

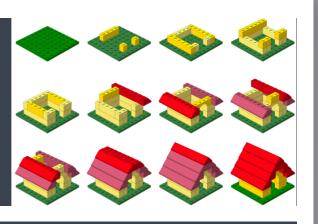
## Introduction to R

Harvard Chan Bioinformatics Core

https://tinyurl.com/intro-to-r-1-5

Sponsored by HMS (TnT) and HSCI

## 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

http://tinyurl.com/HMS-HSCI-introR-exit-survey

## Upcoming free, monthly, 3hr workshops

- December 13th:

Setting up for Success: Introduction to Version Control (Git)

# Bioinformatics Community Networking Breakfast!



- **Free and open** to the LMA community
- Food and seats are first-come-first-served
- Last Wednesday of every month

Date: Dec. 18th, 2019

Time: 9:00 to 10:30 am

More info: <a href="http://bioinformatics.sph.harvard.edu/breakfast/">http://bioinformatics.sph.harvard.edu/breakfast/</a>

### Resources

Resources for RNA-Seq, R, general bioinformatics

Asking for help with R

ggplot2 cheatsheet

**HBC** Training materials

## Contact us!

HBC webpage: <a href="http://bioinformatics.sph.harvard.edu">http://bioinformatics.sph.harvard.edu</a>
HBC training materials: <a href="http://hbctraining.github.io/main">http://hbctraining.github.io/main</a>
HBC workshop listserv: <a href="https://tinyurl.com/hbc-mailing-list">https://tinyurl.com/hbc-mailing-list</a>

Training email: hbctraining@hsph.harvard.edu

Consulting email: bioinformatics@hsph.harvard.edu

**Twitter** 

<u>@bioinfocore</u>