UCLouvain WSbim2115

2019

Protein structure / Function relationships

In view of the health context linked to the spread of the coronavirus, the methods of organisation and evaluation of the learning units could be adapted in different situations; these possible new methods have been - or will be - communicated by the teachers to the students.

4 credits	30.0 h	Q1

Teacher(s)	Bertrand Luc ;Collet Jean-François ;Gatto Laurent ;Laloux Géraldine ;Rider Mark (coordinator) ;				
Language :	English				
Place of the course	Bruxelles Woluwe				
Main themes	Methods of expression, purification an in vitro renaturation of proteins (5h) Protein sequencing (2h) Bioinformatic analysis of proteins (homology searches, alignments, phylogenetic studies, motif and domain searching, structure modelling) (10h) Structure determination by NMR, crystal structures by X-ray diffraction in relation to function (3h) Enzymology (thermodynamics, pre- and steady state kinetics, calculation of kinetic parameters, ligand binding and allosteric enzymes, site-directed mutagenesis, theory of metabolic control) (10 h)				
Aims	To provide Masters students in Biomedical Sciences with the necessary competence to study: techniques of overexpression/purification and structural analysis of proteins structure-function relationships in proteins physiological roles of enzymes The contribution of this Teaching Unit to the development and command of the skills and learning outcomes of the programme(s) can be accessed at the end of this sheet, in the section entitled "Programmes/courses offering this Teaching Unit".				
Evaluation methods	Due to the COVID-19 crisis, the information in this section is particularly likely to change. Written exam of 3 h duration and 6 questions with at least one question from each lecturer. The exam questions will be in English but the students are free to answer either in French or in English. There are no negative points or ponderation according to the questions and course content. However, when a student has a mark between 9/20 et 10/20 after correction, the lecturers re-examine all the points to decide if the mark should be rounded up to 10 or stay at 9. If the answers are insufficient, the final mark will be rounded down.				
Teaching methods	Due to the COVID-19 crisis, the information in this section is particularly likely to change. Formal lectures plus a "workshop" on bioinformatic analysis.				
Content	The aim is to teach to masters students in biomedical sciences (SBIM Master) techniques of enzymology and metabolic control analysis, purification, sequencing, structure determination and bioinformatic analysis of proteins. Content: Expression, purification and renaturation of proteins <i>in vitro</i> (6h) Protein sequencing (3h) Bioinformatic analysis of proteins (3h) Techniques for determining the 3D structures of proteins (3h) Analysis of protein-protein interactions (3h) Kinetics and mechanisms of enzymes, cooperativity, structure-fuction relationships, epresentation of protein structures, metabolic control analysis (12h)				
Inline resources	There is no formal syllabus ! PDF versions of slides presented in the course, which cover the subject in a comprehensive way, will be made available on MoodleUCL (https://moodleucl.uclouvain.be/). In addition, a tablet will be used to explain certain aspects of the course. The "Tablet" PDF versions of the PowerPoint files will also be made available to students via MoodleUCL.				
Bibliography	Les étudiants sont encouragés à lire des revues dans la littérature ! The students are encouraged to do background reading, for example by consulting numerous comprehensive reviews on the subject matter of the course available in the literature (for example via Pub Med).				
Other infos	Teaching of the course material will be in English and the Power Point files will be mostly in English.				

Faculty or entity in	SBIM
charge	

Programmes containing this learning unit (UE)							
Program title	Acronym	Credits	Prerequisite	Aims			
Master [60] in Biomedicine	SBIM2M1	3		٩			
Master [120] in Statistic: Biostatistics	BSTA2M	4		۹			
Master [120] in Biomedicine	SBIM2M	4		۹			
Master [180] in Medecine	MD2M	4		٩			