Description:
This 1.5 day hands-on workshop will introduce participants to using R and RStudio. R is a simple programming environment that enables the effective handling of data, while providing excellent graphical support. RStudio is a tool that provides a user-friendly environment for working with R.

This workshop is intended to provide basic R programming knowledge. Participants should be interested in:
- using R for increasing their efficiency for data analysis
- visualizing data using R (ggplot2)
- learning good data management practices

Workshop segments will address the following:
- R syntax: Understanding the different 'parts of speech' in R; introducing variables and functions, demonstrating how functions work, and modifying arguments for specific use cases.
- Data structures in R: Getting a handle on the classes of data structures and the types of data used by R.
- Data inspection and wrangling: Reading in data from files. Using indices and various functions to subset, merge, and create datasets, including the tidyverse suite of packages.
- Visualizing data: Visualizing data using plotting functions in base R as well as from external packages such as ggplot2.
- Exporting data and graphics: Generating new data tables and plots for use outside of the R environment.

In addition to the above, we will focus on good data management practices, installing and working with data analysis packages from various sources, and the different ways to get help when coding in R. After this workshop participants can expect to be comfortable with the basics of data wrangling and plotting using R.

Prerequisites?
This is one of our basic workshops, and has NO prerequisites and no prior programming experience is required.

This workshop is a prerequisite for advanced workshops introducing the analysis of RNA-seq and ChIP-seq data

Click here to register!
(Please check the eligibility requirements prior to registering)

Who should attend?
Eligible* Harvard researchers who:
1. want to learn how to use R and RStudio for data analysis and visualization
2. are interested in taking an advanced workshop on analysis of sequencing data within the next 6 - 8 months

Eligibility requirements:
* To be eligible to attend this workshop you should fulfill at least one of the following criteria:
- Harvard Medical School researcher in a lab on the Quad, with grants administered by HMS
- Harvard researchers whose PIs are associated with the HSCI as Principal or Affiliated faculty

If you are unsure of your eligibility, please register anyways and we will get back to you

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Schedule:
**Tuesday, March 12th - 9 AM - 5 PM**

**Wednesday, March 13th - 9 AM - 12:30 PM**

There will be one or two short coffee breaks during the day and a longer lunch break (lunch will not be provided).

**Cost and Registration:**

There is a non-refundable $35 registration fee for this workshop.

We will be accepting 25-30 participants on a first-come, first-served basis:

- If you are one of the first 25-30 eligible registrants, you will receive an email within 1 week with a link to pay the (non-refundable) $35 registration fee.
- If you are not among the first 25-30 eligible registrants, you will be added to the waitlist and accordingly notified.

**NOTE:** You will not have a reserved seat for this workshop until you pay the registration fee. Please make sure you pay within the time stated in that email, else you will lose your spot to someone on the waitlist.

**Questions?**

Please contact us at hbctraining@hsph.harvard.edu with any questions.