

# UNIVERSITÀ DEGLI STUDI DI PALERMO

DEPARTMENT	Biomedicina, Neuroscienze e Diagnostica avanzata				
ACADEMIC YEAR	2021/2022				
MASTER'S DEGREE (MSC)	MEDICAL BIOTECHNOLOGIES AND MOLECULAR MEDICINE				
INTEGRATED COURSE	MEDICAL-FORENSIC GENETICS AND MOLECULAR EPIDEMIOLOGY - INTEGRATED COURSE				
CODE	19372				
MODULES	Yes				
NUMBER OF MODULES	2				
SCIENTIFIC SECTOR(S)	BIO/13, MI	ED/42			
HEAD PROFESSOR(S)	SEIDITA C	GREGO	RIO	Ricercatore	Univ. di PALERMO
OTHER PROFESSOR(S)	TRAMUTO SEIDITA O			Professore Associato Ricercatore	Univ. di PALERMO Univ. di PALERMO
CREDITS	9				
PROPAEDEUTICAL SUBJECTS					
MUTUALIZATION					
YEAR	1				
TERM (SEMESTER)	1° semester				
ATTENDANCE	Mandatory				
EVALUATION	Out of 30				
TEACHER OFFICE HOURS	SEIDITA GREGORIO				
	Tuesday	14:30	16:30	Sezione di Biologia e Genetica di Biopatologia e Biotecnologia	
	Wednesday	11:00	13:00	Sezione di Biologia e Genetica di BiND)	a via divisi, 83 (Dipartimento
	TRAMUTO	FABIO			
	Monday	14:00	16:00	Dipartimento di Promozione de Medicina Interna e Specialistic D'Alessandro"AOUP "P. Giaco 133Piano terra	a di Eccellenza "G.
	Wednesday	14:00	16:00	Dipartimento di Promozione de Medicina Interna e Specialistic D'Alessandro"AOUP "P. Giaco 133Piano terra	a di Eccellenza "G.
	Friday	14:00	16:00	Dipartimento di Promozione de Medicina Interna e Specialistic D'Alessandro"AOUP "P. Giaco 133Piano terra	a di Eccellenza "G.

## DOCENTE: Prof. GREGORIO SEIDITA

PREREQUISITES	The student should have the basic knowledge of biology and genetics. It must also know the basic fundamentals of biostatistics and show ability to run at least one operating systems (i.e. Linux, MacOS and or Microsoft Windows) and a internet browser.
LEARNING OUTCOMES	Knowledge and understanding At the end of the course students will be able to develop, analyze and solve problems related personal identification, detection, or the exclusion of family relationships using appropriate techniques of molecular biology; analyze and apply the methods available to study the genetic variability and its relationship with human diseases; experimental use genetic information in a statistical sense and develop comparison skills with international databases; evaluation capacity of a biological evidences. Understanding of the ethical and legal problems associated with the personal identification procedures; At the end of the course students will be expected to have acquired the needed skills for critically knowing and using informatics tools and programs for genomic and proteomic data analysis. The above results will be achieved trough the knowledge of both biological database organization models and search strategies, and pairwise and multiple sequence alignments.
	Graduates will demonstrate knowledge of the molecular epidemiology applications in different health situations, knowing how to interpret and evaluate the results. Recognize the nature of biomedical data in order to be able to properly apply the most appropriate methods for analyzing statistics, and describe and properly detect the presence or absence of relationships between variables. To know and critically use computer tools applied to the analysis of experimental genetic information and epidemiological items.
	Applying knowledge and understanding Students will spend the knowledge acquired during the course in Human Molecular Genetics directly into work (in research or analysis laboratories, public or private). They will know how to apply the new techniques for the investigation of genetic variability, considering also bioethical implications this analysis raise.
	<ul> <li>Graduates must also demonstrate:</li> <li>competence in the design and organization of epidemiological studies appropriate to the various health needs;</li> <li>competence in the application and interpretation of measures of statistical description, of association and of the main descriptive graphics in molecular epidemiology;</li> <li>Knowledge of laboratory techniques and computer aimed at assessing the genetic characteristics and potential epidemiological pattern. Use and search in online biological databases.</li> <li>Use and do research of gene and protein sequences.</li> <li>Use simple data analysis tools.</li> </ul>
	Making judgments Be able to independently meet the professional issues related to knowledge of the course. Students will be able to evaluate in a rational and autonomous basic knowledge provided by the course and will be able to address issues of Human Molecular Genetics through a scientific approach. Autonomy, the application and interpretation of the measures of the main descriptive statistics and graphs in molecular epidemiology. Students will be expected to have acquired autonomous and critic judgment skills for evaluating and analysing significance and limits of the programs and tools produced results.
	Communicative skills Ability to communicate and disseminate the knowledge acquired during the course in their professional responsibilities. Students will gain a communicative methodology of scientific / experimental in the survey genetics and genetic- forensic and data analysis. They will know how to manage the genetic investigations with particular decision-making autonomy to guarantee confidentiality. Gain ability to evaluate the results. Students should be able to clearly report on the topics dealt with during the course in its own professional domain. Following a strong scientific approach, the use of correct technical and computer science based-terms is also required.
	Learning skills Application of knowledge of the course in order to improve the overall approach to their health care professional. Ability to update by consulting their own scientific publications of the Molecular Genetics sector with particular attention to the human and forensic genetics and special network resources. The

	development of students' awareness for scientific publications and for new tool and program position papers will be a further target.
ASSESSMENT METHODS	The Learning evaluation is done by oral examination. The outcome of the evaluation is expressed by a vote ranging from 1-30. The oral exam consists of a conversation usually lasting about 20-30 minutes in order to check the disciplinary knowledge of the curriculum contents. In the assessment of learning it is taking into account the participation and evaluation obtained of test carried out during the course. The following table shows the learning evaluation scheme with the ECTS grades, the equivalent rating of thirty and their significance.
	A – A+ / Excellent 30/30 cum laude Excellent knowledge of teaching contents; students should show high analytical and synthetic capabilities and should be able to apply their knowledge to solve highly complex problems.
	B / Very good 27-29/30 Very good knowledge of the teaching contents and excellent language control; students should show analytical and synthetic skills and be able to apply their knowledge to solve problems of medium and, in some cases, even higher complexity.
	C / Good 24- 26/30 Good knowledge of teaching contents and good language control; the students should be able to apply their knowledge to solve problems of medium complexity
	D / Satisfactory 21-23/30 Average knowledge of the teaching contents, in some cases limited to the main topic; acceptable ability to use the specific discipline language and independently apply the acquired knowledge.
	E / Sufficient 18-20 Minimum teaching content knowledge, often limited to the main topic; modest ability to use the subject specific language and independently apply the acquired knowledge.
	F / Fail Lack of an acceptable knowledge of the main teaching content knowledge; very little or no ability to use the specific subject language and apply independently the acquired knowledge.
	A preliminary and optional examination is carried out during the lessons. This is a test with 6 open questions to be performed in 90 minutes. The exam questions will focus on the lessons contents untile then.
TEACHING METHODS	Lessons in the classroom; practical lessons and exercises in classroom and bioinformatic classroom.

## MODULE MOLECULAR EPIDEMIOLOGY

Prof. FABIO TRAMUTO

 SUGGESTED BIBLIOGRAPHY

 Rothman Kenneth J, Epidemiologia, Idelson-Gnocchi, ISBN-13: 978-8879474580

 Weiss NS, Exercises in Epidemiology, Oxford University Press, ISBN-13: 9780190651510

 AMBIT
 20885-Attività formative affini o integrative

 INDIVIDUAL STUDY (Hrs)
 51

 COURSE ACTIVITY (Hrs)
 24

COURSE ACTIVITY (Hrs) EDUCATIONAL OBJECTIVES OF THE MODULE

The purposes of this module are:

a) to outline the fundamentals of epidemiological data analysis and interpretation.

b) to discuss the theoretical aspects of technics in molecular epidemiology.

SYLLABUS		
Hrs	Frontal teaching	
4	Elements of biostatistics Epidemiological studies: cross-sectional studies, case-control studies, cohort studies	
4	Epidemiological data sources and sampling strategies. Epidemiological bias and confounding factors.	
4	Molecular methods applied to epidemiology. Molecular epidemiology of infectious and chronic diseases	
4	Elements of molecular evolution. Nucleic acids searches in international databases. Multiple alignment: algorithm	
Hrs	Practice	
2	Application of major epidemiological study models, analysis and interpretation of results.	
2	Calculation and interpretation of measures of morbidity (incidence and prevalence). Calculation and interpretation of association measures in epidemiology (odds ratio, relative risk and attributable risk).	
4	Reading and interpretation of methods and results presented in scientific articles of specialized journals	

## MODULE MEDICAL AND FORENSIC GENETICS

#### Prof. GREGORIO SEIDITA

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SUGGESTED BIBLIOGRAPHY		
<ul> <li>Strachan &amp; Read "Genetica Molecolare Umana . Garland Sciences 4 Edizione (2012).</li> <li>Butler J.M. "Fundamental of Forensic DNA typing". Academic Press (2010).</li> <li>Dispense fornite dal docente durante il corso in file pdf (articoli, linee guida e metodologie).</li> </ul>		
АМВІТ	50636-Discipline di base applicate alle biotecnologie	
INDIVIDUAL STUDY (Hrs)	102	
COURSE ACTIVITY (Hrs)	48	
EDUCATIONAL OBJECTIVES OF THE MODULE		
Medical Ecrensic Constics module aims to provide students with:		

Medical-Forensic Genetics module aims to provide students with: -the theoretical basis for understanding problems related to the analysis of genetic variation. It provides guidelines for

applying molecular genetics techniques;
Detailed knowledge of the main techniques of molecular biology and molecular genetics applied in the biomedical field;
Analysis of the main polymorphisms of forensic interest and the organization of a forensic genetics laboratory.

## **SYLLABUS**

Hrs	Frontal teaching
4	Genome structure and function. Genetic variability and measurement of variability. Organization of the Human Genome.
4	Gene and chromosomal mutations. Main techniques for the study of human genetic variations.
4	The main Genetic tests on the individual and their meaning. The distribution of genes in families and populations, mode of transmission of characters, Hardy Weinberg Equilibrium.
4	Counseling in medical genetics. Biological samples of forensic interest, evidence sampling techniques, evidences conservation and characterization. Extraction of DNA from biological samples of forensic interest. Quantization and DNA amplification.
4	Human STR loci: Biology and Genetics. STR forensic features. Capillary electrophoresis. Quality control and validation of methods procedures. The DNA database. Missing persons and mass disaster.
4	DNA Low Copy Number samples. Y chromosome STR markers. Analysis of Mitochondrial genome. Analysis of chromosome markers X. Genetic aspects of the forensic interest DNA polymorphisms (mutation rate; Hardy Weinberg theorem and its applications, Investigation and personal identification, Paternity Investigation).
3	Random Match Probability. limits and boundaries of Forensic Genetics.
3	The judicial inspection for sampling biological evidences. Predictive and related technical tests. DNA extraction from biological traces in Forensic Genetics and Molecular Diagnostics.
2	The Italian DNA Data Bank. Procedures for forensic genetics laboratories accreditation. Biostatistic in forensic genetics. The paternity investigations. Routines and exceptions.
Hrs	Practice
3	Analysis and design software for synthetic DNA sequences.
3	Pubmed and MyNCBI. Databases of public domain DNA. Management software and analysis of nucleotide and amino acid sequences. Analysis of the main information of a biological sequence (BLAST, CLUSTAL,)
3	Genetic counseling. Family trees and Risk-of-recurrence calculation. Tutorials on the Hardy Weinberg equation.
3	Technical investigation of genetic variation
2	DNA extracted genomic characterization: quantity and quality (Real time PCR). Amplification of genomic DNA.
2	Genetic characterization technologies for Low Copy Number DNA samples (low template DNA).