## BIRLA INSTITUTE OF TECHNOLOGY AND SCIENCE, PILANI, HYDERABAD CAMPUS SECOND SEMESTER 20212022 COURSE HANDOUT (PART II)

Date: 15/01/2022

In addition to part I (General Handout for all courses appended to the timetable) this portion gives further specific details regarding the course.

Course NO. : Bio F242

Course Title : Introduction to Bioinformatics

Instructor-in-Charge: SHUVADEEP MAITY (L), VAISHALI KUMAR (T)

## 1. Scope and objective of the Course:

Introduction to genomic & Proteomics, Human Genome and other sequencing projects, Biological databases and data mining, sequence similarity search and sequence alignment, Protein structure predication and structure analysis, use of software package in Bioinformatics. This course designed to impart the beginner with the fundamentals, which would enable understanding of the intricacies and vast scope of Bioinformatics. A sampling of the different areas required for understanding of this upcoming field will be provided along with *in slilico* exercises to familiarize individuals with different program packages.

2. Text Book : "Introduction to Bioinformatics" Arthur M. Lesk; Oxford University Press (2009) (TB)

**3. Reference Books**:1. "Instant Notes in MOLECULAR BIOLOGY" P.C. Turner, A.G. McLennan, A.D. Bates & M.R.H. White, Viva Books Private Ltd, New Delhi. (RB1)

2. "Bioinformatics Genome and sequence Analysis" by David W Mount, CSHL

Press, 2003 (RB2)

## 4.Course Plan:

Lectur	Learning Objectives	Topics to be covered	Chapter in
e			Textbook
No.			
1.	Introduction	What is Bioinformatics, Scope	Lecture Notes
		Nucleic acid; Structure & function	Sec C- RB1
2-6			
	Overview of molecular biology &		
	genetics		
		Protein Structure & function	Sec B- RB1
		Central dogma of life –	Secs E/K/Q-
		Replication/Transcription/Translation	RB1
		Genetic code, Codon bias	Sec P- RB1
7-13.	General overview of different	DNA sequencing, RNA sequencing, PCR,	Class Notes
	techniques to generate	NMR, X-ray crystallography, Micro array,	
	biomolecular information and	Programming language for bioinformatics	
	analysis	(R,Perl)	
14.	Information Networks	WWW, TCP/IP, HTTP, URLs	Chap.2 TB
15-16	Collection and storage of	Data repositories (Genomics &	Chap-2,3,4,5
	sequences	proteomics), Submission of sequences to	(TB)
		the databank, Computer storage of	
		sequences, Web resources in	Class notes
		Bioinformatics	

17-18	Information Resources	Biological databases	Chap.4 TB Class notes
		Primary databases	Chap.3 TB
		Secondary databases	Chap.8 TB
19-25	Sequence Analysis and alignment	Definition of sequence alignment, Method of sequence analysis, Dot-matrix, dynamic programming algorithms for sequence alignment, use of scoring matrix and gap penalties, significance of sequence alignment, Multiple sequence alignment, statistical methods for aiding alignment, Markov models, Hidden Markov models, position-specific scoring matrices.	Chap. 3 and Chap. 4 RB2
26-29	Phylogenetic analysis	Tree building and evaluation methods	Chap. 4 TB
30-38	Next generation sequencing	RNAseq and its variants, Quality control experimental, Transcriptome assembly, data processing, differential expression analysis	Class Notes
39- 41	Analysis Packages	Commercial databases and software	Chap. 3 & 10 TB
42	Bioinformatics Programming	Introduction of different scripting language, Demo with R languages	Class notes

## 5. Evaluation scheme:

EC No.	Evaluation Component	Duration	Weightage %	Date, Time & Venue	Nature of Compone nt
1	4 Announced Quizzes* (Continuous evaluation)	Variable	35% (70M =15M+20M+ 15M+20M)	To be announced	OB# (50%) and Closed book (50%)
2	Mid-Sem	90 Min.	30% (60M)	12/03 11.00am to12.30pm	OB
3	Comprehensive	2 Hrs.	35% (70M)	11/05 AN	Closed Book

<sup>\*</sup>Quizzes will be conducted during lecture hours; two before the mid-semester and two after. Out of 4 quizzes 2 open books and 2 closed books.

- **6. Consultation Hour:** To be announced in the class.
- 7. Notices: Notices will be displayed via CMS.
- 8. Make up Policy: Make up will be given on genuine grounds as determined by the IC.
- **9. Academic Honesty and Integrity Policy:** Academic honesty and integrity are to be maintained by all the students throughout the semester and no type of academic dishonesty is acceptable.

<sup>#</sup> OB- Open Book Only prescribed textbook/Reference book(s), slides and hand written notes (in a binding format, No loose pages) are permitted