Harvard Chan Bioinformatics Core presents:

NGS Data Analysis Course

For all members of the Harvard Stem Cell Institute (HSCI)

This 10-day in-depth course will instruct participants on the theoretical and practical concepts related to Next-Generation Sequencing (NGS) data analysis, enabling them to perform these analyses independently.



Topics include:

- Unix & High-Performance Computing
- NGS data analysis (RNA-Seq, ChIP-Seq, Variant calling)
- Statistical analysis using R
- Functional analysis

Prior sequence analysis or command line experience not required

June 22nd – July 21st, 2016

Course details and application instructions at:

http://tinyurl.com/ngs-course-summer-2016

Pre-registration ends on May 1st, 2016

Sponsored by the Harvard Stem Cell Institute (HSCI)