# CS4465: COMPUTATIONAL BIOLOGY AND BIOINFORMATICS

## **Effective Term**

Semester A 2024/25

# Part I Course Overview

## **Course Title**

Computational Biology and Bioinformatics

# **Subject Code**

CS - Computer Science

## **Course Number**

4465

#### **Academic Unit**

Computer Science (CS)

## College/School

College of Computing (CC)

## **Course Duration**

One Semester

#### **Credit Units**

3

# Level

B1, B2, B3, B4 - Bachelor's Degree

# **Medium of Instruction**

English

## **Medium of Assessment**

English

# Prerequisites

(BMS2801 Molecules and Cells or

BME2106 Introduction to Cellular and Biomolecular Engineering or equivalent)

AND

(CS1102 Introduction to Computer Studies or

CS1302 Introduction to Computer Programming or equivalent)

## **Precursors**

Nil

# **Equivalent Courses**

Nil

## **Exclusive Courses**

Nil

# Part II Course Details

## **Abstract**

This course aims to introduce students concepts and techniques in computational biology and bioinformatics, and to develop practical skills in applying bioinformatics approaches to solve problems in biological studies.

# **Course Intended Learning Outcomes (CILOs)**

	CILOs	Weighting (if app.)	DEC-A1	DEC-A2	DEC-A3
1	Describe the main characteristics of different computational methods in bioinformatics.		X		
2	Explain the effectiveness and efficiency of current computational biology and bioinformatics techniques.			x	Х
3	Identify the problems in biological studies that can be analyzed using computational methods.		X	X	
4	Design bioinformatics solutions for practical problems in biological studies by applying and integrating suitable computational tools.		X	x	x

#### A1: Attitude

Develop an attitude of discovery/innovation/creativity, as demonstrated by students possessing a strong sense of curiosity, asking questions actively, challenging assumptions or engaging in inquiry together with teachers.

## A2: Ability

Develop the ability/skill needed to discover/innovate/create, as demonstrated by students possessing critical thinking skills to assess ideas, acquiring research skills, synthesizing knowledge across disciplines or applying academic knowledge to real-life problems.

# A3: Accomplishments

Demonstrate accomplishment of discovery/innovation/creativity through producing /constructing creative works/new artefacts, effective solutions to real-life problems or new processes.

# **Learning and Teaching Activities (LTAs)**

	LTAs	<b>Brief Description</b>	CILO No.	Hours/week (if applicable)
1	Lecture	Students will engage in formal lectures to gain knowledge about computational biology and bioinformatics techniques, and the features of the main bioinformatics software tools.	1, 2, 3	3 hours per week

2	Laboratory	Students will discover the characteristics of the main computational methods in bioinformatics. They will also perform a critical evaluation of the effectiveness and efficiency of these methods in the form of lab sheets.	1, 2, 3	8 hours per semester
3	Project and Assignments	Students will identify suitable problems in biological studies that are amenable to computational analysis, and create new bioinformatics solutions to address these problems.	2, 3, 4	After class
4	Final Exam	Students will be tested for their understanding and ability to discover and assess bioinformatics techniques. Students will also be assessed for their abilities to identify and create bioinformatics solutions.	1, 2, 3, 4	After class

# Assessment Tasks / Activities (ATs)

	ATs	CILO No.	Weighting (%)	Remarks (e.g. Parameter for GenAI use)
1	Lab Sheets	1, 2, 3	10	
2	Assignments	2, 3, 4	10	
3	Project Report and Deliverables	2, 3, 4	20	

# Continuous Assessment (%)

40

# Examination (%)

60

# **Examination Duration (Hours)**

2

# **Additional Information for ATs**

For a student to pass the course, at least 30% of the maximum mark for the examination must be obtained.

# **Assessment Rubrics (AR)**

# **Assessment Task**

Laboratory Participation

4 CS4465: Computational Biology and Bioinformatics

#### Criterion

1.1 CAPACITY for SELF-DIRECTED LEARNING to understand and apply the principles of bioinformatics methods and tools 1.2 CAPACITY for ASSESSING different computational biology and bioinformatics techniques critically

Excellent (A+, A, A-)

High

Good (B+, B, B-)

Significant

Fair (C+, C, C-)

Moderate

Marginal (D)

Basic

Failure (F)

Not even reaching marginal levels

## **Assessment Task**

Assignments

#### Criterion

- 2.1 ABILITY to EXPLAIN different bioinformatics techniques and discover their characteristics
- 2.2 ABILITY to PERFORM a critical assessment of different computational biology and bioinformatics techniques
- 2.3 ABILITY to EXPLAIN different examples of problems in biological studies, and their ability to identify the suitable class of problems which are amenable to bioinformatics analysis

Excellent (A+, A, A-)

High

Good (B+, B, B-)

Significant

Fair (C+, C, C-)

Moderate

Marginal (D)

Basic

Failure (F)

Not even reaching marginal levels

## Assessment Task

Project

studies

### Criterion

3.1 ABILITY to EXPLAIN in DETAIL and with ACCURACY about different examples of problems in biological studies, and their ability to identify the suitable class of problems which are amenable to bioinformatics analysis
3.2 ABILITY to DISCOVER/INTEGRATE and CREATE new bioinformatics solutions for practical problems in biological

Excellent (A+, A, A-)

High

Good (B+, B, B-)

Significant

Fair (C+, C, C-)

Moderate

Marginal (D)

Basic

Failure (F)

Not even reaching marginal levels

## **Assessment Task**

Final Exam

#### Criterion

4.1 ABILITY to EXPLAIN and COMPARE in DETAIL and with ACCURACY about different bioinformatics methods
4.2 ABILITY to EXPLAIN in DETAIL and with ACCURACY about different examples of problems in biological studies, and
their ability to identify the suitable class of problems which are amenable to bioinformatics analysis

Excellent (A+, A, A-)

High

Good (B+, B, B-)

Significant

Fair (C+, C, C-)

Moderate

Marginal (D)

Basic

Failure (F)

Not even reaching marginal levels

# **Part III Other Information**

# **Keyword Syllabus**

Pairwise sequence alignment, multiple sequence alignment, phylogenetic tree reconstruction, motif identification, RNA informatics, protein domain annotation and prediction, protein structure prediction, contemporary topics in bioinformatics.

## **Reading List**

# **Compulsory Readings**

1 W.K.Sung (2009). Algorithms in Bioinformatics: A Practical Introduction. Chapman and Hall/CRC.

# **Additional Readings**

	Title	
1	J. Pevsner (2015). Bioinformatics and Functional Genomics. Wiley-Blackwell, 3rd edition	
2	A. Lesk (2014). Introduction to Bioinformatics. Oxford University Press, 4th edition.	
3	N. Cristianini and M. Hahn (2007). Introduction to Computational Genomics: A Case Stud University Press.	dies Approach. Cambridge