MASTER 'S PROGRAMME Molecular Genetics ST YEAR OF STUDY, 1ST SEMESTER

COURSE TITLE	GENOMICS	
COURSE CODE	BGM6101	
COURSE TYPE	full attendance	
COURSE LEVEL	2 nd cycle (master's degree)	
YEAR OF STUDY, SEMESTER	1 st year of study, 1 st semester	
NUMBER OF ECTS CREDITS	6	
NUMBER OF HOURS PER WEEK	4 (2 lecture hours + 2 seminar hours)	
NAME OF LECTURE HOLDER	Lucian Gorgan	
NAME OF SEMINAR HOLDER	Lucian Gorgan	
Prereouisites	Advanced level of English	
A GENERAL AND COURSE-SPEC	IFIC COMPETENCES	
General competences:		
\rightarrow The usage of terms and in-depth concepts, working principles and methodologies characteristic		
for the field of study and specialization "Molecular genetics"		
\rightarrow Investigation and interpretation of the molecular basis of organization and functioning of living		
matter for the elaboration of studies / reports that can be published and / or applied in a professional		
The use of mol	acular analysis equinments and tools specific to different professional	
	scular analysis equipments and tools specific to uncrent professional	
Course specific competences:		
Course-specific compete	IILES.	
→ to understand the structures of different genome types and the differences between them		
\rightarrow to differentiate the influence of the main evolution factors on the individual status		
\rightarrow to use genomics-specific scientific language		
\rightarrow to understand the	importance of knowing the mutational processes and the adaptability of the	
organisms		
\rightarrow to know the main n	nethods and techniques for genomic analysis	
B LEARNING OUTCOMES		
\rightarrow Initiation into the kr	nowledge of molecular structures and mechanisms at the genomic level.	
\rightarrow Awareness of the	\rightarrow Awareness of the influence of environmental factors on individual status.	
\rightarrow The development	of the skills needed to analyze nucleic acids and to correlate structures with	
the spatial and tem	poral factors of evolution.	
C LECTURE CONTENT		
The concept of Genome		
The genome in prokaryotes and eukaryotes		
Comparative genomics		
Metagenome		
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Molecular markers	Molecular markers	
Sampling and analysis techniques		
Methods for identifying polymorphisms		
Statistical methods used in comparative genomics		
Databases - DNA sequences, complete genomes		
Gene annotations. BLAST Algorithm Sequence alignment		
Phylogeny and molecular phylogeography.		
Phylogenetic trees, trees calibration		
F RECOMMENDED READING FOR SEMINARS		
1. Nei M., Kumar S., 2000 – Molecular evolution and phylogenetics, Oxford University Press.		
2. Saccone C., Pesole C., 2003 - Handbook of comparative genomics - Principles and Methodology,		
G EDUCATION STVLE		
	systematic exposure: conversation: didactic demonstration	
LEARINING AND TEACHING METHODS		
ASSESSMENT METHODS	Exam	
LANGUAGE OF INSTRUCTION	English	