

Subject card

Subject name and code	Marine OMICS -lecture, PG_00099385						
Field of study	Marine Biotechnology						
Date of commencement of studies	October 2025		Academic year of realisation of subject			2026/2027	
Education level	Master's studies		Subject group			Obligatory subject group in the field of study Subject group related to scientific research in the field of study	
Mode of study	full-time studies		Mode of delivery			at the university	
Year of study	2		Language of instruction			English	
Semester of study	3		ECTS credits			2.0	
Learning profile	academic		Assessment form			exam	
Conducting unit	Intercollegiate Faculty of Biotechnology UG-MUG -> Rector						
Name and surname of lecturer (lecturers)	Subject supervisor		dr hab. Paulina Czaplewska				
	Teachers						
Lesson types	Lesson type	Lecture	Tutorial	Laboratory	Project	Seminar	SUM
	Number of study hours	28.0	0.0	0.0	0.0	0.0	28
	E-learning hours included: 0.0						
Learning activity and number of study hours	Learning activity	Participation in didactic classes included in study plan		Participation in consultation hours		Self-study	SUM
	Number of study hours	28		5.0		20.0	53
Subject objectives	The aim of the course is to familiarize the student with the basics of metabolomics, genomics, transcriptomics and proteomics as techniques used in broadly understood research related to the sea and its ecosystem. Additionally, as part of the course, students will learn next-generation sequencing (NGS) techniques and the analysis of microbiomes in marine environments.						

Learning outcomes	Course outcome	Subject outcome	Method of verification
	[MBMU2-KW03] Knows and understands complex biological phenomena at the molecular level, understands their significance for an organism, marine environment and marine biotechnology	KW_01: Understands complex biological phenomena at the molecular level, knows their importance for biotechnology and their connections with other fields and disciplines of science K_W04: Knows the basic principles of occupational safety, understands the risks associated with laboratory work, knows the risks associated with conducting laboratory research, knows the risks associated with working with pathogenic organisms and GMOs	[SW4] test/exam - oral or written
	[MBMU2-KW01] Knows and broadly understands the value of natural marine resources	KU_01: Has the skills necessary for laboratory work; is able to plan and carry out an experiment, is able to document his own operations and results; in laboratory work, under the supervision of the instructor, uses complex techniques and research tools, is able to use laboratory equipment. Collects and interprets empirical data, uses statistical methods and IT tools in data analysis, draws conclusions based on empirical data (K_U05)	[SW4] test/exam - oral or written
Subject contents	Organization and genetic content of prokaryotic (bacteria, archaea) and eukaryotic (yeast, humans, plants) genomes. The Human Genome Project. The importance of mobile genetic elements for the organization and size of genomes. Genome and mitochondrial genomeplastids. Viral genomes. Comparative genomics. The impact of genomics on medicine and society. Next generation sequencing (NGS) techniques: Collecting environmental samples from the seas, securing samples, isolating genetic material, preparing a library for high-throughput sequencing, conducting the sequencing process. Metagenomic data analysis: Manipulation and analysis of collected metagenomic data, interpretation of results regarding microbiomes of marine environments. The use of ancient DNA in genomics. Evolution of genomes. Introduction to mass spectrometry, physical basis for measuring MS spectra, equipment and basics of recording MS spectra. Qualitative and quantitative analysis in proteomic analysis using mass spectrometry Methods of sample preparation for MS analysis, registration and analysis of peptide and protein spectra (ESI, MALDI) Analysis of post-translational modifications in MS Protein loss before MS analysis, digestion in solution and digestion in gel Analysis of MS data using MS spectra and protein databases. Protein digestion, recording of MS/MS spectra, data analysis. The influence of various factors on changes in the metabolite profile.		
Prerequisites and co-requisites	Formal requirements - no formal requirements It is required to obtain knowledge, skills and competences for specific subjects: Biochemistry (lecture), Organic chemistry (lecture), Biodiversity and basics of taxonomy, Bioinformatic sequence analysis, Molecular biology and genetics After passing the compulsory subjects in the first three semesters, the student has the knowledge and skills that qualify him to participate and pass the course		
Assessment methods and criteria	Subject passing criteria	Passing threshold	Percentage of the final grade
	Exam	51.0%	100.0%
Recommended reading	Basic literature	Scientific publications and studies prepared by the teacher and made available to students during classes. Genomes 3 T.A. Brown, 2007, Garland Science Brown T.A. "Genomes", ed. II, translation edited by P. Węgleński, Wydawnictwo Naukowe PWN, Warszawa 2009. Molecular Biology of the Gene, 7th edition, 2014, Pearson Johnstone Robert A.W. I Malcolm E. Rose, Mass spectrometry, PWN 2001 De Hoffmann, Edmond, Charette, Jean Joseph, Stroobant, Vincent, Mass Spectrometry, Wydawnictwa Naukowo-Techniczne 1998 Materials provided by the teacher	

	Supplementary literature	Primers for Proteomics https://doi.org/10.1142/13595 May 2024 Pages: 250 Edited by: Paulina Czaplewska (University of Gdańsk, Poland & Medical University of Gdańsk, Poland), Katarzyna Macur (University of Gdańsk, Poland & Medical University of Gdańsk, Poland), and Paweł Ciborowski (University of Nebraska Medical Center, USA) Metagenomics: Techniques, Applications, Challenges and Opportunities; Reena Singh Chopra, Chirag Chopra, Neeta Raj Sharma; 2020, Springer https://doi.org/10.1007/978-981-15-6529-8
	eResources addresses	
Example issues/ example questions/ tasks being completed		
Work placement	Not applicable	

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