

**Course name:****BIOINFORMATICS IN ANIMAL HUSBANDRY**

ECTS	3
Course status	obligatory
Course final assesement/evaluation of outcomes	exam
Prerequisites	passing the subject Genetics and Molecular Biology

**Main field of study:****BIOENGINEERING IN ANIMAL SCIENCE**

Profile of study	General-academic
The code of studies (education level)	SM (master)
Semester of studies	summer
Language of instruction	English

**Course offered by:**

Name of faculty offering the course	Faculty of Animal Sciences
Name of department offering the course	Department of Genetics, Animal Breeding and Ethology
Course coordinator	dr inż. Łukasz Migdał

**Learning outcomes of the course:**

Symbol of outcome	Description of learning outcome	Reference to	
		main field of study outcomes	discipline#

**KNOWLEDGE – student knows and/or understands:**

BIER_W01	analyse of the results of experiments in the field of bioengineering and related fields	BIOI2_W01	RZ
BIER_W02	knows the rules for preparation of scientific publication and has the ability to use	BIOI2_W17	RZ

**SKILLS – student is able to:**

BIER_U01	plan and perform experiment, analyze and interpret the results obtained, using	BIOI2_U01	RZ, PB
BIER_U02	skillfully select and modify techniques and technologies in order solutions to specific problems in the field of animal bioengineering and	BIOI2_U07	RZ
BIER_U03	use English in the fields of science and disciplines relevant to the field of study being studied, in accordance with	BIOI2_U11	RZ

**SOCIAL COMPETENCE- student is ready to:**

BIER_K01	understands the need for systematic work on long-term implementation projects and is aware of the responsibility for the team's work results	BIOI2_K02	RZ
----------	--	-----------	----

**Teaching contents:**

<b>Lectures</b>	<b>10</b>	<b>hours</b>
<ol style="list-style-type: none"> <li>1. Bioinformatics databases - Major types of bioinformatics data – 1h</li> <li>2. Dynamic programming (pair -wise sequences alignment) - 2h</li> <li>3. PAM and BLOSUM matrices, multisequence alignment (MSA), Markov models - 2h</li> </ol>		

Topics of the lectures	4. FASTA and BLAST searching - 1h 5. Protein Structures - 2h 6. Introduction to Phylogenetics -1h 7. Exam - 1h
------------------------	---

Accomplished learning outcomes	<i>BIER_W01, BIER_U01, BIER_K01</i>
--------------------------------	-------------------------------------

Verification methods, rules and criteria of outcome assessment	<i>Short question, multiple choice questions (25 - 22 points - 5.0; 21-20 - 4.5; 19-17 - 4.0; 16-15 - 3.5; 14-12 -3.0)</i>
--	--

<b>Classes</b>	<b>30</b>	<b>hours</b>
----------------	-----------	--------------

Topics of the classes	1.NCBI and ENSEMBL databases - BLAST options (2 lectures - 1,5h each - 3h) 2.DNA - Genes, genes regulation, gene prediction and promoter prediction, polymorphisms analysis (6 lectures - 1,5h each - 9h) 3. RNA - RNA structure prediction (2 lectures - 1,5h each - 3h) 4. Proteins - structure, structure classification, prediction and visualization (7 lectures - 1,5h each - 10,5h) 5 Phylogenetics (2 lectures - 1,5h each - 3h) 6. Evaluation of laboratories - ( 1 lecture - 1,5h)
-----------------------	---

Accomplished learning outcomes	<i>BIER_W01, BIER_W02, BIER_U01,BIER_U02, BIER_U03, BIER_K01</i>
--------------------------------	--

Verification methods, rules and criteria of outcome assessment	<i>two tests (70%) and student project evaluation(30%) - for every test there will be 5 works (5 points each) - (25 - 22 points - 5.0; 21-20 - 4.5; 19-17 - 4.0; 16-15 - 3.5; 14-12 -3.0)</i>
--	---

<b>Seminars</b>	<b>...</b>	<b>hours</b>
-----------------	------------	--------------

Topics of the seminars	
------------------------	--

Accomplished learning outcomes	<i>symbol of learning outcomesof the seminars</i>
--------------------------------	---

Verification methods, rules and criteria of outcome assessment	<i>together with participation in the final assesement (in %)</i>
--	---

**References:**

Basic	<i>Bioinformatics: Sequence and Genome Analysis, Mount, CSHLP 2004</i>
Supplementary	<i>Bioinformatics programme instructions(available online)</i>

**Structure of learning outcomes:**

Dyscipline – animal science and fishery (RZ)	3	ECTS**
Discipline: # (provide appropriate symbol - if the course relates to more than one academic discipline )	...	ECTS**

**Structure of student activities:**

Contact hours	40	hours	1,6	ECTS**
including:	lectures	9	hours	
	classes and seminars	28,5	hours	
	consultations	...	hours	
	participation in research	...	hours	

mandatory traineeships	...	hours		
participation in examinations	2,50	hours		
e-learning	0	hours	0	ECTS**
student own work	35	hours	1,4	ECTS**

Syllabus valid from the academic year 2021/2022

\* **where 10 hours of classes = 1 ECTC (in case of 15 h → 2 ECTS)**

\*\* stated with an accuracy to 0.1 ECTS, where 1 ECTS = 25 - 30 hours of classes

# academic discipline code: RZ - animal science and fishery, PB - biological sciences, etc.