

Subject card

Subject name and code	Bioinformatics - applications , PG_00153620						
Field of study	Biotechnology						
Date of commencement of studies	October 2026	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	1	Language of instruction			Polish		
Semester of study	2	ECTS credits			2.0		
Learning profile	academic	Assessment form			credit		
Conducting unit	Intercollegiate Faculty of Biotechnology UG-MUG -> Rector						
Name and surname of lecturer (lecturers)	Subject supervisor		dr hab. Małgorzata Waleron				
	Teachers						
Lesson types	Lesson type	Lecture	Tutorial	Laboratory	Project	Seminar	SUM
	Number of study hours	15.0	0.0	0.0	0.0	0.0	15
	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	15	5.0	30.0	50		
Subject objectives	<p>Getting students acquainted with the basic knowledge in the area of bioinformatics that allows students to consciously use bioinformatics methods and tools applied in biotechnology and other fields of science.</p> <p>Students will widen their knowledge in the related areas and disciplines of science that will allow them to see relationships and dependencies in nature, in particular those essential for biotechnology.</p>						
Learning outcomes	Course outcome		Subject outcome		Method of verification		
	[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.		The student has knowledge in the field of mathematical methods, including statistical ones, applied in biotechnology.		[SW4] test/exam - oral or written		
	[BIOTECHMU2_W02] The graduate has in-depth knowledge of the application of laboratory techniques and methods of genetic modification of cells and organisms and their use in biotechnology.		The student has an extended knowledge of the performance and interpretation of bioinformatics analyses and their use in biotechnology.		[SW4] test/exam - oral or written		

Subject contents	<ol style="list-style-type: none"> 1. Review of sequential databases. Collection, storage and preparation of sequences to be deposited in public databases. 2. Sequence comparison (comparison of pairs of sequences, comparison of many sequences). 3. Search for similar sequences in databases. 4. Chimeric sequences. OTU (operational taxonomic units) determination, Metagenomics 5. Genomics and projects of genome sequencing 6. Genome assembly and annotation 7. Comparative genomics. Phylogenetic profiles and regions of genome plasticity. 8. Pangenomics. Pangenome, core genome and dispensable genes discrimination. 9. Overall Genome Related Index (<i>OGRI</i>) <i>methods</i>: ANI/AAI, DNA-DNA hybridisation in silico. 10. Phylogenetics . Evolution models. Phylogenetic prediction 11. Searching for the gene clusters encoding secondary metabolites 12. Searching for plasmids, prophages, mobile genetic elements in NGS data 		
Prerequisites and co-requisites	<p>Unicellular organisms - Structure, diversity and environment Fundaments (M03_B1)</p> <p>Unicellular organisms - Genetics Methodology (M03_B2)</p> <p>Unicellular organisms - Genetics Metabolism (M03_B3)</p>		
Assessment methods and criteria	Subject passing criteria	Passing threshold	Percentage of the final grade
	Graded test with open questions	51.0%	100.0%
Recommended reading	Basic literature	<ul style="list-style-type: none"> • Applied Bioinformatics : An Introduction, Paperback by Selzer, Paul M. (EDT) Springer 2018 • Bioinformatics: A Practical Guide to the Analysis of Genes and Proteins (Methods of Biochemical Analysis) • Bioinformatics and Molecular Evolution. Paul G. Higgs, Teresa K. Attwood. 	
	Supplementary literature	<ul style="list-style-type: none"> • Bioinformatics. Sequence and genome analysis". D.W. Mount. 2001. • Students individually search for materials concerning classes using electronic resources 	
	eResources addresses		
Example issues/ example questions/ tasks being completed			
Work placement	Not applicable		

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