		Teachin	g Guide				
	Identifyir	ng Data			2021/22		
Subject (*)	Fundamentals of bioinformatics Code 614522008			614522008			
Study programme	Mestrado Universitario en Bioinfo	rmática para C	iencias da Saúd	e			
	<u>'</u>	Desci	riptors				
Cycle	Period Year Type C			Credits			
Official Master's Degre	ree 1st four-month period First Obligatory 6				6		
Language	English						
Teaching method	Hybrid						
Prerequisites							
Department	Ciencias da Computación e Tecr	oloxías da Info	rmaciónComputa	aciónFisioterapia, Medicir	na e Ciencias Biomédicas		
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	Munteanu , Cristian Robert			c.munteanu@ud	c.es		
Web	moodle.udc.es			·			
General description	This course will provide concepts	on the basic p	rinciples of geno	me annotation, sequence	analysis, processing tools of		
	molecular information, tools for d	rug design and	evaluation of tox	cicity, biological database	s, omics and epigenetics, the		
	Human Genome, Exposome and	Variome project	cts, and bioinforr	natics applications in clini	cal practice.		
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	Study programme competences
Code	Study programme competences
A1	CE1 - Ability to know the scope of Bioinformatics and its most important aspects
A6	CE6 - Ability to identify software tools and most relevant bioinformatics data sources, and acquire skill in their use
A7	CE7 - Ability to identify the applicability of the use of bioinformatics tools to clinical areas.
B1	CB6 - Own and understand knowledge that can provide a base or opportunity to be original in the development and/or application of ideas,
	often in a context of research

B2	CB7 - Students should know how to apply the acquired knowledge and ability to problem solving in new environments or little known within
	broad (or multidisciplinary) contexts related to their field of study
В3	CB8 - Students to be able to integrate knowledge and deal with the complexity of making judgements from information that could be
	incomplete or limited, including reflections on the social and ethical responsibilities linked to the application of their skills and judgments
B5	CB10 - Students should possess learning skills that allow them to continue studying in a way that will largely be self-directed or
	autonomous.
B6	CG1 -Search for and select the useful information needed to solve complex problems, driving fluently bibliographical sources for the field
B7	CG2 - Maintain and extend well-founded theoretical approaches to enable the introduction and exploitation of new and advanced
	technologies
B8	CG3 - Be able to work in a team, especially of interdisciplinary nature
C1	CT1 - Express oneself correctly, both orally writing, in the official languages of the autonomous community
C2	CT2 - Dominate the expression and understanding of oral and written form of a foreign language
C3	CT3 - Use the basic tools of the information technology and communications (ICT) necessary for the exercise of their profession and
	lifelong learning
C6	CT6 - To assess critically the knowledge, technology and information available to solve the problems they face to.
C8	CT8 - Rating the importance that has the research, innovation and technological development in the socio-economic and cultural progress
	of society

Learning outcomes				
Learning outcomes		Study programme competences		
To identify the characteristics of the computer science applications in health sciences	AJ1	BJ1		
	AJ6	BJ2		
		BJ3		
To be able to develop a research project in the field of biomedical informatics according to ethical and security health data	AJ7	BJ5	CJ1	
requirements		BJ6	CJ2	
		BJ7	CJ3	
		BJ8	CJ6	
			CJ8	
To know how to identify fields of application of information technologies and communications to improve the delivery of health	AJ7		CJ1	
services to citizens			CJ2	
			CJ3	
			CJ6	
			CJ8	

Contents		
Торіс	Sub-topic	
Basic principles for Genome Annotation		
Sequence analysis		
Processing tools of molecular information		
Tools for drug design and evaluation of toxicity		
Biological databases		
Omics and epigenetics: genomics, proteomics,		
transcriptomics		
Projects: Human Genome, Variome, Exposome		
Bioinformatics applications in clinical practice		

	Planning	g		
Methodologies / tests	Competencies	Ordinary class	Student?s personal	Total hours
		hours	work hours	

ICT practicals	A1 A6 A7 B1 B2 B3	30	30	60
	B5 B6 B7 B8 C1 C2			
	C3 C6 C8			
Oral presentation	A1 C1 C2 C3 C6 C8	5	5	10
Supervised projects	A1 C1 C2 C3 C6 C8	10	10	20
Objective test	A1 A6 A7 B1 B2 B3	1	14	15
	B5 B6 B7 B8 C1 C2			
	C3 C6 C8			
Guest lecture / keynote speech	A1 A6 A7 B1 B2 B3	20	20	40
	B5 B6 B7 B8 C1 C2			
	C3 C6 C8			
Personalized attention		5	0	5
(*)The information in the planning table i	o for guidance only and does not take	o into account the l	notorogonoity of the ct	Idonto

(\*)The information in the planning table is for guidance only and does not take into account the heterogeneity of the students.

	Methodologies
Methodologies	Description
ICT practicals	Laboratory practice can be face-to-face or through computer platforms such as TEAMS.
Oral presentation	Public presentation of the supervised work can be face-to-face or through computer platforms such as TEAMS.
Supervised projects	Practical work on the theoretical content of the course can be face-to-face or through computer platforms such as TEAMS.
Objective test	Exam on the theoretical content and supervised work carried out throughout the course if there is no supervised work. It can
	be face-to-face or through computer platforms such as TEAMS.
Guest lecture /	In the theory sessions, the teacher describes the objectives and contents of the subject, to give a particular view of the subject
keynote speech	to be dealt with and to relate it to others within the subject.
	Then the corresponding topic is developed in the form of a lecture session, using the technical tools available, emphasizing certain issues in which the student must deepen his self-learning.
	The master sessions can be face-to-face or through computer platforms such as TEAMS. It is also possible to include
	explanatory videos of different parts of the theoretical contents.

Personalized attention		
Methodologies	Description	
Supervised projects	To solve the most complex aspects of the course, individual or group tutorials with students will be held.	
Objective test		
Oral presentation		
Guest lecture /		
keynote speech		
ICT practicals		

		Assessment	
Methodologies	Methodologies Competencies Description		Qualification
Supervised projects	A1 C1 C2 C3 C6 C8	The proposed work on the subject will be part of the evaluation.	30
Objective test	A1 A6 A7 B1 B2 B3	If deemed necessary, a test on the theoretical and practical content of the course	30
	B5 B6 B7 B8 C1 C2	(including the topics of the lectures and publicly exposed supervised projects) may be	
	C3 C6 C8	conducted. The teacher can distribute points of this test among other methods if	
		deemed appropriate.	
Oral presentation	A1 C1 C2 C3 C6 C8	The public presentation of the supervised work will be part of the final assessment.	30
ICT practicals	A1 A6 A7 B1 B2 B3	The quality and delivery in time of the practices will be assessed.	10
	B5 B6 B7 B8 C1 C2		
	C3 C6 C8		



## **Assessment comments**

To pass this course, the student needs to obtain a minimum percentage in each of the methodologies.

## Plagiarism:

In any submission in which plagiarism is detected, the submission will be assessed a zero. Plagiarism in the objective test will be penalized in accordance with current university regulations.

	Sources of information
Basic	- Stekel, Dov. (2003). Microarray bioinformatics. Cambridge: Cambridge University Press, 2003
	- Ohlebusch, Enno (2013). Bioinformatics algorithms : sequence analysis, genome rearrangements, and phylogenetic
	reconstruction. Ulm : Oldenbusch Verlag
	- Dan E. Krane, Michael L. Raymer (2003). Fundamental concepts of bioinformatics. San Francisco, California :
	Benjamin Cummings
	- Edward Keedwell and Ajit Narayanan (2005). Intelligent bioinformatics the application of artificial intelligence
	techniques to bioinformatics problems. Chichester : John Wiley & Dons
	Graph-based Processing of Macromolecular Information, Current Bioinformatics 10(5): 606-631 (2016), DOI:
	10.2174/1574893610666151008012438   Cristian R. Munteanu, Vanessa Aguiar-Pulido, Ana Freire, Marcos
	Martínez-Romero, Ana B. Porto-Pazos, Javier Pereira, Julian Dorado   onlineRRegrs: An R package for
	Computer-aided Model Selection with Multiple Regression Models, Journal of Cheminformatics 7(1), 1-16,
	doi:10.1186/s13321-015-0094-2 (2015)   Georgia Tsiliki, Cristian R. Munteanu, Jose A Seoane, Carlos
	Fernandez-Lozano, Haralambos Sarimveis, Egon L. Willighagen   GitHub  10.5281/zenodo.21946   online Bio-AIMS
	Collection of Chemoinformatics Web Tools based on Molecular Graph Information and Artificial Intelligence Models,
	Combinatorial Chemistry & Combinatorial Chem
	González-Díaz, Rafael García, Mabel Loza, Alejandro Pazos   online S2SNet: A Tool for Transforming Characters and
	Numeric Sequences into Star Network Topological Indices in Chemoinformatics, Bioinformatics, Biomedical, and
	Social-Legal sciences, Current Bioinformatics 8(4), 429-437 (2013)   Cristian R. Munteanu, Alexandre L Magalhães,
	Aliuska Duardo Sánchez, Alejandro Pazos, Humberto González-Díaz   onlineTutorial Biopython:
	http://biopython.org/DIST/docs/tutorial/Tutorial.html
Complementary	

Recommendations
Subjects that it is recommended to have taken before
Subjects that are recommended to be taken simultaneously
Subjects that continue the syllabus
Other comments
Materia impartida en inglés

(\*)The teaching guide is the document in which the URV publishes the information about all its courses. It is a public document and cannot be modified. Only in exceptional cases can it be revised by the competent agent or duly revised so that it is in line with current legislation.