M.Sc. BIOINFORMATICS

REGULATIONS AND SYLLABI

(Effective from 2007-2008)



Centre for Bioinformatics SCHOOL OF LIFE SCIENCES PONDICHERRY UNIVERSITY PUDUCHERRY

PONDICHERRY UNIVERSITY SCHOOL OF LIFE SCIENCES

Centre for Bioinformatics

SYLLABUS FOR M. Sc. BIOINFORMATICS

(Academic Year 2007-2008 onwards)

Semester	Course Code	Course Title	Credits
	BINF 401	Cell and Molecular Biology	3
	BINF 402	Physics and Chemistry for Biologists	3
Ι	BINF 403	Mathematics for Biologists	3
	BINF 404	Bioinformatics–I	3
	BINF 405	Data Structures and Programming Concepts	3
	BINF 406		3
	BINF 407	Computers for Biologists (Elective)	3
	BINF 451	Lab - Cell and Molecular Biology	1
	BINF 452	Lab - Biological Databases	1
	BINF 453	Lab - Basics of Computer & Operating Systems	1
	BINF 454	Lab - Programming in C/ C++	1
	BINF 421	Genomics and Proteomics	3
	BINF 422	Biophysical Chemistry	3
	BINF 423	Bioinformatics–II	3
	BINF 424	Statistics for Biologists	3
Π	BINF 425	Programming in Java	3
	BINF 426	Relational Database Management System	3
	BINF 427	Fundamentals of Algorithms	3
	BINF 455		1
	BINF 456		1
	BINF 457	Lab - Biosequence Analysis	1
	BINF 501	Structural Biology	3
	BINF 502	Molecular Modeling and Drug Design	3
	BINF 503	Programming in Perl	3
	BINF 504	Applications of Bioinformatics	3
III	BINF 505	Networks and Distributed Computing	2
	BINF 506	Scientific Presentation	2
	BINF 551	Lab – Structural Biology	1
	BINF 552	Lab – Molecular Modeling and Drug Design	1
	BINF 553	Lab - Programming in Perl	1
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	BINF 520	Bioethics and Intellectual Property Rights	3
IV	BINF 521	Finishing School	2
	BINF 522	Project	4

SEMESTER - I BINF 401 – CELL AND MOLECULAR BIOLOGY

Unit-I

Architecture of prokaryotic and eukaryotic cells and tissues. Dynamics of the eukaryotic cell-Molecules of life- Cellular evolution- assembly of macromolecules and Origin of life- integrated structural organization of prokaryotic and eukaryotic cells- Concept of a composite cell and Molecular composition of cells; Cell division - mitosis and meiosis, eukaryotic cell cycle and its regulation

Unit-II

The mitochondrion – structure, electron transport chain, oxidative phosphorylation, Chemiosmotic process; Chloroplasts - photosynthesis - photosystems, Calvin cycle, Dark reactions

Unit-III

Cell Communication – membrane transport – principles, active transport, ion channels; Protein sorting – an overview of targeting proteins to mitochondria, nucleus, endoplasmic reticulum, lysosome and plasma membrane

Unit-IV

Organisation of eukaryotic genome- components of eukaryotic chromatin- chromatin and chromosome structure- DNA-supercoiling -linking number- satellite DNA-possible functions-Cot curve- C- value paradox.

DNA replication- Prokaryotic and eukaryotic DNA replication, mechanism of replication. Enzymes and necessary proteins in DNA replication. Telomeres, telomerase and end replication. Role of telomerase in aging and cancer.

Unit-V

Transcription- Prokaryotic and eukaryotic Transcription- RNA polymerases- general and specific transcription factors- regulatory elements- mechanism of transcription regulation- Transcription termination; Post transcriptional modification

Translation- Genetic code- Prokaryotic and eukaryotic translation- translational machinery-Mechanism of initiation- elongation and termination- Regulation of translation.

Recommended Texts:

- 1. Karp, G. (2005) "Cell and Molecular Biology: Concepts and Experiments"; Fourth Edition, Wiley Publishing Co. USA
- 2. Lodish, H., Scott, M.P., Matsudaira, P., Darnell, J., Zipursky, L., Kaiser, C.A., Berk, A. and Krieger, M. (2003) "Molecular Cell Biology"; Fifth Edition, W.H. Freeman and Co., New York.

7 Lectures

6 Lectures

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Reference Books

- 1. Alberts, B., Johnson, A., Lewis, J., Raff, M., Roberts, K and Walter, P. (2002) "Molecular Cell Biology of the Cell", Fourth Edition, Garland Science, Taylor and Francis Group, USA.
- 2. Lewin, B. (2004) "Genes VIII"; Eighth edition, Pearson Education International.
- 3. De Robertes and De Robertis (2002) "Cell and Molecular Biology", Saunders College, Philadelphia, USA

5 Lectures

7 Lectures

11 Lectures

5 Lectures

SEMESTER - I BINF 402 – PHYSICS AND CHEMISTRY FOR BIOLOGISTS

Unit-I

8 Lectures Mechanics - Newton's Law of motion - Mechanics of a particle - Work-energy principle -Rotational dynamics - Static and dynamic equilibrium - Conservation laws. Thermodynamics -Concept of temperatures – Thermodynamic equilibrium: Zeroth law – Conservation of energy: first law - Concept of entropy: second law - Absolute zero of temperature: third law of thermodynamics.

Unit-II

Current electricity and circuits - Review of basic electrodynamics - Current and current density - Ohm's law - Conductivity in metals, insulators, semiconductors and ionic liquids -Electrical energy and dissipation – Combinations of resistors – Kirchoff's fules – RC circuits – RL and LC circuits.

Unit-III

Optics and Lasers – Interference: single and multiple beam interference – Diffraction at single and multiple rectangular slits – Polarization – Optical instruments – Resolving power – Introduction to LASERs and their applications.

Unit-IV

Chemistry – Introduction to Organic Functional Groups and ionic reaction mechanisms: Functional Groups, Reactive intermediates - electrophiles nucleophiles, radicals, substitution reactions, addition reactions, elemination reactions, redox reactions. Heterocyclic Chemistry -Introduction to heterocyclic compounds: Nomenclature, Reactivity of five membered and six membered heterocycles

Unit-V

Spectroscopy of organic molecules - Spectroscopic identification of simple organic molecules -Infrared Spectroscopy, nuclear magnetic resonance Spectroscopy, mass spectrometry

Recommended Texts:

- 1. Resnick, R., Halliday, D. and Walker (2001) "Fundamentals of Physics", Sixth edition, John Wiley and Sons, USA.
- 2. Tipler, P.A. (1999) "Physics for Engineers and Scientists"; Fourth edition, W.H. Freeman and Company, USA.

SEMESTER - I BINF 403 – MATHEMATICS FOR BIOLOGISTS

Unit I

Algebra - Logarithms – Quadratic equations – Simple problems on cubic equations – Solution of a system of linear equations using determinants

Unit II

Matrices and Vectors - Different types of matrices – Addition, Subtraction and Multiplication of matrices – Transpose of a matrix – Singular matrix - Inverse of a matrix

The concept of a vector – Dot product – Cross product – Angle between two vectors - Scalar triple product – Divergence – Curl – Equation of a normal

Unit III

8 Lectures Trigonometry and Analytical Geometry - Trigonometric ratios- De Moivre's theorem; Cartesian coordinates in two dimensions - Distance formula - Straight line - Slope of a line -Intercepts of a line - Equation of a line - Intersection of two lines - Angle between two lines -Circle, ellipse and parabola

Unit IV

Calculus: The concept of limit – Derivatives of simple, standard functions – Geometrical application of differentiation - Maxima and minima - Partial differentiation Definite and indefinite integrals – Numerical integration

Unit V

7 Lectures

7 Lectures

Differential equations: First order ordinary differential equations – Laplace transform and its application

Recommended Texts:

- 1. Narayanan, S. and Manicavachaagam Pillai, T.S. (1993) "Calculus, Vol. I and II"; Vishwanathan Printers and Publishers.
- 2. Veerarajan, T. (2003) "Engineering mathematics"; Third Edition, Tata McGraw Hill Publishing Co. Ltd, New Delhi.
- 3. Veerarajan, T. (2003) "Trignometry, Algebra and Calculus"; Third Edition, Tata McGraw Hill Publishing Co. Ltd, New Delhi.

Reference Books

- 1. Sharma, A.K. (2005) "Text Book of Integral Calculus", Discovery Publishing House.
- 2. Grewal, B.S. (2000) "Higher Engineering Mathematics"; Thirty seventh edition, Khanna Publishers, New Delhi.

7 Lectures

SEMESTER - I BINF 404 – BIOINFORMATICS–I

Unit I

Bioinformatics: an overview - Introduction to Computational Biology and Bioinformatics; Emergence of bioinformatics as a separate discipline; some of the biological problems that require computational methods for their solution; Role of internet and www in bioinformatics.

Unit II

Biological Data Acquisition – The form of biological information; DNA sequencing methods – basic DNA sequencing, automated DNA sequencing, DNA sequencing by capillary array and electrophoresis; Types of DNA sequences – genomic DNA, cDNA, recombinant DNA, Expressed sequence tags (ESTs), Genomic survey sequences (GSSs); RNA sequencing methods; Protein structure determination methods; gene expression data.

Unit III

Databases : Format and Annotation – Conventions for databases indexing and specification of search terms; Common sequencing file formats – NBRF/PIR, FASTA, GDE; Files for multiple sequence alignment – multiple sequence format (MSF), ALN format; Files for structural data – PDB format and NMR files; Annotated sequence databases – primary sequence databases (GenBank-NCBI, the nucleotide sequence database-EMBL, DNA sequence databank of Japan-DDBJ; Subsidiary data storage (ESTs, dbESTs, GSSs), unfinished genomic sequence data, organisms specific databases (EcoGene, SGD, MatDB, TAIR, FlyBase, OMIM, etc.); Protein sequence and structure databases (PDB, SWISS-PROT and TrEMBL); List of Gateways (NCBI, GOLD, MIPS, TIGR, UniGene)

Unit IV

Data : Access, Retrieval and Submission – Data access – standard search engines, Data retrieval tools – Entrez, DBGET and SRS (sequence retrieval systems); Software for data building; Submission of new and revised data.

Unit V

Sequence Similarity Searches – Sequence homology as product of molecular evolution; Sequence similarity searches; Significance of sequence alignment; Sequence alignment – global, local and free-space; Alignment scores and gap penalties; Measurement of sequence similarity; Similarity and homology.

Recommended Texts:

- 1. Mount, D. (2004) "Bioinformatics: Sequence and Genome Analysis"; Cold Spring Harbor Laboratory Press, New York. (ISBN 0-87969-712-1)
- 2. Baxevanis, A.D. and Francis Ouellellette, B.F. (1998) "Bioinformatics a practical guide to the analysis of Genes and Proteins"; John Wiley & Sons, UK.

Reference Books

- 1. Pevzner, P.A. (2004) "Computational Molecular Biology"; Prentice Hall of India Ltd, New Delhi.
- 2. Pevsner, J. (2003) "Bioinformatics and Functional Genomics"; John Wiley and Sons, New Jersey, USA.

3 Lectures

7 Lectures

10 Lectures

7 Lectures

SEMESTER - I

BINF 405 – DATA STRUCTURES AND PROGRAMMING CONCEPTS

Unit –I

Concepts in Computing

Overview and functions of computer systems – History of Computers – Storage Devices – Memory – Type of Processing – Types of Computers – Operating System Concepts – Internet and its resources.

Unit – II

Elementary Data Structures

Arrays, Stacks, Queues, Dequeues, Order Lists, Generalized List, Linear List, Linked lists, Circular Linked Lists, Doubly-Linked Lists, Infinite Lists, Hash tables, Hash functions, Recursive functions

Unit – III

Computing Algorithms

Algorithms in Computing, Analyzing algorithms, Designing algorithms – Sorting and Searching techniques: Bubble Sort, Merge Sort and Insertion sort – Binary Search

Unit IV

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Algorithms, flow-charts, programming languages, compilation, linking and loading, testing and debugging, documentation – C programming – variables and identifiers, data types, Conditional statements and loops – if, if-else statements, while, do-while, for loop, switch case, Structured Programming, Library Functions

Unit V

Object Oriented Programming Concepts

C++ – Abstract Data type, Encapsulation, Object, Message, Method, class, Inheritance, Polymorphism, Virtual Functions, Abstract Classes, Interface, Constructors & Destructors, Overloading & Overriding, Copy Instructor

Reference Books

- 1. Lafore, R. (2002) "Object Oriented Programming using C++", Fourth Edition, Sams Publishers.
- 2. Ritchie, D.M. (1996) " The C programming language", Second Edition, Prentice Hall Publishers, USA.
- 3. Balaguruswamy, E. (1985) "Computer Fundamentals and Applications", Second Edition, Tata McGraw-Hill Publishing Co. Ltd., India.
- 4. Fundamentals of Data Structures, E. Horowitz and S. Sahani, Galgotia Booksource Pvt. Ltd., (1999

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SEMESTER - I BINF 406 – GENERAL BIOLOGY (Elective)

Unit I

Molecules and Macromolecules of the Cell - Elements, molecules and macromolecules of the cell; Formation of macromolecules; Types of bonds; Structure of carbohydrates, lipids, DNA, RNA and proteins

Unit II

Cell Structure and Functions – Unity and diversity in life forms; Structure of virus and bacteria; Features of prokaryotic and eukaryotic cells; Levels of biological organization – cell, tissue and organs.

Unit III

Supramolecular Assemblies of the cell – Self assembly of macromolecules – ribosomes, chromosomes, membrane, collagen, actin and cellulose

Unit IV

Energy and Cellular Work – Energy input and output in cell; The role of ATP; Electron transfer reactions; Electron transfer molecules; Electron transport chains; Light-driven electron flow; Catabolism and Metabolism; Metabolic pathways

Unit V

Cell Cycle – Different phases of cell cycle; mitosis and meiosis; Regulation of cell cycle; Apoptosis

Recommended Texts:

- 1. Roberts, M.B.V. and King, T.J. (1987) "Biology: A functional approach"; Nelson Thornes Limited, USA.
- 2. Edwards, G.I. (2000) "Biology: The easy way"; Barrons Educational Series, USA

Reference Books

- 1. Murray, R.K., Mayes, P.A. and Graner, D.K. (1996) "*Harpers* Biochemistry", Appleton and Lange Publishers, USA.
- 2. De Robertes and De Roberties (2002) "Cell and Molecular Biology", Saunders College, Philadelphia, USA

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SEMESTER - I BINF 407 – COMPUTERS FOR BIOLOGISTS (Elective)

Unit I

Computer Organization

Fundamentals of computers – Block diagram of computer (input and output devices) – Generations - Advantages and Limitations of Computers - Basics of operating systems DOS, Windows NT & XP, UNIX – Application Softwares.

Unit II

Network Basics

Communication Technology – Networking – LAN, WAN & MAN, Intranet–Wireless communication – Internet.

Unit III

Introduction to Database systems

Fundamentals of database - Database models (Hierarchical, Network, Relational, Object-Oriented Models) – RDBMS - Database System Applications and Security.

Unit IV

Ms-Office

Introduction to M.S. office package - Word – creating a new document – templates and wizards – scientific data representation and basic calculations with EXCEL - Creating Tables and databases using Access – interactive presentations creating using Power Point

Unit V

Internet Technologies

Web Services – WWW, URL, DNS - Servers-E-mail server, WEB servers, Browsers-IP Addressing, IPV6.

Reference Books

- 1. Sherman, J. (2001) "Basic Computer Skills made easy", Butterworth-Heinemann Ltd, USA
- 2. Balaguruswamy, E. (1985) "Computer Fundamentals and Applications", Second Edition, Tata McGraw-Hill Publishing Co. Ltd., India.
- 3. Horwitz, E. and Sahni, S. (1978) "Fundamentals of Computers", Second Edition, WH Freeman & Co., USA.
- 4. Microsoft Office Manual

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SEMESTER - I BINF 451 – LAB - CELL AND MOLECULAR BIOLOGY

Exercises in Cell Biology

- 1. Paper Chromatography of Chlorophyll pigments
- 2. Estimation of Chlorophyll
- 3. Ascorbic acid estimation in different tissues of plants and animals.
- 4. Growth curve of bacteria.
- 5. Estimation of cell mass of bacteria.

Exercises in Molecular Biology

- 1. Isolation & Purification of genomic DNA from bacteria
- 2. Isolation & Purification of plasmid DNA
- 3. Agarose gel electrophoresis of chromosomal & plasmid DNA
- 4. Restriction Digestion of chromosomal & plasmid DNA
- 5. Isolation of DNA fragment from agarose gel

SEMESTER - I BINF 452 – LAB - BIOLOGICAL DATABASES

Exercises:

- 1. Entrez and Literature Searches.
 - a. PubMed
 - b. PubMed central
 - c. OMIM / OMIA
 - d. Citation matcher
- 2. SRS of Biological Databases
 - a. Nucleotide/ Genome Databases.
 - b. Protein Sequence Database.
 - c. Structure databases.
 - d. Protein Pattern Databases
- 3. File format conversion
 - a. FmtSeq
 - b. ReadSeq
 - c. Sequence manipulation Suite
- 4. Sequence Analysis
 - a. Dot Plot
 - b. Pairwise alignment
 - c. Multiple Sequence Alignment
- 5. Phylogenetic analysis using PHYLIP, Phylodraw, PAUP, Treeview, JalView.
- 6. Softwares
 - a. BioEdit.
 - b. GeneDoc
 - c. ClustalW / X, MEGA, MEME
- 7. Visualization Tool
 - a. RasMol
 - b. Cn3D
 - c. MolMol

SEMESTER- I BINF 453 – LAB – BASICS OF COMPUTER & OPERATING SYSTEMS

Exercises:

- 1. DOS Commands Internal Commands: Viewing a directory, Changing Directory, Renaming a Directory - File operations: Creating files, removing a file, renaming files, viewing a file - External commands: Copying a disk, Comparing disks
- 2. Overview of different versions of Windows –Working with Windows- Desktop Basic Layout, Icons, Opening Windows, Window Characteristics, Window Controls, Resize Windows, Arrange Windows, Taskbar.
- 3. Working with Programs: Basic Program Layout, WordPad Program, Scrolling in Documents, Moving Insertion Point, Delete & Insert Key, Selecting Text, Cut, Copy & Paste, Working with Multiple Programs.
- 4. Files & Folders: Organization, View Folder Structure, Working with Folders Search for Files, Organizing Workspace - Personal Desktop, Shortcuts, Start Menu, Start Properties, Display as Menu, Taskbar, Quick Launch.
- 5. Windows Properties Navigating Control Panel, Changing Theme, Desktop Settings, Screen Saver Settings, Appearance Settings, Display Settings, Mouse Settings
- 6. Working with documents: Creating a document, Manage files and folders for documents, working with icons, editing documents Text formatting and alignment, Indentation.
- 7. Paragraph formatting Margins, tabs and page numbering.
- 8. Working with tables and borders Printing Working with Images and Text Find and replace text Mail merge.
- 9. Creating and formatting a presentation –Creation of a new Presentation, Adding Slides and Text to a Presentation, Editing Slide Text, Saving a Presentation, and Running a Slide Show- Adding Tables and charting data – Modifying objects and adding Images, Preparing to deliver a presentation.
- 10. Creating and modifying a worksheet- Formatting Worksheets Working with multiple worksheets Performing Calculations
- 11. Surfing information using Search Engines, Saving web pages to a disk, Composing Email, Sending E-mail.

SEMESTER- I BINF 454 – LAB - PROGRAMMING IN C/ C++

Exercises:

- Operating System: Overview of Linux Architecture, Installation, Booting and Shutdown Process, System Processes(an overview), User Management- Types of users, Creating Users, Granting Rights, File Quota, File-system Management and Layout, Login Process, Linux shells(bash and tcsh), Shell Programming, Printing.
- 2. C Programming: Flowcharts, Algorithm, Keywords, Identifiers, variables, Constants, Scope of Life of variables- Local and Global variables. Data types, Expressions, Operators - Arithmetic operators, Logical operators, Relational, conditional, Bitwise operators - Input / Output Library functions.

Declaration statement, Conditional statement: If statement, If...Else statement, Nesting of If...Else statement, Switch statement – Iteration statements - Arrays: Concept of Single and Multi-dimensional arrays, Array declaration, and initialization of arrays. Functions: User defined and library functions - File Handling: Opening a file, Closing a file, Reading and Writing into a file, Appending to a file

 C++: I/O statements- Escape sequences- Comment lines - Expressions and Statements-Standard libraries - Prototype of main() function - Data types – Conditional Statements -Functions and variables - Classes and objects – Constructors and Destructors – Inheritance.

SEMESTER - II BINF 421 – GENOMICS AND PROTEOMICS

Unit-I

Organization of the prokaryotic and eukaryotic genomes; Genome maps and types; current sequencing technologies; partial sequencing; gene identification; gene prediction rules and softwares; Genome databases; Annotation of genome. Genome diversity: taxonomy and significance of genomes – bacteria, yeast, *Caenorhabditis, Homo sapiens, Arabidopsis*, etc.

Unit-II

Microarray - Gene Expression, methods for gene expression analysis; DNA array for global expression profile; Types of DNA array, Array databases; Applications of DNA microarray – analysis of gene expression, differential gene expression under different conditions and during development of organisms.

Unit-III

Human Genome - Mapping of Human Genome; Construction of physical maps; Basics of radiation hybrid maps; Sequencing of the entire human genome, annotation and analysis of genome sequences: sequence repeats, transposable elements, gene structure, pseudogenes; Gene analysis; gene order; chromosome rearrangement; compositional analysis; clustering of genes; composite genes. Implications of the Human Genome Project; Basics of Single Nucleotide Polymorphisms, detection and its implications.

Unit-IV

The proteome and Proteome technology – Introduction; Expression proteomics (express profile); Cell map proteomics; Protein separation technology - 2D-Gel Electrophoresis, liquid chromotagraphy, affinity chromatography (for cell map proteomics); mass spectroscopy and its uses in protein identification; Forward and Reverse Proteomics

Unit-V

Protein-Protein Interactions – Yeast two hybrid, Co-Precipitation, Phage Display, Phylogenetic Profile, Domain fusion, Gene Neighborhood, Gene Cluster, Mirror Tree, Analysis of genome wide Protein-Protein Interactions in yeast, Genome wide yeast two hybrid analysis of other organisms, Protein fragment complementation assays.

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Recommended Texts:

- 1. Primrose, S.B. and Twyman, R.M. (2003) "Principles of Genome Analysis and Genomics"; Third edition, Blackwell Publishing Company, Oxford, UK.
- 2. Liebler, D.C. (2002) "Introduction to proteomics Tools for the new biology"; First edition, Human Press Inc., New Jersey, USA.
- 3. Pevsner, J. (2003) "Bioinformatics and Functional Genomics"; John Wiley and Sons, New Jersey, USA.

Reference Books

- 1. Mount, D. (2004) "Bioinformatics: Sequence and Genome Analysis"; Cold Spring Harbor Laboratory Press, New York.
- 2. Lesk, A.M. (2002) "Introduction to Bioinformatics", First edition, Oxford University Press, UK.
- 3. Sensen, C.W. (2002) "Essentials of Genomics and Bioinformatics"; Wiley-VCH Publishers, USA

SEMESTER - II BINF 422 – BIOPHYSICAL CHEMISTRY

Unit-I

Classical and quantum mechanics – Elementary introduction to Lagrangian and Hamiltonian formulation of mechanics; breakdown of classical mechanics; Planck theory of blackbody radiation; photoelectric effect; Bohr model of the atom; atomic spectra; De Broglie theory of matter waves; Schrodinger wave equation; interpretation of wave function; atomic orbitals; molecular orbitals; hybrid orbitals; valency of carbon atom; covalent bond; bond order; resonance structure of benzene; partial double bond; character of peptide bond.

Unit-II

Thermodynamics and energetics – Thermodynamics systems; laws of thermodynamics; statement and applications; concepts of entropy and enthalpy; chemical potentials; free energy; Gibb and Helmholtz free energy; ATP as energy currency in biological systems; free energy of hydrolysis of ATP and other organophosphates.

Unit-III

Molecular Mechanics and Dynamics – Basic principles; molecular representations; force fields; atom; atom pair potentials; bond length and bond angle and torsion angle potential; van der Waals and electrostatic potential – hydrogen bonding terms – MM3, AMBER, GROMOS, ECEPP/3 force fields; minimization techniques; line search and elementary introduction to gradient techniques; concepts of molecular dynamics.

Unit-IV

Protein and Nucleic Acid Structure – Levels of protein structure – primary, secondary, tertiary and quaternary with examples; alpha helix, beta sheet and beta turn; domains and structural motifs; Rossmann fold, Immunoglobulin fold; Double helical structure of DNA – DNA polymorphism; RNA secondary and tertiary structure, with viruses – TMV, TBSV and HIV.

Unit-V

X-Ray Crystallagraphy and Spectroscopy – Elementary description of crystallography; crystal growth, data collection, structure solution, refinement and interpretation; concept of resolution, IR spectroscopy, UV-visible spectroscopy; hyperchromism and hypochromism; Raman Spectroscopy; 'finger printing' using Raman spectra; complementarity of Raman and IR spectroscopy Fluorescence spectroscopy; NMR spectroscopy chemical shift; Fourier transform NMR spectroscopy; protein structure determination using NMR.

Recommended Texts:

- 1. Vasantha Pattabhi and N. Gautham. (2001) "Biophysics"; Narosa Publishing Company, New Delhi, India.
- 2. Narayanan, P (1999) "Introductory Biophysics"; New Age Publishing House, Mumbai, India.

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Reference Books

- 1. Cantor, C.R. and Schimmel, P. (1985) "Biophysical Chemistry Vol. 1 and 2"; W.H. Freeman and Company, New York, US
- 2. Freifelder, D. (1982) "Physical Biochemistry"; W.H. Freeman and Company, New York, USA.
- 3. Banwell, C.N. (1983) "Fundamentals of Molecular Spectroscopy"; Tata McGraw Hill Publishing Company, New Delhi, India.
- 4. Leach, A.R. (1992) "Molecular Dynamics Simulation", John Wiley and Sons, New York, USA.

SEMESTER - II BINF 423 – BIOINFORMATICS–II

Unit I **Sequence Analysis** – Methods of sequence alignment: graphic similarity comparison; Dot plots; Hash tables; Scoring matrices - identify matrix, genetic code matrices (GCM); Substitution matrices, Mutation Data Matrices (MDM), Percentage accepted Mutation (PAM). Block Substitution Matrices (BLOSUM), mutation probability matrices; Sequence similarity searches and alignment tools - dynamic programming algorithms; Needlman-Wunch and Smith Waterman; alignment scores and gap penalties; measurement of sequence similarity; percentage of identically aligned residues; Optimal global alignment and optimal local alignment;

Unit II

Pairwise Sequence Alignment - Concept; Programmes (Dot matrix, Dot plot, Dynamic programming); Similarity Searches; Sequence repeats and inversion; Database searching (BLAST and FASTA).

Unit III

7 Lectures Multiple Sequence alignment (MSA) - significance; softwares (PIMA, Clustal, Pileup, ClustalW, Meme, MACAW); Considerations while choosing a MSA software for analysis; sensitivity and specificity of each software.

Unit IV

Comparative Genome Analysis – Relevance of comparative genomics; orthologs and paralogs; Comparative genomics of prokaryotes; Minimal genome; Vertical and horizontal gene transfer. Comparative genomics of organelles; Comparative genomics of eukaryotes. Differences and similarities in genomes of organisms; Evolution of protein families; Applications of comparative genomics in reconstruction of metabolic pathways.

Unit V

Phylogenetic analysis – Phylogenetics, cladistics and ontology; Phylogenetic representations – graphs, trees and cladograms; Classification and ontologies; Steps in phylogenetic analysis; Methods of phylogenetic analysis – similarity and distance tables, distance matrix method; Method of calculation of distance matrix (UPGMA, WPGMA); The Neighbour Joining Method; The Fitch/Margoliash method; Character-based Methods - maximum parsimony, maximum likelihood; Reliability of Phylogenetic trees; Steps in constructing alignments and phylogenies; Limitations of phylogenetic algorithms; Phylogenetic softwares – PAUP, PHYLIP, MacClade.

7 Lectures

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Recommended Texts:

- 1. Mount, D. (2004) "Bioinformatics: Sequence and Genome Analysis"; Cold Spring Harbor Laboratory Press, New York.
- 2. Baxevanis, A.D. and Francis Ouellellette, B.F. (1998) "Bioinformatics a practical guide to the analysis of Genes and Proteins"; John Wiley & Sons, UK.

Reference Books

- 1. Pevzner, P.A. (2004) "Computational Molecular Biology"; Prentice Hall of India Ltd, New Delhi
- 2. Pevsner, J. (2003) "Bioinformatics and Functional Genomics"; John Wiley and Sons, New Jersey, USA.
- 3. Lesk, A.M. (2002) "Introduction to Bioinformatics", First edition, Oxford University Press, UK.
- 4. Sensen, C.W. (2002) "Essentials of Genomics and Bioinformatics"; Wiley-VCH Publishers, USA

SEMESTER - II BINF 424 – STATISTICS FOR BIOLOGISTS

Unit-I

Measures of central tendency : Tabulation of data - Construction of a frequency table -Measures of central tendency - Arithmetic Mean, Median and Mode - Quartiles, Deciles and Percentiles - Geometric mean and Harmonic mean - Weighted arithmetic average - Crude and standardized death rate

Unit-II

Measures of dispersion : Range – Inter quartile range – Mean deviation - Standard deviation and Coefficient of variation – Application of Lorenz curve

Unit-III

Linear correlation and regression

Scatter diagram - Correlation - Types of correlation - The coefficient of correlation - Properties of the coefficient of correlation - Rank correlation - Estimation using time series data - Linear regression - Fitting a straight line – Multiple regression

Unit-IV

Probability and probability distributions : The concept of probability – Sample space – Independent events - Mutually exclusive events - Addition law of probability - Conditional probability - Bayes formula - Expected value - Variance - Binomial distribution - Poisson distribution – Normal distribution – Chi squared distribution – Students t distribution

Unit-V

Theory of Sampling and Theory of Queues : The concept of sampling – Type s of sampling – Techniques of probability sampling - Techniques of non-probability sampling - Sample size -Sampling error

Queues – Types of queues – Markov chains - The concept of dynamic programming – The use of statistics in information systems - The use of spread sheets

Recommended Texts:

- 1. Zar, J.H. (1984) "Bio Statistical Methods"; Prentice Hall International Edition, USA.
- 2. Warren, J., Gregory, E. and Grant, R. (2004) "Statistical methods in Bioinformatics"; First edition, Springer-Verlag, Berlin.

Reference Books

- 1. Milton, J.S. (1992) "Statistical methods in the Biological and Health Sciences"; Second Edition, McGraw Hill Publishers.
- 2. Rosner, B. (2005) "Fundamentals of Biostatistics"; Duxbury Press.
- 3. Bailey, N.T.J. (1995) "Statistical methods in Biology"; Oxford University Press, UK

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SEMESTER - II BINF 425 – PROGRAMMING IN JAVA

Unit I

Java Basics - Importance and features of java, Modifiers, Access Controls, Data types, Expressions, Declarations, Statements & Control Structures, Program Structures, String handling, Packages, Interfaces, Working with java util Package, Garbage Collection, Object Class

Unit II

Exception Handling,I/O & JDBC – Exception Handling: Fundamentals exception types, uncaught exceptions, throw, throw final, built in exception, creating your own exceptions, Input Stream & Output Stream: Streams, Byte and Character stream, Predefined streams, Reading and Writing from Console and Files, Buffered Reader & Writer, Serialization, Data Compression, Using Standard Java Packages (lang, util, io, net) Database: JDBC Architecture, JDBC Basics, JDBC Drivers, Connecting to Database and accessing databases

Unit III

AWT & Event Handling – Creating User interface with AWT, Applets, Applet Life Cycle, Simple Graphics, Fonts and Colors, Events, Listeners, Components, Containers, Working with Layouts, Image Processing, AWT Exceptions, Delegation Event Model, Event Classes, Event Listener Interfaces, Adapter and Inner Classes

Unit IV

Multithreading and Communication – Java Thread Model: Priorities, Synchronization, Messaging, Life Cycle of Thread, Thread class, Runnable interface, Interthread Communication, Suspending, Resuming and Stopping threads, Multithreading, Synchronization, Scheduling and Priority of Threads.

Unit V

BioJava - Installing BioJava, Symbols, Basic Sequence Manipulation (DNA to RNA, Reverse Complement, motif as regular expression), Translation (DNA to Protein, Codon to amino acid, Six frame translation), Proteomics (Calculate the mass and pI of a peptide), Sequence I/O (File Formats conversions), Locations and Features (PointLocation, RangeLocation, Feature modifications), BLAST and FASTA (Blast and FastA Parser, extract information from parsed results), Counts and Distributions, Weight Matrices and Dynamic Programming, User Interfaces.

Reference Books

- 1. Naughton, P. and Schildt, H. (1999) "Java-2: The complete Reference"; Third Edition, McGraw Hill Publishers.
- 2. Horstmann, C.S. (2000) "Computing Concepts with Java 2 Essentials"; Second Edition, John Wiley Publishers.
- 3. Benjamin, Cummings and Booch, G. (1994) "Object Oriented Design and Applications"; Second edition, Addison Wesley Publishers.

6 Lectures

8 Lectures

7 Lectures

7 Lectures

SEMESTER - II BINF 426 – RELATIONAL DATABASE MANAGEMENT SYSTEM

Unit-I

Introduction - Database & Database users, Characteristics of Database, Database System Applications, Database System Versus File Systems, Concepts and Architecture, Data Models, Schemas & Instances, DBMS architecture and Data Independence, Database Languages & Interfaces, View of Data, Data Models, Database Languages, Database Users and Administrators, Database System Structure

Unit-II

Data models - ER Model: Keys, Constraints, Design Issues, Extended ER features, Reductions of ER Schema to Tables. Relational Model: Structure, Relational Algebra; Hierarchical Model, Network Model, Object Oriented Model

Unit-III

Structured Query Language – Basic Structure, Set Operations, Aggregate Functions, Null Values, Nested Sub queries, Views, Integrity: Domain constraints, Joined Relations, Data-Definition Language, Embedded SQL, Dynamic SQL

Unit-IV

Relational Database and Storage – Pitfalls in Relational Design Database, Functional dependencies, Decomposition Normal Forms - 1NF, 2NF, 3NF & Boyce-Codd NF, Overall Database Design Process, Multivalued Dependencies, Data Storage - Ordered indices, Static Hashing, Dynamic Hashing - Transaction Management - Security and Authorization.

Unit-V

Concurrency control techniques & Information retrieval – Locking techniques, Granularity of Data Items - Database System Architecture and Information retrieval: Centralized and Client-Server Architecture, Distributed DBMS, Data Mining, Data Integration, Data Warehousing

Recommended Texts:

- 1. Silberschatz, A., Korth, H.F. and Sudarshan, S. (2002) "Database system Concepts"; Fourth Edition, McGraw Hill Publishers.
- 2. Date, C.J. (2000) "An introduction to Database systems"; Seventh Edition, Addison Wesley Publishers.
- 3. Elmasri and Navathe (2004) "Fundamentals of Database systems"; Fourth Edition, Addison Wesley Publishers.
- 4. Ullman, J. D. (2001) "Principles of Database systems"; Second Edition, Galgotia Publications.

7 Lectures

6 Lectures

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SEMESTER - II BINF 427 – FUNDAMENTALS OF ALGORITHMS

Unit – I

Computing Algorithms

Algorithms in Computing, Analyzing algorithms, Designing algorithms, Asymptotic notation, Standard notations, Big 'O' notations, Time and space complexity of algorithms and common functions

Sets: Union and Intersections, Differences, Disjoint Sets, Counting Elements, Relations Matrices: Adding and Multiplying, Extracting a sub-matrix, Combining, Inverting

Unit – II

Sorting, Searching & Strings Matching

Sorting: Bubble Sort, Insertion sort, Selection sort, Quick Sort, Radix sort, Exchange sort, Shellsort, Mergesort. External sort (K-way mergesort, balanced mergesort, polyphase mergesort) Sorting in Linear time, Heaps (Binary Heaps, Janus Heap, Heap sort, Binomial Heaps, Fibonacci Heaps)

Searching: Binary Search, Fibonacci Search, Hash Search, Lookup Searches, Generative Searches

String Matching: Naïve algorithm, Boyer-Moore algorithm, Knuth-Morris-Pratt algorithm

Unit – III

Graphs

Representation of Graphs, Breadth First Search, Depth First Search, Topological Sort, Connected Components, Minimum Spanning Tree, Single-Source Shortest Path (Dijkstra's and Bellman Fort Algorithm), All-Pairs Shortest Paths (Floyd-Warshall algorithm), Coloring of Graphs (Kruskal's Algorithm, Prim's Algorithm),

Unit – IV

Trees

Forests, DAGs, Ancestors, and Descendants, Binary Search Trees, Querying a Binary search tree, Insertion and Deletion, Tree Traversals, Red-Black Trees, Properties of Red-Black Trees, AVL-Trees, Rotations, Insertion, Deletion, B+ Tree, B* Trees.

Unit – V

Algorithm Design and Analysis

The substitution method, The iteration method, The master method, Divide and Conquer, Greedy Algorithms, Dynamic Programming (Traveling Sales Person Problem, Hamiltonian Path Problem), Backtracking Algorithms (8-queens Problem, Graph Coloring), Branch and Bound Algorithms

Recommended Texts

- 1. E. Horowitz and S. Sahani, "Fundamentals of Data structures", Galgotia Booksource Pvt. Ltd., (1999)
- 2. Ellis Horwitz, Sartaz Sahani and Sanguthevar Rajasekaran,(1999),"Computer Algorithms", Galgotia Publications
- 3. T.H. Cormen, C. E. Leiserson, R.L. Rivest (2001) "Introduction to Algorithms", 3rd Ed PHI

7 Lectures

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SEMESTER - II BINF 455 – LAB - PROGRAMMING IN JAVA

Exercise in JAVA

- 1. Concepts of Flowcharting, Algorithm Development.
- 2. Writing Pseudo Codes.
- 3. Working with Objects, Arrays, Conditionals and Loops.
- 4. Creating Classes and Applications in Java.
- 5. Java Applets Basics, Graphics, Fonts and Color.
- 6. Simple Animation and Threads, Advanced Animation, Images and Sound.
- 7. Managing Simple Events and Interactivity.
- 8. Creating User Interfaces with AWT, Modifiers.
- 9. Packages and Interfaces, Exception, Multithreading.
- 10. Streams and I/O, Using Native Methods and Libraries.
- 11. Java Programming Tools, Working with Data Structures.

SEMESTER - II BINF 456 – LAB - PROGRAMMING IN RDBMS

Exercise in RDBMS (MYSQL)

Data Definition Language (DDL) statements:

Creating database, Selecting database, Deleting database, Creating table, Modifying Table, Deleting table

Data Manipulation statements:

Inserting, updating and deleting records Retrieving Records Retrieving specific rows and columns Use of MySQL operators – Arithmetic operators, Comparison Operators, Logical operators Math functions, Aggregate functions String operations Limiting, Sorting and grouping query results Handling null values Renaming or aliasing table and column names Using subqueries Using Joins – joining a table to itself, joining multiple tables Use of Indexes Security Management Granting and Revoking rights on tables

SEMESTER - II BINF 457 – LAB - BIOSEQUENCE ANALYSIS

Exercises:

- 1. Sequence Analysis Packages EMBOSS, NCBI ToolKit
- 2. Dynamic programming.
- 3. Analysis of Biological Sequences.
 - a. Basic Blast
 - b. Specialized Blast
- 4. FASTA
- 5. Multiple sequence alignment
- 6. MEME/MAST, eMotif, InterproScan, ProSite, ProDom, Pfam
- 7. Phylogenetic analysis PAUP, PHYLIP, MacClade
- 8. Genome annotation Artemis.
- 9. Hypothetical Protein analysis
- 10. Genome Comparison

Anatomy of proteins; Ramachandran Plot; secondary structures; motifs; domains; tertiary and quaternary structures. Anatomy of DNA; A, B, Z-DNA, DNA bending. Structure of RNA.

7 Lectures

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Methods for prediction of secondary and tertiary structures of proteins – knowledge-based structure prediction; fold recognition; *ab initio* methods for structure prediction, Comparative protein modeling.

Unit-IV

Unit-III

Methods for comparison of 3D structures of proteins; Methods to predict three dimensional structures of nucleic acids, rRNA; Electrostatic energy surface generation.

Unit-V

Molecular Mechanics and Molecular dynamics of Oligopeptides, Proteins, Nucleotides and small molecules – Mechanism and dynamics of bio-macromolecules, Simulation of molecular mechanics and dynamics, Simulations of Free energy changes; Force fields. Molecular interactions of protein-protein, protein-DNA, protein-carbohydrate and DNA-small molecules.

Recommended Texts:

- 1. Andrew R. Leach (2001) "Molecular Modeling Principles and Applications"; Second Edition, Prentice Hall, USA
- 2. Creighton, T.E. (1993) "Proteins: structure and molecular properties"; Second edition, W.H. Freeman and Company, New York, USA.

Reference Books

- 1. Mount, D. (2004) "Bioinformatics: Sequence and Genome Analysis"; Cold Spring Harbor Laboratory Press, New York.
- 2. Lesk, A.M. (2001) "Introduction to Protein Architecture", Oxford University Press, UK.
- 3. Mcpherson, A. (2003) "Introduction of Molecular Crystallography", John Wiley Publications, USA.

7 Lectures

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SEMESTER - III BINF 501 – STRUCTURAL BIOLOGY

Unit-I

Unit-II

Structure of Ribosome.

Structural features of biomolecules; techniques used to determine the structure of biomolecules; Methods for single crystal X-ray Diffraction of macromolecules: molecular replacement method and direct method – Fiber diffraction; analysis of structures and correctness of structures; submission of data to PDB: atomic coordinates and electron density maps.

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SEMESTER - III BINF 502 – MOLECULAR MODELING AND DRUG DESIGNING

Unit-I

Concepts in Molecular Modeling – Introduction; Coordinate System; potential energy surfaces molecular graphics; Computer hardware and software; Mathematical concepts – introduction of molecular mechanics & quantum mechanics.

Unit-II

Molecular Mechanics – Features of molecular mechanics, force fields; Bond structure and bending angles – electrostatic, van der Waals and non-bonded interactions, hydrogen bonding in molecular mechanics; Derivatives of molecular mechanics energy function; Calculating thermodynamic properties using force field; Transferability of force field parameters, treatment of delocaliised pi system; Force field for metals and inorganic systems – Application of energy minimization.

Unit-III

Molecular Dynamics Simulation Methods – Molecular Dynamics using simple models; Molecular Dynamics with continuous potentials and at constant temperature and pressure; Timedependent properties; Solvent effects in Molecular Dynamics; Conformational changes from Molecular Dynamics simulation.

Unit-IV

Molecular Modeling in Drug Discovery – Deriving and using 3D pharmacophore; Molecular Docking; Structure-based methods to identify lead compounds; *de novo* ligand design; Applications of 3D Database Searching and Docking

Unit-V

Structure Activity Relationship - QSARs and QSPRs, QSAR Methodology, Various Descriptors used in QSARs: Electronic; Topology; Quantum Chemical based Descriptors. Use of Genetic Algorithms, Neural Networks and Principle Components Analysis in the QSAR equations.

Recommended Texts:

1. Andrew R. Leach (2001) "Molecular Modeling – Principles and Applications"; Second Edition, Prentice Hall, USA

Reference Books

- 1. Fenniri, H. (2000) "Combinatorial Chemistry A practical approach", Oxford University Press, UK.
- 2. Lednicer, D. (1998) "Strategies for Organic Drug Discovery Synthesis and Design"; Wiley International Publishers.
- 3. Gordon, E.M. and Kerwin, J.F. (1998) "Combinatorial chemistry and molecular diversity in drug discovery"; Wiley-Liss Publishers.
- 4. Swatz, M.E. (2000) "Analytical techniques in Combinatorial Chemistry"; Marcel Dekker Publishers.

8 Lectures

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SEMESTER - III BINF 503 – PROGRAMMING IN PERL

Unit-I Data Structures and modular programming – Basic Operators and Control Flow, Basic Perl Data Types, References, Matrices, Complex/Nested Data Structures, Scope (my, local, our), Function/Subroutines, System and User Function, The local Operator, Variable-length Parameter Lists, Notes on Lexical Variable, File handle and File Tests, stat and lstat Functions, Formats, Directory Access & Manipulation, Process Management, Formatting Data, System Information, Perl Modules, Name Spaces, Packages, Defining Modules, Storing Modules, Using Modules, **CPAN Modules**

Unit-II

Regular expressions and Pattern Matching – Uses of Regular Expressions, Patterns, Single-Character Patterns, Grouping Patterns (Sequence, Multipliers, Parentheses as memory, Alternation) Anchoring Patterns, Precedence, Matching Operators, Ignoring Case, Different Delimiter, Variable Interpolation, Special Read-Only Variables, Substitutions, Split and Join Functions, Dynamic Programming, Approximate String Matching

Unit-III

Common Gateway Interface - Gene.pm, Arrow notation (->), AUTOLOAD, Cleaning unused objects with DESTROY URL Encoding, CGI Environment Variables, Handling forms, Accessing form Input, Extra Path Information, CGI.pm Module, Passing Parameters via CGI, Less Typing, Sever Side Includes, Debugging CGI programs, Stepping through programs, Breakpoints, Line Action

Unit-IV

Relational Databases – DBM Databases and DBM Hashes, Design of DBI, DBI Methods, DBI Environment Variables, DBD Interface Modules, Fixed Length Random-Access Databases, Variable-Length Databases, Win32 Database Interface, Perl Graphics, Using the GD.pm graphics, Animation

Unit-V

BioPerl – Installing Bioperl, General Bioperl Classes, Sequences (Bio::Seq Class, Sequence Manipulation), Features and Location Classes (Extracting CDS), Alignments (AlignIO), Analysis (Blast, Genscan), Databases (Database Classes, Accessing a local database), Implementing REBASE

Reference Books

- 1. Dranell, R. (1994) "HTML4 unleashed"; Second Edition, Techmedia Publications.
- 2. Wall, W., Christiansen, T. and Orwant, J. (2000) "Programming Perl", Third Edition, O'Reilly Publishers.
- 3. Tisdall, J. (2004) "Beginning Perl for Bioinformatics", First Edition, O'Reilly Publishers.

8 Lectures

7 Lectures

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SEMESTER - III BINF 504 – APPLICATIONS OF BIOINFORMATICS

Unit-I

Profile analysis – Expression profile analysis of cells, Mining data from Yeast. Microarray and genome wide expression analysis: transcriptomes, proteome: Genomics in medicine, disease monitoring, profile for therapeutic molecular targeting.

Unit-II

Reconstruction of pathways and annotation – Reconstructing metabolic pathways from sequence and function information in microbial species; statistical profiling and function annotation of genomes with a microbial genome as an example.

Unit III

Drug Designing Related Applications – Finding new drug targets to treat diseases – Pharmacophore identification - Structure based drug design – Molecular Simulations.

Unit-IV

Systems Biology - Objectives of Systems Biology, Strategies relating to *In silico* Modeling of biological processes, Metabolic Networks, Signal Transduction Pathways, Gene Expression Patterns. E-cell and V-cell Simulations and Applications.

Unit-V

Commercial Bioinformatics - Definition of Bioinformatics company. Genome Technology: high throughput sequencing and assembly. Diagnostic drug discovery and genomics. Pharmacogenomics and its application. SNPs and their applications. Proteomics in medicine, Toxicology.

Recommended Texts:

- 1. Hunt, S.P. and Livesey, F.J. (2000) "Functional Genomics a practical approach", Oxford University Press, UK.
- 2. Wilkins, M.R., Wiliams, K.L., Appel, R.D. and Hochstrasser, D.F. (1997) "Proteome Research: New frontiers in Functional Genomics", Springer Verlag, New York, USA.
- 3. Witten, I.H. and Frank, E. (2005) "Data mining: Practical Machine Learning Tools and Techniques", Morgan Kauffman Publishers, USA.

7 Lectures

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SEMESTER - III BINF 505 – NETWORKS AND DISTRIBUTED COMPUTING

Unit – I

Nuts & Bolts in Networks

Reference Model, Network Topologies and Protocols, Types of Networks: Local Area Network (LAN), Wide Area Network (WAN), Metropolitan Area Network (MAN), Network Security (Firewall, Packet Filtering, VPN), Uses of Computer Networks

Unit – II

Network Architecture

OSI & Internet Architecture, IEEE 802 standards, Physical Layer - Transmission Media, Switching. Data Link Layer - Design Issues, Example Data Link Protocols, Data Link layer in the Internet, Media Access Sub layer

Unit – III

Network Layer and Transport Layer

Network Layer - Design Issues, Routing Algorithms, Congestion control algorithm, Router Operation, Router Configuration, Internetworking, IP Addressing, IP Subnet Mask, IPv6 (an overview) Transport Layer – Transport Service, TCP/IP Protocols (TCP, UDP)

Unit – IV

Application Layer

Design Issues, Conventional Encryption, Classical and Modern Techniques, Encryption and Decryption Algorithms (RSA), Confidentiality, DNS, SNMP, RMON, WWW, E-mail, Digital Signatures

Unit – V

Characteristics of Distributed Computing

Introduction to Distributed Computing, Examples, Key Characteristics, Historical background, Basic design issues, User requirements

Introduction to IPC, Building Blocks, Client Server Communication, Group Communication, Remote Procedure Call (RPC).

Recommended Texts:

- 1. Tananbaum A.S.,(1999) "Computer Networks", 3rd Ed, PHI
- 2. Black U.,(1996) "Computer Networks-Protocols, Standards and Interfaces", PHI, 1996
- 3. George Coulouris, Jean Dollimore, Tim Kindsberg,(2000) "Distributed Systems : Concepts & Design" 3rd Ed, Addison Wesley

6 Lectures

8 Lectures

7 Lectures

8 Lectures

SEMESTER - III BINF 506 – SCIENTIFIC PRESENTATION

This course is designed to introduce the bioinformatics students to standard scientific presentation formats and to provide a forum to practice/improve oral and written communication skills.

- 1. Readings, writing exercises and in-class discussion Students (in small groups) will lead in-class discussions on assigned readings and writing exercises.
- 2. **Individual oral presentation** Students select a research project/topic and present results as an oral presentation followed by a Q&A session.
- 3. **Individual Poster presentation** Students select a research project/topic and present results as a poster followed by a Q&A session.

Recommended Texts:

1. Mathews. Successful Scientific Writing: A step-by-step Guide for Biomedical Scientists. 2nd ed. Cambridge University Press, 2001.

SEMESTER - III BINF 551 – LAB - STRUCTURAL BIOLOGY

Exercises

- 1. Advanced Visualization Software and 3D representations.
- 2. Coordinate generations and inter-conversions.
- 3. Secondary Structure Prediction
- 4. Fold Recognition, *ab initio* (Rosetta Server)
- 5. Homology based comparative protein modeling.
- 6. Energy minimizations.
- 7. Validation of models.
 - a. WHATIF
 - b. PROSA
 - c. PROCHECK
 - d. VERIFY 3D
- 8. Protein Structure Alignment.
- 9. Modeller
- 10. Geno-3D
- 11. Discovery Studio Server.

SEMESTER - III BINF 552 – LAB - MOLECULAR MODELING AND DRUG DESIGN

Exercise:

- 1. Binding Site Identification.
- 2. Pharmacophore Identification
- 3. Rigid body Docking using AutoDock and ADT
- 4. Molecular dynamics simulations using Gromacs
- 5. Visual molecular Dynamics (VMD)
- 6. Docking with LigandFit (Discovery studio)
- 7. Receptor and Ligand Optimization.
- 8. Conformational Analysis
- 9. BABEL, MOPAC

SEMESTER - III BINF 553 – LAB - PROGRAMMING IN PERL

Exercise in Structured Programming

Basic Operators and Control Flow, Basic Perl Data Types, References, Matrices, Complex/Nested Data Structures, Scope (my, local, our), Function/Subroutines, System and User Function, The local Operator, Variable-length Parameter Lists, Notes on Lexical Variable, File handle and File Tests, stat and lstat Functions, Formats, Directory Access & Manipulation, Process Management, Formatting Data, System Information

Exercise in Regular Expressions

Uses of Regular Expressions, Patterns, Single-Character Patterns, Grouping Patterns (Sequence, Multipliers, Parentheses as memory, Alternation) Anchoring Patterns, Precedence, Matching Operators, Ignoring Case, Different Delimiter, Variable Interpolation, Special Read-Only Variables, Substitutions, Split and Join Functions, Dynamic Programming, Approximate String Matching

Exercise in CGI

URL Encoding, CGI Environment Variables, Handling forms, Accessing form Input, Extra Path Information, CGI.pm Module, Passing Parameters via CGI, Less Typing, Sever Side Includes, Debugging CGI programs, Stepping through programs, Breakpoints, Line Action

Exercise in CPAN Database Modules

DBM Databases and DBM Hashes, Design of DBI, DBI Methods, DBI Environment Variables, DBD Interface Modules, Fixed Length Random-Access Databases, Variable-Length Databases, Win32 Database Interface, Perl Graphics, Using the GD.pm graphics library

Exercise in Bioperl

Installing Bioperl, General Bioperl Classes, Sequences (Bio::Seq Class, Sequence Manipulation), Features and Location Classes (Extracting CDS), Alignments (AlignIO), Analysis (Blast, Genscan), Databases (Database Classes, Accessing a local database), Implementing REBASE

SEMESTER - III BINF 520 – BIOETHICS AND INTELLECTUAL PROPERTY RIGHTS

Unit-I

The legal and socioeconomic impacts of biotechnology; public education of the process of the processes of biotechnology involved in generating new forms of life for informed decision making

Unit-II

Biosafety regulation and national and international guidelines; rDNA guidelines; Experimental protocol approval; levels of containment

Unit-III

Environmental aspects of biotechnology applications; Use of genetically modified organisms and their release in environment; Special procedures for rDNA-based product production.

Unit-IV

General principles of Intellectual property rights (IPR); Patents and methods; application of patents; Legal implications; International treaties for protection of IP – Bern, Paris, TRIPS, WIPO treaties, Biodiversity convention, etc.

Unit-V

Biodiversity and farmers rights; Beneficial applications and development of research focus to the need of the poor; Identification of directions for yield effect in agriculture, aquaculture, etc; Bioremediation

Recommended Texts:

- 1. Sasson, A. (1988) "Biotechnologies and Development", UNESCO Publications.
- 2. Sasson, A. (1993) "Biotechnologies in developing countries present and future"; UNESCO Publishers.
- 3. Singh, K. "Intellectual Property Rights on Biotechnology"; BCIL, New Delhi.

9 Lectures

7 Lectures

6 Lectures

5 Lectures

SEMESTER - IV BINF 521 – FINISHING SCHOOL

Unit-I Team Work – Interpersonal skills, Behavioural attitude, People management – skills, Personality development, Clean and healthy living tips.	5 Lectures Intrapersonal
Unit-II Organizational Behaviour – Goal setting, Individual goal, Organizational goal.	4 Lectures
Unit-III Time Management – Planning, Scheduling	4 Lectures
Unit-IV Ethics, Values, Attitudes.	4 Lectures
Unit-V Indian Culture and Heritage	4 Lectures

SEMESTER - IV BINF 522 – PROJECT

The course is designed to result in the satisfactory completion and defense of the Masters dissertation.

This process includes

- a) the conceptualization of the independent research that will comprise the dissertation,
- b) the preparation of and satisfactory defense of the dissertation proposal,
- c) the collection, analysis, and interpretation of data,
- d) presentation of findings in the dissertation format, and
- e) oral defense of the dissertation.

Dissertation activity must be completed within prescribed time frame for the semester.

PONDICHERRY UNIVERSITY SCHOOL OF LIFE SCIENCES

Centre for Bioinformatics

SYLLABUS FOR M. Sc. BIOINFORMATICS

(Academic Year 2007-2008 onwards)

Semester	Course Code	Course Title	Credits
	BINF 401	Cell and Molecular Biology	3
	BINF 402	Physics and Chemistry for Biologists	3
Ι	BINF 403	Mathematics for Biologists	3
	BINF 404	Bioinformatics–I	3
	BINF 405	Data Structures and Programming Concepts	3
	BINF 406		3
	BINF 407	Computers for Biologists (Elective)	3
	BINF 451	Lab - Cell and Molecular Biology	1
	BINF 452	Lab - Biological Databases	1
	BINF 453	Lab - Basics of Computer & Operating Systems	1
	BINF 454	Lab - Programming in C/ C++	1
	BINF 408	Genomics and Proteomics	3
	BINF 409	Biophysical Chemistry	3
	BINF 410	Bioinformatics-II	3
	BINF 411	Statistics for Biologists	3
II	BINF 412	Programming in Java	3
	BINF 413	Relational Database Management System	3
	BINF 414	Fundamentals of Algorithms	3
	BINF 455		1
	BINF 456		1
	BINF 457	Lab - Biosequence Analysis	1
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	BINF 501	Structural Biology	3
	BINF 502	Molecular Modeling and Drug Design	3
	BINF 503	Programming in Perl	3
	BINF 504	Applications of Bioinformatics	3
III	BINF 505	Networks and Distributed Computing	2
	BINF 506	Scientific Presentation	2
	BINF 551	Lab – Structural Biology	1
	BINF 552	Lab – Molecular Modeling and Drug Design	1
	BINF 553	Lab - Programming in Perl	1
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	BINF 507	Bioethics and Intellectual Property Rights	3
IV	BINF 508	Finishing School	2
	BINF 596	Project	4

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