BIOINFORMATIC ALGORITHM									
1	Course Title:	BIOINFO	DRMATIC ALGORITHM						
2	Course Code:	BM5130							
3	Type of Course:	Optional							
4	Level of Course:	Second (	Cycle						
5	Year of Study:	1							
6	Semester:	2							
7	ECTS Credits Allocated:	6.00							
8	Theoretical (hour/week):	3.00							
9	Practice (hour/week):	0.00							
10	Laboratory (hour/week):	0							
11	Prerequisites:								
12	Language:	Turkish							
13	Mode of Delivery:	Face to f	ace						
14	Course Coordinator:	Doç. Dr.	GIYASETTİN ÖZCAN						
15	Course Lecturers:								
16	Contact information of the Course Coordinator:	Tel.:+90	ır Müh. Bölüm Binası, 1. kat, oda 107 (224) 294 2792 ozcan at uludag.edu.tr						
17	Website:								
18	Objective of the Course:	To learn methods and algorithms for analyzing high-volume biological data / signals. To learn how to obtain results that have meaning in medical terms with these methods.							
19	Contribution of the Course to Professional Development:	Understanding biological databases, making the algorithm designs necessary to process these databases							
20	Learning Outcomes:								
		1	Students learn the algorithms which are used to analze high volume biological data.						
		2	Students learn to use probabilistic prediction methods to solve problems.						
		3	In terms of medicine, students understand the benefits of the bioinformatics algorithms						
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	Occurs Occuts of	10							
21	Course Content:	una Cantanti							
Mook	Theoretical	Co	Proctice						
vveek 1	Theoretical Fundamental problems of bioinformalits computations	tics and	Practice						
2	Sequence Alignment Algorithms								
3	Short read sequence alignment								
4	Alignment against database, BLAST								
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5	Mult	iple S	Seque	nce A	lignm	ent											
6	Motif search algorithms																
7	Probabilistic algorithms																
8	Phylogeny Algorithms																
9	Next Generation Sequencing																
10	Gen	Genomic Integration															
11	Biological networks																
12	Secondary prediction																
13	Protein structure prediction																
14	Interaction with cancer drugs																
22							Mit	Richard Durbin, Sean R. Eddy, Anders Krogh, Graeme Mitchison Biological Sequence Analysis: Probabilistic Models of Proteins and Nucleic Acids , 1998									
23	Asse	esme	nt														
TERM L	LEARI	NING	ACTI	VITIES			N R	IUMBE	WE	WEIGHT							
Midterr	n Exa	am					1		25	25.00							
Quiz							0		0.0	0.00							
Home	work-	proje	ct				2		15.	15.00							
Final E	' '						60.	60.00									
Total							4		100	0.00							
	Activites						1	Number Duration (hour) Total Work Load (hour									
Theoretical Exam to Success Grade						OV,	3.00				42.00						
Practic	Practicals/Labs								)		0.00			0.00			
Measil	Measurement and evaluation is carried out according to									g to							
	Homeworks								0 0.00 0.00								
Pr <b>2</b> #ect	price ECTS / WORK LOAD TABLE							(	0 0.00				0.00				
Field S	d Studies							C	)			0.00			0.00		
Midterr	erm exams							1	1 30.00				30.00				
Others	rs							C	0 0.00				0.00				
Final E	Exams						1	1 52.00				52.00					
Total V	Work Load							180.00									
Total w	ork lo	oad/ :	30 hr							6.00							
ECTS (	S Credit of the Course												6.00				
25 CONTRIBUTION OF LEARNING OUTCOMES TO PROGRAMME QUALIFICATIONS																	
	ı	PQ1	PQ2	PQ3	PQ4	PQ5	PQ6	PQ7	PQ8	PQ9	PQ1 0	PQ11	PQ12	PQ1 3	PQ14	PQ15	PQ16
ÖK1	į	5	4	3	1	1	1	0	0	0	0	0	0	0	0	0	0
ÖK2	4	4	4	4	1	1	1	0	0	0	0	0	0	0	0	0	0
ÖK3	·	1	1	4	1	1	1	0	0	0	0	0	0	0	0	0	0
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LO: Learning Objectives PQ: Program Qualifications																	

Contrib	1 very low	2 low	3 Medium	4 High	5 Very High
ution					
Level:					