

		Teaching Guide			
	Identifying I	Data		2022/23	
Subject (*)	Protein Structure and Dynamics		Code	610441012s	
Study programme	Máster Universitario en Bioloxía Mol	ecular, Celular e Xenética (semipresencial)		
		Descriptors			
Cycle	Period	Year	Туре	Credits	
Official Master's Degre	ee 2nd four-month period	First	Optional	3	
Language	SpanishEnglish			, ,	
Teaching method	Hybrid				
Prerequisites					
Department	BioloxíaDepartamento profesorado r	náster			
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Web					
General description	This subject pretends to meet and m	anage the theoretical found	lations and the experime	ntal approaches to the analysis of	
	the physical and chemical of biological macromolecules, especially proteins, properties in order to relate their structures				
	with its function and biological activity. We will study the concepts needed for the description of the structures,				
	computational and experimental methods for their study and the theoretical foundations that justify them.				

	Study programme competences / results
Code	Study programme competences / results
A3	Skills of understanding the functioning of cells through the structural organization, biochemistry, gene expression and genetic variability.
A9	Skills of understanding the structure and dynamics of proteins to individual and proteomic level, as well as the techniques that are necessary to analyze them and to study their interactions with other biomolecules.
B2	Skills of decision making for the problem solving: that are able to apply theoretical knowledges and practical acquired in the formulation of biological problems and the looking for solutions.
B3	Skills of management of the information: that are able to gather and to understand relevant information and results, obtaining conclusions and to prepare reasoned reports on scientific and biotechnological questions
B4	Organization and work planning skills: that are able to manage the use of the time as well as available resources and to organize the wor in the laboratory.
C2	Ability to know and use appropriately the technical terminology of the field of knowledge of the master, in the native language and in English, as a language of international diffusion in this field
C3	Using ICT in working contexts and lifelong learning.
C8	Valuing the importance of research, innovation and technological development for the socioeconomic and cultural progress of society.

Learning outcomes			
Learning outcomes	Stud	y progra	amme
	cor	npetenc	es/
		results	
Ability to understand concepts and theories related to the dynamics of proteins in cells	AR3	BR2	CC3
	AR9		CC8
Familiarization with the bibliographic and information sources where you can get updated information	AR3	BR2	CC2
	AR9		CC3
			CC8
Know the systems for the determination of structures by x-ray diffraction	AR9	BR2	CC3
			CC8



Learn different computer programs for the representation of proteins and their use	AR3	BR2	CC3
	AR9		CC8
Learn the techniques to determine interactions between proteins and proteins with other biomolecules and ligands	AR3	BR4	CC8
	AR9		
Ability to interpret critically the data of a structure of a protein in a publication	AR3	BR3	CC2
	AR9		CC3

	Contents		
Торіс	Sub-topic		
Structural classification of proteins.	Structural domains of proteins. Classification of proteins according to its		
	three-dimensional structure. Alpha proteins. Alpha/beta protein. Protein beta.		
	Structural classes of proteins. CATH classification. SCOP classification. DALI		
	classification. SMART classification.		
Criteria for the choice of a method of purification and	Chromatographic techniques: gel filtration, ion exchange, affinity and hydrophobic		
preliminary characterization.	interaction. Purification strategies. Preliminary characterization of the protein		
	conformation: State of aggregation, compactness. Secondary structure and tertiary		
	structure indicators. Quantification of proteins.		
Experimental determination of the structure of proteins using	Crystallization techniques. Tools and strategies for diffraction data. Interpretation of		
diffraction X.	the XRD. Obtaining and refinement of the molecular model. Parameters for calculating		
	the convergence of the model. Modelling.		
Interactions between biomolecules.	Interactions of proteins for the formation of complexes with proteins and other ligands.		
	Experimental methods used to determine these interactions and their structure. The		
	double hybrid method. The split-ubiquitin method. Pull-down. GST-Pull-down. FRET.		
	EMSA trials. CHIP test. Other methodologies.		

Plannin	g		
Competencies /	Teaching hours	Student?s personal	Total hours
Results	(in-person & virtual)	work hours	
A9 C2	1	41	42
A9 B2 B3 B4 C8	1	9	10
A3 C3	1	3	4
A9	2	12	14
	5	0	5
	Competencies / Results A9 C2 A9 B2 B3 B4 C8 A3 C3	Results(in-person & virtual)A9 C21A9 B2 B3 B4 C81A3 C31	Competencies / ResultsTeaching hours (in-person & virtual)Student?s personal work hoursA9 C2141A9 B2 B3 B4 C819A3 C313

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

	Methodologies	
Methodologies	Description	
Guest lecture /	The student will be able to attend the face-to-face sessions synchronously through TEAMS. Sessions will be recorded for	
keynote speech	viewing asynchronously.	
Case study	Case study consists of simulations and works using the problem-based learning methodology.	
ICT practicals	ICT allow display of protein structure models and design interaction experiments.	
Mixed	Combination of multiple choice questions and short of relationship questions	
objective/subjective		
test		

Personalized attention	
Methodologies	Description



Case study	The personalized attention that is described in relation to these methodologies are conceived as moments of face-to-face
ICT practicals	student work with the teacher through TEAMS by involving a compulsory student participation.

		Assessment	
Methodologies	Competencies /	Description	
	Results		
Case study	A9 B2 B3 B4 C8	Students will have to answer questionnaires based on the case study.	15
Mixed objective/subjective test	A9	Test related to theoretical and practical knowledge. Students in blended mode, in addition to passing the test, must submit a series of tasks that will be requested throughout the course.	75
ICT practicals	A3 C3	Students in blended mode will carry out the practices through ICT on their own and will submit a report of the work carried out.	10

Assessment comments

To get honours preference will be given to the students evaluated at the

first opportunity in June.

For the students who request the DECEMBER ADVANCE CALL,

the current regulations will be applied, according to which the teaching guide

of the current course governs.

Implications of PLAGIARISM in the qualification: The

current regulations will be applied, according to which the fraudulent

performance of the tests or evaluation activities will directly imply the

qualification of failure.

	Sources of information
Basic	
Complementary	
	Recommendations
	Subjects that it is recommended to have taken before
Molecular Techniques/610441	02
Advanced Cellular Biology/610	441003

Subjects that are recommended to be taken simultaneously

Recombinant proteins and protein Engineering /610441013

Proteomics/610441014

Bioinformatics and Biomolecular models /610441021

Subjects that continue the syllabus

Project/610441023

Other comments

Green

Campus Faculty of Sciences ProgramTo help

achieve a sustainable immediate environment and comply with point 6 of the

"Environmental Declaration of the Faculty of Sciences (2020)", the

documentary works to be carried out in this subject will be requested in

virtual format and computer support.



(*)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.