

# Genomics and Applied Bioinformatics

## BIOS 4150 / BIOL 6150, 3 credits

Mon & Wed, 2:00 pm - 2:50 pm, Instructional Center 115

Fri, 12:30 pm - 3:15 pm, Instructional Center 115

### Instructor Information

<b>Instructor</b> King Jordan	<b>Email</b> king.jordan@biology.gatech.edu	<b>Drop-in Hours &amp; Location</b> EBB 2115B, 3:00 pm, Mon
<b>Teaching Assistant(s)</b> Shivam Sharma	<b>Email</b> shivamsharma13@gatech.edu	<b>Drop-in Hours &amp; Location</b> Instruc Center 115, 2:00 pm, Fri

### General Information

#### Description

This course provides a practical introduction to foundational analysis techniques in genomics and bioinformatics, using the human genome as a working model. Course topics include: human genome structure and variation, genetic epidemiology, common and rare disease genetics, pharmacogenomics, population genomics, and genetic ancestry. The course will emphasize best practices in next-generation sequence (NGS) analysis and reproducible bioinformatics along with high-performance and cloud computing. Lectures will cover conceptual background and lab sessions will cover the implementation and use of bioinformatics methods.

#### Pre- &/or Co-Requisites

The course assumes prior working knowledge of biology, genetics, scientific computing, and programming in R and Python. The course is considered as complementary to the concurrently offered course BIOL 7200 Programming for Bioinformatics, and it will build on bioinformatics skills taught in BIOL 7200.

#### Course Goals and Learning Outcomes

Upon successful completion of this course, you should be able to:

- Conduct collaborative bioinformatics projects using high-performance and cloud-computing resources
- Conduct and document reproducible bioinformatics analyses using Jupyter Notebook
- Perform foundational techniques for NGS analysis: data access, quality control, read mapping, and variant calling
- Statistically model disease outcomes
- Clinically interpret human genome sequence variation
  - Common and rare diseases
  - Drug response
- Characterize genome-wide patterns of genetic diversity and ancestry

### Course Requirements & Grading

Students will be graded based on ten projects performed using Jupyter Notebook. Jupyter Notebooks will be run and evaluated on the class high-performance compute (HPC) cluster. Projects will be performed in small student groups, and students will have one week to complete each project. The project schedule is shown as part of the class schedule (below).

## Grading Scale

Your final grade will be assigned as a letter grade according to the following scale. Grades will not be curved.

A	90-100%
B	80-89%
C	70-79%
D	60-69%
F	0-59%

## Course Materials

### Course Text

Course readings will be taken from the primary scientific literature and provided to students via Canvas.

### Additional Materials/Resources

Students will be given access to a dedicated compute cluster as part of the Georgia Tech PACE-ICE cloud HPC environment. Students will be provided with markdown tutorials and Jupyter Notebook frameworks for genomics and multi-omics methods.

### Course Website and Other Classroom Management Tools

The course will use the Georgia Tech Canvas classroom management tool.

## Course Expectations & Guidelines

### Academic Integrity

Georgia Tech aims to cultivate a community based on trust, academic integrity, and honor. Students are expected to act according to the highest ethical standards. For information on Georgia Tech's Academic Honor Code, please visit <http://www.catalog.gatech.edu/policies/honor-code/> or <http://www.catalog.gatech.edu/rules/18/>.

Any student suspected of cheating or plagiarizing on a quiz, exam, or assignment will be reported to the Office of Student Integrity, who will investigate the incident and identify the appropriate penalty for violations.

### Accommodations for Students with Disabilities

If you are a student with learning needs that require special accommodation, contact the Office of Disability Services at (404)894-2563 or <http://disabilityservices.gatech.edu/>, as soon as possible, to make an appointment to discuss your special needs and to obtain an accommodations letter. Please also e-mail me as soon as possible in order to set up a time to discuss your learning needs.

### Attendance and/or Participation

Attendance and participation at Monday and Wednesday lecture sessions and Friday bioinformatics lab sessions is mandatory. Illness is considered an excused absence; students are encouraged to stay home when sick in order to take of themselves and each other. Please see <http://www.catalog.gatech.edu/rules/4/> for information about institute expectations and restrictions around attendance, including information about excused absences.

## Collaboration & Group Work

Bioinformatics is a highly collaborative field. Collaboration and group work are encouraged in order to emulate the environment that students will encounter in the working world. Projects will be conducted in student groups. Students are free to use outside resources to support their project work, much as they would do if they were in the working world.

## Extensions, Late Assignments, & Re-Scheduled/Missed Exams

Projects will be submitted as Jupyter notebooks via the Georgia Tech PACE-ICE cloud HPC environment. Missed project deadlines will result in a grade of zero.

Note that Georgia Tech makes some exceptions for “approved Institute activities” (e.g. field trips and athletic events) and religious observances, which will be handled on a case-by-case basis. See <http://www.catalog.gatech.edu/rules/4/> for more information. We will also consider the impact of events like the [All-Majors Career Fair](#), and off-campus interviews, and plan accordingly.

## Student-Faculty Expectations Agreement

At Georgia Tech we believe that it is important to strive for an atmosphere of mutual respect, acknowledgement, and responsibility between faculty members and the student body. See <http://www.catalog.gatech.edu/rules/22/> for an articulation of some basic expectation that you can have of me and that I have of you. In the end, simple respect for knowledge, hard work, and cordial interactions will help build the environment we seek. Therefore, I encourage you to remain committed to the ideals of Georgia Tech while in this class.

## Student Use of Mobile Devices in the Classroom

As research on learning shows, unexpected noises and movement automatically divert and capture people's attention, which means you are affecting everyone's learning experience if your cell phone, tablet, laptop, etc. makes noise or is visually distracting during class. That said, many students find it useful to have a mobile device on hand to access course materials.

With this in mind, students are allowed to take notes or access course materials on mobile devices, with the sound off so that other students' learning is not disrupted. Please refrain from doing anything other than taking notes or looking at course materials on your mobile devices.

## Campus Resources for Students

Georgia Tech provides a number of campus resources in support of students' mental health and emotional well-being, including but not limited to:

- Student Life Mental Health and Well-being <https://studentlife.gatech.edu/services/mental-health-well-being>
- The Center for Mental Health Care & Resources <https://mentalhealth.gatech.edu/>
- Stamps Health Services Psychiatric Clinic <https://health.gatech.edu/psych/>

## Class Schedule

Week	Day	Date	Topic	Instructor	Assignment	Points
1	Mon	19-Aug-24	Introduction & logistics	Jordan & Sharma		
	Wed	21-Aug-24	Reproducible bioinformatics	Jordan & Sharma		
	Fri	23-Aug-24	PACE-ICE & Jupyter Lab	Sharma		
2	Mon	26-Aug-24	Human genome	Jordan & Sharma		
	Wed	28-Aug-24	Reference genome sequence	Jordan		
	Fri	30-Aug-24	PACE-ICE & Jupyter Lab	Sharma		
3	Mon	2-Sep-24	Labor Day holiday	No class		
	Wed	4-Sep-24	Genome sequencing	Jordan		
	Fri	6-Sep-24	Genome data structure lab	Sharma		
4	Mon	9-Sep-24	Data access	Jordan		
	Wed	11-Sep-24	Quality control	Jordan		
	Fri	13-Sep-24	Data access & Quality control lab	Sharma	Project #1	100
5	Mon	16-Sep-24	Make up and review	Jordan		
	Wed	18-Sep-24	Sequence read mapping	Jordan		
	Fri	20-Sep-24	Read mapping lab	Sharma	Project #2	100
6	Mon	23-Sep-24	Variant calling	Jordan		
	Wed	25-Sep-24	Variant data structures	Jordan		
	Fri	27-Sep-24	Variant calling lab	Sharma	Project #3	100
7	Mon	30-Sep-24	Population biobanks	Jordan		
	Wed	2-Oct-24	Electronic health records	Jordan		
	Fri	4-Oct-24	Phecode lab	Sharma	Project #4	100
8	Mon	9-Oct-23	Genetic epidemiology	Jordan		
	Wed	11-Oct-23	Disease modeling	Jordan		
	Fri	13-Oct-23	Modeling lab	Sharma	Project #5	100
9	Mon	14-Oct-24	Fall Break	No class		
	Wed	16-Oct-24	Make up and review	Jordan		
	Fri	18-Oct-24	Review lab	Sharma		
10	Mon	21-Oct-24	Pharmacogenomics	Jordan		
	Wed	23-Oct-24	Pharmacogenomic diversity	Jordan		
	Fri	25-Oct-24	Pharmacogenomics lab	Sharma	Project #6	100
11	Mon	28-Oct-24	Genome-wide association studies	Jordan		
	Wed	30-Oct-24	Polygenic risk scores	Jordan		
	Fri	1-Nov-23	Polygenic risk score lab	Sharma	Project #7	100
12	Mon	4-Nov-24	Mendelian genetics & disease	Jordan		
	Wed	6-Nov-24	Genetic testing	Jordan		
	Fri	8-Nov-24	Clinical genetics testing lab	Sharma	Project #8	100
13	Mon	11-Nov-24	Genetic diversity	Jordan		
	Wed	13-Nov-24	Principal components analysis	Jordan		
	Fri	15-Nov-24	Principal components analysis lab	Sharma	Project #9	100
14	Mon	18-Nov-24	Genetic ancestry	Jordan		
	Wed	20-Nov-24	Admixture	Jordan		
	Fri	22-Nov-24	Admixture lab	Sharma	Project #10	100
15	Mon	25-Nov-24	Make up and review	Jordan		
	Wed	27-Nov-24	Thanksgiving break	No class		
	Fri	29-Dec-24	Thanksgiving break	No class		
16	Mon	4-Dec-24	Course wrap up	Jordan		