

Subject card

Subject name and code	Introduction to bioinformatics, PG_00148916						
Field of study	Medical Biology						
Date of commencement of studies	October 2024	Academic year of realisation of subject			2026/2027		
Education level	undergraduate studies	Subject group			Obligatory subject group in the field of study		
Mode of study	full-time studies	Mode of delivery			at the university		
Year of study	3	Language of instruction			Polish		
Semester of study	5	ECTS credits			2.0		
Learning profile	academic	Assessment form					
Conducting unit	Faculty of Biology						
Name and surname of lecturer (lecturers)	Subject supervisor		dr Beata Guzow-Krzemińska				
	Teachers						
Lesson types	Lesson type	Lecture	Tutorial	Laboratory	Project	Seminar	SUM
	Number of study hours	0.0	0.0	30.0	0.0	0.0	30
	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	30		4.0		16.0	50
Subject objectives	<p>To introduce students to basic bioinformatics tools and methods of sequence analysis.</p> <p>To train students to use modern bioinformatics tools.</p> <p>To obtain skills in acquiring biological data from databases and analyzing these data.</p>						
Learning outcomes	Course outcome	Subject outcome			Method of verification		
	[BIOLMEDL3_K03] is aware of his/her own limitations and knows when to seek expert assistance	is aware of his/her own limitations and knows when to seek expert assistance			[SK2] presentation/project/paper/report [SK8] observation of student's independent or team work		
	[BIOLMEDL3_U14] is able to prioritize and organize the work of a small team and work effectively as part of a team	is able to prioritize and organize the work of a small team and work effectively as part of a team			[SU2] presentation/project/paper/report [SU8] observation of student's independent or team work		
	[BIOLMEDL3_W14] describes the principles of using computer tools to analyze data and interpretation of biological phenomena and processes	knows and describes the principles of using bioinformatics tools for data analysis and interpretation of biological phenomena and processes			[SW4] test/exam - oral or written [SW2] presentation/project/paper/report		
	[BIOLMEDL3_U04] applies basic statistical methods and computer algorithms and techniques to describe phenomena and data analysis	applies basic bioinformatics tools to analyze biological data			[SU2] presentation/project/paper/report [SU4] test/exam - oral or written [SU6] demonstration of practical skills		
	[BIOLMEDL3_U08] can interpret scientific data related to the profession of medical biologist	can analyze sequences of biological molecules and can interpret the results of simple bioinformatics analyses			[SU2] presentation/project/paper/report [SU4] test/exam - oral or written [SU6] demonstration of practical skills		
	[BIOLMEDL3_W13] presents the basic methods of statistical analysis and understands their importance in the interpretation of biological phenomena and processes	Has knowledge of basic techniques and bioinformatics tools and understands their importance in molecular data analysis			[SW4] test/exam - oral or written [SW2] presentation/project/paper/report		

Subject contents	Topics covered include the following: health and safety rules and ergonomics of working at a computer; introduction to databases; acquisition and processing of biological information using biological and biomedical databases - nucleotide and amino acid sequences, structures of biological molecules, literature; introduction to methods of sequence analysis; overview of basic bioinformatics tools: editing and analysis of chromatograms, sequence comparison, primer design, sequence alignment, search and visualization of tertiary structures of biological molecules, in silico cloning.		
Prerequisites and co-requisites	Completed course: Molecular basis of medical biology Good computer skills, including MS Office. At least basic knowledge of molecular biology, types of biological macromolecules and basic mechanisms at the molecular level. Good English language skills enabling one to work with software and databases.		
Assessment methods and criteria	Subject passing criteria	Passing threshold	Percentage of the final grade
	Preparation of group project and presentation of results	51.0%	40.0%
	test	51.0%	50.0%
	class work	51.0%	10.0%
Recommended reading	Basic literature	Lesk A. (2019) Wprowadzenie do bioinformatyki. Wydawnictwo Naukowe PWN, ISBN 9788301208110 Baxevanis A.D., Ouellette B.F. (red.) (2005) Bioinformatyka - podręcznik do analizy genów i białek. PWN, ISBN 83-01-14211-1 Xiong J. (2011) Podstawy bioinformatyki, PWN, ISBN: 9788323505112	
	Supplementary literature	Paul G. Higgs, Teresa K. Attwood (2008) Bioinformatyka i ewolucja molekularna. PWN, ISBN: 978-83-01-15494-3 https://www.ncbi.nlm.nih.gov/books/NBK1762/ https://www.ncbi.nlm.nih.gov/books/NBK143764/ Marta A. Skowron, Beata Guzow-Krzemińska, Sylwia Barańska, Paulina Jędrak, Grzegorz Węgrzyn. 2015. A rapidly progressing, deadly disease of <i>Actias selene</i> (Indian moon moth) larvae associated with a mixed bacterial and baculoviral infection. <i>Journal of Biosciences</i> 40(3):487-495.	
	eResources addresses	Adresy na platformie eNauczanie:	
Example issues/ example questions/ tasks being completed			
Work placement	Not applicable		

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