Syllabus

For

Ph.D. Coursework (Biotechnology)

Session: 2020-2021



P.G. Department of Biotechnology Utkal University Bhubaneswar, Odisha

Outline of Ph.D. Coursework Curriculum P.G. Department of Biotechnology Utkal University

Semester-I			
Sl No	Paper	Subject	Marks
1	I	Research Methodology/ Tools and Techniques in Biotechnology	100
2	II	Biostatistics and Computational Biology	100
3	III	Review of Literature : 50 marks Presentation before SRC : 50 marks	100
4	IV	 Seminar-I: 50 marks Seminar-II: 50 marks 	100
Total			400

Ph.D. coursework Syllabus (Biotechnology)

Paper-I Marks: 100

<u>UNIT -I (RESEARCH METHODOLOGY)</u>

- 1. Research objective and statement of problem, types of research.
- 2. Research proposal designing and formulation.
- 3. Research Methodology.
- 4. Review of literature, concept development, theory, hypothesis and manuscript preparation.

UNITS-II (INSTRUMENTATIONS)

- 1. Principles and applications of spectrophotometer.
- 2. Principles and applications of Chromatography. Types of Chromatography.
- 3. Principles and applications of Electrophoresis.
- 4. Principles and applications of PCR.

UNIT – III (TECHNIQUES IN BIOTECHNOLOGY)

- 1. Isolation and purification of DNA.
- 2. Isolation and purification of RNA.
- 3. Isolation and Purification of Proteins.
- 4. Commonly used vectors for gene-cloning, DNA manipulating enzymes, construction of genomic and cDNA libraries.

UNIT – IV (MICROBIOLOGY)

- 1. Methods in Microbiology: Sterilization, Culture Media, Pure culture technique, enrichment culture technique, Microbial staining methods, Maintenance and preservation of Microorganisms.
- 2. Microbial growth: Growth curve, measurement of growth, growth yields, synchronous growth, continuous culture, growth as affected by environmental factors.
- **3.** Antibiotics: Penicillin and Cephalosporin, broad-spectrum antibiotics, antibiotics from prokaryotes, antifungal antibiotics, mode of action of different antibiotics, resistance to antibiotics, Multiple Drug Resistance.
- **4.** Microbes for industrial use. Fermentation and fermented products, Bio-fertilizers, Microbes in mineral beneficiation and bioremediation.

<u>UNIT-V (IMMUNO-TECHNOLOGY)</u>

- 1. Antigen antibody interaction and their applications in immunodiagnostics.
- 2. Detection of antigen /protein by western blotting.
- 3. Antigen based biosensor, ELISA and CHIP.
- 4. FACS and its applications.

Paper-II Marks: 100

BIOSTATISTICS AND COMPUTATIONAL BIOLOGY

UNIT - I

- 1. Statistics: Definition, functions and limitations.
- 2. Treatment of data: frequency distribution, Graph of Frequency Distribution
- 3. Descriptive Measures: Averages and Dispersions (Grouped and ungrouped).
- 4. Probability: Concepts, definition and elementary problems based on definition.

UNIT - II

- 1. Inference: Definition-parameter, Statistic sampling distributors, standard error,
- 2. Test of Hypothesis, type I and Type II errors.
- 3. Large sample tests: Z tests, small sample tests: t and F tests.
- 4. Chi-square test: Goodness of fit and Test of independence.

UNIT - III

- 1. Curve Fitting: First, Second degree and exponential curve.
- 2. Simple correlation and Regression.
- 3. Concept of multiple correction and Regression.
- 4. Analysis of variance: one way and two way classification.

<u>UNIT – IV</u>

- 1. Fundamentals of Computer: CPU, memory, I/O unit, storage, multimedia. Introduction to Operating Systems: DOS, Windows and Linux Operating Systems. Ideas about Computer Viruses. Use of online resources and the internet communication technology.
- 2. Idea on working with MS-Word, Excel and Power point.
- 3. Programming with C++ and PERL: Introduction, Control Flow: Statement and Block, if, ifelse, Nested if-else statements, For, while, do-while loops, break, switch continue, Statements, go to statement. Functions and Arrays.
- 4. Computation of simple mathematical and statistical formulae using the Programming C++ and PERL.

UNIT - V

- 1. Introduction to Bioinformatics
- 2. Elementary idea about Database management system, e.g. Gene bank, EMBL, Swiss-Prot, Sequence database like FASTA, BLAST algorithm and Bioinformatics tools.
- 3. Pairwise sequence alignment, multiple sequence alignment, Gene prediction and Protein structure prediction.

Paper-III Marks: 100

3. Review of Literature : 50 marks4. Presentation before SRC : 50 marks

Paper-IV Marks: 100

3. Seminar-I: 50 marks4. Seminar-II: 50 marks