



**MASTERS OF PHILOSOPHY (M. PHIL.)**  
**BIOINFORMATICS**  
**DETAILED SYLLABUS**  
**SESSION 2013-14**



## **PAPER I**

### **RESEARCH METHODOLOGY THEORY AND TECHNIQUES**

#### **UNIT - I**

Research: Definition, Importance and Meaning of research, Characteristics of research, Types of Research, Steps in research, Identification, Selection and formulation of research problem, Research questions – Research design – Formulation of Hypo Dissertation, Review of Literature.

#### **UNIT – II**

Sampling techniques: Sampling theory, types of sampling – Steps in sampling – Sampling and Non-sampling error – Sample size – Advantages and limitations of sampling.

Collection of Data: Primary Data – Meaning – Data Collection methods – Secondary data – Meaning – Relevances, limitations and cautions.

#### **UNIT – III**

Statistics in Research – Measure of Central tendency, Dispersion, Skewness and Kurtosis in research, Hypo Dissertation, Fundamentals of Hypo Dissertation testing, Standard Error, Point and Interval estimates, Important Non-Parametric tests: Sign, Run, Kruskal, Wallis tests and Mann, Whitney test.

#### **UNIT – IV**

Para metric tests: Testing of significance, mean, Proportion, Variance and Correlation, testing for Significance of difference between means, proportions, variances and correlation co-efficient. Chi-square tests, ANOVA, One-way and Two-way.

#### **UNIT– V**

Research Report: Types of reports, contents, styles of reporting, Steps in drafting reports, editing the final draft, evaluating the final draft.

#### **Reference Books:**

1. Statistical Methods - S.P. Gupta
2. Research Methodology Methods and Techniques - C.R. Kothari
3. Statistics (Theory and Practice) - B.N. Gupta
4. Research Methodology Methods and Statistical Techniques - Santosh Gupta



## **PAPER-II**

# **BIOLOGICAL DATABASES, DATA MINING AND MICRO ARRAYS**

### **UNIT-I**

Biological database - Sequence databases - Other specialized databases – Microarray databases - Database browsers and search engines.

### **UNIT-II**

Data mining definition – Classification and clustering of data – Association rules – Data visualization.

### **UNIT-III**

Introduction to Microarrays - Oligonucleotide and Spotted cDNA arrays – Design considerations for microarray experiments – Goals of a microarray experiment.

### **UNIT-IV**

Basic research with DNA microarrays – Microarrays and Cancer - Myeloid leukemia (AML) vs. acute lymphoblastic leukemia (ALL) data analysis.

### **UNIT-V**

Use of array analysis programs – SAM - TIGR programs – MEV.

### **References:**

1. Analysis of DNA Microarray Data by Steen Knudsen.
2. Discovering Genomics, Proteomics, and Bioinformatics by A.M. Campbell and L.J. Heyer.



## **PAPER-III**

### **STRUCTURAL BIOINFORMATICS**

#### **UNIT I:**

Structural features of biomolecules – techniques used to determine the structure of biomolecules - geometrical parameters – potential energy surfaces – molecular graphics – hardware and software requirements – Internet – mathematical concepts, molecular file formats

#### **UNIT II:**

Structure prediction – secondary structure – homology modeling, fold recognition and ab initio 3D structure prediction – structure comparison and alignment – prediction of function from structure.

#### **UNIT III:**

Molecular dynamic using simple models – simulations with continuous potentials – advantage of constant temperature and pressure simulation – solvent effects – analysis of conformational changes during molecular dynamic simulation.

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#### **UNIT IV:**

Molecular docking – structure based drug design – de novo approach – molecular similarity – quantitative structure-activity relationship – 3D pharmacophore derivation and matching – importance of molecular modeling in drug discovery, Cheminformatics and its applications, Combinatorial libraries and Chemical diversity

#### **UNIT V:**

Protein stability and folding-SCOP-DALI-assignment of protein structures to genomes-determining gene function through conserved protein structure-prediction of protein function-approaches to protein structural genomics



### References:

1. Mount, D.W. (2001).” **Bioinformatics – Sequence and Genome Analysis**”, 1<sup>st</sup> Edition, Cold Spring Harbor Laboratory Press, New York, USA.
2. Westhead, D.R., Parish, J.H. and Twyman, R.M. (2003). **Instant Notes Series Bioinformatics**, 1<sup>st</sup> Edition, Viva Books Private Limited, New Delhi, India.
3. Ignacimuthu (s.j.), S. (2005). **Basic Bioinformatics**, 1<sup>st</sup> Edition, Narosa Publishing House, New Delhi, India.

### PAPER-IV DISSERTATION

