

**Subject card**

<b>Subject name and code</b>	Mass spectrometry in proteomics , PG_00153633						
<b>Field of study</b>	Biotechnology						
<b>Date of commencement of studies</b>	October 2026	<b>Academic year of realisation of subject</b>			2026/2027		
<b>Education level</b>	Master's studies	<b>Subject group</b>			Obligatory subject group in the field of study Optional subject group Subject group related to scientific research in the field of study		
<b>Mode of study</b>	full-time studies	<b>Mode of delivery</b>			at the university		
<b>Year of study</b>	1	<b>Language of instruction</b>			English		
<b>Semester of study</b>	2	<b>ECTS credits</b>			2.0		
<b>Learning profile</b>	academic	<b>Assessment form</b>			credit		
<b>Conducting unit</b>	Intercollegiate Faculty of Biotechnology UG-MUG -> Rector						
<b>Name and surname of lecturer (lecturers)</b>	<b>Subject supervisor</b>		dr hab. Paulina Czaplewska				
	<b>Teachers</b>						
<b>Lesson types</b>	<b>Lesson type</b>	Lecture	Tutorial	Laboratory	Project	Seminar	SUM
	<b>Number of study hours</b>	10.0	0.0	20.0	0.0	0.0	30
	E-learning hours included: 0.0						
<b>Learning activity and number of study hours</b>	<b>Learning activity</b>	Participation in didactic classes included in study plan		Participation in consultation hours		Self-study	SUM
	<b>Number of study hours</b>	30		5.0		15.0	50
<b>Subject objectives</b>	<ul style="list-style-type: none"> <li>learning and acquiring basic concepts and terminology used in mass spectrometry (K_W01, K_W04)</li> <li>acquiring the knowledge and skills necessary to independently prepare a sample and conduct proteomic analysis of peptides and proteins (K_W01, K_W04, K_U01, K_U05, K_K05)</li> <li>familiarization with methods of identifying peptides and proteins based on ESI and MALDI mass spectra (K_W05, K_U05, K_K05)</li> <li>familiarization with the analysis of fragmentation spectra of peptides (K_W01, K_U05, K_K01)</li> </ul>						

Learning outcomes	Course outcome	Subject outcome	Method of verification
	[BIOTECHMU2_U01] The graduate possesses the skills necessary to design and conduct laboratory research, critically assessing risks, method limitations, and ethical implications of undertaken activities.	Is aware of the importance of occupational safety rules, especially work in the laboratory; applies work safety rules; is responsible for the safety of himself and others; is able to deal with threats (K_K05).	[SU8] observation of student's independent or team work
	[BIOTECHMU2_W05] The graduate possesses in-depth knowledge of methods used in the field of exact and natural sciences, necessary to understand biological phenomena and processes at the molecular level, and the connection of this knowledge with medical sciences.	Understands complex biological phenomena at the molecular level, knows their importance for biotechnology and connections with other fields and disciplines of science (K_W01) Knows the basic principles of occupational health and safety; understands the dangers of working in the laboratory; knows the risks associated with conducting laboratory tests; knows the risks of working with pathogenic organisms and GMOs (K_W04)	[SW1] oral statement/ conversation/discussion
[BIOTECHMU2_U04] The graduate possesses the ability to proficiently use scientific information, including English, regarding biotechnology; critically analyses and selects information; uses electronic sources; has the ability to use appropriate databases.	Has the skills necessary for laboratory work; is able to plan and conduct an experiment; independently documents activities and results; in laboratory work, under the supervision of a supervisor, uses complex research techniques and tools; has the ability to operate laboratory equipment (K_U01) Collects and interprets empirical data; uses statistical methods and IT tools to analyze data; formulates conclusions based on empirical data (K_U05).	[SU2] presentation/project/paper/ report	
Subject contents	<ul style="list-style-type: none"> <li>• Introduction to mass spectrometry, physical basis of MS spectrum measurement, equipment and basics of recording MS spectra.</li> <li>• Qualitative and quantitative analysis in proteomic analysis using mass spectrometry</li> <li>• Sample preparation methods for MS analysis, registration and analysis of peptide and protein spectra (ESI, MALDI)</li> <li>• Analysis of post-translational modifications in MS</li> <li>• Protein depletion before MS analysis, solution digestion and in-gel digestion</li> <li>• Analysis of MS data using MS spectra and protein databases.</li> </ul>		
Prerequisites and co-requisites	No formal requirements. Entrance requirements: It is required to obtain knowledge, skills and competences specified for the following courses: Biochemistry (lecture), Organic Chemistry (lecture).		
Assessment methods and criteria	Subject passing criteria	Passing threshold	Percentage of the final grade
	Final project	51.0%	75.0%
	Report on laboratory classes	51.0%	25.0%
Recommended reading	Basic literature	<a href="#">Johnstone Robert A.W. I Malcolm E.Rose</a> , Spektrometria mas, PWN 2001  <a href="#">De Hoffmann, Edmond, Charette, Jean Joseph, Strobant, Vincent</a> , Spektrometria mas, Wydawnictwa Naukowo-Techniczne 1998	

	Supplementary literature	<p>K. Macur, J. GrzenkiewiczWydra, L. Konieczna, J. Bigda, C. Temporini, S. Tengattini, T. Bączek. A (2018) ProteomicBased Approach to Study the Mechanism of Cytotoxicity Induced by Interleukin1 and Cycloheximide <b>Chromatographia</b> 81:4756 (<a href="https://doi.org/10.1007/s10337-017-3382-3">doi.org/10.1007/s10337-017-3382-3</a>)</p> <p>M. Rafalik, M. Spodzieja, A.S. Kołodziejczyk, S. Rodziewicz-Motowidło, A. Szymańska, A. Grubb, P. Czaplewska (2018) The identification of discontinuous epitope in the human cystatin c monoclonal antibody hCC3 complex. <b>Journal of Proteomics</b> (<a href="https://doi.org/10.1016/j.jprot.2018.04.020">doi.org/10.1016/j.jprot.2018.04.020</a>)</p> <p>M Prądzińska, I Behrendt, M Spodzieja, AS. Kołodziejczyk, S Rodziewicz-Motowidło, A Szymańska, SL. Lundström, RA. Zubarev, K Macur, P Czaplewska (2016). <i>Isolation and characterization of autoantibodies against human cystatin C</i>. <b>Amino Acids</b> (DOI 10.1007/s00726-016-2271-7)</p> <p>AE. Lewandowska, K. Macur, P. Czaplewska, J. Liss , K. Łukaszuk, S. Ołdziej. (2017) <a href="#">Qualitative and Quantitative Analysis of Proteome and Peptidome of Human Follicular Fluid Using Multiple Samples from Single Donor with LC-MS and SWATH Methodology</a>. <b>J Proteome Research</b> (DOI: 10.1021/acs.jproteome.7b00366)</p> <p>Primers for Proteomics <a href="https://doi.org/10.1142/13595">https://doi.org/10.1142/13595</a>   May 2024 Pages: 250 Edited by: <a href="#">Paulina Czaplewska</a> (University of Gdańsk, Poland &amp; Medical University of Gdańsk, Poland), <a href="#">Katarzyna Macur</a> ( University of Gdańsk, Poland &amp; Medical University of Gdańsk, Poland ), and <a href="#">Paweł Ciborowski</a> (University of Nebraska Medical Center, USA)</p>
	eResources addresses	
Example issues/ example questions/ tasks being completed		
Work placement	Not applicable	

Document generated electronically. Does not require a seal or signature.