Lab2 Instruction

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Note

- All R codes are present in boxes with grey background. You can run them in your R Console. Lines leading by ## are outputs of R codes.
- All R codes are run correctly on my own Mac OS X. When you try them on your own computer, please customize your own working directory.
- The functions in **bold** are recommended to use.
- The Lab Instruction in different formats including .Rmd, .md, .html and .pdf are available on my GitHub. For your convenience, the data used and results produced in the Lab are also provided in the Github.

Lab2 contains basics for file and directory manipulation, and R data input and output. Rather than a complete collection of functions, I will introduce the frequenty-used functions from my own R experience.

Part I File and Directory Manipulation

R has a variety of functions for file and directory manipulation. The following are a few examples:

setwd() and getwd(): used to change or determine the current working directory. It's a good habit to set working directory before your data analysis as all results during your data analysis will be stored in the working directory.

list.files() and list.dirs(): returns a character vector of names of files or directories under the given directory.

file.info(): gives file size, creation time, directory vs. ordinary file status, and so on for each file whose name is in the argument, a character vector.

file.create() and dir.create(): creates files or directories with the given names if they do not already exist.

file.exists() and dir.exists(): returns a logical vector indicating whether the given file exists for each name in the first argument, a character vector.

file.copy() and file.rename(): moves files from source path to destination path.

file.remove() and unlink(): deletes the files or directories specified by the first argument, a character vector.

```
# set current working directory to DAE
setwd("/Users/tonytsai/Documents/R/DAE")
getwd()
```

[1] "/Users/tonytsai/Documents/R/DAE"

list all files including directories under current working directory DAE list.files()

```
## [1] "Lab1note.pdf" "Lab2"
## [3] "Lecture1Introduction.pdf" "LICENSE"
## [5] "README.md" "reference"
## [7] "script" "生态数据分析课程大纲.pdf"
```

extract file information for those files file.info(list.files())

```
##
                             size isdir mode
                                                           mtime
                            553464 FALSE 777 2015-09-20 12:59:30
## Lab1note.pdf
                              340 TRUE 755 2015-09-20 20:00:34
## Lab2
## Lecture1Introduction.pdf 1134795 FALSE 640 2015-09-20 12:59:16
                             1077 FALSE 644 2015-09-18 09:40:04
## LICENSE
## README.md
                              244 FALSE 644 2015-09-18 09:58:11
                              170 TRUE 777 2015-09-16 11:22:01
## reference
## script
                              136 TRUE 755 2015-09-20 20:00:26
## 生态数据分析课程大纲.pdf 253722 FALSE 777 2015-09-12 18:07:12
                                        ctime
                                                            atime uid gid
## Lab1note.pdf
                           2015-09-20 14:24:21 2015-09-20 14:24:19 501
## Lab2
                           2015-09-20 20:00:34 2015-09-20 20:00:26 501
## Lecture1Introduction.pdf 2015-09-20 12:59:29 2015-09-20 12:59:15 501
## LICENSE
                           2015-09-18 09:42:06 2015-09-20 12:49:52 501
## README.md
                           2015-09-18 09:58:11 2015-09-18 09:58:04 501
## reference
                           2015-09-16 13:27:58 2015-09-20 20:00:26 501
                           2015-09-20 20:00:26 2015-09-20 20:00:26 501 20
## script
## 生态数据分析课程大纲.pdf 2015-09-16 13:27:58 2015-09-16 13:27:58 501 20
                             uname grname
## Lab1note.pdf
                           tonytsai staff
                           tonytsai staff
## Lecture1Introduction.pdf tonytsai staff
## LICENSE
                           tonytsai staff
## README.md
                           tonytsai staff
## reference
                           tonytsai staff
## script
                           tonytsai staff
## 生态数据分析课程大纲.pdf tonytsai staff
```

list only directories under DAE list.dirs()

```
## [1] "."
## [2] "./Lab2"
## [3] "./Lab2/data"
## [4] "./Lab2/data/CMDSSS"
## [5] "./Lab2/data/CMDSSS/SURF_CLI_CHN_MUL_DAY_V3.0"
## [6] "./Lab2/Lab2"
## [7] "./Lab2/Lab2/data"
## [8] "./Lab2/Lab2/data/CMDSSS"
## [9] "./Lab2/Lab2/data/CMDSSS/SURF_CLI_CHN_MUL_DAY_V3.0"
## [10] "./reference"
## [11] "./script"
```

```
# find all R scripts under DAE and give their full path names (or absolute paths)
list.files(recursive = TRUE, pattern = ".R$", full.names = TRUE)
## [1] "./script/20150916.R"
# create a recursive directory under DAE/Lab2, which stores the TXT data
# that will be read in Part III.
if(!dir.exists("Lab2/data/CMDSSS/SURF_CLI_CHN_MUL_DAY_V3.0"))
  dir.create("Lab2/data/CMDSSS/SURF CLI CHN MUL DAY V3.0", recursive = TRUE)
# create a temporary directory under script
if(!dir.exists("script/tmp")) dir.create("script/tmp")
# create a temporary R script under tmp to say Hello World, Hello R!
file.create("script/tmp/tmp.R")
## [1] TRUE
cat("print('Hello World, Hello R!')", file = "script/tmp/tmp.R")
# excute the R script
source("script/tmp/tmp.R")
## [1] "Hello World, Hello R!"
# copy tmp.R to helloworld.R
file.copy("script/tmp/tmp.R", "script/helloworld.R")
## [1] TRUE
list.files("script", recursive = TRUE)
## [1] "20150916.R"
                      "helloworld.R" "tmp/tmp.R"
# rename helloworld.R to hello.R
file.rename("script/helloworld.R", "script/hello.R")
## [1] TRUE
list.files("script", recursive = TRUE)
## [1] "20150916.R" "hello.R"
                                 "tmp/tmp.R"
# delete all R scripts under script directory except for 20150916.R
# attempt to delete inexistent hellworld.R
file.remove(c("script/hello.R", "script/helloworld.R"))
## Warning in file.remove(c("script/hello.R", "script/helloworld.R")): cannot
## remove file 'script/helloworld.R', reason 'No such file or directory'
## [1] TRUE FALSE
```

```
list.files("script", recursive = TRUE)

## [1] "20150916.R" "tmp/tmp.R"

# delete the temporary directory that is not empty.
unlink("script/tmp", recursive = TRUE)
list.files("script")

## [1] "20150916.R"

To see all the file- and directory-related functions, type the following:
?files
```

Part II Capturing R Console Output

R provides functions to save the results that appear in your R Console into a text file.

sink(): diverts R output to a file connection.

[1] 1 2 3 4 5 6 7 8 9 10

capture.output(): sends R output to a character string or file connection.

```
# type following codes in your Console
# divert R output to CO2.txt under data directory
sink(file = "data/CO2.txt")
# load R built-in dataset CO2. type ?CO2 to see the decription about CO2 dataset.
data("CO2")
# divert the first 7 rows of CO2
head(CO2, 7)
# end the divertion
sink()
# sink() does not work in knitr, because it is already used internally to capture
# results. To make it work, you have to use {} wrap up all aboving code in a
# single expression and print results to Console explicitly.
  sink(file = "data/CO2.txt")
 data("CO2")
 print(head(CO2, 7))
  sink()
# the content of CO2.txt will be checked in Part III.
# generate a vector containing a numeric sequence from 1 to 10
1:10
```

```
# send the output to variable x
x <- capture.output(1:10)
x</pre>
```

```
## [1] " [1] 1 2 3 4 5 6 7 8 9 10"
```

```
# divert the first 7 rows of CO2 dataset to CO2.txt
capture.output(head(CO2, 7), file = "data/CO2.txt")
```

Part III Imports and Exports

```
read.table() and write.table():
read.csv() and write.csv():
read.xls() and write.xls():
read.dbf() and write.dbf():
load() and save():
data():
```

References

The following are materials on R data import/export that you can access on the Web.

• R Data Import/Export