

CSE 308-408

Bioinformatics: Issues and Algorithms

Fall 2007 • TuTh 2:35 pm – 3:50 pm • Packard Lab 208

- Instructor** **Professor Daniel Lopresti**
 Office PL 404B • Ext 85782 • Email dal9@lehigh.edu
 Office Hours 4:00 – 6:00 W (or by appointment)
- Biology Advisor** **Professor Jutta Marzillier**
 Ext 84902 • Email jym2@lehigh.edu
- Grading Assistant** **Wen Cheng**
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- Textbooks** *An Introduction to Bioinformatics Algorithms*, Neil C. Jones and Pavel A. Pevzner
 MIT Press, ISBN 0-262-10106-8. (IBA)
- Bioinformatics Biocomputing and Perl*, Michael Moorhouse and Paul Barry
 John Wiley & Sons, ISBN 047085331X. (BB&P)
- Blackboard** Lecture slides, assignments, etc. will be available @ <http://ci.lehigh.edu>

Grading	CSE 308	CSE 408
• Homework assignments =	25% of grade	20% of grade
• Programming assignments =	25% of grade	20% of grade
• Final project or paper =	50% of grade	50% of grade
• Scribe duty (CSE 408 only) =	n/a	10% of grade

- Notes**
- Homework / programming assignments will generally be posted to Blackboard by 9:00 am and due by 5:00 pm on the specified day. Carefully follow all instructions when naming your programs and submitting your work.
 - Assignments turned in up to one week late will received one-half credit. After that point, no credit will be given. Extensions must be approved by Professor Lopresti.
 - If you already consider yourself proficient in Unix and Perl programming, you may choose to skip the lectures marked with an asterisk (*). Please skim the indicated reading, however, and review the lecture notes online in Blackboard.

Date	Topics	Readings	Other Activities
Tu 8/28	Course introduction		
Th 8/30	Intro to molecular biology <i>genetics and genomes, analyzing DNA</i>	IBA 3, BB&P 1	
Tu 9/4	Intro to algorithms <i>correctness, recursion, iteration, time complexity</i>	IBA 2, BB&P 2	HW #1 out
Th 9/6	Intro to Unix; Perl programming 1 * <i>getting started, control flow, variables, arrays, hashes</i>	BB&P 3-4	
Tu 9/11	Perl programming 2 * <i>subroutines, scoping rules, file I/O</i>	BB&P 5-6	HW #1 due
Th 9/13	Perl programming 3 <i>pattern matching, regular expressions, sorting</i>	BB&P 7-8	HW #2 out
Tu 9/18	Restriction mapping <i>biology, full and partial digests, brute-force and practical algorithms</i>	IBA 4.1-4.3	
Th 9/20	Motifs, search trees <i>regulatory motifs, profiles, search trees, motif-finding</i>	IBA 4.4-4.9	HW #2 due

Date	Topics	Readings	Activities
Tu 9/25	Genomics (Prof. Marzillier)	Reading TBA	HW #3 out
Th 9/27	Proteomics (Prof. Marzillier)	Reading TBA	
Tu 10/2	Genome rearrangements <i>biology, sorting by reversals, greedy and approximation algorithms</i>	IBA 5	HW #3 due
Th 10/4	Sequence comparison & alignment 1 <i>biology, comparison models, dynamic programming, global alignment</i>	IBA 6.1-6.7	HW #4 out
Tu 10/9	No class		Pacing Break
Th 10/11	Sequence comparison & alignment 2 <i>local alignment, gap penalties, multiple alignment</i>	IBA 6.8-6.10	HW #4 due
Tu 10/16	Sequence comparison & alignment 3 <i>saving time and space, divide-and-conquer</i>	IBA 7	HW #5 out
Th 10/18	Sequencing & assembly 1 <i>biology, graph theory, shortest superstrings, sequencing by hybridization</i>	IBA 8.1-8.9	
Tu 10/23	Sequencing & assembly 2 <i>protein sequencing, spectral analysis</i>	IBA 8.10-8.15	HW #5 due
Th 10/25	Genetic pattern matching 1 <i>repeat-finding, suffix trees</i>	IBA 9.1-9.5	HW #6 out
Tu 10/30	Genetic pattern matching 2 <i>heuristics: FASTA and BLAST</i>	IBA 9.6-9.8	
Th 11/1	Tools, datasets, and applications <i>EMBL/GenBank, SWISS-PROT/PIR, ClustalW, BLAST</i>	BB&P 17-18	HW #6 due
Tu 11/6	DNA microarrays	Reading TBA	HW #7 out
Th 11/8	Clustering for expression analysis <i>biology, hierarchical clustering, k-means</i>	IBA 10.1-10.4	
F 11/9			Project proposals due
Tu 11/13	Evolutionary trees 1 <i>biology, distance-based tree reconstruction, additive trees</i>	IBA 10.5-10.7	HW #7 due
Th 11/15	Evolutionary trees 2 <i>character-based tree reconstruction, parsimony (small and large)</i>	IBA 10.8-10.11	
Tu 11/20	RNA and protein structure prediction <i>RNA secondary structure prediction, protein threading</i>	Reading TBA	
Th 11/22	No class		Thanksgiving Break
Tu 11/27	Bioethics	Reading TBA	
Th 11/29	History of the Genetic Code	Reading TBA	
Tu 12/4	Student final project / paper presentations I		
Th 12/6	Student final project / paper presentations II		
F 12/7			Final projects due

Accommodations for Students with Disabilities If you have a disability for which you are or may be requesting accommodations, please contact both your instructor and the Office of Academic Support Services, University Center C212 (610-758-4152) as early as possible in the semester. You must have documentation from the Academic Support Services office before accommodations can be granted.

Academic Integrity The work you submit in CSE 308-408 must be entirely your own. While we encourage you to discuss basic concepts and strategies with friends and classmates, the copying or sharing of solutions to homework or programming assignments, or of final papers or projects, is never acceptable. Such cases will be referred to the University Committee on Discipline and, if found guilty, you may be given the failing grade WF in the course.

You should keep in mind that computer programs exhibit an individual's "style" just as much as other forms of authorship. Changing variable names, editing comments, or making other trivial updates in an attempt to hide plagiarism is rarely effective.

If you have questions about this policy at any point throughout the semester, ask. It is far better to be safe than sorry when your academic career may be on the line.