Optional course – main information		
Academic subject	Plant Molecular Phylogeny	
ECTS credits (CFU)	4	
Compulsory attendance		
Teaching language	Italian	
Accademic Year	2020/2021	

Professor/Lecturer		
Name & SURNAME	Fabrizio Grassi	
email	fabrizio.grassi@uniba.it	
Tel.		
Tutorial time/day	Wednesday (by appointment)	

Course details	Pass-fail exam/Exam with mark out of 30	SSD code	Type of class
Course details	Exam with mark out of 30	Bio/01	Lecture/workshop

	Semester	day and time	room
Teaching schedule	1	Tuesday and Wednesday	
	•	<b>15.00 - 16.40</b> (to be defined with the students)	

Lesson type	CFU/ECTS	Lessons (hours)	CFU/ECTS lab	Lab hours	CFU/ECTS tutorial/workshop	Tutorial/workshop hours	CFU/ECTS field trip	Field trip Hours
	4	32						

Time	Total hours	Teaching hours	Self-study hours
management		32	

Academic	First lesson	Final lesson
Calendar	27/10/2020	23/12/2020

Syllabus				
Course entry requirements	Fundamentals of plant biology			
Expected learning outcomes (according to Dublin Descriptors) (it is recommended that they are congruent with the				
learning outcomes contained in	A4a, A4b, A4c tables of the SUA-CdS)			
Knowledge and understanding	To acquire further knowledge in plant phylogeny and methods to construct phylogenetic trees			
Applying knowledge and understanding	The student should be able to describe and apply the most suitable methods for the study of plant phylogeny			
Making informed judgements and choices	Acquisition of autonomy in the evaluation and interpretation of the scientific letterature			
Communicating knowledge and understanding	To describe the phylogenetic relationships between the main taxonomic groups using a correct scientific terminology			
Capacities to continue learning	Some scientific papers will be commented and critically analyzed to assess the progressive level of learning.			

Sylabus	
Course content	Introduction to the plant phylogeny. The origins of the phylogeny and description of the main applications. Cladistics vs. Phenetics. Description of the main methods for the construction of phylogenetic trees. Description and use of the Maximum Parsimony method and probabilistic methods. Nuclear and plastid DNA, and genes of interest for the molecular phylogeny. Origin of Bryophytes, Tracheophytes and

	Spermatophytes. Evolution of the morphological and molecular traits. Origin and diversification of Angiosperms. Morphological and molecular evidence to support the main monophyletic groups. Basal angiosperms and paleoherbs. The molecular clock. The use of fossils to define the times of speciation. Polyploidy in the evolution of plants. Domestication of plants. The use of molecular markers to distinguish related species and critical groups. Resolution of taxonomic problems through the analysis of morphological and molecular traits. Description of some software for phylogenetic analysis.
Course books/Bibliography	Judd, Plant Systematics: A Phylogenetic Approach, Sinauer Associate
Notes	During the course students will be provided with further bibliographical references.
Teaching methods	Lectures, Power Point, software and database
Assessment methods (indicate	
at least the type written, oral,	Oral examination.
other)	
Evaluation criteria (Explain for	-Acquisition of phylogenetic concepts and description of relationships between
each expected learning	principal taxonomic groups.
outcome what a student has to	- Ability to describe phylogenetic trees and correct use of scientific language.
know, or is able to do, and how	- Ability to describe and apply principal phylogenetic methods.
many levels of achievement	- Capacity to resolve basic phylogenetic problems.
there are	
Further information	