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Subject name and code	High-throughput methods in biotechnology, PG_00069250								
Field of study	Biotechnology								
Date of commencement of studies	February 2025		Academic year of realisation of subject			2025/2026			
Education level	second-cycle studies		Subject group						
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		3.0				
Learning profile	general academic profile		Assessment form		assessment				
Conducting unit	Department of Chemistry Technology and Biotechnology of Food -> Faculty of Chemistry -> Wydziały Politechniki Gdańskiej								
Name and surname	Subject supervisor		prof. dr hab. inż. Agnieszka Bartoszek-Pączkowska						
of lecturer (lecturers)	Teachers								
Lesson types and methods	Lesson type	Lecture	Tutorial	Laboratory	Projec	t	Seminar	SUM	
of instruction	Number of study hours	15.0	15.0	0.0	0.0		15.0	45	
	E-learning hours inclu	ided: 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	45		5.0		25.0		75	
	of life sciences. These methods are increasingly frequently applied in biological sciences, medicin and their practical usefulness is on the constant rise. The subject addresses students of specialisations: BT, Ch, TCh, because all these subjects have essential imput in the creation of multiomic technologies.								
Learning outcomes	Course out	come	Subi	ect outcome			Method of veri	fication	
Esaming Sucomos	[K7_K101] acknowledges the importance of knowledge related to the field of study in solving cognitive and practical problems, critically assessing the information obtained		Students appreciates the scientific and practical value of multiomic technologies for the development of mankind.		[SK2] Assessment of progress of work [SK5] Assessment of ability to solve problems that arise in practice				
	[K7_K02] is aware of the potential risks and opportunities associated with the development of science and technology for the natural environment and society		Student comprehends consequences, both beneficial as well as controvertial, resulting from the accessibility of detailed bioinformatic data concerning e.g. individual person.		[SK2] Assessment of progress of work [SK5] Assessment of ability to solve problems that arise in practice				
	[K7_U02] uses research methods used in biotechnology and related fields		Student knows informatic and bioinformatic methods applied in analysis of data obtained from omic technologies.		[SU2] Assessment of ability to analyse information				
	[K7_W01] defines the phenomena, processes and laws of living nature applied to the production of useful goods and the carrying out of services		Student exhibits the basic knowledge of biology and chemistry needed to understand biochemical phenomena and processes utilized in omic technologies.		[SW1] Assessment of factual knowledge				

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Historical skotch from Mendel to Human Genome Project, the birth of epigenomics and discovery of small regulatory RNA molecules, variety of "omes" and "omits" in life science studies; big genomic projects and their inheticular and practical significance; the difference between genomics and transcriptomics and the opportunities of couples and practical significance; the plans; high frincipling methods in recognition of personalized nutrition, proteomics and its variants in medical and nutritional studies; scope and applications of metabolomics, increasional and methods are scope and applications of metabolomics, increasional and nutritional studies; scope and applications of metabolomics, increasional and nutritional research. Topics realized during seminars During seminars, students are expected to present talks based on the most recent scientific publications connected with subjects discussed during lectures and recommended by tutors. The aim is to support totures with the revest information. Topics realized during exercises aimed at making students familiar with technical side of high throutopput methods: 1/2. Strategies of synthesis of polynucleotides used in genomic experiments 3. Methods employed upon Human Genome Project realisation. 4. Methods developed as a result of Human Genome Project realisation, including bioinformatic tools. 5/6. Technological approaches employed in production of DNA arrays; design of sequences. 7/8. Research strategies and methods used in transcriptomic experiments, RT-PCR as an example. 9/10. Methods applied in proteomic studies; spectroscopic and immunological. 11. Methods used in studies on human metabolome; purpose decides 12/13. Research strategies and technologies used in metagenomic atudies. 14/15. Presentation of big data derived on high throughput studies. Prerequisites Accomplished courses in organic chemistry and biochemistry. Subject passing criefis Passing threahold Percentage of the final grade Lecture attendance and exam 50.0% 20.0% 20.0% 20.0% 20.0%	Subject contents	Topics realized during lectures						
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Example issues/ example questions/ tasks being completed	What are scopes of genomics, transcriptomics and proteomics?			
	What is the difference between the impact of epigenetic vs. genetic mechanisms on transcriptome function?			
	What is named "multiomic studies"?			
Work placement	Not applicable			

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