

## Syllabus

**Course:** PCB4917- Modeling Rare Genetic Diseases.

**Date:** Jan 12<sup>th</sup> to February 11<sup>th</sup>, 2026

**Time and Location:** 9:30 am to 12:35 pm; 1:55 pm to 4:55 pm, CGRC Rm 438 and Rm 436.

**Contact information:** Dr. Xiaofei Bai

**Email:** [baixiaofei@ufl.edu](mailto:baixiaofei@ufl.edu)

**Office hours:** CGRC Room 438 by appointment

## Introduction

Welcome to the immersion course. This section will provide hands-on experiences using the nematode model system *Caenorhabditis elegans*, which contains approximately 3000 disease-associated genes. You will receive instruction on designing and applying the novel and powerful gene-editing technique CRISPR/Cas9, focusing on establishing precision disease modeling in *C. elegans*.

### What will you learn when finishing the course?

- 1) Basic molecular biology techniques, including PCR, restriction enzyme digestions, and electrophoresis.
- 2) Basic strategies to design CRISPR/Cas9 gene editing.
- 3) The *C. elegans* maintenance and operation.

### What will you need for the class?

The course will be textbook-free. However, it will be helpful if you can read the literature below before attending the classes. Download the literature using the UF network.

[http://www.wormbook.org/chapters/www\\_strainmaintain/strainmaintain.html](http://www.wormbook.org/chapters/www_strainmaintain/strainmaintain.html)  
<https://www.sciencedirect.com/science/article/pii/S1046202316302857?via%3Dihub>

### Learning assessment during the class.

- 1) **Participation and discussion:** the class attendees will be requested to actively participate in all class activities and meetings.
- 2) **Exam:** There will be no exam for the course. However, everyone will be required to submit a final report by the end of the course, which will include their CRISPR/Cas9 design and final data.
- 3) **Presentation:** You will give a final presentation in your project groups at the end of the classes.

### Grade components and scale:

The participation of the class: 20% of the total grade

Final report: 50% of the total

Presentation: 30% of the total grade.

**Scale:** A  $\geq$  90.00%; A-  $\geq$  86.66%; B+  $\geq$  83.33%; B  $\geq$  80.00%; B-  $\geq$  76.66%; C+  $\geq$  73.33%; C  $\geq$  70.00%; C-  $\geq$  66.66%; D+  $\geq$  63.33%; D  $\geq$  60.00%; D-  $\geq$  56.66%; F < 56.66%

**Illness and unforeseen things will be excused to miss the class. However, proper documentation will be required to explain the absence. You can find UF policy for excused absence here:** <https://catalog.ufl.edu/UGRD/academic-regulations/attendance-policies/>

#### **Academic honesty**

Any cheating, plagiarism, or other forms of academic dishonesty will result in a 0 grade. For additional information on Academic Honesty, please visit the UF Student Honor Code and Student Conduct Code at <https://sccr.dso.ufl.edu/policies/student-honor-code-student-conduct-code/>

**If you have any questions, concerns, or comments on the course, please feel free to contact me by email** [baixiaofei@ufl.edu](mailto:baixiaofei@ufl.edu) **or by phone 352-294-8447.**

#### **Schedule**

<b>Date</b>	<b>Time</b>	
Jan 12 <sup>th</sup> to Jan 19 <sup>th</sup>	9 am to 12 pm	Location is CGRC Rm436 Course introduction and discussion Introduction of disease modeling in the model systems Introduction of <i>C. elegans</i> biology and genetics
Jan 12 <sup>th</sup> to Jan 19 <sup>th</sup>	1 pm to 5 pm	Location is CGRC Rm 438 <i>C. elegans</i> maintenance and operation
Jan 20 <sup>th</sup> to Jan 27 <sup>th</sup>	9 am to 12 pm	Location is CGRC Rm436 CRISPR/Cas9 gene editing introduction and designing
Jan 27 <sup>th</sup> to Feb 9 <sup>th</sup>	9 am to 12 pm	Location is CGRC Rm 438 Genotyping CRISPR edited animals and phenotype the <i>C. elegans</i> .
Feb 11 <sup>th</sup>	9 to 12 pm	Location is CGRC Rm436 Last presentations.
Feb 11 <sup>th</sup>		Deadline to submit the final report.

**Note: There will be no lecture classes on weekends, but the students will be approved and get access to the lab by maintaining the *C. elegans* or performing experiments if needed.**