genomics. CO5K5

Part C

5 x 12 = 30

Answer ALL Questions

Each answer should not exceed 800 words or four pages

- 16.a. Describe the methods and advantages of different genome sequencing techniques in detail. CO1K2
(or)
- 16.b. Summarize the salient features of Human genome project and explain its notable findings in detail. CO1K2
- 17.a. Discuss in detail about principle and method of gene expression analysis using microarrays. CO2K2
(or)
- 17.b. Illustrate with examples about transcriptomics study. CO2K3
- 18.a. Assess the applications of GC/MS and NMR in metabolomics. CO3K5
(or)
- 18.b. Point out the different tools and databases used in Metabolomics and explain in detail. CO3K4
- 19.a. Compare and contrast the various protein sequencing methods in detail. CO4K4
(or)
- 19.b. Distinguish between the experimental and computational methods of determining protein-protein interactions. CO4K4
- 20.a. Discuss in detail about the ethical, legal and Social issues involved in personal genomics. CO5K2
(or)
- 20.b. Employ the techniques involved in integrating Omics data and explain in detail. CO5K3
