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SC-103 508.286.3970 Office Hours: MW 11:30-12:30, 2-3

Meeting Times: Monday, Wednesday, Friday 10:30 - 11:20 Room 154, SciCenter or csLab-A118b

Text: **Perl for Exploring DNA** by LeBlanc and Dyer (Oxford University Press 2007)



10 GOALS for this course

- (0) You are at a cocktail party and the topic of genomes comes up. You are able to recall significant phrases, terms, and techniques and your understanding of the main ideas and concepts enables you to lead the conversation for a while. Your friends raise their eyebrows.
- (1) You learn to **identify** and **classify** problems that are candidates for a computer to handle.
- (2) You demonstrate the ability to think algorithmically, breaking what originally seems like an overly complicated problem into a series of smaller, manageable tasks.
- (3) You learn to craft creative solutions by "writing software" ("to program", "to script").
- (4) You appreciate when to **include external modules** of previously written software.
- (5) You design experiments to first solve small computationally-intensive tasks (e.g. on one gene sequence) and then scale your solutions to very large sets of data (e.g. all genes in a genome).
- (6) You **apply** problem solving strategies previously learned to novel situations.
- (7) You learn to professionally **document** your software and produce quality summaries, graphs, and reports of your results.
- (8) You begin to appreciate genome-wide **microarray experiments** and the accompanying software that is required to analyze results.
- (9) You feel empowered to **evaluate** the ethical implications of your work and learn to **appraise**, **critique**, and **defend** your own as well as the work of others.

Connections: The Bigger Picture

This course is part of the *connection "Genes in Context"* with Philosophy 111 (Ethics). Throughout the semester, students will be exposed to the ethical aspects of living in a post-genomic world and the increasing use and challenges of sequenced genomes as applied to "**personalized medicine**". A series of assignments and events with students from other classes will include:

- A showing of the movie GATTACA and follow-up discussion.
- Talks and discussions with a genetic counselor and professor of bioethics.
- Student-produced, one-minute YouTube "commercials" of companies currently promoting and selling medical profiles based on individual genomes. The commercials will be framed from one of two points of view: (i) from the point of view of the company (e.g., 23andME, deCODEme, Navigenics, etc) or (ii) from a consumer advocacy point of view.

DNA

Catalog Description

An amazing blend of science, computing, and mathematics emerges when considering the molecule "Deoxyribonucleic Acid" (DNA). DNA is the blueprint of life for all organisms on Earth and throughout evolutionary time. Its distinctive and beautiful physical nature, a double helix of four bases, maps onto its functionality as a



ature, a double helix of four bases, maps onto its functionality as a bearer of information, generation after generation. Fully sequenced genomes including the human genome and hundreds of microbial genomes have become the starting point for attempts to answer a wide range of biological and quantitative questions.

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This DNA-242 course satisfies the Quantitative Analysis (QA) requirement.

Exact pages to reading and use of web sites will be **assigned in class**. URLs are listed as assigned/needed in the onCourse (Moodle) website.

Web Resources:

From the PBS show NOVA: "Cracking the Code of Life" Watch the program in 16 online video segments http://www.pbs.org/wgbh/nova/genome/program.html

From the Dolan DNA Learning Center & Cold Spring Harbor Laboratory: "DNA From the Beginning". DNA from the Beginning is organized around key concepts. http://www.dnaftb.org/dnaftb/

DNA interactive: http://www.dnai.org/a/index.html

NCBI: National Center for Biotechnology Information NCBI creates public databases, conducts research in computational biology, develops software tools for analyzing genome data, and disseminates biomedical information. http://www.ncbi.nlm.nih.gov/

A Perl FAQ

http://theoryx5.uwinnipeg.ca/CPAN/perl/pod/perlfaq1.html

Notes on grading, due dates, and submission procedures

Because the lab and class preparation is costly both in money (\$) and time, you must attend all classes and labs. When **homework** is "due in class", this means at the beginning of the class on that particular day, e.g., we will collect your homework at the start of class. Homeworks that arrive after the class and up to one day late will lose 10%. No homeworks are accepted after one-day late.

Programs are due on various dates (see detailed syllabus); however, since we know from experience that many students like to use the last night for testing, we will allow you to submit your programs until 5am the following day. For example, Program #1 is due THUR, Sept 17th, but you can submit it electronically until 5am FRI, Sept 18th -- Careful! The course website (onCourse) makes it appear as if the program is due on Friday, but remember, Friday at 5am! Programs submitted after this time will lose 20% each day it is late upto two days late. A program submitted more than two days late will not be accepted and will receive a grade of zero. In addition to **electronically submitting** your Perl programs (more instructions will be given in class), you must submit a **hardcopy printout** of your program in class after it is due. Note that Perl programs must *always* be **printed** in **landscape mode** and your source listing must be **stapled** to your sample output and documentation (README) file. (Your professor will say @%^\$#&* and will deduct points if you do not staple.)

- based on attendance / participation in all sessions

- see detailed schedule (usually at start of class)

- one-minute "personalized medicine" video

- installation of Perl environment

- chalking RegEx in sidewalk art

- RegEx (Part I)

It is expected that you spend at least 2 hours on reading and practice problems for every hour of lecture. This computes to at least 6 hours of work outside of class per week. This should be done throughout the semester. Please assume all deadlines are fixed. Obviously, see your professor if you know of a conflict beforehand.

Honor Code Revisited: It goes without saying that all submitted work will be the student's own, in keeping with the Wheaton Honor Code. For labs, you may get help from fellow classmates, but remember that all submitted work must be your own. All homeworks, Perl programs, and other hand-written work must be your own from beginning to end unless otherwise noted in the instructions (e.g., paired work).

Guest Lectures

Jenny Lanni, Biology	"The Central Dogma", Sept. 4 th
Teresa Celada, Philosophy	"Designer Babies", Oct. 2 nd
Mike Kahn, Statistics	"Randomness", Oct. 30 th

YOUR GRADE

5% ATTENDANCE/PARTICIPATION

10% QUIZZES (5)

15% HOMEWORKS

Sept 4	Homework #1 – 1 point
Sept 11	Homework #2 – 2 points
Sept 30	Homework #3 – 10 points
Oct 02	Homework #4 – 2 points

50% PROGRAMS (six)

Sept 17	Program #1 (5%)	- String Play
Sept 24	Program #2 (5%)	- Chargaffin Counts
Oct 2	Program #3 (10%)	- Gene Finder
Oct 16	Program #4 (10%)	- "eLmer" a motif finder
Oct 30	Program #5 (10%)	- Fuzzy Olfactory Gene Counter
Nov 18	Program #6 (10%)	- Building a Concordance
	-	-

20% Final **Project**

Nov 23	Written Proposal (Intro & Methods)	(5%)
Dec 7, 9, 11	Oral Presentation	(5%)
Dec 11	Final Paper, Software, and Results	(10%)

Throughout the semester, you will have the opportunity to show Superior Effort, for example, a homework or programming assignment has additional steps. Remember, when determining your final, overall grade: an 'A' is superior, 'B' is above average, 'C' is average effort, etc.

As you can see, the final projects are a significant part of your final grade (20%). Your professor will determine the team pairs. Each pair of students will be given a certain number of points and it will be up to the pair to determine how those points should be allocated. For example, if the pair was given 150 points, then the pair could decide that Person A, who did more of the work, should get 80 and Person B should get 70. If they pair felt they each did the same amount of effort, then each would get 75 points. Each of the three parts (Proposal, Oral Presentation, Final Paper including your software and results) will receive a separate grade.

Note: Bonus Points will be awarded to students who discover errors (especially technical and factual errors) in the book. The number of points awarded will be determined by the instructor. Before you report an error, please check the Errata to see if someone else has already found this error (see onCourse link to the Errata).

Sept 02, WED print "hello DNA";

Reading:

Text: Acknowledgements, Preface, Chapters 1 and 2. Reading on Central Dogma (read before class on Friday); see onCourse **Homework #1** (Due at *start* of class, Friday Sept 4): Install Perl summary.

Sept 04, FRI

Guest Lecture: Dr. Jenny Lanni, Biology – "The Central Dogma" **Reading:** Text: Ch. 4 p47-61 (you need to read this **before** next WED's lab)

Bring your textbook to lab next WED.

Week 2

Sept 07, MON – Labor Day <no class>

Sept 09, WED (meet in csLab)

Quiz #1 – Collectively we give a brief Central Dogma lecture. (See Homework #1).

csLab: Playing with **Reg**ular **Ex**pressions ("**Regex**"):

Regex: a tool for searching for patterns in DNA sequences—Be sure to have the book with you; we will be using Chapter 4 "String Play with Regular Expressions".

Homework #2 (Due at start of class, Friday Sept 11): Finish Part I handout on Regex

Opportunity for <u>Superior Effort</u>: If you really get into regex's, complete Part II ... and even make yourself a regex bookmark.

Fun Readings:

Agee, J. (1996). <u>So many dynamos! And other palindromes</u>. Farrar, Straus, Giroux Publishers, New York, NY.

Agee, J. (2000). <u>Elvis Lives and other anagrams.</u> Farrar, Straus, Giroux Publishers, New York, NY.

Sept 11, FRI

Homework #2 (Playing with Regex – Part I) due at start of class.

Good practices when printing output, using variables, and built-in functions; Introduction to "genic" and "intergenic" regions and codons.

Intro to Perl: "hello DNA Land", print, variables, built-in functions

length, lc, uc, reverse, index, substr, tr, s
Reading:

Text: Ch. 3 p17-46 and some reading on "algorithms" p203-211

a1 Specification is ready: (see onCourse for Starter Kit)

a1 algorithm is due MON, Sept 14

a1 program is due Thursday, Sept 17

Sept 14th MON a1 algorithm is due in class

csLab - playing with Perl ... strings, strINGS, STRINGS ... index, substr, reverse, lc, uc and string operators tr and concatenation (.)

Sept 16th WED

a1 submission is due tomorrow, THUR Sept 17 (really, Friday by 5am)

Assign one-minute videos of "personalized medicine" companies e.g., deCODEme, 23andMe, Knome, Illumina, Pathway Genomics, etc. (Due: Wed, Sept 30 in class)

more Perl

Sept 18th FRI

a2 Specification is ready: (see onCourse for Starter Kit)
a2 algorithm is due MON, Sept 21
a2 program is due Thursday, Sept 24

Using Perl to do calculations: arithmetic expressions

Specifications "Specs" for Program #2 "Chargaffian Counts" and instructions about completing the program.

Reading:

Text: Ch. 5 (all) and Ch. 8 Reading from Files p135-142

Week 4

Sept 21st MON

a2 algorithm is due in class

csLab: good practices of using arithmetic operators, precedence, math functions, and formatted output with printf

Using a subroutine written by someone else (e.g., readInDNA)

Quiz #2 next time on Wed, Sept 23rd at start of class

Sept 23rd WED Quiz #2 at start of class a2 due on Thursday, Sept 24

conditional control (if-elsif-elseif-else) **Reading:** Text: Ch. 6 p93-98, 103-106 and Ch. 7 (all)

Sept 25th FRI

a3 Specification is ready: (see onCourse for Starter Kit)a3 algorithm is due MON, Sept 28a3 program is due Friday, Oct 2

one-minute videos due next Wed, Sept 30

Transcription - Translation ... and a3 specification

Week 5

Sept 28th MON

a3 algorithm is due in class

Good practices of using if-else; using if and index together; Using subroutines revisited

Specifications "Specs" for Program #3 "Gene Finder" and instructions about completing the program.

Sept 30th WED

one-minute videos due today

Regex Chalk: due Friday morning around the dimple (rain date, Sat morning)

csLab – practice with if and index

Reading (in preparation for Friday's guest lecture): TBA

Oct 2nd FRI

Regex Chalk: <u>Due in morning</u> before classes

a3 due today

Guest Lecture: Dr. Teresa Celada, Philosophy -- "Designer Babies"

Oct 5th MON

a4 Specification is ready: (see onCourse for Starter Kit) a4 algorithm is due FRI, Oct 9th a4 program is due Friday, Oct 16th

while loops, greediness in regular expressions, commenting those tough regex

Introduction to gene regulation

Reading:

Text: Ch. 6 p98 – 107 (if-elsif-else) and while loops, p107-115

Oct 7th WED

Quiz #3

csLab - p114 #1,2,3,6

Oct 9th FRI -

a4 algorithm is due

More looping, more looping, more looping, more looping

Week 7

Oct 12th MON and TUES 13th "Fall Break"

Oct 14th WED

Introduction to modules and BioPerl

Oct 16th FRI a4 due today

More BioPerl

Oct 19th MON

a5 Specification is ready: (see onCourse for Starter Kit) a5 algorithm is due FRI, Oct 23rd a5 program is due Friday, Oct 30th

csLab -- BLAST Lab (Homework #6; due at the end of lab)

Specifications "Specs" for Program #5 "Fuzzy Olfactory Gene Finder" and instructions about completing the program

Olfaction and seven trans-membrane proteins (Part I)

Reading:

Text: 13 (all) - Randomness - "So, like totally random, dude"

Oct 21st WED Quiz #4

Using your BLAST output as input ... Olfaction and seven trans-membrane proteins (Part II)

Oct 23rd FRI

a5 algorithm is due

csLab: more subroutines and Perl's rand

Week 9

Oct 26th MON

Tweeking regex

Reading:

Text: 12 (all) - Regex Revisited

Oct 28th WED

Introduction to arrays and Perl's split and join functions Reading: Text: 9 – Arrays, p151-168

Heads/Tails homework - due at start of guest lecture on Friday, 30th

Oct 30th FRI

a5 due today

Guest Lecture: Dr. Mike Kahn -- "Randomness"

Week 10

Nov 02nd MON

csLab: Intro to microarray analysis - MagicTool software

Nov 04th WED

csLab: Continuation of microarray analysis - MagicTool software

Nov 5th THUR

Superior Effort: Join us in csLab to help the Genetics class use Magic Tool, 2-5pm.

Nov 6th FRI

a6 Specification is ready: (see onCourse for Starter Kit)
 a6 algorithm is due WED, Nov 11th
 a6 program is due Wed, Nov 18th

Sorting with arrays and hash tables **csLab** – Practice with arrays and hash tables

Reading:

Text: Ch. 9 – Arrays, p168-177 and Ch. 10 Hash Tables, p179-192

Week 11

Nov 9th MON

More hash tables and help on Program #6 Lots(!) of Reading: Text: split – p163-167, join p167-168, sorting p168-173, hashes p179-192 Counting motifs p192-202 Breaking Sequence into Motifs p214-216

Nov 11th WED

a6 algorithm is due

csLab - Entering a valid motif \$MIN <= \$size <= \$MAX</pre>

Nov 13th FRI

csLab: Work on a6

Nov 16th MON

Final Project Specification for Individual ProjectsWhat might *you* do?How will you present your results?How do you write the <u>Methods</u> section of a paper?What is expected of you?

Proposals due MON, Nov 23rd

Nov 18th WED

a6 due today

Class time to search for appropriate datasets for your final project: Go to NCBI, select sequences, store them for future use.

Nov 20th FRI Quiz #5

Samples of written and oral presentations—an analysis of examples

Week 13

Nov 23rd MON

Final Project Proposal due: type-written Title, Introduction, and Methods due by the beginning of class

csLab: Open lab time for individual and group help on final projects

Nov 25th WED – Nov 27th FRI Thanksgiving Break

Week 14

Nov 30th MON [open TBA]

Dec 2nd WED [open TBA]

Dec 4th FRI

Lab time for final presentations

Dec 7th MON Oral Presentations on Projects

Dec 9nd WED

Oral Presentations on Projects

Dec 11th FRI

Oral Presentations on Projects

Experimental program and paper are due at the **BEGINNING of CLASS**

Course Evaluations and Farewells