SARDAR PATEL UNIVERSITY T.Y.B.Sc (BIONFORMATICS)

BNF-301: Immunology

Effective from June- 2008

Three periods per week

External Marks: 80 Internal Marks: 40 Total Marks: 120

Unit –I:

Introduction to immunoloty, Types of immunity- Innate & Acquired. Humoral & Cell mediated, Active & Passive immunity, Antigens- Antigenic determinants, haptens components of Innate & Acquired immunity, Cells of the immune system, primary & secondary lymphoid organs (Bone marrow, Thymus, Lmyph nodes, spleen. MALT & GALT)

Unit - II:

Antibiotics – Structure, types & function. Monoclonal antibodies- Hybridoma technology (HAT selction) Application of monoclonal antibodies, B-Lymphocytes, BCR, B-cell development (Maturation, activation & differentiation), T-lymphocytes, TCR, T-cell development (Maturation, activation & differentiation)

Unit III:

Antigen processin & presentation- Endogenous and exogenous antigens. MHC-Structures, types & functions, Rearrangement & Expression of immunoglobulin genes, Generation of Antibody diversity.

Unit IV:

Antigen- Antibody interactions- Agglutination. Precipitation, Immunodiffusionsigle & double, Immuno- Electrophoresis, Complement Fixation Test, ELISAprinciple, variants (indirect, sandwich & competitive) and applications, RIA.

Unit V:

• Compliment system- Introductions & functions, Classical, Alternative & lectin pathway, Cytokines- introduction & types, Role of cytokines in immune response, Transplantation Localised & systemic. Proposed mechanism of generation fo auto immunity.

Unit VI:

Immunodeficiency- Primary & Secondary. (AIDS and SCID). Hypersensitivity-Indroduction & Types. Apoptosis- Definition, mechanism & significance cell signaling- overview & significance. Role of Bioinformatics in immunology & vaccine development.

Reference Books:

- 1. Immunology by Janis Kuby
- 2. Immunology by Ivan Roits.
- 3. Text Book of Immunology by C.V Rao

SARDAR PATEL UNIVERSITY T.Y.B.Sc (BIONFORMATICS) BNF-302: Genetic Engineering

Effective from June- 2008

Three periods per week

External Marks: 80 Internal Marks: 40 Total Marks: 120

Unit – 1:-

- R-DNA technology, Restriction enzymes, DNA-RNA modifying enzymes
- Cloning and expression vectors- plasmids (pBR322, pUC), cosmids,
- Phagemids(replacement and insertional vector),
- Eukaryotic expression vector(SV-40,retroviral vector)
- yeast plasmid based vectors (YAC,YEP,YIP) shuttle vectors and Ti- plasmid based vector. (Co-integrate and Binary reactor)

Unit –2

- Transformation and Transfection various methods of Recombinant selection & screening,
- construction of C- DNA & genomic DNA Library.
- In situ hybridization, methods and application
- Reporter genes-scorable reporter genes (LAC,LUX,GUS) Selectable reporter genes (CAT,NPT,Kan^r).

Unit - 3

- Electrophoresis Agarose and Polyacrylamide,
- DNA foot printing, Gel filtration, assay
- HPLC, Ion exchange and affinity chromatography.
- PCR technology Introduction and overview of classical and RT PCR,
- Application of Sequencing technologies Automated and chemical methods of DNA and Protein sequencing .

Unit -4

- Hybridization technique -
- Southern, Northern & Western Blotting techniques,
- dot-Blot hybridization ,Nucleic acid probes and its applications, Radioactive and non radioactive probes and their application
- Autoradiography principle and application

Unit – **5**

- Molecular Mapping RFLP , RAPD , AFLP , STS , EST, repetitive DNA Micro and mini satellite ,
- SNPs and their application.
- Genomics Organization of human genome,
- DNA finger printing and its application

Unit – 6

- Gene transfer to animals and plants, Electoporation , Micro injection, Particle bombardment , Ca++ Phosphate Mediated, –Retroviral mediated and liposomes
- Application of transgenics, animals & plants
- Stem cell -Potential and applications.
- Gene Therapy-types, methods and application
- Site directed mutagenesis methods and application

Reference Book

- 1. Principle of Genetic Manipulation by Old and Primrose
- 2. Genomes by T. A. Brown
- 3. Gene Cloning T.A.Brown
- 4. An introduction to Genetic Analysis Griffith Miller, Suzuki Levonting

SARDAR PATEL UNIVERSITY T.Y.B.Sc (BIONFORMATICS)

BNF – 303 : Bioinformatics Applications Effective from June- 2008

- **Unit 1:- Genomics:** Genome Anatomy- Prokaryotic genomes structures, Eukaryotic genomes structures , Human Genome Project overview & applications, Genome sequencing, Comparative genomics.
- **Unit 2:- Gene Prediction:** Gene structure in Prokaryotes and Eukaryotes, Gene prediction methods: Neural Networks, Pattern Discrimination methods, Signal sites Predictions (Promoter, Splice, UTR, CpG-islands), Evaluation of Gene Prediction methods.
- **Unit 3 :- Multiple Sequence Alignment :** Significance of MSA, Various approaches for MSA (Progressive & Iterative), Profile analysis, Block analysis, Pattern searching, Motif analysis. Sequence logo, Statistical methods for aiding alignment Markov Models, PSI/PHI-BLAST
- **Unit 4:- Microarray Informatics:** Microarray Introduction, Microarray Fabrication, Microarray hybridization and detection, Microarray Image Processing and analysis, Expression ratios, Transformations of the Expression ratio, Data Normalization, Distance Measures: Euclidean distance, Pearson Correlation Coefficient. Clustering Methods: Hierarchical (Single linkage, Complete linkage, Average linkage, Centroid Linkage), Non-Hierarchical (K-means, SOMs) Clustering, Applications of Microarray.
- **Unit 5 :- Phylogenetic & molecular evolution :** molecular basis of evolution, Phylogenetic trees & different methods for phylogenetic inference, Construction-Maximum Parsimony Method, Maximum likelihood method and Distance Methods & significance of phylogenetic tree
- Unit 6:- Drug design & discovery: Contour of Drug, steps in development of new drugs, chemical & physiochemical parameters in drug-designing, drug metabolism, Interaction energy calculations, structure based drug designing, Combinatorial chemistry, virtual & high throughput screening, Docking.

Reference Books-

- 1. Genomes T.A. Brown
- 2. Bioinformatics: Methods & Protocols Misener & Krawetz
- 3. Genomics and its applications- David Mount
- 4. Setubal.J., Meidanis J.-Introduction to computational molecular biology.
- 5. Essentials of Genomics and Bioinformatics C. W. Sensen (Editor), 2002, John Wiley & Sons
- 6. Bioinformatics: From Genomes to Drugs, Thomas Lengauer (Editor), 2001, John Wiley & Sons

T.Y.B.Sc. (Bioinformatics)

BNF-304: WebDesigning and PERL Effective from June – 2008

Three periods per week

External Marks: 80 Internal Marks: 40 Total Marks: 120

Unit 1:- Introduction to HTML

- Introduction to Markup Language
- Creating Simple Webpage
- ➤ Linking to other Web Page
- > Text Allignment and Lists
- Text Formatting and Font Control
- Images
- > Introduction to tables
- Introduction to Forms
- > Introduction to Frames

Unit 2:- Introduction to XML

- > XML : Difference between HTML & XML
- Creating your own XML Files
- > XSL

Unit 3:- Perl

- Getting started with Scripting Language
- Sequences and String

Unit 4:-

- Motifs and Loops
- > Subroutine and bugs

Unit 5:-

> Mutations and Randomization

Unit 6:-

> The Genetic Code

Reference Books

- 1. Beginning Perl for Bioinformatics : James Tisdall (O' Reilly)
- 2. Teach Yourself XML: Sandra E. Eddy & John E. Schnyder
- 3. Sams Teach Yourself HTML in 24 hours : Dick Oliver (Techmedia)

T.Y.B.Sc. (Bioinformatics)

BNF-305: Object Oriented Programming and Data Structure Effective from June – 2008

Three periods per week: External Marks: 80

Internal Marks: 40 Total Marks: 120

Unit-1: Concept of Procedural, structured and object oriented programming

- Concept of Encapsulation, Data hiding, Inheritance and Polymorphism
- History of Object oriented Programming techniques
- Classes and Objects
- Advantages of object oriented programming over procedural languages
- **Unit- 2 :-** Data types, Variables and Constants, Expression and statements, logical, relational, mathematical operators, turnery operator
 - Simple I/O statements reading and writing, Statements for formatted I/O usage of header files using INCLUDE statement
 - Looping : while, Do...While, For loop, Continue and break statement, Switch statement, IF statement, IF..ELSE statement
 - Array: Initializing one dimensional and two dimensional, Multi dimensional array, Passing arrays to functions.
- Unit- 3:- Structures and Enumerated Data types: Declaration of structure, initialization of structure, Array of structure and pointers to structure, structure within structures, Enumerated Data types
 - Classes: Creating new type of class, Classes and members, Accessing class members, Implementing class methods, Constructors and Destructors, Private and public class.

Unit- 4:- Function Definition

- Declaring and defining function, passing of parameters, passing structure variables as argument, passing address (passing by reference), Function with arguments, Overloading function, Inline Function, Storage classes and Static Storage Classes
- Pointer
- Concept of a pointer variable and its declaration, Pointer arithmetic, Pointers in string handling, Pointer to pointer, Arrays of pointers, Pointers to array, Dynamic memory allocations, Pointers to objects

Unit- 5 :- Array & Stack

- Storage structure for arrays
- Definition of stacks, operations on stacks (PUSH, POP, PEEP, CHANGE)
- Application of Stack
- Recursion(Factorial)
- Polish Notation

- Conversion of infix expression to polish notation
- Conversion of polish notation to code

Unit-6:- Queues

- Operation on queue (insert, delete)
- Circular queue (insert, delete)
- Linked List
 - Linked list (insert, delete & storage representation)
 - Single Linked List
 - Circular Linked List
- Trees: Introduction to Trees

Text Books:

- 1. Balaguruswamy: Programming in Ansi C, Tata McGraw Hill Publication
- 2. Trembley & Sorenson: An Introduction to Data Structure with Application, Tata McGraw Hill Publication

Reference Books:

- 1. Henry Mullish & Herbert L. Cooper: The Spirit of C, Jaico Publication
- 2. Bhagat Singh & Thomas L. Naps: Introduction to Data Structure, Tata McGraw Hill Publication

Main Reference Books:

1. Rajaraman : Object Oriented Programming & C++, New age international publishers.

SARDAR PATEL UNIVERSITY T.Y.B.Sc (BIONFORMATICS)

BNF – 306: Structural Bioinformatics & Bio-database and RDBMS

Effective from June- 2008

Three periods per week External Marks: 80

Internal Marks: 40 Total Marks: 120

Unit 1:- Macromolecular Structures : DNA: Types of Double helices (A,B,Z). Structural and Geometrical parameters of each and their comparison, RNA: Secondary structures, Tertiary structures, t-RNA tertiary structure. RNA secondary structure prediction methods and its limitations (mfold method of Zuker), Applications of RNA structure modeling

Unit 2 :- Anatomy of proteins : Primary Structure, Secondary Structure, Super Secondary Structures, Tertiary & Quarternary structure of proteins. Protein – protein interaction. Principle of Protein folding & Unfolding: Levinthal Paradox, Protein folding models, Protein Unfolding, Disorders/Diseases of Protein Misfolding (Alzheimer's disease, Cystic fibrosis, Mad Cow disease, Prions)

Unit 3:- Protein Structure Prediction & Comparison: Methods for protein secondary and tertiary structure prediction-Algorithms of Chou Fasman, GOR, PHD, PSI-PRED and Neural network method; Steps involved in Homology Modeling, Ramachandran Plot, Transmembrane prediction methods. Concepts in 3D structure comparison, Algorithms for structure comparison (VAST & DALI).

Unit 4:- SQL and Constraints: Fundamentals of RDBMS Fundamentals of RDBMS:

- ➤ Normalization, CODD's Principles
- ➤ Introduction to SQL syntax
- Components of SQL
- > Oracle Data Types: Char, Varchar2, Varchar, number, date, raw, long raw,
- Command: Create, insert, update, delete, select, alter, drop
- ➤ Use of Distinct and Order By clause
- ➤ Select *from Tab, Use of Desc keyword
- > Create the table from another table
- > Inserting data into table from another table
- ➤ Data constraints : column level, table level, Null value, primary key, unique key, foreign key, Default
- > Check Integrity Constraints
- > Drop constraints
- ➤ Logical Operators : AND, OR, NOT
- Range Searching: Between, Not Between
- > Pattern Matching : Like, In

Unit 5:- Oracle Functions, Joins and Sub queries:

- Group / Aggregate functions : AVG, MIN, MAX, COUNT, SUM
- Numeric functions: ABS, POWER, ROUND, TRUNC, SQRT, GREATEST, LEAST, MOD, FLOOR, CEIL, EXP
- ➤ Character Functions: UPPER, LOWER, INITCAP, LENGTH, SUBSTR, LPAD, RPAD, LTRIM, RTRIM, TRIM, INSTR
- Date functions: ADD_MONTHS, LAST_DAY, MONTHS_BETWEEN, NEXT_DAY
- Special Date Formats using To_Char
- ➤ Conversion Functions : TO_NUMBER, TO_CHAR, TO_DATE,
- > GROUP BY, HAVING clauses
- > Sub-Queries, nested sub-queries
- > Joins of two or more tables, self join, outer join, inner join
- Subqueries: Union, Intersect, Minus Clause Oracle transactions: Commit, Rollback, SavePoint

Unit 6 :- Oracle Transaction and PL/SQL

- ➤ Indexes creating, dropping, uses
- > Sequences creating, dropping, altering, using (CURRVAL, NEXTVAL)
- ➤ Views creating, dropping, using updateable and non-updateable views
- Granting and Revoking Permissions
- ➤ Introduction to PLSOL
- > Control Structures: If, Loop, While, For
- > Stored procedure and Stored Functions
- ➤ Data warehousing, Data Mining, DDBMS (Only Definitions)

Reference Books:

- 1. AN INTRODUCTION TO COMPUTATIONAL BIOCHEMISTRY by C. Stan Tsai, A JOHN WILEY & SONS, INC., PUBLICATION
- 2. Introduction to Bioinformatics by Arthur M. Lesk
- 3. SQL, PL/SQL The Programming Language of ORACLE, By Ivan Bayross, BPB Publications, 3rd Revised Edition

Sardar Patel University T.Y.B.Sc. (Bioinformatics)

BNF-307 : Practicals based on BNF – 301 and BNF-302 Effective from June – 2008

Three periods per week: External Marks: 80

Internal Marks: 40 Total Marks: 120

- 1. Isolation of genomic DNA from plant cells.
- 2. Isolation of RNA from mammalian cell.
- 3. Isolation of Plasmid DNA.
- 4. Preparation of competent cell. Restriction digestion, ligation and Transformation
- 5. Gel electropheresis- Agarose (for separation of DNA & RNA)
- 6. SDS PAGE. (Demo)
- 7. Radial Immunodiffusion,
- 8. ELISA.
- 9. PCR amplification of target DNA.(Demo)
- 10. Study of chromosomal Abberrations using colchicine.
- 11. Study and identification of amino acid by paper chromatography.
- 12. Estimation of DNA by DPA method.
- 13. Estimation of protein by Folin Lowry method.
- 14. Bacterial cell culture, media preparations and inoculation(spread plate, streak plate and pore plate).
- 15. Estimation of antibiotic activity by Disc method.
- 16. Replica plating.
- 17. Enzyme assay of pH, temperature and incubation time.

T.Y.B.Sc. (Bioinformatics)

BNF-308 : Practicals based on BNF – 303 and BNF-304 and 306 (structural bioinformatics part)

Effective from June - 2008

External Marks: 80 Internal Marks: 40 Total Marks: 120

Part 1: Practical based on 303 & 306 (50%)

- 1. Retrieve DNA and Protein sequence, do MSA, using clustal W (from EBI), draw Dendogram and Jalview.
- 2. Advance Quering of databases- use of PSI BLAST (NCBI)
- 3. Identification of gene using Genscan.
- 4. Proteomics analysis (physiochemical property of protein) using Protparam.
- 5. 3D structure visualization and analysis using Rasmol,
- 6. Structure alignment visualization using Cn3D and structure comparison.
- 7. ORF finder (NCBI)
- 8. Motif and domain search (ExPasy server), Prosite database
- 9. Sequence logo.
- 10. Phylogenetic analysis: phylodraw, treecon
- 11. RNA structure prediction using Mfold
- 12. Add more tool based on paper Bioinformatics applications and structural bioinformatics.
- 13. A MINI PROJECT.

Part 2: Practical based on 304 (50%)

- 1. Programs based on scripting languages (PERL)
- 2. Programs based on HTML
- 3. Creating Forms, tables, frames, images etc
- 4. Program based on XML

T.Y.B.Sc. (Bioinformatics)

BNF-309: Practical based on BNF – 305 and BNF-306 Effective from June – 2008

External Marks: 80 Internal Marks: 40 Total Marks: 120

Part 1: Develop & process the programs for data structures using C++ language (50%)

- Simple program using object oriented programming language
- Programs based on arrays
- Programs based on Structures
- Programs based on Classes
- Programs based on Data Structures

Part 2: Using SQL and PL/SQL (50%)

- Creation and manipulation of a database table
- Simple database queries
- Database Queries based on joins
- Database Queries based on Group by and Having
- Simple PL/SQL blocks