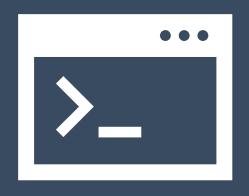
Current Topics in Bioinformatics

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



Harvard Chan Bioinformatics Core



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







Training

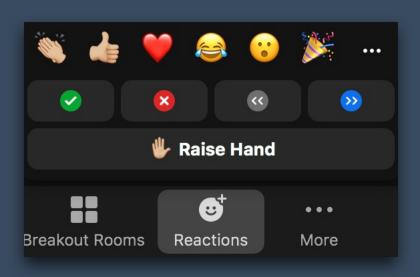
- Basic Data Skills
 - Shell
 - **❖** R
- Advanced Topics: Analysis of highthroughput sequencing data
 - Chromatin Biology
 - Bulk RNA-seq
 - Differential Gene Expression
 - scRNA-seq
 - Variant Calling
- **Current Topics in Bioinformatics**

Consulting

- Transcriptomics: RNA-seq, small RNA-seq, scRNA-seq
- Epigenetics: ChIP-seq, genome wide methylation, ATAC-seq
- DNA Variation: WGS, resequencing, exome-seq, CNV studies
- Functional enrichment analysis
- Experimental design help
- Grant support

Odds & Ends

- Quit/minimize all applications that are not required for class
- Are you all set?
 - = "agree", "I'm all set"
 - = "disagree", "I need help"



Odds & Ends

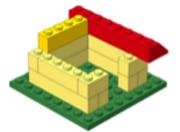
- Questions for the presenter?
 - Post the question in the Chat window OR
 - ♣ Raise Hand when the presenter asks for questions
 - Let the Moderator know
- Technical difficulties with software?
 - Start a private chat with the Troubleshooter with a description of the problem

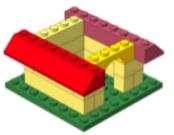
Exit survey

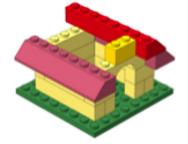
http://tinyurl.com/hbc-modules

Keep building!









Topic	Pre-requisites	Date/Time	Time	Registration
Publication Perfect II	R basics	8/21/24	1 – 4pm	Coming soon!
RShiny	R basics	9/18/24	1 – 4pm	Coming soon!

Join us for HBC Community Breakfast!

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



Thursday September 5th, 2024 9:00 to 10:30am

More Info:

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