General Information	
Academic subject	Plant Genetics
Degree course	Environmental and forest Sciences
Curriculum	Genetics
ECTS credits	4 ECTS Lectures + 2 ECTS Laboratory
Compulsory attendance	No
Language	Italian

Subject teacher	Name Surname	Mail address	SSD
	<b>.</b>	rosanna.simeone@uniba.it	AGR/07
	Simeone		

ECTS credits details	Торіс	SSD	Credits
Basic teaching activities	Principal aspects	AGR/07	6
	of genetics and		
	biology of agro-		
	forestry plants.		

Class schedule	
Period	II semester
Year	2017-18
Type of class	Lecture- workshops

Time management	
Hours	150
In-class study hours	60
Out-of-class study hours	90

Academic calendar	
Class begins	5th March, 2018
Class ends	22nd June, 2018

Syllabus	
Prerequisites/requirements	Knowledge of inorganic and organic chemistry and biology
Expected learning outcomes (according to	<ul> <li>Knowledge and understanding</li> </ul>
Dublin Descriptors) (it is recommended	Knowledge on the principal genetic methodologies
that they are congruent with the learning	
outcomes contained in A4a, A4b, A4c	<ul> <li>Applying knowledge and understanding</li> </ul>
tables of the SUA-CdS)	Applying and understanding the principal genetic methodologies for agro-forestry species.
	<ul> <li>Making informed judgments and choices</li> <li>Capacity of applying the principal genetic methodologies for agro-forestry species.</li> </ul>
	<ul> <li>Communicating knowledge and understanding Capacity of identifying the principal genetic methodologies for agro-forestry species.</li> </ul>
	<ul> <li>Capacities to continue learning Capacity of communicate and continue learning the principal genetic methodologies for agro-forestry species.</li> </ul>

<ul> <li>and in eukaryotes. Mitosis. Meiosis. Life cycles in organisms of interest for genetic studies.</li> <li>Mendelian analysis. Mendel's experiments. Genotype and phenotype. The chromosome theory of heredity. Multiple alleles. Epistatic genes. Statistical analyses.</li> <li>Linkage and eukaryotic chromosome mapping. Crossing-over and recombination. Linkage maps. Two point test-crosses.</li> <li>The genetic material. The structure of DNA and RNA. Replication of DNA. The chromosome structure. Geneprotein relation-ships. Gene function.</li> <li>Transcription and translation. Transcription. Eukaryotic RNA. Mechanism of gene splicing. The genetic code. Translation.</li> </ul>	Contents	Genetics and the organism. Chromosomes in prokaryotes
<ul> <li>Mendelian analysis. Mendel's experiments. Genotype and phenotype. The chromosome theory of heredity. Multiple alleles. Epistatic genes. Statistical analyses.</li> <li>Linkage and eukaryotic chromosome mapping. Crossing-over and recombination. Linkage maps. Two point test-crosses.</li> <li>The genetic material. The structure of DNA and RNA. Replication of DNA. The chromosome structure. Geneprotein relation-ships. Gene function.</li> <li>Transcription and translation. Transcription. Eukaryotic RNA. Mechanism of gene splicing. The genetic code. Translation.</li> </ul>		• • • • • • • • •
<ul> <li>phenotype. The chromosome theory of heredity. Multiple alleles. Epistatic genes. Statistical analyses.</li> <li>Linkage and eukaryotic chromosome mapping. Crossing-over and recombination. Linkage maps. Two point test-crosses.</li> <li>The genetic material. The structure of DNA and RNA. Replication of DNA. The chromosome structure. Geneprotein relation-ships. Gene function.</li> <li>Transcription and translation. Transcription. Eukaryotic RNA. Mechanism of gene splicing. The genetic code. Translation.</li> </ul>		interest for genetic studies.
<ul> <li>phenotype. The chromosome theory of heredity. Multiple alleles. Epistatic genes. Statistical analyses.</li> <li>Linkage and eukaryotic chromosome mapping. Crossing-over and recombination. Linkage maps. Two point test-crosses.</li> <li>The genetic material. The structure of DNA and RNA. Replication of DNA. The chromosome structure. Geneprotein relation-ships. Gene function.</li> <li>Transcription and translation. Transcription. Eukaryotic RNA. Mechanism of gene splicing. The genetic code. Translation.</li> </ul>		Mondolian analysis Mondol's experiments Constype and
alleles. Epistatic genes. Statistical analyses. Linkage and eukaryotic chromosome mapping. Crossing-over and recombination. Linkage maps. Two point test-crosses. The genetic material. The structure of DNA and RNA. Replication of DNA. The chromosome structure. Gene- protein relation-ships. Gene function. Transcription and translation. Transcription. Eukaryotic RNA. Mechanism of gene splicing. The genetic code. Translation.		
Crossing-over and recombination. Linkage maps. Two point test-crosses. <b>The genetic material.</b> The structure of DNA and RNA. Replication of DNA. The chromosome structure. Gene- protein relation-ships. Gene function. <b>Transcription and translation.</b> Transcription. Eukaryotic RNA. Mechanism of gene splicing. The genetic code. Translation.		
test-crosses. <b>The genetic material.</b> The structure of DNA and RNA. Replication of DNA. The chromosome structure. Gene- protein relation-ships. Gene function. <b>Transcription and translation.</b> Transcription. Eukaryotic RNA. Mechanism of gene splicing. The genetic code. Translation.		Linkage and eukaryotic chromosome mapping.
Replication of DNA. The chromosome structure. Gene- protein relation-ships. Gene function. <b>Transcription and translation.</b> Transcription. Eukaryotic RNA. Mechanism of gene splicing. The genetic code. Translation.		
protein relation-ships. Gene function. <b>Transcription and translation.</b> Transcription. Eukaryotic RNA. Mechanism of gene splicing. The genetic code. Translation.		The genetic material. The structure of DNA and RNA.
RNA. Mechanism of gene splicing. The genetic code. Translation.		
Translation.		Transcription and translation. Transcription. Eukaryotic
<b>DNA recombinant.</b> Restriction enzymes. Recombinant		
		DNA recombinant. Restriction enzymes. Recombinant
DNA methodology. Cloning vectors. Gene libraries. Clone		DNA methodology. Cloning vectors. Gene libraries. Clone
identification. Polymerase chain reaction (PCR). DNA polymorphism. Genetic transformation in forest trees.		
Mutations. Origin of mutations. Gene mutations. Changes in		Mutations. Origin of mutations. Gene mutations. Changes in
chromosome structure. Aneuploidy. Auto- and allo- polyploidy. Interspecific hybridization in forest trees.		chromosome structure. Aneuploidy. Auto- and allo-
Quantitative genetics. Qualitative and quantitative		<b>Quantitative genetics.</b> Qualitative and quantitative
characters. Continuous variation. Genetic and environment variation. Heritability. Mapping of quantitative traits in plants.		characters. Continuous variation. Genetic and environment
<b>Population genetics.</b> Hardy-Weinberg equilibrium. Changes		<b>Population genetics</b> Hardy-Weinberg equilibrium Changes
from Hardy-Weinberg equilibrium: migration, mutation		
selection, inbreeding depression, heterosis.		
Introduction to plant breeding. Racial selection. Selection		Introduction to plant breeding. Racial selection. Selection
of plus trees. Half-sib and full-sib selection. Clonal selection.		•
Phenotypic selection. Seed orchard. Seedling seed orchard.		Phenotypic selection. Seed orchard. Seedling seed orchard.
Course program	Course program	
Bibliography • Russel P.J., Wolfe S.L., Hertz P.E., Starr C., McMillan B.		
2016. Genetica Agraria. EdiSES S.r.I. Ed.		0
<ul> <li>Lorenzetti F., Ceccarelli S., Rosellini D., Veronesi F. 2011.</li> <li>Genetica agraria. Patron Ed.</li> </ul>		
Barcaccia G., Falcinelli M. 2005. Genetica e Genomica.		
Liguori Ed.		Liguori Ed.
• Figliuolo G. 2012. Genetica vegetale. Favia Ed.		• Figliuolo G. 2012. Genetica vegetale. Favia Ed.
Further readings:		Further readings:
• Griffiths A.J. F., Gelbart W. M., Miller J. H., Lewontin R. C.		•
2004. Genetica moderna. Zanichelli, Vol. I-II.		
Chrispeels M. J., Sadava D. E. 2005. Genetica, Biotecnologie     e agricoltura sostenibile. Idelson-Gnocchi		

	Notes from classes
Notes	
Teaching methods	Lectures will be presented through PowerPoint and overhead
Assessment methods (indicate at least the type written, oral, other)	A midterm write exam is scheduled for students enrolled to the Course. This exam will test the first half of the course's information in the middle of the semester and the rest of the course's information at the final exam. The final exam will consist on an oral test, as reported in the Guidelines of the Degree of Environmental and Forest Sciences. (art.10) and in the Annex A. The evaluation of the student will be based on established criteria, as explained in the Annex A of the Degree of Environmental and Forest Sciences. The final grade will be an average of both the midterm and final exam. For foreign the exam consists of an oral test with questions related to the course's information.
Evaluation criteria (Explain for each	• Knowledge and understanding
expected learning outcome what a	Knowledge on the principal genetic methodologies
student has to know, or is able to do, and	5 1 1 5 5
how many levels of achievement there	• Applying knowledge and understanding
are.	Applying and understanding the principal genetic
	methodologies for agro-forestry production
	• Making informed judgments and choices
	Capacity of apply the principal genetic methodologies for agro-forestry production
	o Communicating knowledge and understanding Capacity of identify the principal genetic methodologies for agro-forestry production
	o Capacities to continue learning Capacity of communicate and continue learning the
Official visiting hours.	principal genetic methodologies for agro-forestry production Monday to Friday, 10.30-13.00 a.m. (Define the appointment
0	by e-mail).