

### AYDIN ADNAN MENDERES UNIVERSITY COURSE INFORMATION FORM

Course Title		Introduction to	o Bioinformatio	~~					
Course Thie		millouucion	Domonnau						
Course Code		BIS529		Couse Leve	el	Second Cycle	e (Master's D	Degree)	
ECTS Credit	4	Workload	98 (Hours)	Theory	2	Practice	0	Laboratory	0
Objectives of	the Course							aspects of informa ptive repositories.	atics
Course Content								s, the analysis of	
			n interaction a	ind networks	, phylogene			next-generation se prediction, molect	
Work Placeme	ent	protein-protei	n interaction a	ind networks	, phylogene				
		protein-protei dynamics, ge	n interaction a netic linkage,	nd networks gene express	, phylogene sion.	etics, protein st	ructure and		

## **Assessment Methods and Criteria**

Method	Quantity	Percentage (%)
Midterm Examination	1	40
Final Examination	1	60

# **Recommended or Required Reading**

1	Claverie, J. M., & Notredame, C. (2006). Bioinformatics for dummies. John Wiley & Sons.	
2	Baxevanis, A. D., & Ouellette, B. F. (2004). Bioinformatics: a practical guide to the analysis of genes and proteins (Vol. 43). John Wiley & Sons.	
3	Zvelebil, M. J., & Baum, J. O. (2007). Understanding bioinformatics. Garland Science.	
4	Rashidi, H. H., & Buehler, L. K. (1999). Bioinformatics basics: applications in biological science and medicine. CRC press.	

Week	Weekly Detailed Course Contents			
1	Theoretical	Introduction to the course: DNA Databases		
2	Theoretical	Protein Databases		
3	Theoretical	DNA Sequencing and Assembly		
4	Theoretical	Translation, Mapping and Primers		
5	Theoretical	Protein Structure Web Tools		
6	Theoretical	Pairwise Comparison Scoring Matrices		
7	Theoretical	Database Similarity Searching		
8	Intermediate Exam	Midterm exam		
9	Theoretical	Multiple Alignment		
10	Theoretical	Motifs, Patterns, and Profiles		
11	Theoretical	Genomics - How were genomes sequenced? Genome Databases and the UCSC Browser		
12	Theoretical	Genome Browsers 2 Comparative Genomics		
13	Theoretical	Dealing with Gene Lists: Gene Ontology		
14	Theoretical	Introduction to Phylogenetic Analysis		
15	Theoretical	Literature review and discussion		
16	Final Exam	Final exam		

#### **Workload Calculation**

Activity	Quantity Preparation		Duration	Total Workload
Lecture - Theory	14	0	2	28
Reading	14	0	1	14
Individual Work	14	0	2	28
Midterm Examination	1	10	1	11



				Course mormation Form
Final Examination	1	15	2	17
		Тс	otal Workload (Hours)	98
		[Total Workload (	Hours) / 25*] = <b>ECTS</b>	4
*25 hour workload is accepted as 1 ECTS				

Learn	ning Outcomes
1	To be able to comprehend the importance of bioinformatics in the field of health care
2	Understand the basics of biological databases and use them in molecular biology and genetics
3	Identify comparison tools for two or more series and discuss similarity / noncompliance
4	To be able to understand the flow of information between protein-RNA and DNA and to analyze large-scale biological data using various bioinformatics based tools
5	Be able to discuss genomics, proteomics and pharmacogenetic approaches using bioinformatics tools

#### Programme Outcomes (Biostatistics Master)

1	To be able to understand the interdisciplinary interaction releated with biostatistics.
2	to be able to use Theoretical and practical knowledge at the level of expertise.
3	To be able to nterpret the information by integrating information from different disciplines and create new information
4	To be able to nalyze the problems encountered by using research methods
5	to be able to conduct a study as an independent specialist
6	To be able to formulate solutions for complex unpredictable problems encountered by developing new approaches and taking responsibility.
7	To be able to resolve problems in environments that require leadership.
8	To be able to evaluate and direct knowledge and skills with a critical approach at the level of expertise.
9	To be able to to give statistical advise at the begining stages of preparing health related projects
10	To be able to get the knowledge and the ability of using statistical packages

# Contribution of Learning Outcomes to Programme Outcomes 1: Very Low, 2: Low, 3: Medium, 4: High, 5: Very High

	L1	L2
P1	5	5
P2	3	5
P3	3	5
P4	3	3
P5	4	4
P6	3	5
P7	4	4
P8	3	4
P9	3	5
P10	3	4

