

# Bioinformatic resources

## In-person courses

**Harvard Chan Bioinformatics Core - Bioinformatics for the Harvard Community**  
(courses range from beginner to advanced); cost ~ \$65 (04/2024)

<https://bioinformatics.sph.harvard.edu/training/>

Online tutorials facilitate learning at your own convenience:

<https://github.com/hbctraining>

## Online courses

**Harvard Edx - Data analysis for Life Science (Beginner)**

Cost including certificate \$876 (can be reimbursed through BWH benefits) (04/2024).

Lab members who have completed this course in the past: Frederike Kramer

<https://www.edx.org/certificates/professional-certificate/harvardx-data-analysis-for-life-sciences>

**Harvard Edx - Data analysis for Genomics (Advanced)**

Cost including certificate \$657 (can be reimbursed through BWH benefits) (04/2024).

Lab members who have completed this course in the past: Merle Riedemann

<https://www.edx.org/certificates/professional-certificate/harvardx-data-analysis-for-genomics>

## Free resources in-person

**Broad CodeRATS**

When? Mondays and Tuesday @ 11am-12pm

Where? 75 Ames, 4th Floor Connector Kitchen

*“Come by if you need coding advice, or help debugging anything. Work or personal projects welcome. We have snacks :)”*

(info last updated 04/2024)

<https://sites.google.com/broadinstitute.org/coderats/home>

**Meetup Group Bioconductor for Genomics**

<https://www.meetup.com/Boston-R-Bioconductor-for-genomics/>

**Meetup Group Boston Women's Bioinformatics**

<https://www.meetup.com/boston-area-womens-bioinformatics/>

## **Harvard Statistics Consulting Service**

When? Mondays, Tuesdays & Fridays from 1pm to 3pm

Where? Science Center, room 400N

*“Note that this service is for statistical issues (e.g., framing a scientific question as a statistical hypothesis, choosing an appropriate statistical method, or even choosing the fastest computational method for deploying that method) and we do not provide assistance on pure programming issues.”*

(last updated 04/2024)

<https://statistics.fas.harvard.edu/harvard-statistics-consulting-service>

## Free resources online

### ***R for absolute beginners***

#### **Swirl**

<https://swirlstats.com/students.html>

### ***Applied R for Biologists/MDs***

#### **YouTube videos explaining gene expression analysis**

<https://www.youtube.com/@Bioinformagician/featured>

#### **Introduction to Data Science by Rafael Irizarry**

*Rafalab, Dana-Farber Cancer Institute*

Online Part 1 <https://rafalab.dfci.harvard.edu/dsbook-part-1/>

Online Part 2 <https://rafalab.dfci.harvard.edu/dsbook-part-2/index.html>

Pdf <https://leanpub.com/datasciencebook>

#### **Data Analysis for the Life Sciences by Rafael Irizarry**

Pdf <https://leanpub.com/dataanalysisforthelifesciences>

Github <https://github.com/rafalab/rafalab.github.io/blob/master/pages/harvardx.md>

#### **Anything scRNAseq**

*Satija Lab, NYU*

<https://satijalab.org/seurat/>

## **Bioinformatics Training at the Harvard Chan Bioinformatics Core**

Lots of free courses on many bioinformatic topics

<https://hbctraining.github.io/main/>

### ***Online books***

Very helpful for creating plots in R

<https://r-graphics.org/>

Bible of R

<https://r4ds.had.co.nz/>

Advance bible of R

<http://adv-r.had.co.nz/Introduction.html>

A collection of pre-processing and visualization scripts for single-cell multi-omics data

<https://bookdown.org/ytlIU13207/SingleCellMultiOmicsDataAnalysis/>

Starter codes for Linux

<https://www.putorius.net/standard-wildcards-globbing-patterns-in.html>