

**Course name: Plant Genomics**

ECTS	3
Course status	<i>facultative</i>
Course final assessment /evaluation of outcomes	<i>Exam</i>
Prerequisite	<i>plant biology, basic biochemistry, conventional and molecular genetics</i>

**Main field of study:**

Agriculture and Horticulture, Biology and Biotechnology (Erasmus+)

Educational profile	General academic
Code of studies and education level	bachelor/engineer (SI) or master of science (SM)
Semester of studies	Winter or Summer
Language of instruction	English

**Course offered by:**

Name of faculty offering the course	Biotechnology and Horticulture
Name of department offering the course	Plant Biology and Biotechnology
Course coordinator	Prof. dr hab. Dariusz Grzebelus

**Learning outcomes:**

Symbol of outcome	Description of the learning outcome	Reference to main field of study outcomes	Area symbol*
KNOWLEDGE – student knows and understands			
PLG_W1	Key issues in structural, functional and comparative genomics	EPB2_W02	R, P
PLG_W2	Basic methods of genome analysis	EPB2_W01 EPB2_W02	R, P
PLG_W3	Principles and methods of DNA sequencing	EPB2_W01 EPB2_W02	R, P
PLG_W4	Basic principles of plant genome evolution	EPB2_W02	R, P
SKILLS – student is able to			
PLG_U1	Utilize online genomic resources and databases of DNA and protein sequences	EPB2_U03 EPB2_U04	R, P
PLG_U2	Perform basic analyses of DNA sequences in silico	EPB2_U01 EPB2_U04	R, P
PLG_U3	Align DNA and protein sequences and evaluate their similarity	EPB2_U01 EPB2_U04	R, P
SOCIAL COMPETENCIES – student is ready to:			

**Teaching contents**

Lectures		15 hours
Topics	<ol style="list-style-type: none"> <li>1. Structural and functional genomics.</li> <li>2. <i>Arabidopsis thaliana</i> as a model plant genomes.</li> <li>3. Methods of genome analysis, genome annotation.</li> <li>4. Evolution of plant genomes, mobile genetic elements.</li> <li>5. Comparative genomics.</li> </ol>	
Accomplished learning outcomes		EPB2_W01, EPB2_W02

Means of verification, rules and criteria of assessment	<i>Test exam, single choice (51%)</i>
Classes:	15 hours
Topics	1. Exploration of online sequence databases (DNA and protein). 2. In silico analysis of DNA sequences. 3. Sequence alignment and similarity, identification of polymorphisms.
Accomplished learning outcomes	EPB2_U01, EPB2_U03, EPB2_U04
Means of verification, rules and criteria of assessment	<i>Case report, demonstration of practical skills (49%)</i>

### References:

Basic	<i>Meksem K, Kahl G (eds.), 2005. The Handbook of Plant Genome Mapping. Wiley-VCH, Weinheim. Sensen CW (ed.), 2005. Handbook of Genome Research. Wiley-VCH, Weinheim, vol. 1 and 2. Lankenau D-H, Voff J-N (eds.), 2009. Transposons and the Dynamic Genome. Springer, Dordrecht.</i>
Supplementary	<i>The Arabidopsis Genome Initiative, 2000. Analysis of the genome sequence of the flowering plant Arabidopsis thaliana. Nature 408: 796-815. Varshney RK, Tuberosa R (eds.), 2007. Genomics-Assisted Crop Improvement. Springer, Dordrecht, vol. 1 and 2.</i>

### Structure of learning outcomes

Area of academic study: agriculture and horticulture	2 ECTS **
Area of academic study: biological sciences	1 ECTS**

### Structure of student activity

Contact hours	34 hrs.	1.4 ECTS**
Including:		
lectures	15 hrs.	
classes and seminars	15 hrs.	
consultations	2 hrs.	
participation in research	- hrs.	
obligatory traineeships	- hrs.	
participation in examination	2 hrs.	
e-learning	- hrs.	- ECTS**
student own work	41 hrs.	1.6 ECTS**

\*areas of academic study in the fields of: P – biological sciences; R – agriculture and horticulture

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