



Democritus University of Thrace, Kavala, Greece

School of Science Department of Informatics

Department of European and International Programmes – Erasmus+ Agios Loukas, 654 04, Kavala University Campus, Greece

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Proposed Course for incoming Erasmus students¹

Responsible for the course	Professor Stergios Papadimitriou
(lecturer)	0030 2510 462 323
(name, phone number, e-	sterg@cs.duth.gr
mail address)	_
Title of the Course	Introduction to Bioinformatics
ECTS credits	5
Short contents of the	
course	Introduction to Bioinformatics algorithmic techniques and computer software tools.
	Study of Biological Databases, their organization, and the data retrieval process. Example databases as NCBI, EBI, KEGG and PDB will be used for web based data retrieval and for studying their programmatic interface.
	Another subject of study is the important one of sequence alignment and biological database searching. Dynamic programming based approaches will be introduced. The proper definition of distance metrics between biological sequences will also be elaborated. Next, approximation based heuristic suboptimal searching algorithms, as the BLAST one will be studied.
	Also, the next generation sequencing technologies and the DNA microarrays will be another important subject of study. The important novel domains of rational drug discovery and personalized medicine based on the individual genome analysis will also be discussed.
Aim of the course and target audience	 The course will introduce students to Bioinformatics Target audience: Undergraduate students of Informatics/ Computer Science OR Education
Teaching Methods duration	Lectures: 26 hours
and Evaluation	Hands-on exercises: 26 hours
	Evaluation: 100% Individual AND/OR Group Assignments
Offered Period	Spring semester

Indicative bibliography	1. Introduction to Bioinformatics, Neil C. Jones and Pavel A. Pevzner, Addison Wesley, 2003
	2. Bioinformatics: An active learning approach, Pavel A. Pevzner, Addison Wesley, 2015, Vol. 1 and 2
	3. Arthur M. Lesk, Introduction to Bioinformatics, Fourth Edition, Oxford, 2014
	4. Bioinformatics and Functional Genomics, third edition, Jonathan Pevzner, WILEY Blackwell, 2015
	5 Paul A Gagniuc PhD Algorithms in Riginformatics WILEY 2021

- 5. Paul A. Gagniuc, PhD, Algorithms in Bioinformatics, WILEY, 2021
- 6. Hannes Hauswedell, Sequence Analysis and Modern C++, Springer 2022
- 7. Basant K. Tiwary, Bioinformatics and Computational Biology, Springer 2022

¹ Could be easily used and offered for TS movement to our Erasmus partners