BHARATHIAR UNIVERSITY-COIMBATORE-641046

M.Phil Bioinformatics Degree Course, University Department

(For candidates admitted during the academic year 2006-2007 batch onwards)

PART I – SYLLABUS

PAPER I – RESEARCH METHODOLOGY

UNIT I:

Research Methodology:Introduction ;meaning of research;objectives of research;types of research; research approaches; significances of research; research methods vs methodology; research and scientific method; importance of knowing how research is done; research process; criteria of good research; problem encountered by researchers in india; Defining the research problem; What is the research problem? Selecting the problem; Techniques involved in defining the problem; Research design; Need for research Design; Features of good Design, important concepts relating to design; different research designs; basic principles of experimental designs.

UNIT II:

Hypothesis testing: What is Hypothesis? Basic concepts concerning testing of hypothesis; procedure for hypothesis testing; Probability; Markov models and Hidden Markov Models; Probability distribution; Binomial; Poisson; Normal distribution and Multiple testing Methods ANOVA; Test of significance-t-test; F-test

UNIT III:

Interpretation and Report writing; Meaning of Interpretations; Techniques of interpretation; precautions of interpretations; significances of report writing; Different steps in report writing; layout of the research project; types of report; oral presentation; mechanics of writing a research project; precautions for writing research reports; conclusions

UNIT IV:

Elements of C Programming;Features of C;Variables;Constants;keywords;Data types;operators;statements;loops – simple programs using Loops,Arrays – integer arrays – character arrays – simple programs using arrays;Introductions to functions – simple programs using functions – Introduction to pointers,structures string Manipulations using pointers and arrays;Files;Defining and opening a file,Closing a file,input/output operations on files PERL:Basic syntax-I/O – Variables,strings & arrays-control structures – regular expressions – simple programs

UNIT V:

Algorithms in computer sciences inspired by biology genetic algorithms, Neural networks and path optimization

Reference:

- 1 Kothari. C.R. 2004 Research Methodology Methods and Techniques, New Age International (P)Ltd
- 2 E Balagurusamy.Programming in ANSI C Tata Mc Graw Hill
- 3 Randa L.Schwartz,tom phoenix,learning perl,third edition

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PART I – SYLLABUS

PAPER II – ADVANCES IN BIOINFORMATICS

UNIT I

High throughput genome sequencing and genome assembly, Gene finding algorithms, DNA Microarrays and large gene expression data sets, clustering algorithms

UNIT II

Protein and Nucleic acid sequence alignments, Sequence databases, the use of algorithm BLAST, Multiple sequence alignments

UNIT III

Protein Structure analysis; Protein structure databases; Protein Structure comparison; Fold Recognition; 3D – ID Profiles; Threading; Comparitive Structure Modeling

UNIT IV

Phylogeny(evolutionary trees)biological networks;pathway analysis

UNIT V

Emerging new ideas on treating biological systems; Pharmacogenesis and its applications; SNPs and their applications

Reference:

- 1 Andreas D Baxevanis and BF Francis Oueliene 2001 Bioinformatics A Practical Guide to the analysis of Genes and Proteins, A John wiley & sons, INC, Pub
- 2 David W Mount,2003 Bioinformatics Sequence and Genome Analysis,CBS Publishers,Ian Korf,Mark Yandell & Joseph Bedell,2003
- 3 Ian Korf.Mark Yandel & Joseph Bedell.2003 BLAST(O' RELLY)SPD Pvt Ltd
