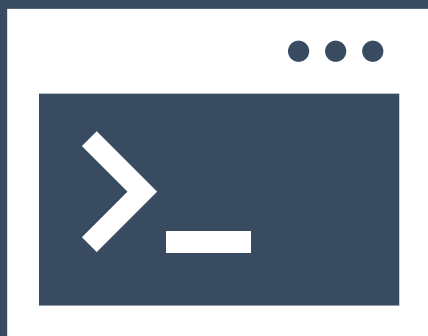


Introduction to Single-cell RNA-seq analysis

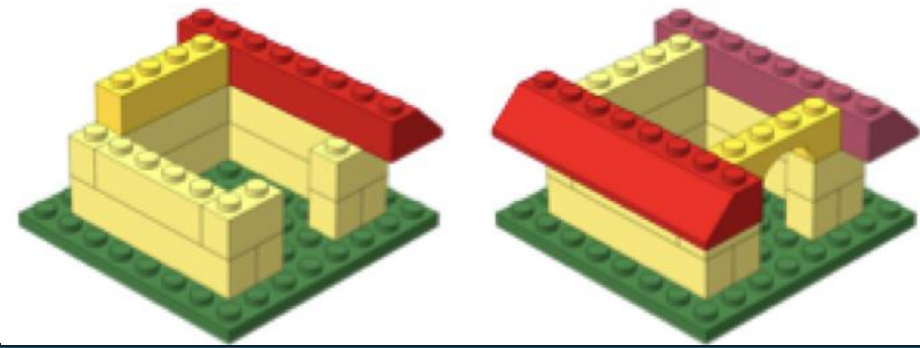
<https://tinyurl.com/hbc-scrnaseq-oct2024>



Harvard Chan Bioinformatics Core



Workshop Scope

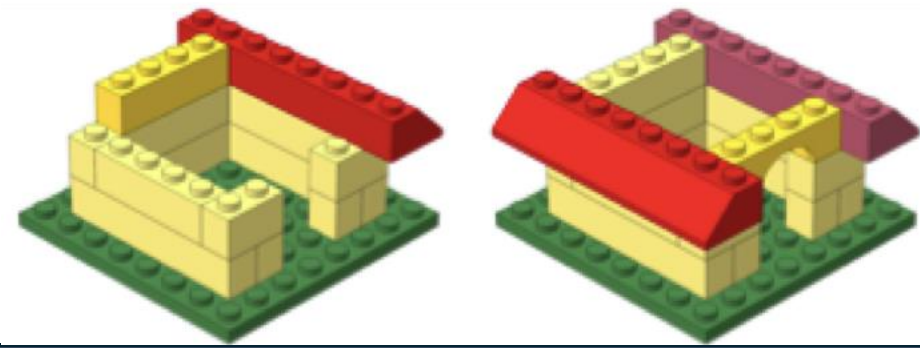


- ❖ Describe best practices for designing a single-cell RNA-seq experiment
- ❖ Describe steps in a single-cell RNA-seq analysis workflow
- ❖ Use Seurat and associated tools to perform analysis of single-cell expression data, including data filtering, QC, integration, clustering, and marker identification
- ❖ Understand practical considerations for performing scRNA-seq, rather than in-depth exploration of algorithm theory

Exit survey

<https://tinyurl.com/scRNAseq-online>

Useful resources



❖ 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:

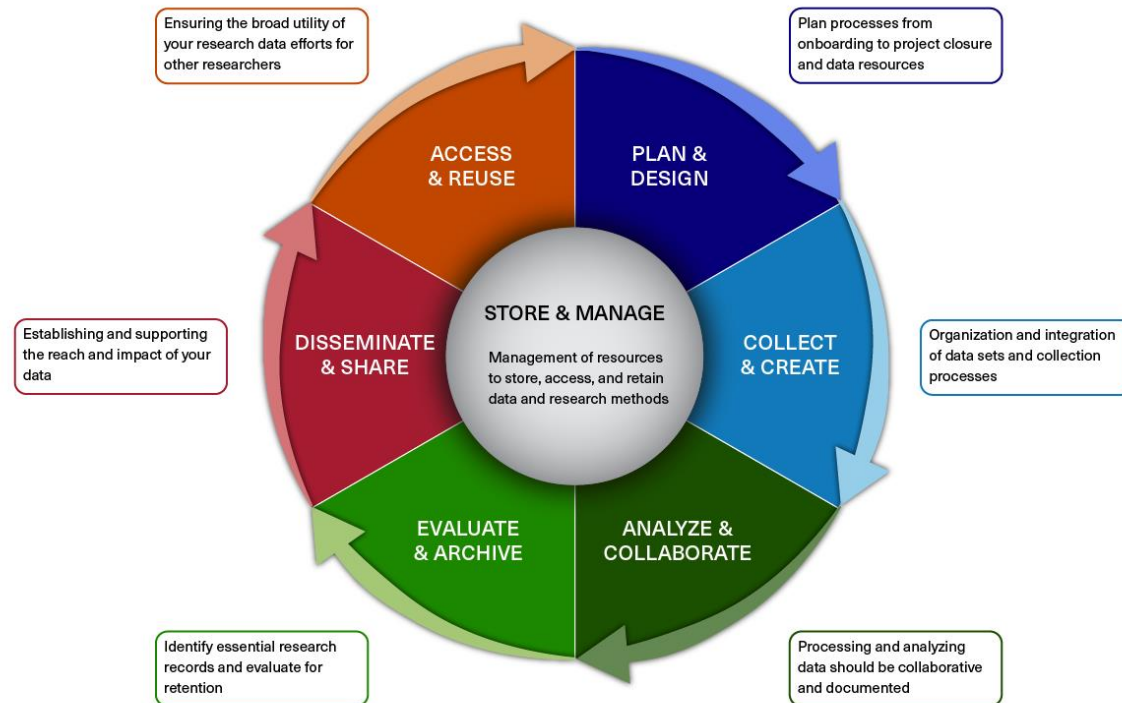
- ❖ <https://hemberg-lab.github.io/scRNA.seq.course/>
- ❖ <https://github.com/SingleCellTranscriptomics>

❖ Resources for scRNA-seq Sample Prep:

- ❖ <https://www.protocols.io/>
- ❖ <https://support.10xgenomics.com/single-cell-gene-expression/sample-prep>
- ❖ <https://community.10xgenomics.com/>

Research Data Management (RDM)

BIOMEDICAL RESEARCH DATA LIFECYCLE



Better RDM practice benefits you

- ❖ **HMS Data Management LMA**

- ❖ **Webpage:** <https://datamanagement.hms.harvard.edu>

- ❖ **Sign up** for quarterly email updates

- ❖ **Harvard-wide Research data Management**

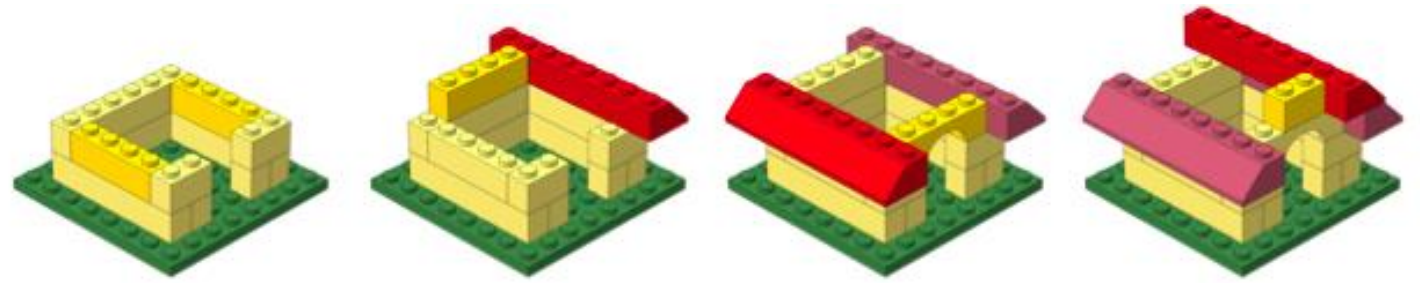
- ❖ <https://researchdatamanagement.harvard.edu/>

NOVEMBER 2024

	Date	Time	Event	Location
<input type="checkbox"/>	Nov 1	10:30am	HKS Data + Donuts: Teddy Svoronos on the Science & Implications of Generative AI	Hybrid
<input type="checkbox"/>	Nov 6	12pm	Data Discussions: Let's Name Your Data	Zoom
<input type="checkbox"/>	Nov 7	10am	LMA Research Data Management Working Group Monthly Meeting	Zoom
<input type="checkbox"/>	Nov 8	10:30am	HKS Data + Donuts: Kate Robb on Leveraging Data to Improve Neighborhoods	Hybrid
<input type="checkbox"/>	Nov 11	2pm	protocols.io Webinar: Introduction	Zoom
<input type="checkbox"/>	Nov 12	11am	Intro to NVivo (for Windows and Mac)	Lamont Library, Room 310
<input type="checkbox"/>	Nov 12	3pm	Managing and Storing Qualitative Data	Zoom
<input type="checkbox"/>	Nov 14	10am	Data Literacy: Principles of Finding and Citing Data	Zoom
<input type="checkbox"/>	Nov 14	10am	HMS RC: Intro to Python	Zoom
<input type="checkbox"/>	Nov 15	10:30am	HKS Data + Donuts: Emerging Tech Fellows on Language Access in Boston	Hybrid
<input type="checkbox"/>	Nov 19	1pm	Better Research II: Writing a Data Management and Sharing Plan	Countway Library Classroom 102 & 103

<https://datamanagement.hms.harvard.edu/about/news-events/rdmwg-calendar>

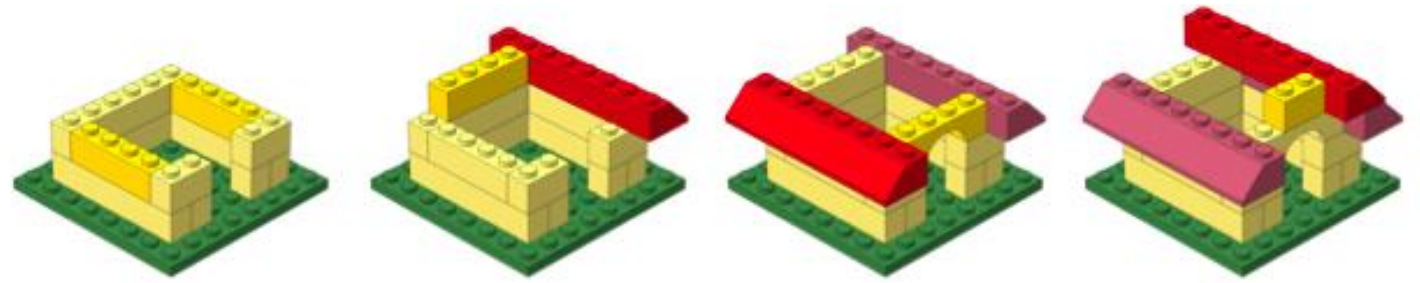
Keep building!



2024 schedule:

Topic	Pre-requisites	Date/Time	Time	Registration
Basic Shell	None	11/20/24	1 – 4pm	Register!
Tips and Tricks on O2	Shell	12/11/24*	1 – 4pm	Register!

Keep building!



Topic	Category	Date	Duration	Prerequisites
Pseudobulk	Advanced	November 12, 15, 19	Three 2.5h sessions	R
Peak analysis	Advanced	December 3, 6, 10	Three 2.5h sessions	R

Talk to us early!

Involvement in study design to optimize experiments



Thanks!

❖ **Dr. Arpita Kulkarni** – Associate Director, HMS Single Cell Core

More Information

- ❖ *HBC training materials: <https://hbctraining.github.io/main>*
- ❖ *HBC website: <http://bioinformatics.sph.harvard.edu>*

Contact Us

Sign up for our mailing list:

<https://tinyurl.com/hbc-training-mailing-list>

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