## UCL Université de Louvain UNIVERSITÉ VFARM2501 2012-2013 Chimie pharmaceutique avancée et drug design

3.0 credits

20.0 h + 10.0 h

Teacher(s) :	Labar Geoffray (compensates Lambert Didier) ; Lambert Didier ; Poupaert Jacques ; Muccioli Giulio (coordinator) ;
Language :	Français
Place of the course	Bruxelles Woluwe
Prerequisites :	organic chemistry and medicinal chemistry
Main themes :	The teacher(s) will discuss first the different techniques allowing the discovery of novel therapeutic targets (lipidomic ' proteomic ' deorphanization), second the methods allowing the identification of hits for a given molecular target (high-throughput screening ; computer assisted de novo drug design); finally the strategies allowing to optimize a hit ('hit to lead') will be discussed (structures-activity relationships ; docking; ')
Aims :	At the end of the activity the student will be able to
	 Interpret, based on what was discussed in class, the results presented in a scientific paper dealing with the development of a novel drug
	 Suggest a strategy allowing to identify novel therapeutic targets
	 Suggest a strategy allowing to identify novel lead compounds for a given target (enzyme, receptor, ')
	Suggest a strategy allowing to optimize the activity of a drug towards its target 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 :	a written exam and a presentation of a case-study based on a scientific paper
Content :	 Lipidomics applied to the discovery of novel targets
	Exploring the proteome looking for novel targets : the enzymes ' the receptors
	Proteins as therapeutic target
	Determination of their 3D structure
	Role of protein's 3D models in drug design
	The interest of establishing structure-activity relationships in drug design
Cycle and year of study :	> Master [120] in Pharmacy
Faculty or entity in charge:	FARM