


Due to the COVID-19 crisis, the information below is subject to change, in particular that concerning the teaching mode (presential, distance or in a comodal or hybrid format).

3 credits	22.5 h + 7.5 h	Q1
-----------	----------------	----

Teacher(s)	Fustin Charles-André ;
Language :	English
Place of the course	Louvain-la-Neuve
Main themes	1. Basic definitions 2. Ionization modes 3. Analyzers 4. Chromatographic couplings 5. Spectral data interpretation 6. Introduction to the identification and sequencing of proteins and peptides by mass spectrometry.
Aims	<p>1 This course covers technical aspects of mass spectrometry and interpretation of spectral data.</p> <p>-----</p> <p><i>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".</i></p>
Evaluation methods	<p>Due to the COVID-19 crisis, the information in this section is particularly likely to change.</p> <p>The participation to all the activities of the course is mandatory to validate the course and be allowed to present the exam. The exam consist in a personnel work based on the scientific literature to be presented to all the other students following the course.</p>
Content	<p>After a brief survey of the basic definitions, the various mode of ionization (EI, CI, FAB, ESI, APCI, APPI, DESI, DAPCI, DAPPI) will be described in details. The different analyzers (Quad, triple Quad, Traps, TOF, Orbitrap, FTICR) will be presented together with their possible scanning modes and their combinations. Couplings with Gc and HPLC will be presented. The interpretation of spectral data will first emphasize the differences between low resolution and low accuracy versus high resolution and high accuracy data. The importance of the isotopic cluster will be demonstrated. The principal rules of fragmentation of radical-cations will be presented together with some basic rules for the fragmentation of ions with even-number of electrons. Selected examples and exercises will be explicitly studied in this part of the course. At the end of the course, several examples of the utility of mass spectrometry for different domains will be seen through the presentations of the students.</p>
Inline resources	A copy of the slides has been deposited on Moodle
Other infos	Prerequisite: - Basic knowledge of chemistry and physics - CHM1251C course.
Faculty or entity in charge	CHIM

Programmes containing this learning unit (UE)				
Program title	Acronym	Credits	Prerequisite	Aims
Master [120] in Chemistry	CHIM2M	3		
Master [60] in Chemistry	CHIM2M1	3		