

The Data Scientist's Toolbox

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CLI (Command Line Interface)

- `/` = root directory
- `~` = home directory
- `pwd` = print working directory (current directory)
- `clear` = clear screen
- `ls` = list stuff
 - `-a` = see all (hidden)
 - `-l` = details
- `cd` = change directory
- `mkdir` = make directory
- `touch` = creates an empty file
- `cp` = copy
 - `cp <file> <directory>` = copy a file to a directory
 - `cp -r <directory> <newDirectory>` = copy all documents from directory to new Directory *
 - `-r` = recursive
- `rm` = remove
 - `-r` = remove entire directories (no undo)
- `mv` = move
 - `move <file> <directory>` = move file to directory
 - `move <fileName> <newName>` = rename file
- `echo` = print arguments you give/variables
- `date` = print current date

GitHub

- **Workflow**
 1. make edits in workspace
 2. update index/add files
 3. commit to local repo
 4. push to remote repository
- `git add .` = add all new files to be tracked
- `git add -u` = updates tracking for files that are renamed or deleted
- `git add -A` = both of the above
 - *Note: `add` is performed before committing*
- `git commit -m "message"` = commit the changes you want to be saved to the local copy
- `git checkout -b branchname` = create new branch
- `git branch` = tells you what branch you are on
- `git checkout master` = move back to the master branch
- `git pull` = merge you changes into other branch/repo (pull request, sent to owner of the repo)
- `git push` = commit local changes to remote (GitHub)

Markdown

- **##** = signifies secondary heading (bold big font)
- **###** = signifies tertiary heading (slightly smaller font than secondary, not bold)
- * = bullet list item

R Packages

- Primary location for R packages → CRAN
- `available.packages()` = all packages available
- `head(rownames(a), 3)` = returns first three names of a
- `install.packages("nameOfPackage")` = install single package
- `install.packages(c("nameOfPackage", "nameOfPackage", "nameOfPackage"))` = install multiple package
- Bioconductor Packages:
 - `source("https://bioconductor.org/biocLite.R")`
 - `biocLite()` = install bioconductor packages
- `library(packagename)` = load package
- `search()` = see all functions in package after loading

Types of Data Science Questions

- in order of difficulty: *Descriptive* → *Exploratory* → *Inferential* → *Predictive* → *Causal* → *Mechanistic*
- **Descriptive analysis** = describe set of data, interpret what you see (census, Google Ngram)
- **Exploratory analysis** = discovering connections (correlation does not = causation)
- **Inferential analysis** = use data conclusions from smaller population for the broader group
- **Predictive analysis** = use data on one object to predict values for another (if X predicts Y, does not = X cause Y)
- **Causal analysis** = how does changing one variable affect another, using randomized studies, Strong assumptions, golden standard for statistical analysis

- **Mechanistic analysis** = understand exact changes in variables in other variables, modeled by empirical equations (engineering/physics)

Data

- **Data** = values of qualitative or quantitative variables, belonging to a set of items (usually population)
- **Variables** = measurement/characteristic of an item (qualitative vs quantitative)
- **Data** = not always structured, usually raw file, different formats
- Most important thing is question, then it is data
- **Big data** = now possible to collect data cheap, but not necessarily all useful (need the right data)