

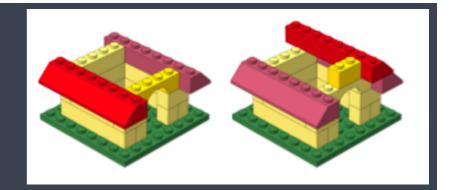
Introduction to Variant Analysis

Harvard Chan Bioinformatics Core

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

Sponsored by DF/HCC and HMS Foundry

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

Exit survey

Interested in additional training?

https://hbctraining.github.io/Training-modules/

Short workshops: Current Topics in Bioinformatics

These workshops are free and open to all researchers at Harvard University and affiliated institutions.

- Workshops on bioinformatics methods & related skills.
- · Once a month for 3 hours
- · Hands-on workshops be prepared with your MAC or Windows computer
- Free and open to everyone at Harvard University and its affiliates
- · Will meet the first Wednesday of the month (with one exception) online via Zoom
- Sign up at the links below to receive the workshop Zoom link

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Current Topics in Bioinformatics workshops 2023 Schedule (1pm - 4pm):

Topic and Link(s) to lessons	Prerequisites	Date	Registration
Introduction to Shell	None	1/18/2023	Sign up!
Intermediate Shell	None	2/15/2023	Sign up!
Advanced Shell	None	3/13/2023	Sign up!
Git/Github	Introduction to Shell	4/19/2023	Sign up!
Basics of Python	None	6/21/2023	Coming soon

Data Management Short Workshops

Bell 12pm 8 Connecting GitHub and Data (libcal.countway.harvard.edu...)
Repositories Zoom

Mar 12pm 2 Ask an Expert: NIH Data Management and Sharing Policy Virtual Office Hours (libcal.countway.harvard.edu...)

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

Get (stay) in touch with us!

Sign up for our mailing list:

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

Training email: hbctraining@hsph.harvard.edu

Consulting email: bioinformatics@hsph.harvard.edu

Twitter: @bioinfocore