

# Computational Approaches to Biology (BIOL 3070)

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# Learning objectives

1. Get to know the R interface and basic commands
2. Learn about variables, functions, vectors, logic, and missing data

# Today's outline

1. General housekeeping items
2. Live coding demonstration
3. Quiz exercise (non-graded)

# Course laptops/R installation

1. ~55 students and ~35 laptops
  - a. If anyone has R (or can get it easily) and a decent computer, we can locally install the needed packages
  - b. Will post a copy/paste tutorial to do so

# Office hours

1. Monday, 12:30-3:30pm
2. Will post announcement with location
3. Good place to install course packages if you want to

# Assigned seats

1. Randomized assigned seating starting next week
2. Groups will be mostly even
3. You'll probably make new friends

# R learning resources

1. `swirl()` Package in R
2. Data Carpentry
  - a. <https://datacarpentry.org/R-genomics/index.html>
3. Learning R: A Step-by-Step Function Guide to Data Analysis (Richard Cotton, 2013)

# Introduction to R

Find the handout via the course schedule. If you have R on your laptop, open so you can enter the commands as we go. If not, just follow along with the handout.

# Practice quiz

Head to the Quizizz link on the schedule for today and get ready to answer some questions about what you learned!