

Course goals: The goal of this course is to provide an understanding of the application of next-generation sequencing approaches to the study of evolutionary phenomena. The students will be expected to understand the molecular techniques involved, the statistical issues associated with these large datasets, and the implications of these datasets with respect to the evolutionary process.

Grading: Letter grades will be assigned based as follows: active participation: 50%; take-home essay exam: 50%.

Grade scale: 90-100 A; 80-89 B; 70-79 C; 60-69 D; < 60 F

Americans with Disabilities Act (ADA): The Americans with Disabilities Act (ADA) is a federal anti-discrimination statute that provides comprehensive civil rights protection for persons with disabilities. Among other things, this legislation requires that all students with disabilities be guaranteed a learning environment that provides for reasonable accommodation of their disabilities. If you believe you have a disability requiring an accommodation, please contact Disability Services, in Cain Hall, Room B118, or call 845-1637. For additional information visit <http://disability.tamu.edu>.

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“An Aggie does not lie, cheat, or steal, or tolerate those who do.”

REQUIRED READINGS

Required readings will originate from the primary literature and will be assigned by email or during class.

LECTURES

1. Comparative genomics and methods.
2. Population genomics.
3. Signatures of selection and adaptation.
4. Phylogenomics.
5. Genome structure and evolution.

6. Speciation genomics and sex-chromosome evolution.

Take-home essay exam due by email at 4 pm the day after lecture 6. *One letter grade will be deducted for each day past the deadline!*