

ORIGINAL COURSE IMPLEMENTATION DATE:

**REVISED COURSE IMPLEMENTATION DATE:** 

September 1995 September 2022 January 2028

**COURSE TO BE REVIEWED** (six years after UEC approval): Course outline form version: 05/18/2018

# **OFFICIAL UNDERGRADUATE COURSE OUTLINE FORM**

Note: The University reserves the right to amend course outlines as needed without notice.

Course Code and Number: BIO 406		Number of Credits: 3 Course credit policy (105)				
Course Full Title: Advanced Genetics						
Course Short Title:						
(Transcripts only display 30 characters. Departments may recommend a short title if one is needed. If left blank, one will be assigned.)						
Faculty: Faculty of Science		Department (c	Department (or program if no department): Biology			
Calendar Description:						
A number of emerging areas of interest in genetics will be covered, including the genetics of human complex disease, epigenetics and environmental influences on our genes, genomics and personalized medicine, the genetics of infectious disease, legal and ethical issues arising from advances in genetics, and quantitative and evolutionary genetics of populations.						
Prerequisites (or NONE):	BIO 202 an	d BIO 220.				
Corequisites (if applicable, or NONE):						
Pre/corequisites (if applicable, or NONE):						
Antirequisite Courses (Cannot be taken for additional credit.)			Special Topics (Double-click on boxes to select.)			
Former course code/number:		This course is offered with different topics:				
Cross-listed with:		🛛 No	No Yes (If yes, topic will be recorded when off			
Dual-listed with:			Independent Study If offered as an Independent Study course, this course may			
Equivalent course(s):						
(If offered in the previous five years, antirequisite course(s) will be included in the calendar description as a note that students with credit			be repeated for further credit: (If yes, topic will be recorded.)			
for the antirequisite course(s) cannot take this course for further credit.)			$\square$ No $\square$ Yes, repeat(s) $\square$ Yes, no limit			
	,	Transfer Credit				
Typical Structure of Instructional Hours			Transfer credit already exists: (See bctransferguide.ca.)			
Lecture/seminar hours		45	🖾 No	Yes		
Tutorials/workshops				outline for (re)articulation:		
Supervised laboratory hours		🖾 No	Yes (If yes, fill in trans	fer credit form.)		
Experiential (field experience, practicum, in	ternship, etc.)	)	Gradin	g System		
Supervised online activities			🛛 Lette	er Grades 🗌 Credit/No 🤇	Credit	
Other contact hours:			Maxim	um enrolment (for inform	nation only): 24	
Total hours 45			Expected Frequency of Course Offerings:			
Labs to be scheduled independent of lecture	o 🗌 Yes	-	annually (Every semester, Fall only, annually, etc.)			
Department / Program Head or Director: G	Bregory Schma	altz	• 	Date of meeting:	October 1, 2021	
Faculty Council approval				Date of meeting:	November 5, 2021	
Undergraduate Education Committee (UE	C) approval			Date of meeting:	January 28, 2022	

# Learning Outcomes:

Upon successful completion of this course, students will be able to:

- 1. Assess current literature and critically discuss most recent data on select areas of genetic research.
- 2. Present a critical review on a specific area of interest based on topics discussed in class.
- 3. Discuss the difficulties associated with the identification of genes responsible for disease.
- 4. Give examples of how the analysis of the human genome sequence can be used to identify genes associated with complex diseases.
- 5. Describe how technological development is changing the face of medicine and the ethical and social issues that this is generating.
- 6. Evaluate, at a genetic level, how selective pressures such as the host immune response or the overuse of antibiotics is driving changes in microbial populations, and how this is resulting in newly emerging infectious disease.
- 7. Discuss how natural selection and other mechanisms such as mutations, migration, and genetic drift alter allele frequencies to bring about evolutionary divergence in populations.

# Prior Learning Assessment and Recognition (PLAR)

🛛 Yes 🗌 No

□ No, PLAR cannot be awarded for this course because

**Typical Instructional Methods** (*Guest lecturers, presentations, online instruction, field trips, etc.; may vary at department's discretion.*) A combination of lectures and small group tutorials emphasizing problem-solving.

# NOTE: The following sections may vary by instructor. Please see course syllabus available from the instructor.

Typical Text(s) and Resource Materials (If more space is required, download Supplemental Texts and Resource Materials form.)

	Author (surname, initials)	Title (article, book, journal, etc.)	Current ed.	Publisher	Year
1.	Klug W. et al	Concepts of Genetics	$\boxtimes$	Pearson	2019
2.	Schuettenngruber B. et al	nuettenngruber B. et al Genome Regulation by Polycomb and Trithorax: 70 Years and Counting		Cell	2017
3.	Nussenzweig P. et al.	Molecular Mechanisms of CRISPR-Cas Immunity in Bacteria		Ann Rev Genetics	2020
4.	Uffelmann E. et al.	Genome Wide Association Studies		Nature Reviews	2021
5.	Gibson G. and Lacek K	Canalization and Robustness in Human Genetics and Disease		Ann Rev Genetics	2020

Required Additional Supplies and Materials (Software, hardware, tools, specialized clothing, etc.)

## Typical Evaluation Methods and Weighting

Final exam:	50%	Assignments:	20%	Field experience:	%	Portfolio:	%
Midterm exam:	30%	Project:	%	Practicum:	%	Assigned Problems:	%
Quizzes/tests:	%	Lab work:	%	Shop work:	%	Total:	100%

## Details (if necessary):

## Typical Course Content and Topics

Human cytogenetics Developmental genetics Genetics of infectious disease Epigenetics Medical genetics/genetics of human complex disease Bioethics DNA barcoding Allele frequency changes in Populations

Genetics of Human Complex Disease/Genome Wide Association Studies (GWAS)

- Single nucleotide polymorphisms
- The human haplotype map
- Linkage for common disease
- Common disease common variant hypothesis
- Linkage disequilibrium
- Indirect association

#### Epigenetic Inheritance: A Contributor to Species Differentiation? • The concept of the epigenome

- What is epigenetic variation?
- Environmental influences on gene expression
- How multiple epigenetic states can be associated with the same genome
- The transmission of epigenetic states through the germline
- Can epigenetic variation be stable enough to underlie species characteristics?
- Examples of epigenetic inheritance which may mediate Darwinian evolution
- Somatic methylation states as they relate to methylation states of the germline
- Paramutation and genomic imprinting

#### Genetics of infectious disease

1. Antibiotic resistance:

- Resistance mechanisms to circumvent the toxic action of antimicrobials
- Mutation of normal cellular genes, the acquisition of foreign resistance genes, or a combination of these two mechanisms
- · The spread of mobile genetic elements and the acquisition of multidrug resistance in a single genetic event
- · The need for infection control in both human and veterinary medicine

#### 2. Antigenic variation

- Mechanisms of antigenic variation; an overview
- · Selective pressures imposed on the microbial world
- Phase variation in Helicobacter pylori lipopolysaccharide
- Genetic variation in the pathogenic Neisseria species
- Trypanosome antigenic variation
- Surface antigenic variation in Giardia lamblia
- Emerging infectious disease; the impact of antigenic variation on pathogen population structure, fitness, and dynamics

#### Genetics in society:

- The development of human genetic and reproductive technologies
- · Legal, ethical, and social issues arising from the development of these technologies
- Bioethics associated with research which include areas such as the use of human subjects in the field of genomics, genomic
  data sharing policy, and how the FDA proposes oversight of laboratory developed tests.