

# INTRODUCTION TO BIOINFORMATICS ALGORITHMS

<b>1</b>	Course Title:	INTRODUCTION TO BIOINFORMATICS ALGORITHMS	
<b>2</b>	Course Code:	BMB4017	
<b>3</b>	Type of Course:	Optional	
<b>4</b>	Level of Course:	First Cycle	
<b>5</b>	Year of Study:	4	
<b>6</b>	Semester:	7	
<b>7</b>	ECTS Credits Allocated:	5.00	
<b>8</b>	Theoretical (hour/week):	3.00	
<b>9</b>	Practice (hour/week):	0.00	
<b>10</b>	Laboratory (hour/week):	0	
<b>11</b>	Prerequisites:		
<b>12</b>	Language:	Turkish	
<b>13</b>	Mode of Delivery:	Face to face	
<b>14</b>	Course Coordinator:	Doç. Dr. GIYASETTİN ÖZCAN	
<b>15</b>	Course Lecturers:		
<b>16</b>	Contact information of the Course Coordinator:	Bilgisayar Müh. Bölüm Binası, 1. kat, oda 107 Tel.:+90 (224) 294 2792 email: gozcan at uludag.edu.tr	
<b>17</b>	Website:		
<b>18</b>	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.	
<b>19</b>	Contribution of the Course to Professional Development:	To understand biological databases, to be able to design the necessary algorithms to process these databases	
<b>20</b>	Learning Outcomes:		
		1	Students learn the algorithms which are used to analyze 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
		4	
		5	
		6	
		7	
		8	
		9	
		10	
<b>21</b>	Course Content:		
		<b>Course Content:</b>	
Week	Theoretical	Practice	
1	Fundamental problems of bioinformatics and its computations		
2	Sequence Alignment Algorithms		
3	Short read sequence alignment		
4	Alignment against database, BLAST		

5	Multiple Sequence Alignment	
6	Motif search algorithms	
7	Probabilistic algorithms	
8	Phylogeny Algorithms	
9	Next Generation Sequencing	
10	Genomic Integration	
11	Biological networks	
12	Secondary prediction	
13	Protein structure prediction	
14	Interaction with cancer drugs	

22	Textbooks, References and/or Other Materials:	1. Richard Durbin, Sean R. Eddy, Anders Krogh, Graeme Mitchison Biological Sequence Analysis: Probabilistic Models of Proteins and Nucleic Acids , 1998
----	---	---

23	Assesment	
----	-----------	--

TERM LEARNING ACTIVITIES		NUMBER	WEIGHT
Midterm Exam		1	25.00
Quiz		0	0.00
Home work-project		2	15.00
Final Exam		1	60.00
Total		4	100.00

Activites	Number	Duration (hour)	Total Work Load (hour)
Contribution of Final Exam to Success Grade	60.00		
Theoretical	14	3.00	42.00
Practicals/Labs	0	0.00	0.00
Measurement and Evaluation Techniques Used in the Course	0	0.00	0.00
Self study and preparation	0	0.00	0.00
Homeworks	0	0.00	0.00
Projects	0	0.00	0.00
Field Studies	0	0.00	0.00
Midterm exams	1	50.00	50.00
Others	0	0.00	0.00
Final Exams	1	58.00	58.00
Total Work Load			200.00
Total work load/ 30 hr			5.00
ECTS Credit of the Course			5.00

24	<b>ECTS / WORK LOAD TABLE</b>
----	-------------------------------

25	<b>CONTRIBUTION OF LEARNING OUTCOMES TO PROGRAMME QUALIFICATIONS</b>															
	PQ1	PQ2	PQ3	PQ4	PQ5	PQ6	PQ7	PQ8	PQ9	PQ10	PQ11	PQ12	PQ13	PQ14	PQ15	PQ16
ÖK1	5	5	4	4	3	2	1	1	1	1	2	3	0	0	0	0
ÖK2	5	5	4	4	3	2	1	1	1	1	2	3	0	0	0	0
ÖK3	5	5	4	4	3	2	1	1	1	1	2	3	0	0	0	0

**LO: Learning Objectives PQ: Program Qualifications**

<b>Contribution Level:</b>	<b>1 very low</b>	<b>2 low</b>	<b>3 Medium</b>	<b>4 High</b>	<b>5 Very High</b>
----------------------------	-------------------	--------------	-----------------	---------------	--------------------