

# CMSE 491/890 - Bioinformatics & Computational Biology

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# Lecture 1: Introduction and Overview

Course overview

- Introductions
- Course website
- Communication
- Course activities
- Tentative Schedule
- Course topics
- Wrap-up

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- Assistant Professor
  - Dept. Computational Mathematics, Science, and Engineering
  - Dept. Biochemistry and Molecular Biology
- Research Interests: Computational genomics, Biomedical data science, Biological networks, Data integration, Machine learning

# What you should get out of this course

## How to become a practicing computational biologist in 60+h?

- An understanding of the major research directions in computational biology.
- Familiarity with the algorithms and statistical techniques for addressing these problems.
- At the end you should be able to:
  - Critically read bioinformatics / computational-biology literature.
  - Apply the methods you have learned to other problems both within and outside of bioinformatics.

# Modules & Topics

Module	Topics
Genome assembly, alignment, & annotation	de Bruijn graphs; Suffix trees; Hidden Markov models
Sequence alignment & pattern finding	Dynamic programming; Substitution matrices; BLAST
Comparative genomics; Phylogenomics	Molecular evolution; Tree construction
Genetic variation & quantitative genetics	LD, GWAS, Regularized linear regression; Statistical inference, Multiple testing
Regulatory genomics	Gibbs sampling; Expectation-Maximization
Functional genomics	Differential expression; Functional enrichment analysis; Clustering; Intro to ML
DataSci Primers	Data wrangling & visualization in R/Python; Exploratory data analysis
ML Primers	Machine learning; Deep learning; Applications
Single-cell genomics	Missing value imputation; Dimensionality reduction; Trajectory inference; Spatial reconstruction
Molecular dynamics; Protein structure prediction	Molecular simulation; Maximum entropy modeling
Modeling cellular pathways; Digital evolution	Dynamical simulation, State Space, Bifurcation; Linear programming; Artificial life
Biological networks	Measuring associations; Network inference; Graph theory, Label propagation
Cancer genomics	Overview
Genome engineering	Overview
Personal genomics	Overview

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Each topic:

- Lecture
- Paper discussion

Arjun presents


You present  
as team of  
two.

# Prerequisites & Expectations

- CMSE 201 and two semesters of introductory biology (LB 144 and 145 OR BS 161 and 162 OR BS 181H and 182H, or equivalent).
  - Statistics at the level of STT 231 is strongly recommended.
- Basically, it would be assumed that you:
  - know how to code in one of the mainstream languages like Python or R,
  - have an understanding of basic statistics and probability, and
  - have studied basic genetics, molecular biology, and cellular biology.

# Course website

[github.com/krishnanlab/teaching/tree/master/2019-spring\\_compbio](https://github.com/krishnanlab/teaching/tree/master/2019-spring_compbio)

- Contact information
  - Course outline and materials 
  - Schedule, location, calendar, and office hours
  - Website and communication
  - Course activities
  - Grading information
  - Attendance, conduct, honesty, and accommodations
- Lecture slides
  - Learning materials
  - Assignments
  - Scribe notes



[cmse-compbio-ss19.slack.com](https://cmse-compbio-ss19.slack.com)

- The primary mode of communication in this course (including major announcements) will be the course Slack account.
- All of you should have invitations to join this account in your MSU email.

#syllabus-schedule

#assignments

#announcements

#semester-projects

#lectures

#random

# Tell us about you

- Name
- Major or graduate program
- Research interests and/or topics you're especially interested in learning about
- Background in programming, algorithms, statistics, biology
- Office hours

Survey: [bit.ly/ss19-compbio\\_incoming-survey](https://bit.ly/ss19-compbio_incoming-survey)

# My office: 2507H Engineering Building (2nd floor)



# Course activities

- Assignments: ~35%
- Class participation: ~15%
- Scribing: ~10%
- Project: ~40%

# Assignments

- For each topic, you will be assigned a paper after the topic's "Lecture" class that you are required to read, summarize, and critique.
- Submit a report before the topic's "Paper discussion" class.
- See class website for details on what this report needs to contain.

# Class participation

- Do the assignments and additional readings.
- Show up to class.
- Work in groups during in-class discussion sessions (Bring your laptops)
- No one will have the perfect background.
  - [Ask questions](#) about computational or biological concepts.
- Correct me when I am wrong.
- Paper presentation (see the class website).

# Scribing

- Each lecture will have one dedicated scribe who will take notes on the lecture.
- Complete draft of scribe notes due in 3 days after lecture.
  - I will read and give comments/suggestions.
- Final scribe notes due in 6 days after lecture.
- We will use [stackedit.io](https://stackedit.io) + Google Drive, and then publish on GitHub.
  - <https://writing.stackexchange.com/a/32393>

# Project

- A major goal of this course is to prepare your ability to perform original research in computational biology, and to present your ideas and research.
- Can be one of:
  - Design and implement a new computational method for a task in biology
  - Improve an existing method
  - Perform an evaluation of several existing methods
  - Develop a fully-reproducible codebase for an existing analysis



# Project

Item	Due date
Project profile	W Jan 16
Project topic	F Feb 01
Project pre-proposal	W Fri 08
Project proposal	W Feb 20
Proposal reviews	F Mar 01
Mid-term project proposal presentations	M Mar 11
	W Mar 13
	F Mar 15
Review response	Th Mar 14
	Sa Mar 16
	M Mar 18
Mid-course project report	F Mar 29
Final project report	F Apr 26
Final project presentations	M Apr 29

# Course reading, External resources, Relevant seminars

- Mostly papers: primary articles and reviews
- Seminar
  - Science at the Edge <https://web.pa.msu.edu/seminars/edge/>
  - ML Seminar (CSE)
  - CSME Colloquium

# Groups @ MSU doing (a sigf. amount of) Bioinfo & Compbio

- Jianrong Wang
- Yuying Xie
- Leslie Kuhn
- Alex Dickson
- Michael Feig
- George Mias
- Ana Vasquez
- Gustavo de los Campos
- Eran Andrechek
- Robin Buell
- Erik Goodman
- Kevin Liu
- Mark Reimers
- Sudin Bhattacharya
- Jiayu Zhou
- Shinhan Shiu
- **MANY OTHERS**

You are welcome to work with me as well!

# What you need to do before the next class

- Read the course website:  
[github.com/krishnanlab/teaching/tree/master/2019-spring\\_compbio](https://github.com/krishnanlab/teaching/tree/master/2019-spring_compbio)
- Fill out the incoming survey: [bit.ly/ss19-compbio\\_incoming-survey](https://bit.ly/ss19-compbio_incoming-survey)
- Sign-up for scribing and paper presentation: check Slack for link

# Lecture 2: Introduction and Overview

Getting started

- Some history & high-level view of the field
- Choosing a good problem
- Organizing a computational biology project
- Reading journal articles | Supplementary materials
- Programming languages and other logistics
- Managing data and code
- Getting help