



Academic year	2016-17
Subject	11261 - Bioinformatics
Group	Group 1, 2S
Teaching guide	A
Language	English

Subject identification

Subject	11261 - Bioinformatics
Credits	0 de presencials (0 hours) 3 de no presencials (75 hours) 3 de totals (75 hours).
Group	Group 1, 2S
Teaching period	Second semester
Teaching language	English

Professors

Lecturers	Horari d'atenció als alumnes					
	Starting time	Finishing time	Day	Start date	Finish date	Office
Gerard Pujadas Anguiano						You need to book a date with the professor in order to attend a tutorial.

Contextualisation

The students of this course have different background and interests for bioinformatics. Therefore, we have designed the course with the goal on mind that you have to learn bioinformatics skills that can be useful for all of you. Following with this idea, the course have two chapters where you will learn how to search and how to analyze biologic information. In the first chapter, you will learn to use the SRS (Sequence Retrieval System). This is a web interface to a large number of biological databases and analysis tools. In the second chapter you will learn to search into microarray results databases.

Requirements

No special skills

Skills

The competences are:

Specific

- * G13 - Knowing the capabilities and potential of ICT (Information and Communication) in the bioinformatics area..

Generic

- * Known the capabilities of the new technologies and their application to the nutrition and nutrigenomics fields.
- * CB7 - That students can apply their knowledge and ability in problem solving environments new or unfamiliar within broader (or multidisciplinary) contexts related to their field of study.



Teaching guide

Basic

* You may consult the basic competencies students will have to achieve by the end of the Master's degree at the following address: http://estudis.uib.cat/master/comp_basiques/

Content

The contents of the course is:

Theme content

- 1.. Sequence analysis of proteins and nucleic acids
 - Sequence databases for proteins and nucleic acids.
 - Sequence searching in databases and analysis of the retrieved information.
 - Sequence comparison
- 2.. Analysis of protein structures and nucleic acids
 - 3D structural databases for proteins and nucleic acids
 - Looking for 3D structures of proteins and nucleic acids and visual analysis of their main features
 - Protein structure comparison
- 3.. Predicting bioactivity of ligands
 - 3.1. Databases ligands nutritional interest
 - 3.2. Fundamentals of virtual screening
 - 3.3. Virtual screening in the absence of the 3D structure of the target
 - 3.4. Virtual screening in the presence of the 3D structure of the target

Teaching methodology

The methodology used will be:

Workload

The activities of the course will be:

In-class work activities

Modality	Name	Typ. Grp.	Description	Hours
Other	There is no activity of this type	Large group (G)	a	0

At the beginning of the semester a schedule of the subject will be made available to students through the UIBdigital platform. The schedule shall at least include the dates when the continuing assessment tests will be conducted and the hand-in dates for the assignments. In addition, the lecturer shall inform students as to whether the subject work plan will be carried out through the schedule or through another way included in the Campus Extens platform.

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Distance education work activities

Modality	Name	Description	Hours
Individual self-study	Exam	Exam	2
Individual self-study	Exercises	To analyze the sequences and structures of nucleic acids and proteins by using software and specialized databases in the field of Molecular Biology. Predict and analyze specific capabilities as bioactive ligands based on their structure and composition.	73

Specific risks and protective measures

The learning activities of this course do not entail specific health or safety risks for the students and therefore no special protective measures are needed.

Student learning assessment

The evaluation will be done according to the following criteria:

Exam

Modality	Individual self-study
Technique	Short-answer tests (retrievable)
Description	Exam
Assessment criteria	
Final grade percentage:	50%

Exercises

Modality	Individual self-study
Technique	Student internship dissertation (non-retrievable)
Description	To analyze the sequences and structures of nucleic acids and proteins by using software and specialized databases in the field of Molecular Biology. Predict and analyze specific capabilities as bioactive ligands based on their structure and composition.
Assessment criteria	
Final grade percentage:	50%

Resources, bibliography and additional documentation



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The materials used (videos, web pages, etc..) will be available through the course moodle space

