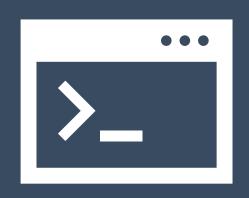


Introduction to Variant Calling

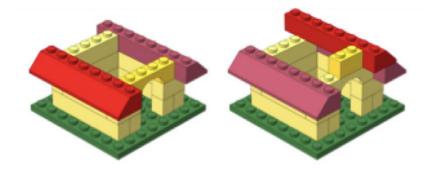
https://tinyurl.com/Intro-to-variant-analysis



Harvard Chan Bioinformatics Core



Learning Objectives

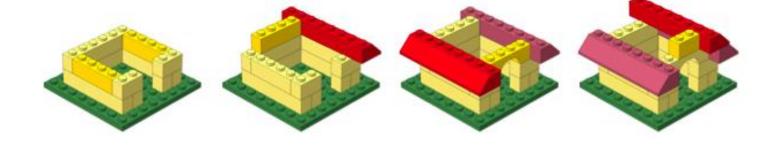


- Evaluate QC metrics for variant calling
- Call variants using GATK
- Filter variants to retain only high-quality variant calls
- Annotate variants using SnpEff and dbSNP
- Prioritize variants by their impact
- Visualize variants in IGV
- Implement cBioPortal to analyze variants

Exit survey

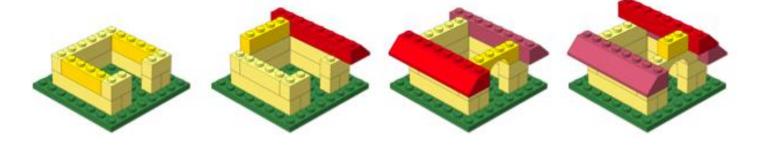
https://tinyurl.com/hbc-varcalling-exit-survey

Keep building!



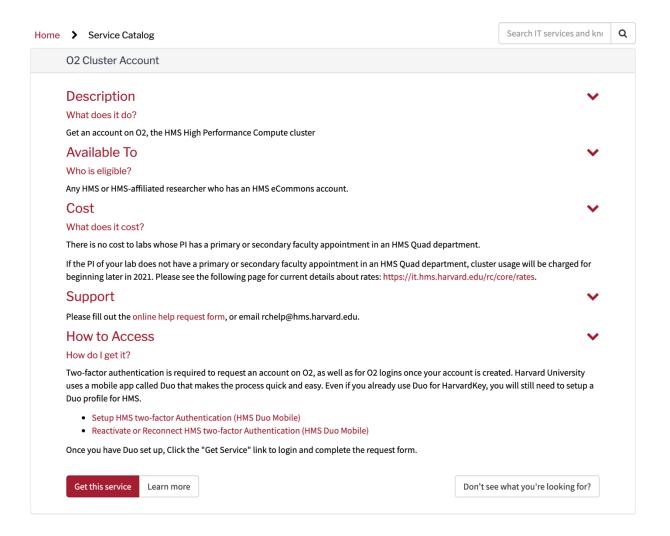
Topic	Category	Date	Duration	Prerequisites
Introduction to R	Basic	October 8, 11, 15, 18	Four 2h sessions	None
Introduction to SingleCell RNA-seq	Advanced	October 25, 29, November 1	Three 2.5h sessions	R
Pseudobulk	Advanced	November 12, 15, 19	Three 2.5h sessions	R
Peak analysis	Advanced	December 3, 6, 10	Three 2.5h sessions	R

Keep building!



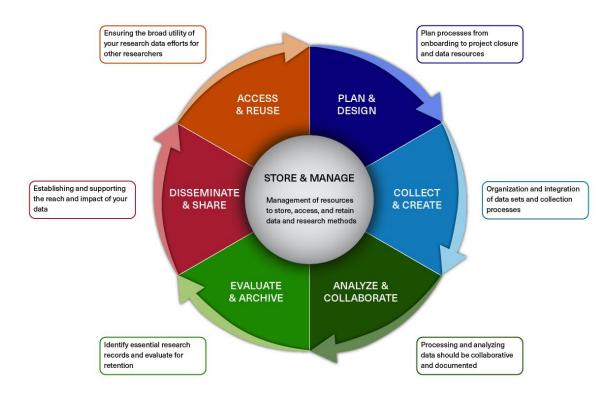
- Introduction to scRNA-seq and data pre-processing
 - October 23rd 1-3pm
 - Shell and R pre-requisite
- Introduction to shell
 - November 20th 1-4pm
- Tips and Tricks on O2
 - December 11th 1-4pm
 - Shell pre-requisite

Get an O2 account!



Research Data Management (RDM)

BIOMEDICAL RESEARCH DATA LIFECYCLE



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

October 16

scRNA pre-preocessing

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

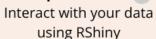
December 17



Managing Electronic Records: Shared Drives and Emails

Share & Publish

September 18



November 14



Principles of Finding and Citing Data

November 20 💂

GitHub Part I

December 3 🚇



Research Management: Tools for Open Science

December 12 💂



Data Sharing in Repositories

In-person

🖳 Virtual

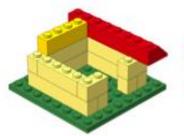


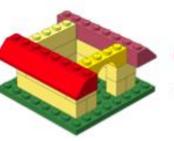
Learn More & Register bit.ly/rdmwg-calendar

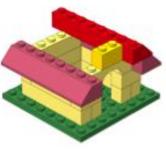


Data Discussions









- Let's Get Your Data Organized
 - October 9th 12-12:30pm
- Let's Name Your Data
 - November 6th 12-12:30pm
- Let's Write a README
 - December 11th 12-12:30pm

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/

Join us for HBC Community Breakfast!

- An opportunity to get to know others in the community
- Free food and beverages
- Great conversations



January 9th 2025 9-10:30am

More Info:

http://bioinformatics.sph.harvard.edu/breakfast/

Thanks!

- Kathleen Chappell and Andy Bergman from HMS-RC
- Dr. Tali Mazor from DFCI
- Data Carpentry

These materials have been developed by members of the teaching team at the <u>Harvard Chan Bioinformatics</u> <u>Core (HBC)</u>. These are open access materials distributed under the terms of the <u>Creative Commons</u> <u>Attribution license (CC BY 4.0)</u>, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

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