**Day 1 - Intro to R and Bioconductor**

*Date:* July 13, 2015, 9 am – 4 pm  
*Instructors:* Jenny Drnevich (HPCBio), Jessica Kirkpatrick (HPCBio), Neal Davis (CSE)  
*Location:* 607 IGB

**Workshop Materials:**
- Materials for Day 1
- R Installation guide
- Continued Use of R and Bioconductor guide
- R swirl guide
- Understanding R code cheat sheet
- CSAMA2015 Labs - Day 1
- Morning video
- Afternoon video

**Additional Resources:**
- Why learn R and Bioconductor:
  - Nature Toolbox: Adventures with R
  - Why you should learn R first for data science

**Day 2 - Intro to Linux & Cluster Usage; Data Management**

*Date:* July 14, 2015, 9 am – 4:30 pm  
*Instructors:* Jenny Drnevich (HPCBio), Jessica Kirkpatrick (HPCBio), Neal Davis (CSE)  
*Location:* 607 IGB

**Linux Materials:**
- Presentation
- Linux cheat sheet
- Getting on a UNIX-based terminal
- Linux / R comparison

**Git Materials/Resources:**
- Data management github repository
- Git Tutorial
- Git cheat sheet

**Additional Linux Resources:**
- Beginner Linux - self-guided tutorials for extra practice
  - Tutorial from the book *UNIX and Perl to the RESCUE!*
  - Parts 1 and 2 cover UNIX
  - [http://korflab.ucdavis.edu/Unix_and_Perl/current.html#part2](http://korflab.ucdavis.edu/Unix_and_Perl/current.html#part2)
• Orchestrating high-throughput genomic analysis with Bioconductor

How to learn R:

• The R manuals
• swirl interactive lessons (see R homework above for guide on how to get started)
• UIUC Intro to R tutorial
• Jared Knowles’ R Bootcamp
• IDRE, UCLA: Resources to help you learn and use R
• Software Carpentry’s Programming with R
• John Hopkins University’s MOOCs on Coursera.org:
  • The Data Scientist's Toolbox
  • R programming
• Code School: Try R (warning: free but must register; they send a lot of spam e-mails)
• Thomas Girke’s R & Bioconductor Manual
• Learning Statistics with R
• R reference card (good to print out)
• Writing an R package from scratch (intermediate R users)
• RStudio’s cheatsheets and online learning resources

R blogs:

• R-bloggers
• hipsteR
• RStudio blog
• R/Bioconductor blog by Sean Davis

• SIB tutorial
  • If it asks you to log in, just log in as a “guest user”
  • Click on “UNIX elearning module” at the bottom to get started
  • http://edu.isb-sib.ch/course/view.php?id=82
• University of Surrey tutorial
  • http://www.ee.surrey.ac.uk/Teaching/Unix/
• Ryan's Tutorials
  • http://ryanstutorials.net/linuxtutorial/

• Linux Books
  • UNIX and Perl to the RESCUE! by Keith Bradnam & Ian Knorf
  • Developing Bioinformatics Computer Skills by Cynthia Gibas & Per Jambeck
  • Linux Pocket Guide by Daniel J. Barrett
  • Beginning Shell Scripting by Eric Foster-Johnson, John C. Welch, & Micah Anderson
  • Learning the bash Shell by Cameron Newham
  • bash Cookbook by Carl Albing, JP Vossen, & Cameron Newham

• Linux-related Software
  • PuTTY: Allows remote access from a Windows computer to a Linux based terminal.
    • http://www.putty.org/
  • Mac already has a UNIX-based terminal pre-installed. You can find it in Applications -> Utilities -> Terminal

• Cluster Resources
  • Biocluster help page
  • Campus cluster help page
  • How to submit a cluster job
  • qsub script generator