# MPP-E1180 Lecture 2: Introduction to the R Programming Language

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### Objectives for the topic

- Basics of object-oriented programming in R
- Simple R data structures
- ► Simple descriptive statistics and plotting with Base R

#### What is R?

**Open source programming language**, with a particular focus on statistical programming.

**History:** Originally (in 1993) an implementation of the S programming language (Bell Labs), by **R**oss Ihaka and **R**obert Gentleman (hence  $\mathbf{R}$ ) at University of Auckland.

Currently the R Foundation for Statistical Computing is based in Vienna.

#### RStudio:

RStudio is an Integrated Developer Environment (IDE) that makes using R and other reproducible research tools easier.

### Growing popularity

R can be easily expanded by  ${\bf user}$  created packages hosted on GitHub and/or CRAN.

#### How to Cite R

##

##

##

##

##

#### citation()

## To cite R in publications use:

address = {Vienna, Austria},

 $year = \{2015\},\$ 

```
##
## R Core Team (2015). R: A language and environment for
## statistical computing. R Foundation for Statistical Co
## Vienna, Austria. URL https://www.R-project.org/.
##
## A BibTeX entry for LaTeX users is
##
## @Manual{,
## title = {R: A Language and Environment for Statistic
## author = {{R Core Team}},
```

organization = {R Foundation for Statistical Comput:

# Fundamentals of the R language

R is object-oriented.

Objects are R's nouns. They include (not exhaustive):

- character string (e.g. words)
- number
- vector of numbers or character strings
- matrix
- data frame
- list

#### **Assignment**

You use the **assignment operator** (<-) to assign character strings, numbers, vectors, etc. to object names

```
## Assign the number 10 to an object called number
number <- 10
number</pre>
```

```
## [1] 10
```

```
# Assign Hello World to an object called words
words <- "Hello World"</pre>
```

```
## [1] "Hello World"
```

# Assignment

You can also use the equality sign (=):

```
number = 10
number
```

## [1] 10

Note: it has a slightly different meaning.

See StackOverflow discussion.

# Special values in R

- NA: not available, missing
- NULL: does not exist, is undefined
- ► TRUE, T: logical true. **Logical** is also an object class.
- ► FALSE, F: logical false

# Finding special values

Function	Meaning
is.na	Is the value NA
is.null	Is the value NULL
isTRUE	Is the value TRUE
!isTRUE	Is the value ${\tt FALSE}$

```
absent <- NA
is.na(absent)</pre>
```

```
## [1] TRUE
```

Operator	Meaning		
<	less than		
>	greater than		
==	equal to		
<=	less than or equal to		
>=	greater than or equal to		
! =	not equal to		
a   b	a or b		
a & b	a and b		
_			

#### Classes

Objects have distinct classes.

```
# Find the class of number
class(number)
## [1] "numeric"
# Find the class of absent
class(absent)
## [1] "logical"
```

# Naming objects

- Object names cannot have spaces
  - Use CamelCase, name\_underscore, or name.period
- Avoid creating an object with the same name as a function (e.g. c and t) or special value (NA, NULL, TRUE, FALSE).
- Use descriptive object names!
  - ▶ Not: obj1, obj2
- ► Each object name must be **unique** in an environment.
  - Assigning something to an object name that is already in use will overwrite the object's previous contents.

# Finding objects

```
# Find objects in your workspace
ls()
```

```
## [1] "absent" "number" "words"
```

### Style Guides

As with natural language writing, it is a good idea to stick to one style guide with your R code:

- Google's R Style Guide
- ► Hadely Wickham's R Style Guide

#### Vectors

A vector is an **ordered collection** of numbers, characters, etc. of the **same type**.

Vectors can be created with the c (combine) function.

```
# Create numeric vector
numeric_vector <- c(1, 2, 3)
# Create character vector
character_vector <- c('Albania', 'Botswana', 'Cambodia')</pre>
```

#### Factor class vector

Categorical variables are called **factors** in R.

```
# Create numeric vector
fruits \leftarrow c(1, 1, 2)
# Create character vector for factor labels
fruit_names <- c('apples', 'mangos')</pre>
# Convert to labelled factor
fruits factor <- factor(fruits, labels = fruit names)</pre>
summary(fruits_factor)
```

```
## apples mangos
## 2 1
```

#### **Matrices**

Matrices are collections of vectors with the same length

```
# Combine numeric_vector and character_vector into a matrix
combined <- cbind(numeric_vector, character_vector)
combined</pre>
```

```
## numeric_vector character_vector
## [1,] "1" "Albania"
## [2,] "2" "Botswana"
## [3,] "3" "Cambodia"
```

Note: In addition to cbind you can rbind new rows onto a matrix.

#### Data frames

Data frames are collections of vectors with the same length.

Each column (vector) can be of a **different class**.

#### Lists

A list is a vector containing other objects.

The objects can have different lengths and classes.

```
## $countries
## [1] "Albania" "Botswana" "Cambodia"
##
## $not_there
## [1] NA NA
##
## $more_numbers
## [1] 1 2 3 4 5 6 7 8 9 10
```

#### **Functions**

Functions do things to/with objects. Functions are like **R's verbs**.

When using functions to do things to objects, they are always followed by parentheses (). The parentheses contain the **arguments**. Arguments are separated by commas.

```
# Summarise combined_df
summary(combined_df, digits = 2)
```

```
##
   numeric vector character vector
##
   Min. :1.0
                 Length:3
##
   1st Qu.:1.5
                 Class:character
   Median :2.0
##
                 Mode :character
##
   Mean :2.0
##
   3rd Qu.:2.5
##
   Max. :3.0
```

### Functions help

Use ? to find out what arguments a function can take.

?summary

The help page will also show the function's **default argument** values.

# Component selection (\$)

The \$ is known as the component selector. It selects a component of an object.

```
combined_df$character_vector
```

```
## [1] "Albania" "Botswana" "Cambodia"
```

# Subscripts []

You can use subscripts [] to also select components.

For data frames they have a [row, column] pattern.

```
# Select the second row and first column of combined_df
combined_df[2, 1]
## [1] 2
```

```
# Select the first two rows
combined_df[c(1, 2), ]
```

# Subscripts []

```
# Select the character_vector column
combined_df[, 'character_vector']
```

```
## [1] "Albania" "Botswana" "Cambodia"
```

# Assigment with elements of objects

You can use assignment with parts of objects. For example:

```
combined_df$character_vector[3] <- 'China'
combined_df$character_vector</pre>
```

```
## [1] "Albania" "Botswana" "China"
```

You can even add new variables:

```
combined_df$new_var <- 1:3
combined_df</pre>
```

#### **Packages**

You can greatly expand the number of functions by installing and loading user-created packages.

```
# Install dplyr package
install.packages('dplyr')

# Load dplyr package
library(dplyr)
```

You can also call a function directly from a specific package with the double colon operator (::).

```
Grouped <- dplyr::group_by(combined_df, character_vector)</pre>
```

#### R's build-in data sets

List internal data sets:
data()
Load <b>swiss</b> data set:
data(swiss)
Find data description:
?swiss

#### R's build-in data sets

Find variable names:

## Franches-Mnt

```
names(swiss)
```

```
## [1] "Fertility" "Agriculture" "Examination"
## [4] "Education" "Catholic" "Infant.Mortal
```

See the first three rows and four columns

```
head(swiss[1:3, 1:4])
```

92.5

##	Fertility	Agriculture	Examination	Education
## Courtelary	80.2	17.0	15	1:
## Delemont	83.1	45.1	6	9

39.7

4 D > 4 B > 4 B > 4 B > 9 Q (2)

5

# What all the cool kids are doing: piping

**Pipe**: pass a value forward to a function call.

#### Why?

- Faster compilation.
- Enhanced code readability.

In R use %>% from the magrittr (or dplyr) package.

%>% passes a value to the **first argument** of the next function call.

#### Simple piping example

Not piped:

```
values <- rnorm(1000, mean = 10)
value_mean <- mean(values)
round(value_mean, digits = 2)</pre>
```

## [1] 10

Piped:

```
library(magrittr)
rnorm(1000, mean = 10) %>% mean() %>% round(digits = 2)
```

```
## [1] 10.04
```

### **Creating Functions**

You can create a function to find the sample mean  $(\bar{x} = \frac{\sum x}{n})$  of a vector.

```
fun_mean <- function(x){
    sum(x) / length(x)
}
## Find the mean
fun_mean(x = swiss$Examination)</pre>
```

```
## [1] 16.48936
```

# Why functions?

#### Functions:

- Simplify your code if you do repeated tasks.
- Lead to fewer mistakes.
- Are easier to understand.
- Save time over the long run—a general solution to problems in different contexts.

#### Descriptive statistics: review

Descriptive Statistics: describe samples

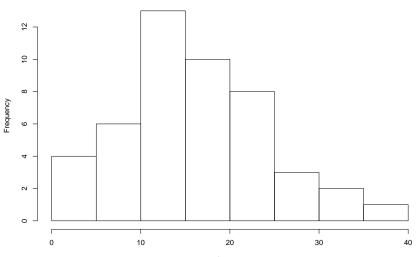
Stats 101: describe samples **distributions** with appropriate measure of

- central tendancy
- variability

# Histograms

#### hist(swiss\$Examination)

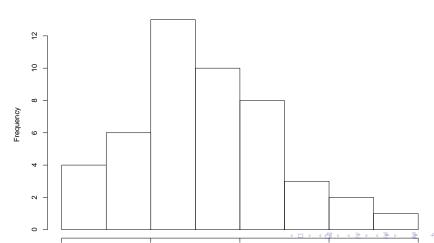
#### Histogram of swiss\$Examination



# Histograms: styling

```
hist(swiss$Examination,
    main = 'Swiss Canton Draftee Examination Scores (1888)
    xlab = '% receiving highest mark on army exam')
```

#### Swiss Canton Draftee Examination Scores (1888)



## Finding means

```
(or use the mean function in base R)
mean(swiss$Examination)
## [1] 16.48936
If you have missing values (NA):
mean(swiss$Examination, na.rm = TRUE)
```

## Digression: Loops

You can 'loop' through the data set to find the mean for each column

```
for (i in 1:length(names(swiss))) {
    swiss[, i] %>%
    mean() %>%
    round(digits = 1) %>%
    paste(names(swiss)[i], ., '\n') %>% # the . directs the cat()
}
```

```
## Fertility 70.1
## Agriculture 50.7
## Examination 16.5
## Education 11
## Catholic 41.1
## Infant.Mortality 19.9
```

# Other functions for central tendency

#### Median

```
median(swiss$Examination)
```

## [1] 16

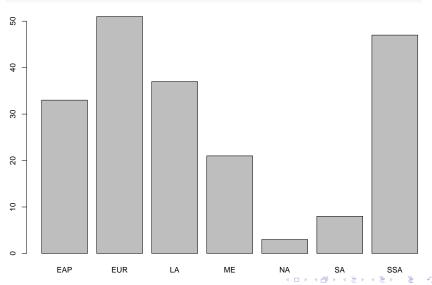
#### Mode

mode is not an R function to find the statistical mode.

Instead use summary for factor nominal variables or make a bar chart.

# Simple bar chart for nominal

```
devtools::source_url('http://bit.ly/OTWEGS')
plot(MortalityGDP$region, xlab = 'Region')
```



### Variation

#### Range:

```
range(swiss$Examination)
```

```
## [1] 3 37
```

#### Quartiles:

#### summary(swiss\$Examination)

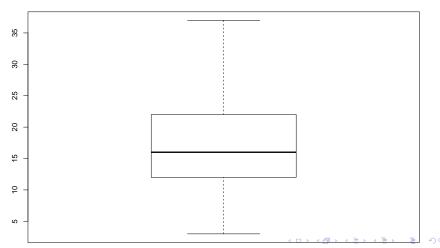
```
## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 3.00 12.00 16.00 16.49 22.00 37.00
```

## Variation

#### **Boxplots:**

boxplot(swiss\$Examination, main = '% of Draftees with High

#### % of Draftees with Highest Mark



## Variation

Interquartile Range ( $IQR = Q_3 - Q_1$ ):

IQR(swiss\$Examination)

## [1] 10

### Variation: standard deviation

Sum of squared deviations:

Sum of Squares = 
$$\sum (x - \bar{x})^2$$

Degrees of freedom (number of values that are free to vary):

$$df = n - 1$$

Variance  $(s^2)$ :

$$s^2 = \frac{\text{Sum of Squares}}{\text{Degrees of Freedom}} = \frac{\sum (x - \bar{x})^2}{n - 1}$$

**Standard deviation (s)** (in terms of the mean):

$$s = \sqrt{s^2}$$

### Variation: Standard Error

The **standard error** of the mean:

If we think of the variation as around a central tendancy as a measure of **unreliability** then we want the measure to **decrease as the sample size goes up**.

$$\mathrm{SE}_{\bar{x}} = \frac{s}{\sqrt{n}}$$

## Variation: Variance and Standard Deviation

#### Variance:

```
var(swiss$Examination)
```

## [1] 63.64662

#### Standard Deviation:

```
sd(swiss$Examination)
```

## [1] 7.977883

### Variation: Standard Error

#### **Standard Error:**

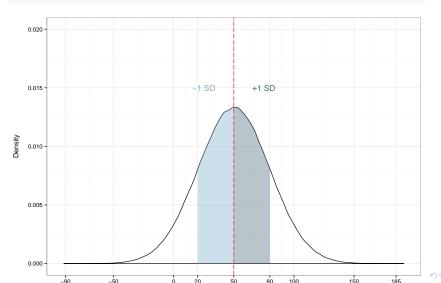
```
sd_error <- function(x) {
    sd(x) / sqrt(length(x))
}
sd_error(swiss$Examination)</pre>
```

```
## [1] 1.163694
```

## Playing with distributions

Simulated normally distributed data with SD of 30 and mean 50

Normal30 <- rnorm(1e+6, mean = 50, sd = 30)

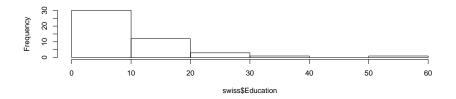


#### Transform skewed data

Highly skewed data can be transformed to have a normal distribution.

Helps correct two violations of key assumptions: (a) non-linearity and (b) heteroskedasticity.

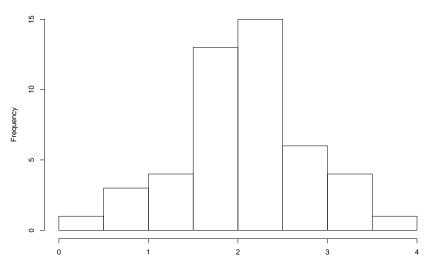
hist(swiss\$Education, main = '')



# Natural log transformed skewed data

log(swiss\$Education) %>% hist()





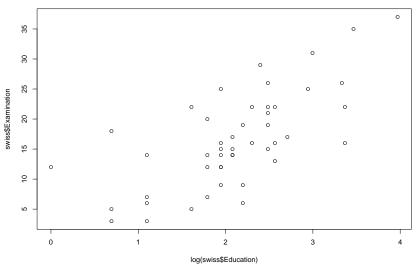
#### **Transformations**

The natural log transformation is only useful for data that **does not contain zeros**.

See http://robjhyndman.com/hyndsight/transformations/ for suggestions on other transformations such as Box-Cox and Inverse Hyperbolic Sine.

### Joint distributions

### plot(log(swiss\$Education), swiss\$Examination)



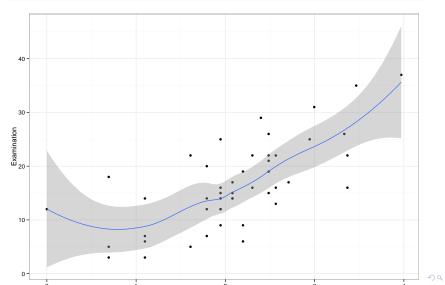
### Summarise with correlation coefficients

cor.test(log(swiss\$Education), swiss\$Examination)

```
##
    Pearson's product-moment correlation
##
##
## data: log(swiss$Education) and swiss$Examination
## t = 6.4313, df = 45, p-value = 7.133e-08
## alternative hypothesis: true correlation is not equal to
## 95 percent confidence interval:
## 0.5053087 0.8168779
## sample estimates:
##
         cor
## 0.6920531
```

### Summarise with loess

```
ggplot2::ggplot(swiss, aes(log(Education), Examination)) +
   geom_point() + geom_smooth() + theme_bw()
```



# Programming Hint (1)

### Always close!

In R this means closing:

- **)**
- **[**]
- **▶** {}
- **▶** 1
- **▶** II I

# Programming Hint (2)

Make your code as **simple as possible**.

- ▶ Easier to read.
- Easier to write (ultimately).
- Easier to find mistakes.
- Often computationally more efficient.

One way to do this is to **define things once**—e.g. use variables to contain values and custom functions to contain multiple sequential function calls.

# Programming Hint (2)

#### Bad

```
mean(rnorm(1000))

## [1] 0.01963857

sd(rnorm(1000))

## [1] 1.019639
```

# Programming Hint (2)

#### Good

```
rand_sample <- rnorm(1000)</pre>
mean(rand_sample)
## [1] -0.01687148
sd(rand_sample)
## [1] 0.9936785
```

# Seminar: Start using R!

- Access R data sets
- Explore the data and find ways to numerically/graphically describe it.
- ► Find and use R functions that were **not covered** in the lecture for exploring and transforming your data.
- Create your own function (what it does is open to you).