		Teaching	g Guide		
	Identifyi	ng Data			2020/21
Subject (*)	Genomics Code 614522006				614522006
Study programme	Mestrado Universitario en Bioinfe	ormática para Ci	iencias da Saúd	е	
		Descri	iptors		
Cycle	Period Year Type			Credits	
Official Master's Degre	e 1st four-month period First Optional 6				6
Language	Spanish				
Teaching method	Face-to-face				
Prerequisites					
Department	Bioloxía				
Coordinador	Vila Taboada, Marta		E-mail	marta.vila.taboa	da@udc.es
Lecturers	Becerra Fernandez, Manuel		E-mail	manuel.becerra	@udc.es
	Cerdan Villanueva, Maria Espera	anza		esper.cerdan@u	ıdc.es
	Vila Taboada, Marta			marta.vila.taboa	da@udc.es
	Vizoso Vázquez, Ángel José			a.vizoso@udc.e	S
Web		ı			
Contingency plan	including interactions of those genes with each other and with the person's environment. However, Genomics also with the genomes of other organisms as well as their evolution. Genomics includes knowledge procuded by Genet Molecular Biology, Biochemistry, Computer Science, Statistics, Maths, Physics and so on. Classic Genetics used to start with a mutant and then search for the gene or genes responsible for that particular phenotype. By contrast, Genomics aims at predicting the function of genes from their sequence and/or their interaction other genes. OMIC sciences (Genomics, Proteomics, Metabolomics) are top science at the moment, particularly because of bioinformatics and the new DNA-sequencing tecnologies. In case of another lockdown becaus of covid19:			edge procuded by Genetics, . nsible for that particular uence and/or their interaction w	
	1. Contents will be the same. 2. In-person instruction will chan using MS TEAMS. 3. Tutoring sessions and any oth MS TEAMS. 4. The only change in the assess	ner communication	on will take plac at all students wi	e by means of email, vide	eocalls or chat as implemented i
	5. The recommended reference list will remain the same. If needed, instructors will provide with any reading and/or course resources to the students.				

	Study programme competences / results
Code	Study programme competences / results
A8	CE8 - Understanding the basis of the information of the hereditary material, its transmission, analysis and evolution
A9	CE9 ? To understand the benefits and the problems associated with the sequencing and the use of biological sequences, as well as
	knowing the structures and techniques for their processing
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

B5	CB10 - Students should possess learning skills that allow them to continue studying in a way that will largely be self-directed or
	autonomous.
В6	CG1 -Search for and select the useful information needed to solve complex problems, driving fluently bibliographical sources for the field
В7	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
C7	CT7 ? To maintain and establish strategies for scientific updating as a criterion for professional improvement.
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 /		
		results		
Knowledge about the molecular tools used in genomics				
	AJ9			
Knowledge about structural, functional and evolutionary genomics		BJ1	CJ8	
		BJ2		
To set up experiments and analyse and interpret data using DNA microarrays		BJ6	CJ2	
		BJ7	CJ3	
Knowledge about the mechanisms involved in the evolution of genomes and the molecular and bioinformatic tools used in that		BJ5	CJ1	
kind of studies		BJ8	CJ7	

	Contents
Topic	Sub-topic
Introduction: from Molecular Genetics to Genomics	Molecular markers
	Applications ot recombinant DNA technologies
	PCR and real-time quantitative PCR
	Sanger sequencing
	DNA editing techniques
The Human Genome Project	Approaches for whole genome sequencing
Next Generation Sequencing (NGS)	Platforms
	Paired-end libraries
	Data files
Whole genome sequencing	Mate-pair libraries
	Annotation
	Comparative genomics
	Palaeogenomics
Metagenomics	Application
Clinical Genomics	Amplicon-seq
	Panel-seq
	Exome-seq
	Pharmacogenomics
Single Nucleotide Polymorphisms (SNPs)	Genome wide association studies (GWAS)
	Digital genetic testing
Functional Genomics	Transcriptome analysis: microarrays and NGS

Hands on	Introduction to the Integrative Genomics Viewer (IGV)
	Solving exercises using GALAXY
	Gene expression analysis using BABELOMICS
	Pharmacogenomic analysis using PHARMGKB

etencies / esults 5 B8 C3	Teaching hours (in-person & virtual)	Student?s personal work hours	Total hours
	· · · /		
5 B8 C3	24		
J DO 00	_ Z1	42	63
2 C1 C2 C3	2	8	10
1 B6 B7 C1	21	52.5	73.5
C7 C8			
	3.5	0	3.5
2	2 C1 C2 C3 1 B6 B7 C1 C7 C8	2 C1 C2 C3 2 1 B6 B7 C1 21 C7 C8 3.5	2 C1 C2 C3 2 8 1 B6 B7 C1 21 52.5 C7 C8

(*)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	Hands on: students solve exercises using their own laptop.
Mixed	Assessment of the learning process. Tests may include multiple choice questions, problem solving and computer exercises.
objective/subjective	Instructors will decide whether scheduling a separate test for the computer exercises depending on the progress of the group.
test	
Guest lecture /	Each instructor will explain the basic contents of each topic interacting as much as possible with the students.
keynote speech	

	Personalized attention
Methodologies	Description
ICT practicals	The instructors will carefully supervise the student's work during the hands-on sessions.
	In the event of having officially certified "part-time" students, the instructors will take the appropriate measures so that their
	scores are not affected.

		Assessment	
Methodologies	Competencies /	Description	Qualification
	Results		
Guest lecture /	A8 A9 B1 B6 B7 C1	In order to pass the subject, all students will have to take a multiple choice test and/or	70
keynote speech	C2 C7 C8	short-answer questionnaire.	
ICT practicals	B2 B5 B8 C3	All students will have to submit several reports following the guidelines provided by each instructor. In these reports, students will answer questions and/or solve exercises using their own computer and the software introduced during the computer labs.	30

Assessment comments



Students scoring at least 50 (out of 100) points but not reaching the minimum thresholds (ICT practicals: 15 out of 30 points; Objective test; 28 out of 70 points) will be awarded a 4.5 (out of 10) score. When resitting, they can choose to take both parts or only the failed one.

Mark "A with distinction" will only be awarded to outstanding students passing the subject in May.

Students will be scored as "ABSENT" (Non presentado) only when not involved in any of the assessed activities.

In the case of exceptional circumstances, lecturers may assist the student to improve his/her learning process and/or catch up on missed work/assessments. The student is responsible for liaising with his/her lecturer to organise this assistance by e.g. applying for: an extended deadline to present his/her work or taking an exam in a different date. The coordinator can request evidence about the reason for such an application.

In the event of having officially certified "part-time" students, the instructors will take the appropriate measures so that their scores are not affected.

	Sources of information
Basic	- Campbell, AM & Discovering Genomics, Proteomics & Discovering Genomics & Discovering Genom
	Cummings
	- Robison PN, Piro RM, Jäger M (2018). Computational Exome and Genome Analysis. CRC Press, Taylor & Computational Exome and Genome Analysis.
	Francis Group
	- Kulkarni S, Pfeifer J (2015). Clinical Genomics. A guide to Clinical NGS. Academic Press, Elsevier
	- Brown TA (2018). Genomes4. Garland Science, Taylor & Eroup
	- Pevsner J (2015). Bioinformatics and Functional Genomics. Wiley Blackwell
Complementary	

Recommendations
Subjects that it is recommended to have taken before
ntroduction to molecular biology/614522004
Genetics and molecular evolution/614522005
Subjects that are recommended to be taken simultaneously
Subjects that continue the syllabus
Fundamentals of bioinformatics/614522008
Other comments
Do not take this course unless your level of English is B1 or higher.

(*)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.