



Introduction to R

Harvard Chan Bioinformatics Core

Learning Objectives



- ✓ 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
- ✓ Visualize data using *ggplot2*

Exit survey

[https://harvard.az1.qualtrics.com/jfe/form/
SV_9HRpofBfjb2vgTX](https://harvard.az1.qualtrics.com/jfe/form/SV_9HRpofBfjb2vgTX)

Resources

[Resources for RNA-Seq, R, general bioinformatics](#)

[Asking for help with R](#)

[ggplot2 cheatsheet](#)

Contact us!

HBC webpage: <http://bioinformatics.sph.harvard.edu>

HBC training materials: <http://hbctraining.github.io/main>

Training email: hbctraining@hsph.harvard.edu

Twitter

[@bioinfocore](#)