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"Tradition and Quality"

Detailed Course Description - Course Plan Development and Updating Procedures/ Pharmacy Department				QF02/0408-3.0E
Faculty Pharmacy Department Pharmacy				
Course number	201742	Course title	Advar Chem Desig	nced Medicinal istry and Drug n
Number of credit hours	3	Pre-requisite/co- requisite		

## **Brief course description**

This course is designed to impart the knowledge in computational methods and drug design approaches. It aims to build students' knowledge in theoretical chemistry and its application in drug design. It is proposed to provide students with an understanding of hit discovery, lead identification, lead optimization, target selection, and molecular recognition employing computer-aided drug design software. And, it will shed the light on computer-based methods, combinatorial chemistry, highthroughput screening, and database mining.

	Course goals and learning outcomes			
Goal 1	To distinguish between bioinformatics and cheminformatics methods and their recruitment in drug design.			
	1.1 Recognize theoretical strategies and their classifications.			
	1.2 Describe the drug design pipeline and understand where computational			
	chemistry fits in.			
	1.3 Discuss informatics approaches to the prediction of chemical properties.			
	1.4 Understand the importance of drug-like properties and their prediction.			
	1.5 Describe the use of lead candidates and database representations.			
	1.6 Understand the use of classifier algorithms and quantum/classical descriptors.			
Learning	1.7 Describe relations between thermodynamic properties and protein-ligand binding			
outcomes	and structure.			
	1.8 Describe protein-ligand docking and the empirical/knowledge-based scoring			
	functions.			
	1.9 Discuss empirical scoring, de-novo design and virtual screening.			
	1.10 Describe simulations of ligand binding thermodynamics.			
	1.11 Appreciate protein sequence searches, homology and loop modelling, protein-			
	protein docking, and describe biologics design.			
	1.12 Describe the relation between Re <sub>50</sub> and Rd, and discuss orophysical methods. 1.13 Know how to use software such as MOE, MAESTRO, AMBER, and PYMOL.			
	1. The Organic Chemistry of Drug Design and Drug Action, 2nd edition,			
	Richard B. Silverman, Elsevier, 2004.			
Textbook	<ol> <li>Foye's Principles of Medicinal Chemistry, 6th edition, Thomas L. Lemke and David A. Williams, Lippincott Williams &amp; Wilkins, 2008.</li> </ol>			



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Detailed Cours	QF02/0408-3.0E	
	<ol> <li>Wilson and Gisvold's Textbook of Organic Medicinal a Chemistry, 12<sup>th</sup> edition, J. N. Delgado and W. A. Remer 2011.</li> </ol>	nd Pharmaceutical s, Lippincott-Raven,
	<ol> <li>Burger's Medicinal Chemistry and Drug Discovery, 6th 2003.</li> </ol>	edition, M. E. Wolff,
	<ol> <li>An Introduction of Medicinal Chemistry, 4<sup>th</sup> edition, Gra University Press, 2008.</li> </ol>	aham Patrick, Oxford
Supplementary references	<ul><li>2- The Organic Chemistry of Drug Synthesis, Vol. 1-6, D. Mitscher, John Wiley and Sons.</li></ul>	Lednicer and L. A.
	3- Computational Chemistry and Drug design Journals.	

Course timeline				
Week	Number of hours	Course topics	Pages (textbook)	Notes
01	1 1 1	<ul> <li>Computational Methods</li> <li>Potential energy.</li> <li>Molecular mechanics.</li> <li>Quantum Mechanics.</li> <li>Geometry Optimization.</li> <li>First Order Minimization.</li> <li>Second Order Minimization.</li> </ul>	Textbooks 1-3/ Drug Design Part	
02	1 1 1	<ul> <li>Conformational Analysis.</li> <li>Molecular Dynamic (MD) Simulation.</li> <li>Monte Carlo Method.</li> <li>Metropolis Method.</li> <li>NPT Model</li> <li>NVT Model.</li> </ul>	Textbooks 1-3/ Drug Design Part	
03-04	2 2 2	<ul> <li>X-ray crystallography</li> <li>Homology Modeling</li> <li>Bioinformatics.</li> <li>Structure-Based Drug Design</li> <li>Molecular Docking</li> <li>Binding Free Energy.</li> </ul>	Textbooks 1-3/ Drug Design Part	



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	Detail	s/ QF02/0	0408-3.0E		
	05-06	2 2 2	<ul> <li>Ligand-Based Drug Design.</li> <li>Cheminformatics.</li> <li>Pharmacophore Modeling</li> <li>Pharmacophore Searching</li> <li>Druggability</li> <li>Lipinski's Rule of Five</li> <li>Database Mining</li> <li>Virtual Screening</li> </ul>	Textbooks 1-3/ Drug Design Part	
	07-08	2 2 2	<ul> <li>Ligand-Based Drug Design</li> <li>Quantitative Structure-Activity Relationship</li> <li>Equations and Graphs.</li> <li>Physicochemical Properties.</li> <li>Hydrophobicity.</li> <li>Electronic Property.</li> <li>Steric Factor.</li> <li>Craig Plot.</li> <li>Topliss Scheme.</li> <li>Hansch Equation.</li> <li>Topliss operational schemes.</li> <li>QSAR: 3D-QSAR (CoMFA).</li> </ul>	Textbooks 1-3/ Drug Design Part	
	09	1 1 1	<ul> <li>Combinatorial Chemistry.</li> <li>Parallel Synthesis.</li> <li>Solid Phase Technique.</li> <li>High-Throughput Screening.</li> </ul>	Textbooks 1-3/ Drug Design Part	
10 1 - Case Study I 1		- Case Study I	Textbooks 1-3/ Drug Design Part		
	11	1 1 1	- Case Study II	Textbooks 1-3/ Drug Design Part	
	12	1 1 1	Article Presentation and Discussion.	Computational Chemistry and Drug design Journals.	
	13	1 1 1	Molecular Modeling Practical	Modeling Lab Applications	



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Detailed Course Desc	QF02/0408-3.0E				
Theoretical course	Participation = 10%	Practical (clinical)	Semester students		
evaluation methods	First exam 20%	course evaluation	work = 50%		
and weight	Second exam 20%	methods	(Reports, research,		
	Final exam 50%		quizzes, etc.)		
			Final exam = 50%		

Approved by head of department	Date of approval	

Extra information (to be updated every semester by corresponding faculty member)

Name of teacher	Dima A. Sabbah, Ph.D.	Office Number	227
Phone number (extension)	31	Email	dima.sabbah@zuj.edu.jo
	Sun-Thurs (4-5 pm)		
Office hours			