

# Introduction to Single-cell RNA-seq analysis

Harvard Chan Bioinformatics Core



<https://tinyurl.com/hbc-intro-to-scrnaseq>

# Learning Objectives



- ✓ Describe best practices for designing a Single-cell RNA-seq experiment
- ✓ Describe steps in a Single-cell RNA-seq analysis workflow.
- ✓ Use Seurat and associated tools to perform analysis of single-cell expression data, including data filtering, QC, clustering, and marker identification

# Useful Resources

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## Computational packages for single-cell analysis:

<http://bioconductor.org/packages/devel/workflows/html/simpleSingleCell.html>

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

<https://scanpy.readthedocs.io/>

<https://github.com/seandavi/awesome-single-cell>

## Online courses:

<https://hemberg-lab.github.io/scRNA.seq.course/>

<https://github.com/SingleCellTranscriptomics>

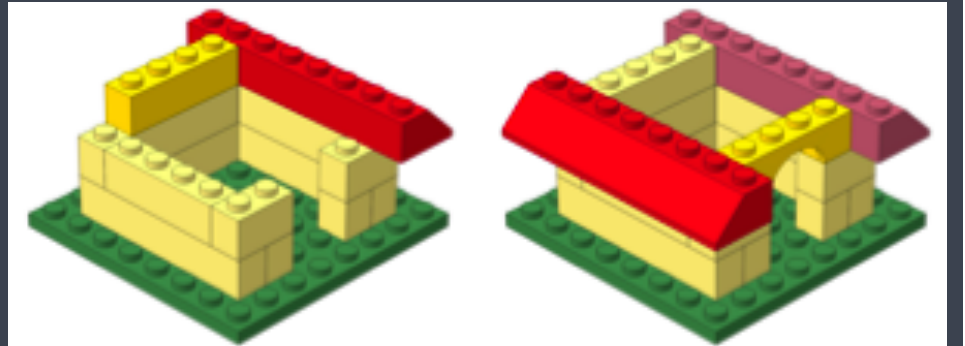
## Resources for scRNA-seq Sample Prep:

<https://www.protocols.io/>

<https://support.10xgenomics.com/single-cell-gene-expression/sample-prep>

<https://community.10xgenomics.com/>

# Survey



<https://tinyurl.com/scrnaseq-exit>

# 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	Location
March 25th	HMS' Jeffrey Modell Center for Immunology, Room 100A
April 29th	HMS' Jeffrey Modell Center for Immunology, Room 100A
May 27th	HMS' Jeffrey Modell Center for Immunology, Room 100A
June 24th	HMS' Jeffrey Modell Center for Immunology, Room 100A



# Upcoming short workshops

## HBC's monthly short workshops

- Workshops on bioinformatics methods & related skills
- Once a month for 3 hours
- First-come-first-served, no registration required
- Free and open to everyone at Harvard University and its affiliates

### Spring 2020 Schedule:

Topic	Date	Time	Location	Pre requisites
<u>Introduction to R</u>	January 15th	1 PM	HSPH Kresge G1	None
<u>Introduction to tidyverse and Data Visualization with ggplot2</u>	February 12th	1 PM	HSPH Kresge G1	<u>Introduction to R</u>
<u>Gene Annotations and Functional Analysis of Gene Lists</u>	March 18th	1 PM	HSPH Kresge G1	<u>Introduction to R</u>
<u>Generating Data Analysis Reports with RMarkdown</u>	April 15th	1 PM	HSPH Kresge G1	<u>Introduction to R</u>

# Thanks!

- Sarah Boswell, HMS
- HBC members

**<https://tinyurl.com/r-on-02>**

# Contact us!

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