

### Current Topics in Bioinformatics

presented by the

Harvard Chan Bioinformatics Core

Workshop materials: <a href="https://hbctraining.github.io/Training-modules/">https://hbctraining.github.io/Training-modules/</a>

HBC training team: <a href="mailto:hbctraining@hsph.harvard.edu">hbctraining@hsph.harvard.edu</a>
HBC consulting: <a href="mailto:bioinformatics@hsph.harvard.edu">bioinformatics@hsph.harvard.edu</a>







#### **Training**

- Basic Data Skills
  - Introduction to command line (Unix) and high-performance computing
  - Introduction to R
- Advanced Analyses of NGS Data
  - Bulk RNA-seq
  - Single-cell RNA-seq
  - Chromatin biology (ChIP-seq/ATACseq)
- Monthly, short workshops on various bioinformatics topics

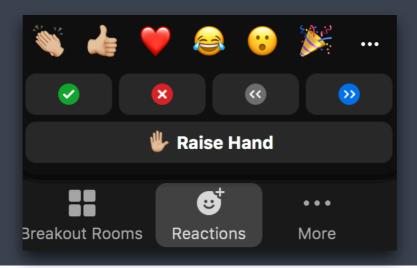
#### Consulting

- Transcriptomics: RNA-seq, small RNA-seq, scRNA-Seq
- Epigenetics: ChIP-seq, genome-wide methylation, ATAC-Seq
- DNA Variation: WGS, resequencing, exome-seq and CNV studies
- Functional enrichment analysis
- Exp. design help & grant support

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

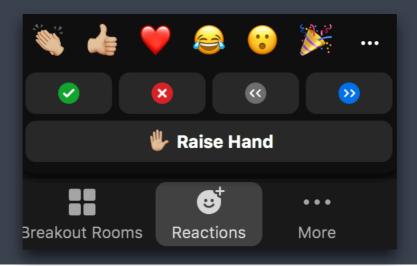
## Odds and Ends (1/2)

- Quit/minimize all applications that are not required for class
- Captioning is available upon request
- Are you all set?
  - = "agree", "I'm all set" (equivalent to a green post-it)
  - = "disagree", "I need help" (equivalent to a red post-it)



## Odds and Ends (2/2)

- Questions for the presenter?
  - Post the question in the Chat window OR
  - Raise Hand when the presenter asks for questions
  - Let the Moderator know



# Upcoming HBC workshop & events

Topic and Link(s) to lessons	Prerequisites	Date	Registration
Publication Perfect: Part II	Publication Perfect: Part I	10/18/2023	Sign up!
Rmarkdown	R Basics or Online R course - Harvard Catalyst	11/15/2023	Sign up!

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

# Exit Survey

http://tinyurl.com/hbc-modules