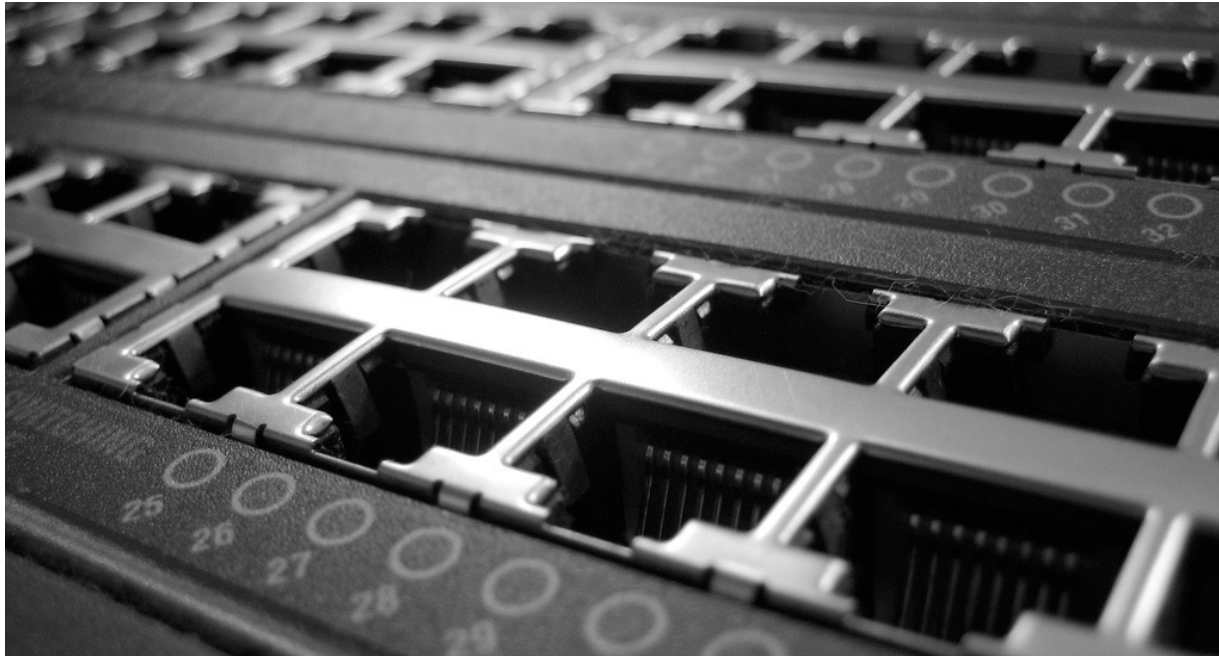**Sign-in:** https://tinyurl.com/modules-sign-in

Current Topics in Bioinformatics

presented by the

Harvard Chan Bioinformatics Core

Workshop materials:

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

HBC training team: hbctraining@hsph.harvard.edu

HBC consulting: bioinformatics@hsph.harvard.edu

Training

- Introduction to **command line** (Unix) and **high-performance computing**

- **Introductory R** and **differential gene expression** analysis

- **In-depth course**: Unix & R, RNA-Seq, ChIP-Seq, and variant calling

- 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/

Exit Survey

http://tinyurl.com/hbc-modules