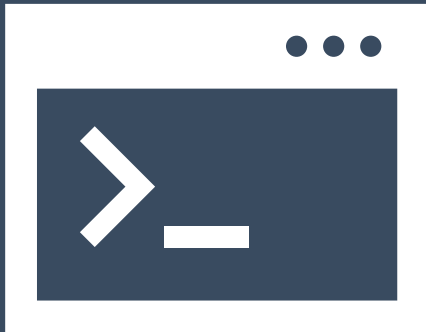


Introduction to R

<https://tinyurl.com/hbc-r-revised>



Harvard Chan Bioinformatics Core



Introductions!





Shannan Ho Sui
Director



Meeta Mistry
Associate Director



Lorena Pantano
*Director of Bioinformatics
Platform*



John Quackenbush
Faculty Advisor



Upen Bhattarai



Heather Wick



Will Gammerdinger



Noor Sohail



Alex Bartlett



Elizabeth
Partan



Emma Berdan



James Billingsley



Zhu Zhuo



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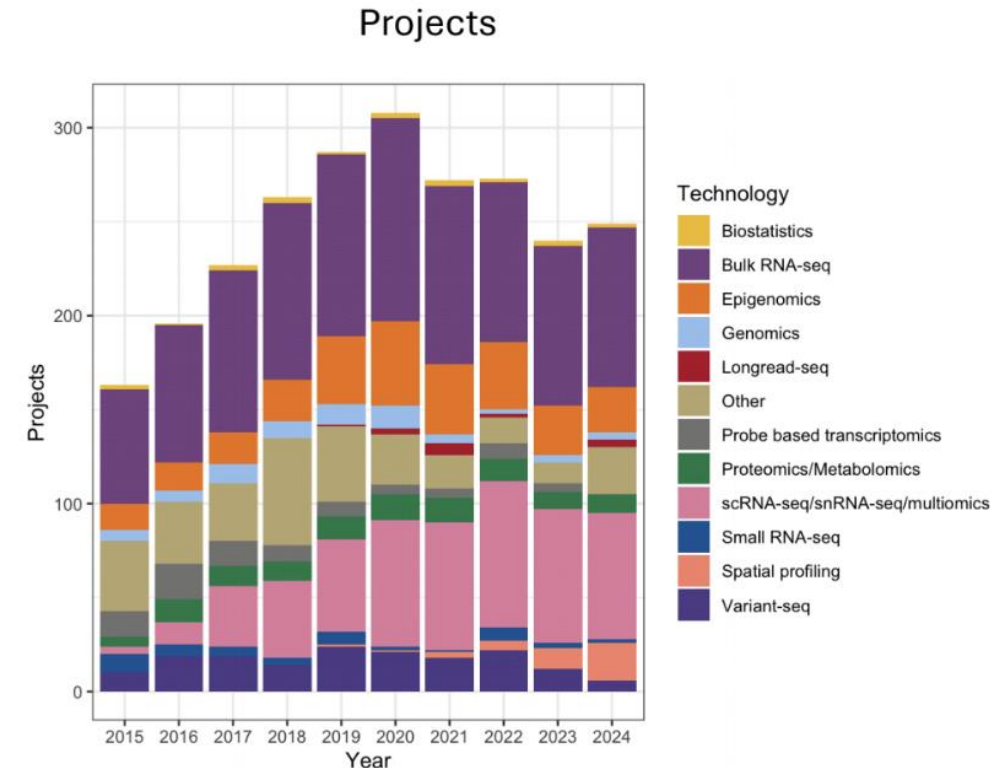
Zhu Zhuo



Maria Simoneau

Consulting

- ❖ Transcriptomics: Bulk, single cell, small RNA
- ❖ Epigenomics: ChIP-seq, CUT&RUN, ATAC-seq, DNA methylation
- ❖ Variant discovery: WGS, resequencing, exome-seq and CNV
- ❖ Multiomics integration
- ❖ Spatial biology
- ❖ Experimental design and grant support



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THE HARVARD CLINICAL
AND TRANSLATIONAL
SCIENCE CENTER



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MEDICAL SCHOOL

Training

- ❖ Hands-on workshops design to reflect best practices, reproducibility and an emphasis on experimental design

- ❖ Basic Data Skills

- ❖ Shell

- ❖ R

- ❖ Advanced Topics: Analysis of high-throughput sequencing data

- ❖ Chromatin Biology

- ❖ Bulk RNA-seq

- ❖ Differential Gene Expression

- ❖ scRNA-seq

- ❖ Variant Calling

- ❖ Current Topics in Bioinformatics

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

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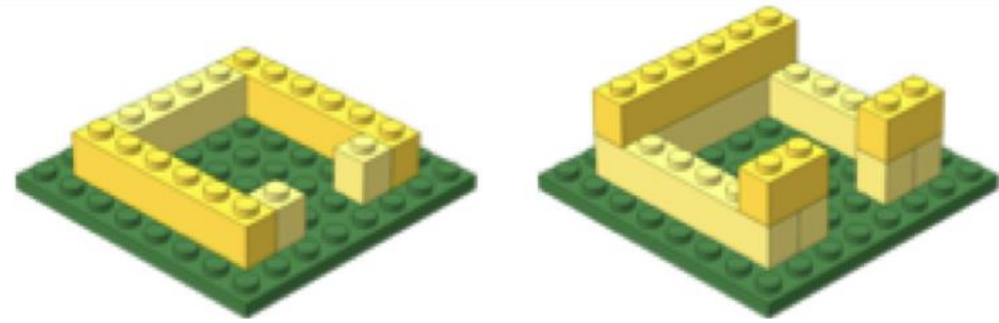
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Workshop scope

Workshop Scope



- ❖ Comfortably use RStudio (a graphical interface for R)
- ❖ Fluently interact with R using RStudio
- ❖ Become familiar with R syntax
- ❖ Understand data structures in R
- ❖ Inspect and manipulate data structures
- ❖ Install packages and use functions in R

CRAN



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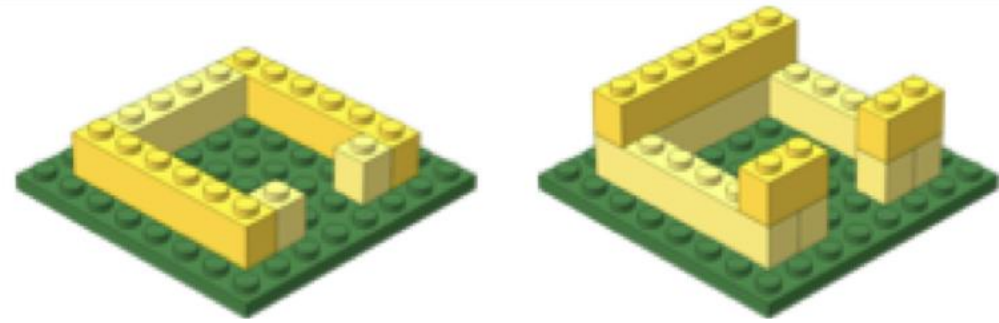
Available CRAN Packages By Name
[A](#)[B](#)[C](#)[D](#)[E](#)[F](#)[G](#)[H](#)[I](#)[J](#)[K](#)[L](#)[M](#)[N](#)[O](#)[P](#)[Q](#)[R](#)[S](#)[T](#)[U](#)[V](#)[W](#)[X](#)[Y](#)[Z](#)

Accurate, Adaptable, and Accessible Error Metrics for Predictive Models
Access to Abbyy Optical Character Recognition (OCR) API
Tools for Approximate Bayesian Computation (ABC)
Computed ABC Analysis
Data Only: Tools for Approximate Bayesian Computation (ABC)
ABCDE_FBA: A-Biologist-Can-Do-Everything of Flux Balance Analysis with this package
Implementation of Artificial Bee Colony (ABC) Optimization
Approximate Bayesian Computational Model for Estimating P2
Approximate Bayesian Computation via Random Forests

- ❖ Comprehensive R Archive Network
- ❖ The main repository for R packages
- ❖ Easy to install

- ❖ An alternative package repository; “..provides tools for the analysis and comprehension of *high-throughput genomic data*.”
- ❖ Includes (but is not limited to) tools for:
- ❖ Performing statistical analysis
- ❖ Accessing public datasets
- ❖ Open source and open development
- ❖ Free

Workshop Scope





- ❖ Comfortably use RStudio (a graphical interface for R)
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- ❖ Become familiar with R syntax
- ❖ Understand data structures in R
- ❖ Inspect and manipulate data structures
- ❖ Install packages and use functions in R
- ❖ Visualize data using ggplot2
- ❖ Utilize pipes, tibbles and functions from the Tidyverse package suite


Logistics



Course schedule

Harvard Chan Bioinformatics Core



 hbctraining/Intro-to-R-m...
☆ 0 🗨 0

Home Workshop Schedule

Table of contents
Day 1
Before the next class
Day 2
Before the next class
Day 3
Before the next class
Day 4
Additional exercises and answer keys
Additional resources

Workshop Schedule

Day 1

Lesson	Overview	Instructor	Time
Workshop Introduction	Welcome and housekeeping	Elizabeth	10:00-10:30
Intro to R and RStudio	Introduction to R and RStudio	Noor	10:30-11:45
Self learning materials	Overview of self-learning materials	Elizabeth	11:45-12:00

<https://tinyurl.com/hbc-r-revised>

Course materials

- ❖ We continuously update our materials to reflect changes in the field/software

HBC
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Home Workshop Schedule

Search

hbctraining/Intro-to-R-m...
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Table of contents

- Learning Objectives
- What is R?
- Why use R?
- What is RStudio?
- Creating a new project directory in RStudio
- What is a project in RStudio?
- RStudio Interface
- Organizing and Setting up RStudio
 - Viewing your working directory
- Structuring your working directory
 - Setting up
- Interacting with R
 - Console window

Introduction to R and RStudio

Approximate time: 45 minutes

Learning Objectives

- Describe what R and RStudio are.
- Interact with R using RStudio.
- Familiarize various components of RStudio.
- Employ variables in R.

What is R?

The common misconception is that R is a programming language but in fact it is much more than that. Think of R as an environment for statistical computing and graphics, which brings together a number of features to provide powerful functionality.

<https://tinyurl.com/hbc-r-revised>

Course participation

- ❖ Mandatory review of self-learning lessons and assignments
- ❖ Attendance required for all classes
- ❖ Your questions and active participation drive learning
- ❖ **We look forward to all of your questions!**



Course participation

- ❖ At-home lessons and exercises after each session
- ❖ Cover material not previously discussed
- ❖ Provides us feedback to help pace the course appropriately
- ❖ 3-5 hours to complete
- ❖ Homework load is heavier in the beginning of this workshop series and tapers off

Using AI for Assignments



❖ Do

- ❖ Try to resolve error messages with it
- ❖ Test code written by AI on a dataset where you have expected results
- ❖ Take the time to review the generated code line-by-line

❖ Don't

- ❖ Implement it in replacement to learning
- ❖ Write code that you don't understand
- ❖ Assume the output from an AI process is correct

Odds & Ends

- ❖ Quit/minimize all applications that are not required for class
- ❖ Name tags
- ❖ Post-its
 - ❖  green - I am all set
 - ❖  red - I need time/help
- ❖ Phones on vibrate/silent
- ❖ Bathrooms

Contact Us

- ❖ *HBC training team:* hbctraining@hsph.harvard.edu
- ❖ *HBC consulting:* bioinformatics@hsph.harvard.edu