# BIRLA INSTITUTE OF TECHNOLOGY AND SCIENCE, PILANI, HYDERABAD CAMPUS SECOND SEMESTER 2018-2019 COURSE HANDOUT (PART II)

07-01-2019

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: Ramakrishna Vadrevu

Instructor : I. Shivakumar

## 1. Course Description:

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.

# 2. Scope and objective of the Course:

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.

- **3. Text Book** : "Introduction to Bioinformatics" Arthur M. *Lesk*; Oxford University Press (2008, Third Edition) (TB)
- **4. 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."GENES VII "Benjamin Lewin. (RB2)

3. "Bioinformatics Genome and sequence Analysis" by David W

Mount, CSHL Press, 2003 (RB3)

#### 5.Course Plan:

Lecture No.	Learning Objectives	Topics to be covered	Chapter in the Text Book
1.	Introduction	What is Bioinformatics, Scope	Lecture Notes
2-6	Overview of molecular biology & genetics	Nucleic acid; Structure & function	Sec C- RB1; Lecture Notes
		Protein Structure & function	Sec B- RB1;

		DNA replication	Sec E- RB1;
		Transcription	Sec K- RB1;
			Chap2, TB
		Translation	Sec Q- RB1;
			Chap2 TB
		Genetic code, Codon bias	Sec P- RB1
7-13.	General overview of different	DNA sequencing, Genome	Chap7 TB
	techniques to generate	sequencing, PCR, NMR, X-ray	Lecture
	biomolecular information and	crystallography, Micro array,	Notes
	analysis	Proteomics	
14.	Information Networks	WWW, TCP/IP, HTTP, URLs	Chap2 TB
15-16	Collection and storage of	Submission of sequences to the	Chap2,3,4,5
	sequences	databank, Computer storage of	(TB)
		sequences, Web resources in	
		Bioinformatics	
17-18		Biological databases	Chap3/4 TB
	Information Resources		(Hands-on)
		Primary databases	Chap3 TB
		Secondary databases	Chap4 TB
19-33		Definition of sequence alignment,	Chap5 TB
		Method of sequence analysis, Dot-	RB3
		matrix, dynamic programming	
		algorithms for sequence alignment,	
		use of scoring matrix and gap	
	Sequence Analysis and	penalties, significance of sequence	
	alignment	alignment, Multiple sequence	
		alignment, statistical methods for	
		aiding alignment, Markov models,	
		Hidden Markov models, position-	
		specific scoring matrices.	
34-35	Phylogenetic analysis	Tree building and evaluation methods	Chap5 TB
36-38	Protein structure prediction	Homology modeling, abinitio	Chap. 6
		structure prediction, Threading	TB/Class
		method	Notes
39- 41	Analysis Packages	Commercial/Public Domain databases	Chap. 3
		and softwares,	Hands-on
42	Bioinformatics Programming	Introduction of different scripting	Class notes
		language	

### **6.** Evaluation scheme:

Components	Duration	Date	Time	Weightage	Nature of
				(%)	Component
Midsem	90 min	11/3	11.00 -12.30	30%	Closed Book
			PM		
Surprise and or	Diverse			30%	Open Book (20%) /
announced					Closed Book(10%)
quiz(s)/Open					
assignment(s)					
either take home					
or in the class					
(surprise/announ					
ced) and or					
Hands-on testing					
sessions					
(computational)					
Comprehensive	3 Hours	01/05	AN	40%	Closed Book
examination					

- 7. Chamber Consultation Hour: To be announced in the class.
- 8. Notices: Notices, if any concerning the course will be displayed on the notice Board of Biology Dept. and on CMS.
- 9. Make up Policy: Only in case of hospitalization for the exams. No Make up for class/take home assignments/announced quizzes. (all non-exam components).
- **10.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.

Instructor-in-charge Bio F242