# Life Sciences Practicals II

**12 Credits** 

## Coordinator: Dr. Niti Puri

**LS431A** 

Participating Faculty Members (Module-wise):

I Maharahar Dishara Dash Don data Dash K. Natara

- I. Molecular Biology: Prof. P.C. Rath\*, Prof. K. Natarajan
- II. Plant Physiology: Prof. A.K. Nandi\*, Prof. S. Chakraborty, Dr. Amarjeet Singh
- III. Animal Physiology: Prof. A.C. Mondal\*, Dr. S.K. Jha, Prof. R.P. Singh
- IV. Developmental Biology: Prof. S. Saran\*, Prof. A.K. Sarkar
- V. Immunology: Dr. N. Puri\*
- VI. **Computational Biology and Bioinformatics**: Dr. D. Kaur, Prof. S. Gourinath, Prof. N. Ramchiary\*, Dr. K. Kar

	S.No.	Торіс	Faculty Name/ Contact Hours
	1.	DNA Structure: Preparation of a ball and stick model of B- DNA. Draw the chemical structures of A,T and G,C base pairs, show different groups/ bonds in the DNA and study the structural features.	PCR/10* (*overlapping with other experiments)
	2.	Genomic DNA isolation from yeast <i>Candida albicans</i> / mouse cell line and DNA quantification	KN/2
(	3.	Primer designing for PCR-based cloning of genes for expression in <i>E.coli</i> - use of various tools and genome sequence databases	KN/2
credits	4.	Titration of template concentrations and PCR amplification; Quantification of amplicons by gel electrophoresis and clean-up of DNA.	KN/1
logy (2	5.	<u>Preparation of Insert DNA</u> : Restriction digestion of PCR fragment with six-base cutter/s, gel analysis and clean-up and quantify DNA. DNA ready for ligation.	KN/1
Molecular Biology (2 credits)	6.	<u>Preparation of vector DNA:</u> Restriction digestion of pET28b vector with compatible six-based cutter/s; verification of RE digestion by gel analysis; purification of vector DNA and quantification by gel analysis.	KN/1
Mole	7.	Preparation of LB medium, LB agar plates with antibiotics and other reagents.	PCR/1
	8.	Streaking of <i>E. coli</i> DH5α strain on LB+Agar+ Nalidixic acid plate	PCR/1
	9.	Preparation of competent <i>E. coli</i> DH5α cells	PCR/2
	10.	Transformation of <i>E. coli</i> DH5α cells with pBluescript, pET28b plasmid DNA and recombinant DNA clones	PCR/2
	11.	Isolation of plasmid DNA from overnight culture of the transformed colony/ clone.	PCR/1
	12.	Agarose gel electrophoresis of plasmid DNA preparation	PCR/1

	13.	Restriction enzyme digestion of the plasmid DNA and clone verification         Restriction enzyme digestion of bacteriophage λ DNA/         genome by Hind III and Hind III+EcoR I; preparation of restriction map for the double digest.	PCR/1
	14.	Blue/ white screening of the recombinant DNA clones in <i>E.coli</i> .	PCR/2
S.	1.	Regeneration and transformation of tobacco using <i>Agrobacterium tumefaciens</i> .	SC/6
olog ts)	2.	Study of photosynthetic efficiency in normal and stressed plants	AJS/1
Plant Physiology (2 credits)	3.	Separation of Chlorophyll pigments and study of absorption spectra	AJS/2
ant (2	4.	Effect of plant hormones on growth and physiology	AJS/4
Pl	5.	Stress-associated cell death and reactive oxygen species accumulation studies	AKN/ 2
	1.	Histology of liver, kidney & brain; Histology of normal and carcinomas of different human organs	RPS/1
	2.	Tissue processing for histological preparations	ACM/1
	3.	Section cutting; Microtomy (Rotary), Vibrotomy, Cryo- sectioning	ACM/1
<b>s</b> )	4.	Staining: Haematoxyline/ Eosin staining for different organs and tissues	ACM/2
credits hysiology	5.	Preparation of permanent slides	ACM/1
(2 cro al Physi	6.	Identifying characters for sections of different tissues or organs stained with H&E.	ACM/1
logy z Anim	7.	Histology of Blood: Preparation & staining of human blood film	ACM/1
Animal Physiology (2 Animal Histology & Animal Pl	8.	<ul> <li>(i) Determination of blood group</li> <li>(ii) Study of haemin crystal</li> <li>(iii) Measurement of arterial blood pressure</li> </ul>	ACM/1
Animal Physiology Animal Histology & Anim	9.	Immunohistochemistry of brain sections and cancer tissues:Staining, microscopic evaluation and quantification ofbiomarkers and determination of proliferation index	RPS/2
7	10.	Demonstration and application of rodent's stereotaxic instruments.	SKJ/1
	11.	Demonstration of recorded brain waves. Identification of vigilant states in the recorded brain waves.	SKJ/1
	12.	Identification of alpha, beta, delta, theta and gamma brain waves in recorded the EEG.	SKJ/1

	Plant Developmental Biology				
Developmental Biology (2 credits) Plant and Animal Developmental Biology	1.	Embryogenesis: sample preparation and microscopic visualization of developmental stages of <i>Arabidopsis</i> embryos.	AS/1		
	2.	Shoot and root meristems: sample preparation, morphological and microscopic visualization of embryonic and post-embryonic shoot/root meristems	AS/1		
	3.	Root growth and branching: morphological and microscopic observation of root growth and number or lateral roots (LRs); microscopical observation of stages of LR development in Arabidopsis	AS/1		
	4.	Auxin localization in shoot and root using DR5:GUS reporter. GUS- staining of <i>Arabidopsis</i> seedlings, analysis and imaging of staining pattern in shoot and root tissue	AS/1		
	5.	Analysis of root developmental/auxin pathway mutant for defect in root, comparison with wild type	AS/1		
	6.	Leaf polarity- analysis of dorsoventral features. Observation of number and density/pattern of stomata and trichomes on leaf surfaces. Tissue fixation, processing, embedding and Histological analysis leaf tissues	AS/1		
ntal	Animal Developmental Biology				
pme t and A	7.	Day 1: Preparation of media, buffer and reagents for experiments	SS/1		
Develo <sub>Plan</sub>	8.	Day 2: Studying growth and development of <i>Dictyostelium discoideum</i>	SS/1		
	9.	Day 3: Continue with development of <i>D.discoideum</i>	SS/1		
	10.	Day 4: To study pattern formation in Dictyostelium using neutral red dye	SS/1		
	11.	Day 5: LacZ staining of few mutants of <i>Dictyostelium</i>	SS/1		
	12.	Day 6: Observing developmental mutants of Drosophila	SS/1		
	13.	Day 7: Observing different model systems used in Developmental Biology	SS/1		
	Immu	nology			
×	1.	Counting of immune cells: Differential Leukocyte count; Total Leukocyte count	NP/1		
Immunology (2 credits)	2.	Innate Immune Response 1: Phagocytosis of Fl-labeled microbes or damaged cells by phagocytes followed by assay of ROS production (Fluorescence microscopy/Flowcytometry)	NP/2		
Imm (2 c	3.	Innate Immune Response 2: Exocytosis/Release of inflammatory mediators by Mast cells in response to allergen challenge (Mediator release assessment by colorimetric assay)	NP/2		
	4.	Activation of complement by classical pathway: Lysis of antibody coated cells	NP/1		

	5.	Adaptive humoral immune responses: Checking antibody titer in immunized mouse serum by indirect ELISA (comparison of	NP/2
	5.	immunized versus unimmunized)	111/2
	6.	Use of Antigen-antibody interaction in immunological techniques	NP/1
		for antigen detection and quantitation (Immunodiffusion/	
		haemagglutination; immunoprecipitation/western blotting)	
	7.	Tissue typing (MHC-typing) by Mixed lymphocyte reaction	NP/1
	<i>.</i>	(MTT/CFSE-FACS)	111/1
	8.	T-cell subset determination (CD4 and CD8) by	NP/2
		Immunophenotyping experiment/labeling/ running sample on	
		Flowcytometry followed by analysis	
	9.	T-cell activation by mitogen (Con A). Study proliferation (MTT/CFSE-FACS) and detection of cytokines/chemokines	NP/2
		secretion by activated immune cells using ELISA (sandwich	
		ELISA).	
	10	NK cell-mediated killing of tumor cell targets by MTT assay	NID /1
	10.	(non-radioactive method)	NP/1
	Comp	outational Biology & Bioinformatics	
	1.	Database searching using Boolean operators (Advanced database	DV /1
tici		searching) and data retrieval	DK/1
na	2.	Sequence analysis of DNA for base compositions and Codon	DK/3
OLT		usage analysis, Use of Pattern finding algorithm to find	
ioinfo			DK/3
	2.	patterns/motifs and statistical relevance using Python or in-silico	DK/3
ioi		tools.	DK/3
ż Bioi	3.	tools. Pair-wise alignment and MSA with DNA/Protein sequence	DK/3 NR/KK/2
y & Bioinformatics its)		tools. Pair-wise alignment and MSA with DNA/Protein sequence dataset, Phylogenetic tree overview	
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utational Biology & Bioi (2 credits)	3.	<ul> <li>tools.</li> <li>Pair-wise alignment and MSA with DNA/Protein sequence dataset, Phylogenetic tree overview</li> <li>Query search against different databases, understanding search set options, program selection and different algorithm parameters, Overview of BLAST suite of programs – Computational approach – Statistical significance</li> <li>Predicting RNA secondary structure and Protein structures using in-silico tools (Homology Modelling, Ab-initio and Threading concepts).</li> </ul>	NR/KK/2 DK/2
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Computational Biology & Bioi (2 credits)	3. 4. 5.	<ul> <li>tools.</li> <li>Pair-wise alignment and MSA with DNA/Protein sequence dataset, Phylogenetic tree overview</li> <li>Query search against different databases, understanding search set options, program selection and different algorithm parameters, Overview of BLAST suite of programs – Computational approach – Statistical significance</li> <li>Predicting RNA secondary structure and Protein structures using in-silico tools (Homology Modelling, Ab-initio and Threading concepts).</li> </ul>	NR/KK/2 DK/2 DK/4

## **Further Reading:**

### 1.

**Molecular Biology** Molecular Cloning- A LABORATORY MANUAL 4<sup>th</sup> Edition Green, M.R. and Sambrook, J.

#### 2. Plant Physiology

Concerned literature/handout will be given by individual faculty members.

#### 3. Animal Physiology

- 1. diFiore's Atlas of Histology with Functional Correlations by Victor P Eroschenko
- 2. Theory and Practice of Histological Techniques Eds. John D Bancroft, Marilyn Gamble
- 3. Atlas of Histology with Functional Correlations Victor P. Eroschenko
- 4. Histology for Pathologists. Stacey E. Mills (Ed.)

#### 4. Developmental Biology

- 1. Plant Physiology and Development, 6th Ed. by Eduardo Zeiger and Lincoln Taiz et al.
- 2. Arabidopsis: A laboratory manual (CSHL press). Detlef Weigel and Jane Glazebrook
- 3. Handouts for experiments will be given in the class.

#### 5. Immunology

Concerned literature/handout will be given by the faculty member

#### 6. Computational Biology & Bioinformatics

#### **Textbook:**

- 1. Jonathan Pevsner. Bioinformatics and functional genomics. Wiley Blackwell, Third Edition, 2015
- 2. Des Higgins and William Taylor. Bioinformatics: Sequence, Structure, and databanks. Oxford University Press, 2001

#### **Reference Book:**

 Richard Durbin, Sean R. Eddy, Anders Krogh, Graeme Mitchison. Biological Sequence Analysis: Probabilistic Models of Proteins and Nucleic Acids. Cambridge University Press, 1998