

**ADVANCED TECHNIQUES IN STRUCTURAL DETERMINATION  
2016-2017**

Master Degree:	CHEMISTRY AND BIOTECHNOLOGY	760M
Course title:	ADVANCED TECHNIQUES IN STRUCTURAL DETERMINATION	5119
Year/Semester:	1st/2nd	ECTS Credits: 5

**DEPARTMENT**

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**CONTENTS**

The course is divided in three parts oriented to the structural determination of organic, inorganic and organometallic compounds using X-ray diffraction methods (solid state), nuclear magnetic resonance techniques (solution state) and computational methods, especially molecular dynamics. These complementary techniques help to the scientist to have a vision close to the reality about how the atoms of the molecules are arranged in 3D. This information is critical given the close relationship between structure and activity (SAR).

**PART A. DIFFRACTION METHODS**

**UNIT1.** Data collection: geometric data collection, data collection based on intensity (space group determination)

**UNIT2.** Data processing: Lorentz and polarization theories, data reduction and absorption correction, structure factor, Friedel's law, Fourier synthesis

**UNIT3.** Structural resolution: The phase problem, direct methods, Patterson methods, heavy atom methods

**UNIT4.** Refinement and results: Software (Wingx, SHELX, DIRDIF, SIR, PLATON, ORTEP, MERCURY, ENCIFER...), crystallographic files, data bases, validation of results

**PART B. NUCLEAR MAGNETIC RESONANCE**

**UNIT1.** Pulsed nuclear magnetic resonance

**UNIT2.** Chemical shift and coupling constants

**UNIT3.** Relaxation and NOE effect

**UNIT4.** Carbon-13 nuclear magnetic resonance and assignment techniques (Decoupling, attached proton test and DEPT)

**UNIT5.** 2D-Nuclear magnetic resonance: Proton-carbon heteronuclear correlations (HMBC, HSQC, HMQC), proton-proton homonuclear correlations (COSY, TOCSY, NOESY)

**UNIT6.** Dynamic nuclear magnetic resonance: concept, calculation of equilibrium constants, study of ligand-protein interactions using saturation transfer (STD-experiments, tr-NOESY experiments, Diffusion (DOSY).

**PART C. COMPUTATIONAL CALCULATIONS. MOLECULAR DYNAMICS, QM/MM**

**UNIT1.** General aspects of the molecular dynamic (MD) and molecular mechanical (MM)

**UNIT2.** MM and MD applications to the conformational analysis of interesting biological entities

## REFERENCES

### Title

Basic one- and two-dimensional NMR Spectroscopy

NMR spectroscopy and computer modeling of carbohydrates

The art of Molecular Dynamics Simulations

Introduction to Practice of Molecular Simulation

X-ray structure determination. A practical guide

Fundamentals of crystallography

Internet sources:

NMR on-line: <https://qshare.queensu.ca/Users01/sauriof/www/webcourse/index.htm>

Molecular dynamics tutorials (AMBER): <http://ambermd.org/tutorials/>

GLYCAM web: <http://glycam.org/>

International Union of Crystallography: <http://www.iucr.org/>

## EVALUATION SYSTEM

Tasks before final exam referred to resolution of structures from X-ray data (30%, unrecoverable)

Final exam (60%, recoverable)

Observation techniques (10%, unrecoverable)