Program in Biological Sciences Biol. Sci. 393: Genetic Analysis

Lectures: MWF 2:00 - 2:50 PM, Tech M128

Discussion F 3:00 - 3:50 PM, Tech M128

Exams: Midterm (May 1, 2-4 PM), Final (June 10, 3-5 PM)

Instructor: Erik Andersen, Department of Molecular Biosciences, Cook 3125,

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Textbook: Introduction to Genetics Analysis by Griffiths *et al.* 10th edition

Website: www.andersenlab.org/bio393

Course Format

This course will have lectures on Mondays and Wednesdays. On Fridays, the class will alternate between quizzes and problem-solving sessions. Participation in lecture and Friday discussions is required. The course will have four quizzes, four problem sets, one midterm, and a final examination.

Problem sets and guizzes

Problem sets will be distributed on Mondays nearly two weeks before the problem-solving session. Students must turn in their completed problem sets and be prepared to work through the problems in class on Fridays. Problem-solving sessions are also meant to discuss questions about the lecture material. Quizzes will assess student progress through basic concepts. Students can take each of the quizzes twice to improve scores. Students will receive their graded guizzes in the following lecture.

Grading

Each examination and quiz will cover material from the preceding lectures, quizzes, and problem sets. Successive exams and quizzes assume you remember concepts and methods discussed earlier in the course. No make-up exams or quizzes will be given. If you get 80% or greater on a quiz, you receive all 20 points. If you get less than 80% on a quiz the first time through, you may take the quiz again on the next scheduled quiz day. If you receive less than 80% the second time through, you will receive that score for the quiz. Problem sets are not graded. Full points are awarded for completed problem sets due on Friday and the student participates in the class discussion on Friday.

Point distribution		
Problem sets	15%	60 points (15 pts each)
Quizzes	20%	80 points (20 pts each)
Participation	5%	20 points
Midterm	30%	120 points
Final	30%	120 points

Any student with a disability requesting accommodations is required to register with AccessibleNU (<u>accessiblenu@northwestern.edu</u>; <u>847-467-5530</u>) and present an accommodation letter from AccessibleNU to the course instructor, preferably within the first two weeks of class. All information will remain confidential.

All work submitted for this class must be your own. Suspected violations of academic integrity will be reported to the Dean's Office. For more information on Northwestern policies on academic integrity, see http://www.weinberg.northwestern.edu/handbook/integrity/index.html.

BIOL SCI 393 (Spring 2015) Genetic Analysis

Date	Lecture topic	
Mon. March 30	Mendelian Inheritance, Basic probability, PS#1 out	
Wed. April 1	Chromosome theory, mitosis, and meiosis	
Fri. April 3	QUIZ #1	
Mon. April 6	Recombination and mapping	
Wed. April 8	Screens, selections, mutants, and dosage	
Fri. April 10	PROBLEM SET #1 Due	
Mon. April 13	Complementation, PS#2 out	
Wed. April 15	Genetic interactions: epistasis	
Fri. April 17	QUIZ #2	
Mon. April 20	Genetic interactions: enhancement and suppression	
Wed. April 22	Principles and methods of genetic analysis I	
Fri. April 24	PROBLEM SET #2 Due	
Mon. April 27	Principles and methods of genetic analysis II	
Wed. April 29	Principles and methods of genetic analysis III	
Fri. May 1	MIDTERM EXAMINATION	
Mon. May 4	Developmental genetics I, PS#3 out	
Wed. May 6	Developmental genetics II	
Fri. May 8	QUIZ #3	
Mon. May 11	Behavioral genetics	
Wed. May 13	Model organisms in genetics (from cells to humans)	
Fri. May 15	PROBLEM SET #3 Due	
Mon. May 18	Human variation and allele frequency spectrum, PS#4 out	
Wed. May 20	Pedigrees and linkage mapping	
Fri. May 22	QUIZ #4	
Mon. May 25	NO CLASS - Memorial Day	
Wed. May 27	Linkage disequilibrium and population structure	
Fri. May 29	PROBLEM SET #4 Due	
Mon. June 1	Complex traits and GWAS	
Wed. June 3	Epigenetics and transgenerational effects	
Fri. June 5	Genome sequencing, exome sequencing, and what do we do with these data	