Course code								
Type and description	EC - elective course for Chemical engineering							
ECTS credit	1							
Course name	Application of - <i>omics</i> techniques for identification of microorganisms and chemical compounds in environmental systems							
Course name in Polish	Zastosowanie technik - omicznych do identyfikacji mikroorganizmów I związków chemicznych w systemach środowiskowych							
Language of instruction	English							
Course level	8 PRK							
Course coordinator	Prof. dr hab. Beata Gutarowska							
Course instructors	Prof. dr hab. Beata Gutarowska							
Delivery methods and course duration		Lecture	Tutorials	Laboratory	Project	Seminar	Other	Total of teaching hours during semester
	Contact hours	0	0	0	5	0	0	5
	E-learning Assessment criteria (weightage)	no 0	no 0	no 0	no 100%	no 0	no 0	no 0
Course objective	The aim of the course is to learn about "omics" new generation techniques, mainly metagenomics and metabolomics, which are applicable to the systemic identification of microorganisms and chemical compounds in environmental samples.							
Learning outcomes	The student is able to discuss the metabolomic and metagenomic methods and their importance The student is able to describe bioinformatics tools, databases, statistical tools for analysing of the omics results The student is able to interpret the results of omics analyzes The student is able to present the problem using the literature databases and multimedia presentation tools.							
Assessment methods	1. Discussion, 2. Discussion, 3. Case study presentation, discussion 4. Multimedia presentation							
Prerequisites	not required							
Course content with delivery methods	 Introduction to the analysis of biology and system chemistry called <i>omics</i>. Definitions, types and importance of new generation methods compared to classical methods, environmental sampling - lecture 1 hour. New generation metagenomic methods: DNA isolation, primers, commercial kits, PCR, qPCR, RTPCR, sequencing with proton detection in integrated circuit (Ion Torrent Sequencing), pyrosequencing 454, sequencing by synthesis (Illumina MiSeq), nanopore sequencing, comparison of methods, apparatus - presentation 1 hour Targeted and untargeted metabolomic methods: high resolution mass spectrometry (HRMS) coupled with liquid (LC) or gas chromatography (GC), NMR (nuclear magnetic resonance), infrared IR spectroscopy, UV spectroscopy, fluorescence spectroscopy, metabolomic imaging MS mass spectrometry with laser desorption / ionization (SALDI) and assisted surface, for example plates with nanoparticles AgNPET or AuNPET - presentation 1 hour Bioinformatics analysis, databases, statistical analysis, software, identification of microorganisms, chemical compounds and metabolic pathways in environmental samples (eg surfaces of historic objects), case study – Project Base Learning PBL 2 hours. 							

Basic reference materials	 Proteomika i metabolomika, Kraj Agnieszka, Drabik Anna, Silberring Jerzy, 2019, wyd Uniwersytetu Warszawskiego, Warszawa, ISBN: 9788323507659 Bioanalityka w nauce i życiu. Cz.1 i 2. Irena Staneczko-Baranowska, Bogusław Buszewski, 2020, wyd. PWN, Warszawa, ISBN:9788301212810
Other reference materials	Manuscripts in scientific journals on metabolomic and metagenomic tools – case studies
Average student workload outside classroom	15 hours
Comments	summer semester
Last update	18.11.2020