# MQIM | R Workshop | Part 3

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## **Outline**



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# **R** Objects and Functions

## R Objects (frequently used)

Vectors

Туре	Example
Doubles	die <- c(1:6)
Characters	text <- c('R', 'Workshop')
Logicals	logic <- c(TRUE, FALSE, TRUE)

#### Matrices

```
> die <- c(1:6)</pre>
```

```
> mtx <- matrix(die, nrow =2, byrow = TRUE)</pre>
```

- Arrarys
- > ary <- array(mtx, dim=c(2,3,3))

# R Objects (frequently used)

Lists

```
> list <- list(die, mtx, ary)</pre>
```

 DataFrame is the two-dimensional version of a list. It is a very useful storage structure for data analysis. You can think of a dataframe as R's equivalent to the Excel spreadsheet.

Try for youself by running rObj.r.

Let's build a simple slot machine that you can play by running an R function. When you're finished, you'll be able to play it like this:

```
> play()
```

```
## [1] "You win $0"
```

## [1] "DD" "O"

The **play** function will need to do two things. First, it will need to randomly generate three symbols; and, second, it wil need to calculate a prize based on those symbols.

• Define your slot machine symbols and then randomly generate three symbols with the **sample** function.

```
## [1] "O" "B" "B"
```

Write a program that can take the output of get\_symbols and calculate the correct prize based on the prize rule, the payout scheme is set as:

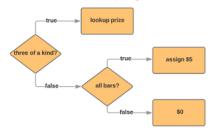
Combination	Prize
DD DD DD	100
777	80
BBB BBB BBB	40
BB BB BB	25
BBB	10
Any combo of bars	5

We call this function **score**, such that score(c("DD", "DD", "DD")) will give you \$100.

Out them together to create the full slot machine, like:

```
> play <- function(){
+    symbols <- get_symbols()
+    print(symbols)
+    score(symbols)
+ }</pre>
```

Now let's write the score function, the flow chart below describes the payout scenarios, diamond shape symbolize an *if else* decision.



```
score <- function(symbols){</pre>
    # identify case
    same <- symbols[1] == symbols[2] && symbols[2] == symbols[3]
    bars <- symbols %in% c("B","BB","BBB")</pre>
    # get prize
    if(same){
+
      payouts \leftarrow c("DD" = 100, "7" = 80, "BBB" = 40, "BB" = 25,
+
                     "B" = 10, "C" = 10, "0" = 0)
+
      prize <- unname(payouts[symbols[1]])</pre>
    } else if(all(bars)){
      prize <- 5
    } else {
      prize <- 0
+
    }
    # return prize
    prize <- paste0("You Win $", prize)</pre>
    return(prize)
+ }
```

Once the **score** function is defined, the **play** function will work as well:

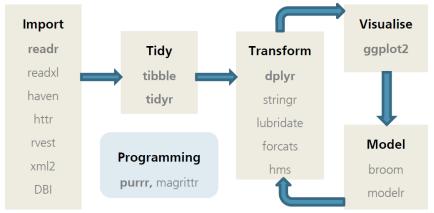
```
> play <- function(){</pre>
    symbols <- get_symbols()</pre>
    print(symbols)
    score(symbols)
> play()
## [1] "BB" "O"
## [1] "You Win $0"
> play()
## [1] "0" "0" "0"
```

[1] "You Win \$0"

## **Data manipulation and Visualization**

## tidyverse

The *tidyverse* is a set of packages for doing data science, it loads six "core" libraries that provide tolls for importing(readr), tidying(tidyr,tibble), manipulation(dplyr) and visualizing(ggplot) data, as well as support for functional programming(purrr).



For more details on using tidyverse for data science see http://r4ds.had.co.nz/.

# Data Manipulation(dplyr)

#### key verbs:

- select for subsetting variables (columns)
- mutate for creating new variables from existing ones
- filter for subsetting observations (rows)
- arrange for rearranging/ordering observations
- summarise for computing various summary statistics

Along with the **group\_by**, combining multiple simple pieces with the pipe operator %>% is a powerful way of solving complex problems.

Let's see a simple example by loading data\_df.RData, which contains information about 40 large-cap companies.

- > library(tidyverse)
- > load("data df.RData")

# Data Manipulation(dplyr)

Suppose we have the following task: Calculate the mean ROE by sector and 12-month momentum (positive or negative), making sure that there are at least 3 observations per group. Finally, arrange the data in descending order of ROE. What would you do it without advanced package except base R?

```
## Sector Momentum ROE

## 3 Consumer Staples Positive 59.76600

## 2 Consumer Discretionary Positive 39.30333
```

# Data Manipulation(dplyr)

It's more clean and straight forward by using dplyr

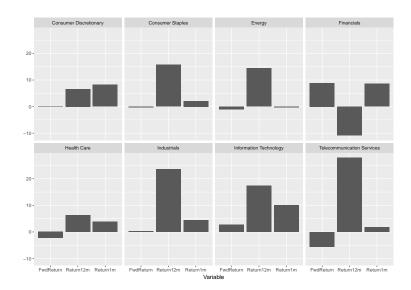
## # A tibble:  $3 \times 3$ 

# Data Visualization(ggplot)

Charts are great tools to help you understand the data from the initial data research. If you don't want to spend too much time writing your own data manipulating and plotting functions, there is a nice and quick way to do it by combining *dplyer* and *ggplot*. Here is an example to visualize the sector mean returns of the data:

```
> myplot <- data_df %>%
+ group_by(Sector) %>%
+ summarise_at(vars(contains("Return")), funs(mean)) %>%
+ tidyr::gather(Variable, Return, -Sector) %>%
+ ggplot(aes(x = Variable, y = Return)) +
+ geom_bar(stat = "identity") + ylab("") +
+ facet wrap(~ Sector, ncol = 4)
```

# Data Visualization(ggplot)



## R Markdown, R Projects and Github

### R Markdown

#### Why R Markdown?

- R Markdown provides various great formats for communicating, presenting as well as publishing rearch results.
- It enables the final report to include code chunks and output (table, plot etc.)
   while executing the codes.
- In addition, it makes typing compliated formulas and equations much less painful.
- It also ensures reproducibility and consistency. You can keep your code, notes, graphs and relevant links all in one place.
- Of course, a great way to submit your assignment.

## R Markdown

#### How it works?

- create .Rmd file, select File > New File > R Markdown. in the menubar.
   RStudio will launch a wizard that you can use to pre-populate your file with useful content that reminds you how the key features of R Markdown work.
- knit the document, R Markdown sends the .Rmd file to knitr, http://yihui.name/knitr/. R Markdown executes all of the code chunks and creates a new markdown (.md) document which includes the code and its output.
- The markdown file generated by knitr is then processed by pandoc, http://pandoc.org/. R Markdown is responsible for creating the final file.



Figure 1

## R Markdown

#### R Markdown formats:

- Documents:
  - html
  - pdf
  - word
  - github document (designed for sharing on GitHub)
- Presentations:
  - ioslides HTML presentation with ioslides
  - slidy HTML presentation with W3C Slidy
  - beamer PDF presentation with LaTeX Beamer.

Let's see a couple of examples.

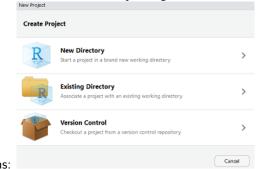
## **Rstudio Projects**

It is a good practice to create a projects for an ongoing research process with kept developing codes. The beauty of using projects are:

- Holds all the files relevant to that particular piece of work in a folder in your computer in a desired project directory. Set unite working directory to Project directory, save the trouble for typing \*setwd("C:/Users/path)\* and rm(list = ls()) for each r script.
- Dedicated R process. File browser pointed at Project directory.
- The way to create RStuodio Project is quite flexible, you can create it in a new folder, in a existing folder, or link it to a version control repository (we will talk about GitHub here).

# **Rstudio Projects**

To create a new Rstudio Projects, go to RStudio -> New Project, you will see



options:

Click **New Directory** if you'd like to make a new folder as project, or **Existing Directory** to make existing folder into an RStudio Project.

#### What is it?

- Git is an open-source version control system that was started by Linus Trovalds, whom created Linux.
- When developers create something (an app, for example), they make constant changes to the code, releasing new versions up to and after the first official release.
- Version control systems keep these revisions straight, storing the modifications in a central repository. This allows developers to easily collaborate, as they can download a new version of the software, make changes, and upload the newest revision. Every developer can see these new changes, download them, and contribute.
- People who have nothing to do with the development of a project can still download the files and use them.
- "Hub" part in GitHub is https://github.com/

- Repository: a repository("repo") is a location where all the files for a particular
  project are stored. Each project has its own repo, and you can access it with a
  unique URL.
- Forking a Repo
  - create a new project based off of another project that already exists.
  - fork the repo that you'd like to contribute to, make the changes you like and release the revised project as a new repo.
- You can fork the R\_Workshop repo from my github to yours to access some R files and data.

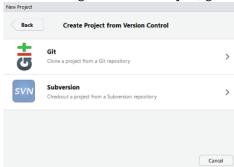
### Create a new repository A repository contains all the files for your project, including the revision history. Owner Repository name ☐ git4casey <del>-</del> r\_Workshop\_2018 Great repository names are short and memorable. Need inspiration? How about jubilant-computing-machine. Description (optional) Public Anyone can see this repository. You choose who can commit. You choose who can see and commit to this repository. ☐ Initialize this repository with a README This will let you immediately clone the repository to your computer. Skip this step if you're importing an existing repository. Add a license: None ▼ Add .gitignore: None ▼ Create repository

Figure 2

## Link Rstudio Project to Github

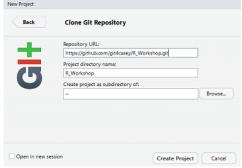
How to use it along with Rstudio Project?

When creating the Rstudio Project, go to the third option Version Control



# Link Rstudio Project to Github

- Select Git, and paste the repo URL you'd like to work on. For example, it can be the forked R\_Workshop repo on your github.
- Select the preferred the directory in your computer by Browse..., then click Create Project. Now you should be able to see all the files in the repo from Rstudio, and edit and push to update your gihub repo.



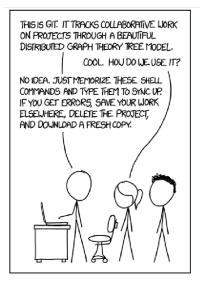
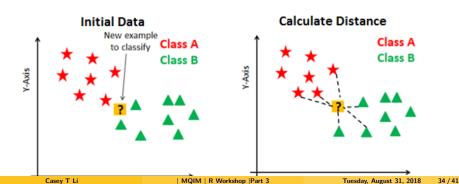


Figure 3

# Using R for k-Nearest Neighbors (KNN)

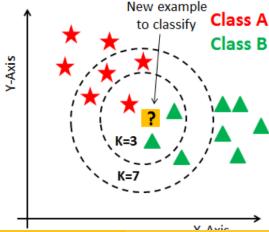
## **KNN** concepts

- The KNN or k-nearest neighbors algorithm is one of the simplest machine learning algorithms and is an example of instance-based learning, where new data are classified based on stored, labeled instances.
- More specifically, the distance between the stored data and the new instance is calculated by means of some kind of a similarity measure.
- This similarity measure is typically expressed by a distance measure such as the Euclidean distance, cosine similarity or the Manhattan distance.



## **KNN**

The number of neighbors(K) in KNN is a hyperparameter that you need choose at the time of model building. Look at the case below: The question is how to choose the optimal number of k neighbors?



## **KNN**

No unified answer that suits all kind of data sets. In general, requires test on different k and validate the performance. Research has also shown that

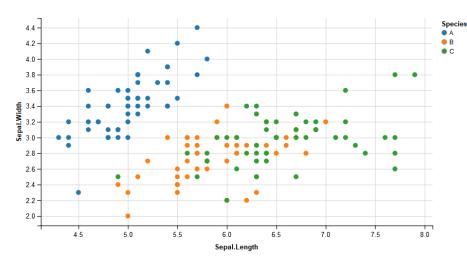
- a small amount of neighbors are most flexible fit which will have low bias but high variance
- a large number of neighbors will have a smoother decision boundary, which implies lower variance but higher bias.

Let's use famous iris dataset to understand how knn works in R. To inspect the data, some simple summary:

```
##
     Sepal.Length
                     Sepal.Width
                                      Petal.Length
                                                       Petal.Width
##
    Min.
           :4.300
                    Min.
                            :2.000
                                     Min.
                                            :1.000
                                                     Min.
                                                             :0.100
##
    1st Qu.:5.100
                    1st Qu.:2.800
                                     1st Qu.:1.600
                                                      1st Qu.:0.300
##
    Median :5.800
                    Median :3.000
                                     Median :4.350
                                                      Median :1.300
##
    Mean
           :5.843
                    Mean :3.057
                                     Mean :3.758
                                                     Mean
                                                             :1.199
##
    3rd Qu.:6.400
                    3rd Qu.:3.300
                                     3rd Qu.:5.100
                                                      3rd Qu.:1.800
##
    Max.
           :7.900
                    Max.
                            :4.400
                                     Max.
                                            :6.900
                                                     Max.
                                                             :2.500
##
          Species
              :50
##
    setosa
##
    versicolor:50
##
    virginica:50
##
##
```

##

To visulize if there is any correlation between variables.



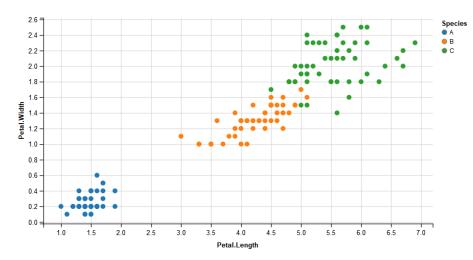


Figure 6: flowchart

- Divide the data set into two parts: a training set and a test set. (2/3, 1/3 split)
- Using the knn() function, which uses the Euclidian distance measure in order to find the k-nearest neighbours to your new, unknown instance.

```
> library(class)
> iris_pred <- knn(train = iris_training, test = iris_test, cl = ir:</pre>
```

Compare the predicted class to the test labels.

```
## [1] "Accuracy: 91.6666666666667%"
```