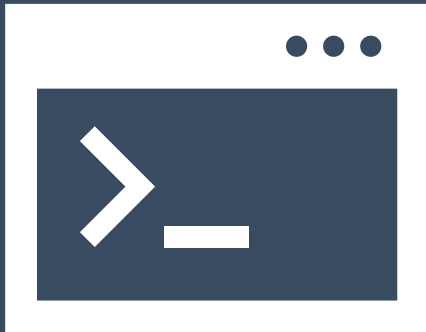


Introduction to Peak Analysis

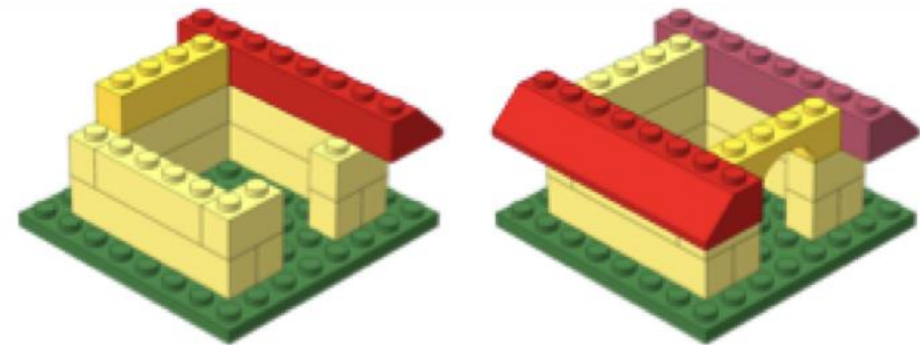
<https://tinyurl.com/hbc-peak-analysis>



Harvard Chan Bioinformatics Core



Workshop Scope



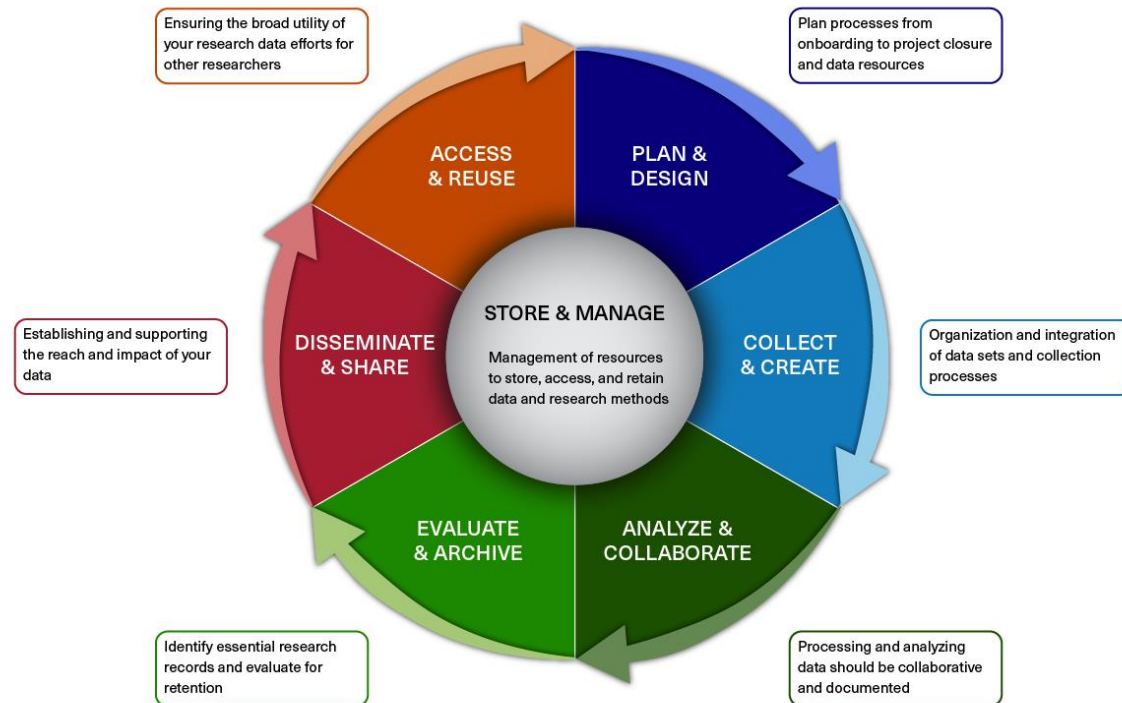
- ❖ Describe peak data and different file formats generated from peak calling algorithms
- ❖ Assess various metrics used to assess the quality of peak calls
- ❖ Compare peak calls across samples within a dataset
- ❖ Create visualizations to evaluate peak annotations
- ❖ Evaluate differentially enriched regions between two sample groups

Exit survey

<https://tinyurl.com/hbc-peak-analysis-exit-survey>

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/>

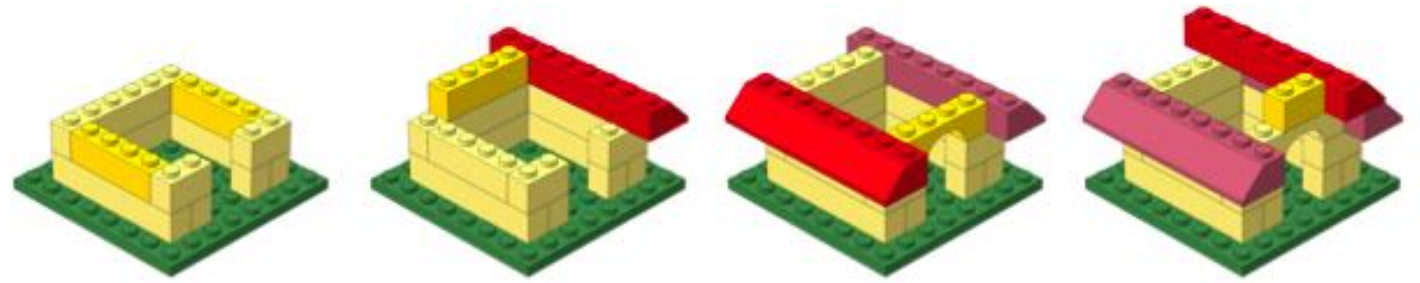
AUGUST 2025

| Date | Time | Event | Location |
|--------|------|--|----------|
| Aug 25 | 10am | Managing Your Electronic Records: Shared Drives and Email | Zoom |

SEPTEMBER 2025

| Date | Time | Event | Location |
|--------|--------|---|--------------------------------|
| Sep 4 | 10am | LMA Research Data Management Working Group Monthly Meeting | Zoom |
| Sep 5 | 1:30pm | CGA Workshop: Introduction to GIS | CGIS South Building, Room S020 |
| Sep 8 | 11am | Getting Started on the OSF: A Hands-on Guide | Zoom |
| Sep 12 | 9am | HKS: Data Carpentry | HKS Library |
| Sep 19 | 9am | HKS: Data Carpentry | HKS Library |
| Sep 25 | 10am | Which transcriptomics approach is right for you? Navigating bulk, single cell and spatial technologies | Hybrid |

Keep building!

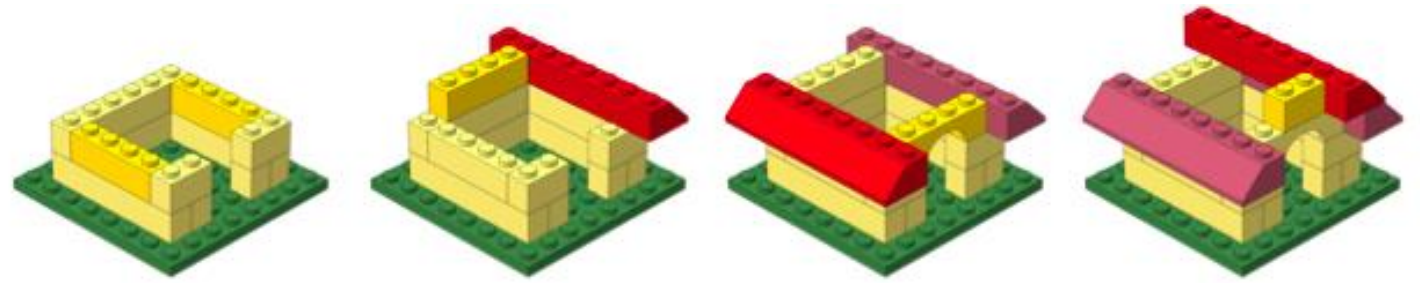


2025 schedule:

| Topic | Pre-requisites | Date | Time | Registration |
|------------------------------------|----------------------|----------|----------|-------------------------------|
| Foundations in Shell | None | 9/17/25 | 1 - 4 pm | Register now! |
| Shell - Needle in a Haystack | Foundations in Shell | 10/15/25 | 1 - 4 pm | Register now! |
| Tips and Tricks for the O2 Cluster | Foundations in Shell | 11/19/25 | 1 - 4 pm | Register now! |

<https://bioinformatics.sph.harvard.edu/current-bioinformatics-topics-workshops>

Keep building!



| Topic | Category | Date | Duration | Prerequisites |
|--|----------|-------------------------|---------------------|-------------------|
| Introduction to single-cell RNA-seq | Advanced | September 9, 12, 16 | Three 2.5h sessions | R |
| Pseudobulk and related approaches for scRNA-seq analysis | Advanced | October 21, 24, 28, 31 | Four 2.5h sessions | R |
| Introduction to Differential Gene Expression Analysis | Advanced | November 14, 18, 21, 25 | Four 2h sessions | R |

<https://bioinformatics.sph.harvard.edu/upcoming-workshops>

Talk to us early!

Involvement in study design to optimize experiments



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