Computing for Big Data (BST-262)

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			$\frac{10}{20}$
			$\frac{19}{19}$
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Introduction

1.1 Prerequisites

For BST262 (Computing for Big Data), we assume familiarity with the material covered in BST260 (Introduction to Data Science).

We will use R to present concepts that are mostly language-agnostic. We could have used Python, as in BST261 (Data Science II).

1.2 Syllabus

Week 1 - Basic tools

- Lecture 1. Unix scripting, make
- Lecture 2. Version control: Git and GitHub (guest lecture: Ista Zhan)

Week 2 - Creating and maintaining R packages

- Lecture 3. Rationale, package structure, available tools
- Lecture 4. Basics of software engineering: unit testing, continuous integration, code coverage

Week 3 - Software optimization

- Lecture 5. Measuring performance: profiling and benchmarking tools
- Lecture 6. Improving performance: an introduction to C/C++, Rcpp

Week 4 – Databases

- Lecture 7. Overview of SQL (SQLite, PostgreSQL) and noSQL databases (HBase, MongoDB, Cassandra, BigTable, ...)
- Lecture 8. R database interfaces (in particular through dplyr)

Week 5 - Analyzing data that does not fit in memory

- Lecture 9. Pure R solutions (sampling, ff and ffbase, other interpreters). JVM solutions (h20, Spark)
- Lecture 10. An introduction to parallel computing; clusters and cloud computing. "Divide and Conquer" (MapReduce approaches)

Week 6 – Visualization

- Lecture 11. Principles of visualization (guest lecture: James Honaker)
- Lecture 12. Maps and GIS: principles of GIS, using R as a GIS, PostGIS

Weeks 7 & 8 - Guest lectures (order and precise schedule TBD)

- Software project management (Danny Brooke)
- R and Spark (Ellen Kraffmiller and Robert Treacy)
- Advanced GIS and remote sensing (TBD)
- Cluster architecture (William J. Horka)

1.3 Evaluation

Grades will be based on **two mandatory problem sets**. Each problem set will correspond to 50% (= 50 points) of the final grade. The first problem set will be available by the end of week 3 and the second problem set by the end of week 6.

You will be required to submit problem set solutions within two weeks. Grades, and feedback when appropriate, will be returned two weeks after submission.

You will submit a markdown document that combines commented code for data analysis and detailed and structured explanations of the algorithms and software tools that you used.

1.4 Software tools and packages

We will mostly use R in this course. Some examples will be run in Python.

In general, we will use free and open-source software programs such as PostgreSQL / PostGIS or Spark.

1.5 Datasets

1.5.1 MovieLens

MovieLens by Harper and Konstan (2015, https://grouplens.org/datasets/movielens/) collects datasets from the website https://movielens.org/.

There are datasets of different sizes. We will use:

- 1. Small (1MB): https://grouplens.org/datasets/movielens/latest/
- 2. Benchmark (~190MB zipped): https://grouplens.org/datasets/movielens/20m/

1.5.2 Airlines data

The airlines dataset comes from the U.S. Department of Transportation and were used in the 2009 Data Expo of the American Statistical Association (ASA).

We will use a version curated by h20o: https://github.com/h2oai/h2o-2/wiki/Hacking-Airline-DataSet-with-H2O.

1.5.3 Insurance claims

Claims data contain Protected Health Information (PHI). There are strong privacy restrictions to store, use and share this type of data.

We will use synthetic data (Sample 1) from the Centers for Medicare and Medicaid Services (CMS).

1.5.4 Census

Census data is commonly merged with administrative claims data such as Medicare. We will use data from the Census Bureau.

1.5.5 PM_{2.5} exposure

We will use $PM_{2.5}$ exposure data from the EPA Air Quality System (AQS) to illustrate GIS linkage concepts.

1.5.6 Methylation

If there is enough interest, we might present methylation examples.

1.6 Contributing with GitHub

If you have suggestions, you can open an issue on GitHub at https://github.com/cchoirat/bigdata17/issues. If you want to contribute, we welcome pull requests.

Basic tools

In this Chapter, we present basic tools that will be important when interacting with big data systems, most importantly the command-line interface (CLI) in a Unix shell and several utilities (less, vi, awk and make).

2.1 Command line tools

We assume some familiarity with the Unix shell, for example as in http://swcarpentry.github.io/shell-novice/.

We also assume that you have access to a shell, either because you use Linux or OS X or because you have the right tools on Windows (for example Cygwin or the Bash shell in Windows 10).

2.1.1 Why use the command line?

- Batch processing
- Cluster and cloud computing

2.1.2 Basic Unix commands

2.1.3 Useful tools

- 2.1.3.1 less
- 2.1.3.2 vi
- 2.1.3.3 awk

2.1.4 Example

Let's apply some of the techniques described in Blackwell and Sen (2012).

2.2 Makefiles

2.3 Git and GitHub

Guest lecture by Ista Zahn.

2.4 Buffer

```
ratings <- read.csv("~/Dropbox/data17/ml-latest-small/ratings.csv")</pre>
head(ratings)
     userId movieId rating timestamp
## 1
          1
                31
                      2.5 1260759144
## 2
          1
               1029
                        3.0 1260759179
## 3
          1
               1061
                     3.0 1260759182
## 4
               1129
                        2.0 1260759185
## 5
                        4.0 1260759205
          1
               1172
               1263
                        2.0 1260759151
links <- read.csv("~/Dropbox/data17/ml-latest-small/links.csv")</pre>
head(links)
     movieId imdbId tmdbId
##
## 1
           1 114709
                       862
## 2
           2 113497
                       8844
## 3
           3 113228 15602
## 4
           4 114885 31357
## 5
           5 113041 11862
## 6
           6 113277
                        949
movies <- read.csv("~/Dropbox/data17/ml-latest-small/movies.csv")</pre>
head(movies)
##
     movieId
                                            title
## 1
                                 Toy Story (1995)
## 2
                                   Jumanji (1995)
## 3
                         Grumpier Old Men (1995)
                        Waiting to Exhale (1995)
## 4
## 5
           5 Father of the Bride Part II (1995)
## 6
                                      Heat (1995)
                                            genres
## 1 Adventure | Animation | Children | Comedy | Fantasy
                       Adventure | Children | Fantasy
## 3
                                    Comedy | Romance
## 4
                             Comedy | Drama | Romance
## 5
                                            Comedy
## 6
                            Action|Crime|Thriller
tags <- read.csv("~/Dropbox/data17/ml-latest-small/tags.csv")</pre>
head(tags)
##
     userId movieId
                                          tag timestamp
## 1
                 339 sandra 'boring' bullock 1138537770
                                      dentist 1193435061
## 2
         15
                1955
```

2.4. BUFFER 11

##	3	15	7478	Cambodia	1170560997
##	4	15	32892	Russian	1170626366
##	5	15	34162	forgettable	1141391765
##	6	15	35957	short	1141391873

Packages

3.1 Why?

- Organize your code
- Distribute your code
- Keep versions of your code

3.2 Structure

- Folder hierarchy
 - NAMESPACE: package import / export
 - DESCRIPTION: metadata
 - R/: R code
 - man/: object documentation (with short examples)
 - tests/
 - data/
 - src/: compiled code
 - vignettes/: manual-like documentation
 - inst/: installed files
 - demo/: longer examples
 - exec, po, tools

3.3 Building steps

- R CMD build
- R CMD INSTALL
- R CMD check

$3.4\,$ R CMD build

R CMD build --help

Build R packages from package sources in the directories specified by 'pkgdirs'

3.5 R CMD INSTALL

```
R CMD INSTALL --help
```

Install the add-on packages specified by pkgs. The elements of pkgs can be relative or absolute paths to directories with the package sources, or to gzipped package 'tar' archives. The library tree to install to can be specified via '-library'. By default, packages are installed in the library tree rooted at the first directory in .libPaths() for an R session run in the current environment.

3.6 R CMD check

```
R CMD check --help
```

http://r-pkgs.had.co.nz/check.html

Check R packages from package sources, which can be directories or package 'tar' archives with extension 'tar.gz', 'tar.bz2', 'tar.xz' or 'tgz'.

A variety of diagnostic checks on directory structure, index and control files are performed. The package is installed into the log directory and production of the package PDF manual is tested. All examples and tests provided by the package are tested to see if they run successfully. By default code in the vignettes is tested, as is re-building the vignette PDFs.

3.7 Building steps with devtools

devtools::builddevtools::installdevtools::check

• and many others: load_all, document, test, run_examples, ...

3.8 Creating an R package

3.8.1 utils::package.skeleton

```
package.skeleton() # "in "fresh" session ("anRpackage")
package.skeleton("pkgname") # in "fresh" session

set.seed(02138)
f <- function(x, y) x+y
g <- function(x, y) x-y
d <- data.frame(a = 1, b = 2)
e <- rnorm(1000)
package.skeleton(list = c("f", "g", "d", "e"), name = "pkgname")</pre>
```

3.8.2 devtools::create

```
devtools::create("path/to/package/pkgname")
```

3.9 Submitting to CRAN

http://r-pkgs.had.co.nz/release.html

3.10 Using GitHub

http://r-pkgs.had.co.nz/git.html

- 3.11 RStudio and GitHub integration (1 / 7)
- 3.12 RStudio and GitHub integration (2 / 7)
- 3.13 RStudio and GitHub integration (3 / 7)
- 3.14 RStudio and GitHub integration (4 / 7)
- 3.15 RStudio and GitHub integration (5 / 7)

3.16 Command line

```
git init
git add *
git commit -m "First commit"
git remote add origin git@github.com:harvard-P01/pkgtemplate.git
git push -u origin master
```

- 3.17 RStudio and GitHub integration (6 / 7)
- 3.18 RStudio and GitHub integration (7 / 7)
- 3.19 Installing from GitHub

3.20 .gitignore (RStudio default)

```
.Rproj.user
.Rhistory
.RData
```

3.21 .gitignore (GitHub default)

```
# History files
.Rhistory
.Rapp.history

# Example code in package build process
*-Ex.R

# RStudio files
.Rproj.user/

# produced vignettes
vignettes/*.html
vignettes/*.pdf
```

3.22 RStudio projects

- .Rproj file extension, in our example pkgtemplate.Rproj
- A project has its own:
 - R session
 - .Rprofile (e.g., to customize startup environment)
 - .Rhistory
- Default working directory is project directory
- · Keeps track of project-specific recent files

3.23 Project options

```
Version: 1.0

RestoreWorkspace: Default
SaveWorkspace: Default
AlwaysSaveHistory: Default

EnableCodeIndexing: Yes
UseSpacesForTab: Yes
NumSpacesForTab: 2
Encoding: UTF-8

RnwWeave: knitr
```

```
LaTeX: pdfLaTeX

AutoAppendNewline: Yes
StripTrailingWhitespace: Yes

BuildType: Package
PackageUseDevtools: Yes
PackageInstallArgs: --no-multiarch --with-keep.source
```

3.24 Package documentation

• Functions and methods

```
VignettesPDFknitr (or Sweave)
```

3.25 Process example

Creating R Packages: A Tutorial (Friedrich Leisch, 2009)

 $\bullet \ \ https://cran.r-project.org/doc/contrib/Leisch-CreatingPackages.pdf$

3.26 Adding linreg.R in R/ directory

```
linmodEst <- function(x, y) {</pre>
  ## compute QR-decomposition of x
  qx \leftarrow qr(x)
  ## compute (x'x)^{(-1)} x'y
  coef <- solve.qr(qx, y)</pre>
  ## degrees of freedom and standard deviation of residuals
  df \leftarrow nrow(x) - ncol(x)
  sigma2 \leftarrow sum((y - x \%*\% coef)^2) / df
  ## compute sigma^2 * (x'x)^{-1}
  vcov <- sigma2 * chol2inv(qx$qr)</pre>
  colnames(vcov) <- rownames(vcov) <- colnames(x)</pre>
    coefficients = coef,
    vcov = vcov,
    sigma = sqrt(sigma2),
    df = df
  )
}
```

3.27 Running our function

3.28 And compare with lm (1 / 2)

3.29 And compare with lm (2 / 2)

```
vcov(lm1)

## (Intercept) Bwt

## (Intercept) 0.4792475 -0.17058197

## Bwt -0.1705820 0.06263081

summary(lm1)$sigma

## [1] 1.452373
```

3.30 Adding ROxygen2 documentation

```
#' Linear regression
#'
```

```
#' Runs an OLS regression not unlike \code{\link{lm}}
#'
\#' Oparam y response vector (1 x n)
\#' @param X covariate matrix (p x n) with no intercept
#'
#' Oreturn A list with 4 elements: coefficients, vcov, sigma, df
#'
#' @examples
#' data(mtcars)
#' X <- as.matrix(mtcars[, c("cyl", "disp", "hp")])
#' y <- mtcars[, "mpg"]
#' linreg(y, X)
#'
#' @export
#'
linmodEst <- function(x, y) {</pre>
  ## compute QR-decomposition of x
  qx \leftarrow qr(x)
  ## compute (x'x)^(-1) x'y
  coef <- solve.qr(qx, y)</pre>
  ## degrees of freedom and standard deviation of residuals
  df \leftarrow nrow(x) - ncol(x)
  sigma2 \leftarrow sum((y - x %*% coef) ^ 2) / df
  ## compute sigma^2 * (x'x)^{-1}
  vcov <- sigma2 * chol2inv(qx$qr)</pre>
  colnames(vcov) <- rownames(vcov) <- colnames(x)</pre>
    coefficients = coef,
   vcov = vcov,
   sigma = sqrt(sigma2),
    df = df
  )
}
```

3.31 Configure Build Tools

3.32 man/linmodEst.Rd

```
% Generated by roxygen2 (4.1.1): do not edit by hand
% Please edit documentation in R/linmodEst.R
\name{linmodEst}
\alias{linmodEst}
\title{Linear regression}
\usage{
linmodEst(x, y)
}
\arguments{
\item{y}{response vector (1 x n)}}
\item{X}{covariate matrix (p x n) with no intercept}
}
```

```
\value{
A list with 4 elements: coefficients, vcov, sigma, df
}
\description{
Runs an OLS regression not unlike \code{\link{lm}}
}
\examples{
data(mtcars)
X <- as.matrix(mtcars[, c("cyl", "disp", "hp")])
y <- mtcars[, "mpg"]
linmodEst(y, X)
}</pre>
```

3.33 Formatted output

3.34 DESCRIPTION

```
Package: pkgtemplate
Type: Package
Title: What the Package Does (Title Case)
Version: 0.1
Date: 2015-10-24
Author: Who wrote it
Maintainer: Who to complain to <yourfault@somewhere.net>
Description: More about what it does (maybe more than one line)
License: What license is it under?
LazyData: TRUE
```

3.35 NAMESPACE

export's automatically generated when parsing ROxygens2 snippets
export(linmodEst)

3.36 S3 basics

[1] "hi"

```
hello <- function() {
    s <- "Hello World!"
    class(s) <- "hi"
    return(s)
}
hello()
## [1] "Hello World!"
## attr(,"class")</pre>
```

3.37. S3 BASICS 21

3.37 S3 basics

```
print.hi <- function(...) {
  print("Surprise!")
}
hello()
## [1] "Surprise!"</pre>
```

3.38 S3 and S4 generics

```
linmod <- function(x, ...)
    UseMethod("linmod")

linmod.default <- function(x, y, ...) {
    x <- as.matrix(x)
    y <- as.numeric(y)
    est <- linmodEst(x, y)
    est$fitted.values <- as.vector(x %*% est$coefficients)
    est$residuals <- y - est$fitted.values
    est$call <- match.call()
    class(est) <- "linmod"
    return(est)
}</pre>
```

3.39 print

```
print.linmod <- function(x, ...) {
  cat("Call:\n")
  print(x$call)
  cat("\nCoefficients:\n")
  print(x$coefficients)
}</pre>
```

3.40 print

```
x <- cbind(Const = 1, Bwt = cats$Bwt)
y <- cats$Hw
mod1 <- linmod(x, y)
mod1

## Call:
## linmod.default(x = x, y = y)
##
## Coefficients:
## Const Bwt
## -0.3566624 4.0340627</pre>
```

3.41 Other methods

```
summary.linmod
print.summary.linmod
predict.linmod
plot.linmod
coef.linmod, vcov.linmod, ...
```

3.42 Formulas and model frames

```
linmod.formula <- function(formula, data = list(), ...) {
    mf <- model.frame(formula = formula, data = data)
    x <- model.matrix(attr(mf, "terms"), data = mf)
    y <- model.response(mf)
    est <- linmod.default(x, y, ...)
    est$call <- match.call()
    est$formula <- formula
    return(est)
}</pre>
```

3.43 Unit tests and testthat

```
http://r-pkgs.had.co.nz/tests.html
In package directory:
```

```
devtools::use_testthat()
```

pre-populates test/testthat/

Test files should start with test to be processed.

3.44 test_coef.R

```
data(cats, package = "MASS")
11 <- linmod(Hwt ~ Bwt * Sex, data = cats)
12 <- lm(Hwt ~ Bwt * Sex, data = cats)

test_that("same estimated coefficients as lm function", {
    expect_equal(l1$coefficients, l2$coefficients)
})
==> devtools::test()

Loading pkgtemplate
Loading required package: testthat
Testing pkgtemplate
.
Woot!
```

3.45. VIGNETTES 23

3.45 Vignettes

http://r-pkgs.had.co.nz/vignettes.html
devtools::use_vignette("linmod")

https://github.com/harvard-P01/pkgtemplate/blob/master/vignettes/linmod.Rmd

Optimization

In this Chapter, we are going to see how to measure and improve code performance

- 4.1 Measuring performance
- 4.1.1 Profiling
- 4.1.2 Benchmarking
- 4.2 Improving performance
- ${\bf 4.2.1} \quad {\bf Introduction \ to \ C/C++}$
- 4.2.2 Rcpp

Databases

- 5.1 Overview
- 5.2 SQL
- 5.3 noSQL
- 5.4 R interfaces

Big data

In this Chapter, we are going to review different approaches to handle and perform analyses on data that does not fit in memory.

- 6.1 Reading big data (that fits in memory)
- 6.1.1 R package comparison
- 6.1.2 Python
- 6.2 Sampling (can be read, not analyzed easily)
- 6.3 Pure R solutions
- 6.4 JVM solutions
- 6.4.1 h20
- 6.4.2 Spark

Visualization

We have finished a nice book.

Bibliography

Blackwell, M. and Sen, M. (2012). Large datasets and you: A field guide. *The Political Methodologist*, 20(1):2–5.

Harper, F. M. and Konstan, J. A. (2015). The movielens datasets: History and context. $ACM\ Trans.\ Interact.\ Intell.\ Syst.,\ 5(4):19:1-19:19.$