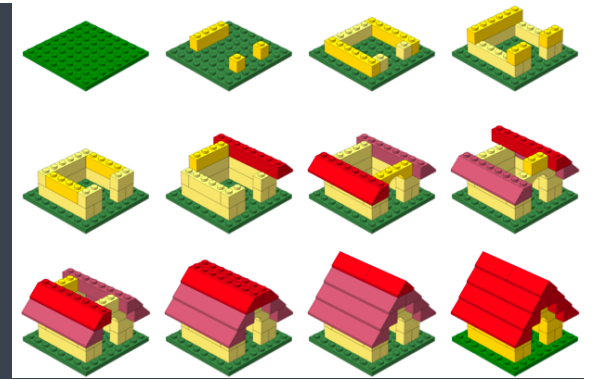


# Introduction to R

Harvard Chan Bioinformatics Core

<https://tinyurl.com/introR-feb2018>

# Learning Objectives



- ✓ Become comfortable with 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 simple and complex plotting methods

# Exit survey

<https://tinyurl.com/R-exit-survey>

# Resources

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

[Asking for help with R](#)

[ggplot2 cheatsheet](#)

# Contact us!

*Training questions?* [hbctraining@hsph.harvard.edu](mailto:hbctraining@hsph.harvard.edu)

*Consulting questions?* [bioinformatics@hsph.harvard.edu](mailto:bioinformatics@hsph.harvard.edu)

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

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

*HBC workshop listserv:* <https://tinyurl.com/hbc-mailing-list>

*Twitter:* [@bioinfocore](https://twitter.com/bioinfocore)