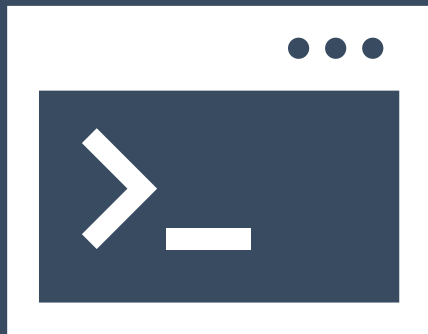


Peak Analysis

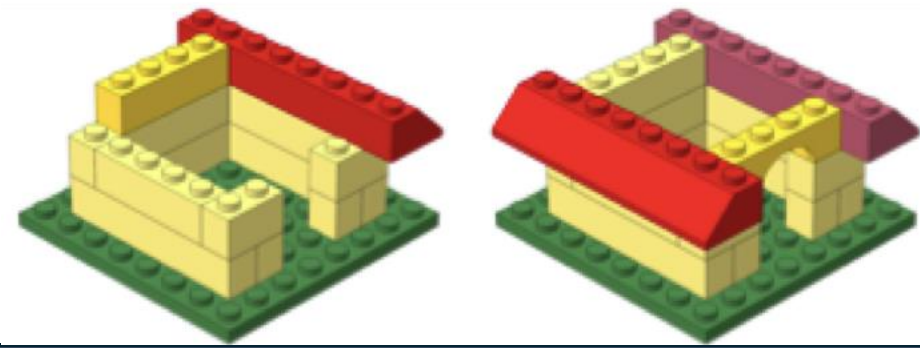
<https://tinyurl.com/Peak-analysis>



Harvard Chan Bioinformatics Core



Workshop Scope



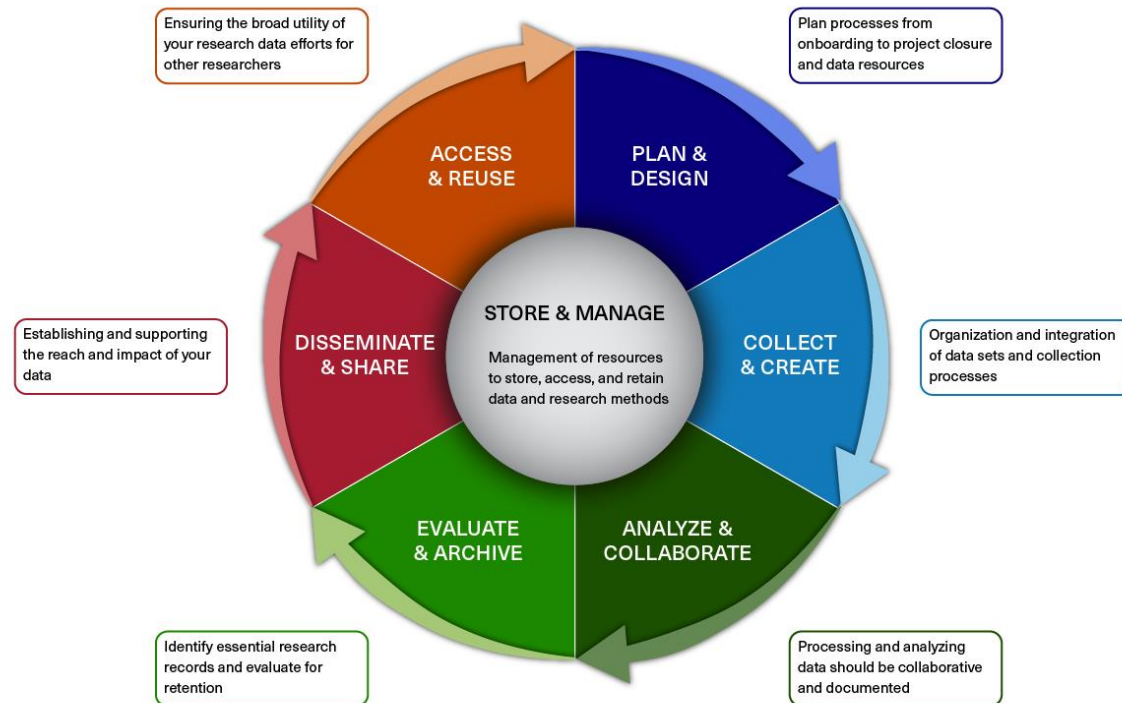
- ❖ Describe peak data and different file formats generated from peak calling algorithms
- ❖ Assess various metrics used to assess the quality of peak calls
- ❖ Compare peak calls across samples within a dataset
- ❖ Create visualizations to evaluate peak annotations
- ❖ Evaluate differentially enriched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

Exit survey



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/>


Fall 2024 Data Lifecycle Training

Plan & Design


September 24 
Managing Research
Data Efficiently


September 26 
Project and Lab
Onboarding


October 31 
Data Horror Stories:
Avoid the Nightmare

November 19 
Writing a Data Management
and Sharing Plan


Collect & Analyze

September 19 
Intro to MATLAB


October 10 
Research Computing:
Intro to Python


November 20 
Basic Shell


November 21 
Research Computing:
Intro to O2


December 5 
RCBio: easy and quick HPC
pipeline builder & runner

Store & Evaluate


October 22 
Introduction to the
General Records Schedule


October 24 
Computing Strategies
and Resources


November 22 
Managing Paper Records:
Off-Site Records Storage


December 17 
Managing Electronic
Records: Shared Drives
and Emails



Share & Publish

September 18 
Interact with your data
using RShiny

November 14 
Principles of Finding
and Citing Data

December 3 
Research Management:
Tools for Open Science

December 12 
Data Sharing in
Repositories

 In-person
 Virtual

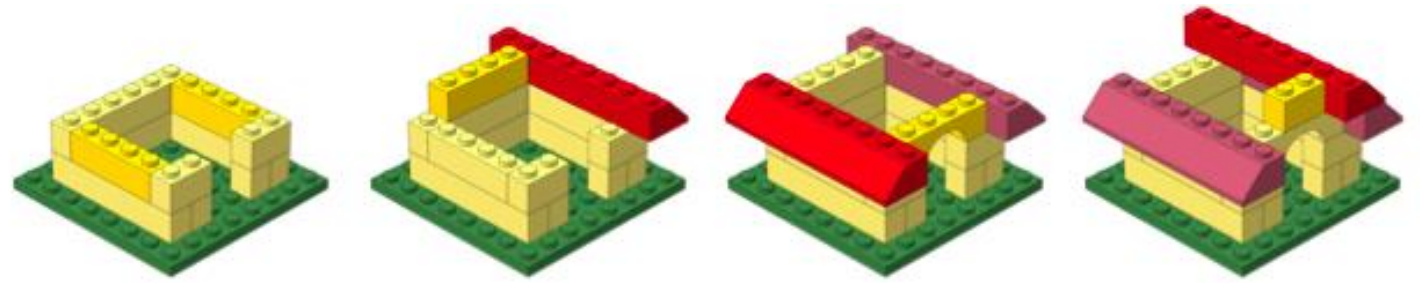


Learn More & Register
bit.ly/rdmwg-calendar

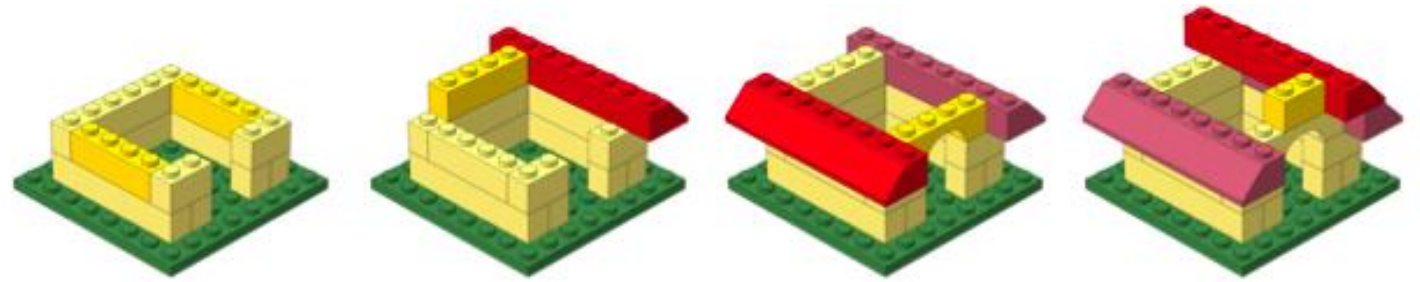


<https://datamanagement.hms.harvard.edu/about/news-events/rdmwg-calendar>

Keep building!



Keep building!

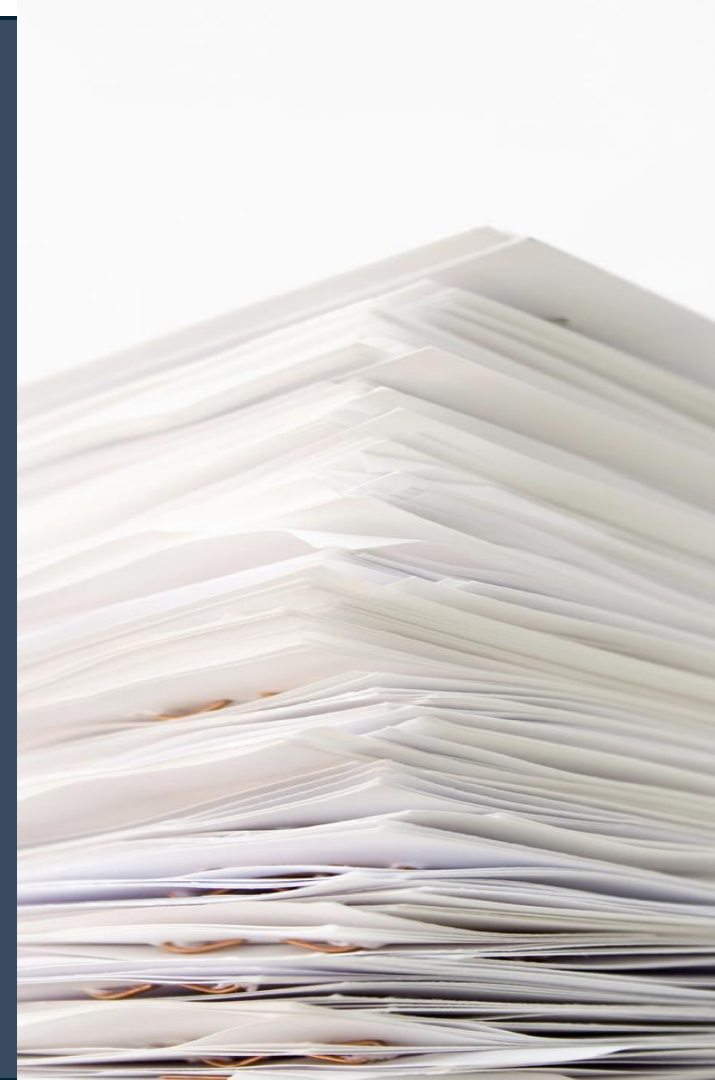


Shell for Bioinformatics	Basic	January 21, 24, 28	Three 2.5h session	None
Introduction to bulk RNA-seq data analysis Part I	Advanced	February 4, 7, 11	Three 2.5h session	Shell for Bioinformatics
Introduction to R	Basic	February 18, 21, 25 and 28	Four 2h sessions	None
Introduction to SingleCell RNA-seq	Advanced	March 4, 7, 11	Three 2.5h session	R
Introduction to Differential Gene Expression (DGE) Analysis	Advanced	March 18, 21, 25, 28	Four 2h sessions	R
Pseudobulk	Advanced	April 4, 8, 11	Three 2.5h 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>

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