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Fall 2010

### CS/BIO 471/671: Algorithms for Bioinformatics

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# CS/BIO 471/671 – ALGORITHMS FOR BIOINFORMATICS

*Fall, 2010*

## Course Description

Theory-oriented approach to the application of contemporary algorithms to bioinformatics. Graph theory, complexity theory, dynamic programming and optimization techniques are introduced in the context of application toward solving specific computational problems in molecular genetics. 4 credit hours.

## Meeting Time and Place

12:15 – 1:20      Monday and Wednesday      155 Russ Engineering Center (RC)

## Textbooks

J. Setubal and J. Meidanis (1997), *Introduction to Molecular Biology*, Brooks/Cole Publishing, ISBN: 0-534-95262-3

D. Krane and M. Raymer (2003), *Fundamental Concepts of Bioinformatics*, Benjamin Cummings, ISBN: 0-8053-4633-3

## Instructors and Office Hours

Dr. Michael Raymer  
391 Joshi  
775-5110

[michael.raymer@wright.edu](mailto:michael.raymer@wright.edu)  
<http://www.wright.edu/~michael.raymer>  
Office hours: Mon & Wed, 4:00 – 6:00 pm  
or by appointment.

Dr. Dan Krane  
225B BH  
775-2257

[dan.krane@wright.edu](mailto:dan.krane@wright.edu)  
Office hours: Mon & Wed, 8:45 – 10:45 am  
or by appointment.

## Course Web Page

The course web page will be the primary method for distributing important announcements, course material, class notes, etc. Please check the page often. Login to the campus WebCT system using your CATS username and password. The URL is: <http://wisdom.wright.edu>

You can find an archive of the course materials at: <http://birg.cs.wright.edu/cs471>

## Grading

Course grades will be determined as follows:

25% Midterm Exam  
40% Final Exam  
35% Project/homework assignment(s)

*Final grades will be based on the standard university-wide score divisions (i.e. 90%, 80%, 70%, etc.). However, the instructors may choose to curve the final grades depending on the distribution of scores at the end of the term.*

## **Tentative Lecture Schedule**

### **Week 1 – Review: Molecular biology and problem solving methods**

- Tools and methods of molecular biology
- Exhaustive, greedy, divide-and-conquer, and other approaches

### **Week 2 – Paper discussion: Aerts; Paper presentation assignments**

### **Week 3 – Sequence alignments and dynamic programming**

- Recursion
- Dynamic programming
- Sub-problem optimality

### **Week 4 – Paper discussions and term project planning**

### **Week 5 – Algorithm complexity, P and NP (Class notes)**

- Complexity of algorithms
- Big-O notation
- Complexity classes, P and NP
- Some classic algorithms

### **Week 6 – Sequencing and Comparative Genomics (Ch4. Setubal & Meidanis)**

- Sequence annotation
- Synteny
- Models and tools for fragment assembly

### **Week 7 – Mapping of DNA (Ch. 5, Setubal & Meidanis)**

- Genetic mapping
- Restriction site mapping, hybridization mapping
- LOD Scores, QTLs, Sulston scores

### **Week 8 – Phylogenetics (Chs. 4 & 5, Krane & Raymer)**

- Trees and distances
- Distance-based and character-based approaches
- Bootstrapping and confidence
- Selective forces and population size

### **Week 9 – Suffix tries and inverted repeats (Class notes)**

- Trees, tries, and PATRICIA trees
- Inverted repeats and microRNA

### **Week 10 – Project results presentations/discussion**

## **Policies & Notes**

**Homework assignments** are due by the end of the lecture period on the assigned due date. Late homework submissions cannot be accepted.

**Collaboration** in learning is encouraged, as discussion of the course contents with other students is an important part of the learning process. However, it is expected that **course assignments will be completed on an individual basis unless the assignment states otherwise.**

Students may not, under any circumstances, work together in actual implementation of any course assignment unless the assignment specifically allows group submissions. Do not allow other students to view or copy your code or writing. Code sharing, including code from previous quarters, is strictly disallowed. Copying or significant collaboration on any graded assignment will be considered a violation of university guidelines for academic integrity and reported to the Office of Judicial Affairs. The Code of Student Conduct can be viewed at <http://www.wright.edu/students/judicial/conduct.html> or a hand copy can be obtained from the Office of Student Judicial Services in W035 Student Union. If you have any questions about these policies, it is your responsibility to discuss them with the instructor of the course or a representative of the Office of Judicial Affairs as soon as possible.

If the same work is turned in by two or more students, all parties will be held equally accountable for violation of academic integrity. In other words, *you are responsible for ensuring that other students do not have access to your work.* If you suspect that your work material has been compromised, notify an instructor immediately.