

#### OLAH DATA KUANTITATIF MENGGUNAKAN BAHASA R (R-LANGUAGE)

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#### Outline

- Pengenalan Penelitian Kuantitatif dan Applied Statistics
- Pengenalan Bahasa Pemrograman R
- Applied Statistics di R
  - Visualisasi Data di R
  - Descriptive Statistics di R
  - Inference Statistics di R
- Wawasan: Penelitian dengan R untuk implementasi dan model komputasi.

## Penelitian Kuantitatif dan Applied Statistics

#### Metode Kuantitatif

 Quantitative research is defined as a systematic investigation of phenomena by gathering quantifiable data and performing statistical, mathematical, or computational techniques.





#### Quantitative Designs:

• **Experimental:** To determine whether an activity or materials **make a difference** in results for participants.

#### • Correlational: To

examine/measure the **association** or relation of one or more variables than in testing the impact of activities or materials.

• **Survey:** to describe trends in a large population of individuals by giving **questionnaire** to a small group of people

Flow of the Research Process through Quantitative and Qualitative Research



#### Statistics in Research

 Research is an endeavour to discover answers to intellectual and practical problems through the application of scientific method.

 Research is a systematized effort to gain **new knowledge**.



#### Tipe Statistics

- 1. Applied statistics: Descriptive statistics and the application of inferential statistics.
- 2. Mathematical statistics: The manipulation of probability distributions necessary for deriving results related to methods of estimation and inference, various aspects of computational statistics and the design of experiments.
- **3.** Theoretical statistics: The logical arguments underlying justification of approaches to statistical inference, as well as encompassing mathematical statistics.

#### **Applied Statistics**

- Descriptive Statistics: to **describe** the characteristics of the sample in an accurate and unambiguous fashion in such a way that the information will be easily **communicated** to others → Distribution (frequency distribution), Central tendency (mean, median, mode), Dispersion (range and std deviation), and their visualizations.
- Inferential Statistics: working with a sample we introduce some unknown amount of error due to the effects of chance. Inferential statistics allow conclusions about a population based on data from a sample.

Inferential statistics consist of techniques that allow us to study samples and then make generalizations about the populations from which they were selected.

# Relationship between population and sample

- Specifically, when a researcher finishes **examining the sample**, the goal is **to generalize** the results back to the entire population.
- Remember that the research started with a general question about the population.
- To answer the question, a researcher studies a sample and then generalizes the results from the sample to the population.



### Sampling Error

**Sampling error** is the naturally occurring discrepancy, or error, that exists between a sample statistic and the corresponding population parameter.

#### FIGURE 1.2

A demonstration of sampling error. Two samples are selected from the same population. Notice that the sample statistics are different from one sample to another and all the sample statistics are different from the corresponding population parameters. The natural differences that exist, by chance, between a sample statistic and population parameter are called sampling error.



Population of 1000 college students

Population Parameters Average Age = 21.3 years Average IQ = 112.5 65% Female, 35% Male

Sample #2

Tom Kristen Sara Andrew John

Sample Statistics Average Age = 20.4 Average IQ = 114.2 40% Female, 60% Male



#### Step 3

Inferential statistics: Interpret results The sample data show a 4-point difference between the two methods of studying. However, there are two ways to interpret the results.

- There actually is no difference between the two studying methods, and the sample difference is due to chance (sampling error).
- There really is a difference between the two methods, and the sample data accurately reflect this difference.

The goal of inferential statistics is to help researchers decide between the two interpretations.

#### FIGURE 1.3

The role of statistics in experimental research.

#### Reproducibility and Replicability

- **Reproducibility** is the ability to get the same research results using the raw data and computer programs provided by the researchers.
- **Replicability** is the ability to independently achieve similar conclusions when differences in sampling, research procedures and data analysis methods may exist.
- **Reproducibility and replicability** together are among the main principles of the scientific method.

## Introduction to R Programming Language

#### What is R?

- R is a programming language and software environment for statistical computing and graphics.
- R is an implementation of the **S programming** language combined with lexical scoping semantics inspired by Scheme.
- R was created by Ross Ihaka and Robert Gentleman at the University of Auckland, New Zealand.
- R is a GNU project/Open Source.
- R is an **interpreted language**; users typically access it through a command-line interpreter.

### Why do we use R?

- It has two repositories (>16.000 R Packages):
  - 1. Comprehensive R Archive Network (CRAN, <u>https://cran.r-</u> project.org/)
  - 2. The Bioconductor project (https://www.bioconductor.org
- The quality of many packages is backed through the following highly reputed academic journals: Journal of Statistical Software, the R Journal, and **Bioinformatics**.



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#### CRAN Task Views

CRAN task views aim to provide some guidance which packages on CRAN are relevant for tasks related to a certain topic. They give a brief overview of the included packages and can be automatically installed using the cty package. The views are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are intended to have a sharp focus so that it is sufficiently clear which packages should be included (or excluded) - and they are included (or excluded) - and t

- · To automatically install the views, the ctv package needs to be installed, e.g., via

and then the views can be installed via install.views or update.views (where the latter only installs those packages are not installed and up-to-date), e.g.

- The task views are maintained by volunteers. You can help them by suggesting packages that should be included in their task views. The contact e-mail addresses are listed on the
  - · For general concerns regarding task views contact the ctv package maintaine

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<u>temPhys</u>	Chemometrics and Computational Physics
inicalTrials	Clinical Trial Design, Monitoring, and Analysis
uster	Cluster Analysis & Finite Mixture Models
atabases	Databases with R
fferentialEquations	Differential Equations
stributions	Probability Distributions
onometrics	Econometrics
vironmetrics	Analysis of Ecological and Environmental Data
perimentalDesign	Design of Experiments (DoE) & Analysis of Experimental Data
ttremeValue	Extreme Value Analysis
nance	Empirical Finance
inctionalData	Functional Data Analysis
enetics	Statistical Genetics
aphics	Graphic Displays & Dynamic Graphics & Graphic Devices & Visualization
ghPerformanceComputing	High-Performance and Parallel Computing with R
drology	Hydrological Data and Modeling
achineLearning	Machine Learning & Statistical Learning
edicalImaging	Medical Image Analysis

### Survey 2018

• <u>https://www.kdnuggets.</u> <u>com/2018/05/poll-tools-</u> <u>analytics-data-science-</u> <u>machine-learning-</u> <u>results.html</u> Top Analytics, Data Science, Machine Learning Tools



#### R Installation

- R installer: Go to https://cran.rstudio.com/
- rStudio: Go to https://www.rstudio.com/pr oducts/rstudio/download/
- Follow the instructions.
- Open R or rStudio.

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#### Simple Commands in R

• At the R prompt we type expressions. The <- symbol is the assignment operator.

```
> x <- ]
> print(x)
[1] 1
> x
[1] 1
> msg <- "hello"
> msg
[1] "hello"
> x <- c(0, 2, 5, 8, 9)
> sum(x)/length(x)
[1] 4.8
> mean(x)
[1] 4.8
```

### R Objects

- R has 5 basic or atomic classes of objects:
  - character,
  - numeric (real numbers),
  - integer,
  - complex,
  - logical (True/False).
- Vector, Matrix, List, Data Frame



#### Functions in R





#### Data from/to a file

```
> install.packages("xlsx")
```

>library(xlsx)

```
>my_data1 <- read.xlsx("test.xlsx", sheetIndex =
1, header=TRUE)</pre>
```

```
> my_data2 <- data.frame(ind = c(1,2,3),
initial=c("LSR", "BD", "AL"))
```

> write.xlsx(my data2, "test1.xlsx")

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Task Views

<u>R Homepage</u> The R Journal CRAN Task Views

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- To automatically install the views, the <u>ctv</u> package needs to be installed, e.g., via
- install.packages("ctv")

and then the views can be installed via install.views or update.views (where the latter only installs those packages are not installed and up-to-date), e.g., ctv::install.views("Econometrics")

- ctv::update.views("Econometrics")
- The task views are maintained by volunteers. You can help them by suggesting packages that should be included in their task views. The contact e-mail addresses are listed on the individual task view pages.
- For general concerns regarding task views contact the <u>ctv</u> package maintainer.

Topics

Bayesian	Bayesian Inference
ChemPhys	Chemometrics and Computational Physics
ClinicalTrials	Clinical Trial Design, Monitoring, and Analysis
Cluster	Cluster Analysis & Finite Mixture Models
Databases	Databases with R
<b>DifferentialEquations</b>	Differential Equations
Distributions	Probability Distributions
<b>Econometrics</b>	Econometrics
Environmetrics	Analysis of Ecological and Environmental Data
ExperimentalDesign	Design of Experiments (DoE) & Analysis of Experimental Data
<u>ExtremeValue</u>	Extreme Value Analysis
Finance	Empirical Finance
<b>FunctionalData</b>	Functional Data Analysis
Genetics	Statistical Genetics
Graphics	Graphic Displays & Dynamic Graphics & Graphic Devices & Visualization
HighPerformanceComputing	g High-Performance and Parallel Computing with R
<u>Hydrology</u>	Hydrological Data and Modeling
MachineLearning	Machine Learning & Statistical Learning
MedicalImaging	Medical Image Analysis

• R Package == Software Library

> install.packages("car")

- > install.packages(c("car", "MASS"))
- > library(car)

Install R

Package

## Applied Statistics di R

- Data Visualization di R
- Descriptive Statistics di R
- Inference Statistics di R

#### Data Visualization in R

- Packages for visualization: "ggplot2", "tidyverse".
   > install.packages("tidyverse")
- To make it available for use, we code:
  - > library(tidyverse)
- R Package: plot()

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### Plotting

> z = rnorm(1000) > w = rnorm(1000) > plot(w, z) > plot(w, z, main = "Plot of W and Z", xlab="z for absicca axis", ylab="w for ordinate axis", pch = 19) > pdf(file = "plot1.pdf", width = 12, height = 17, family = "Helvetica") > plot(w, z)

> dev.off()

```
> tiff("Plot3.tiff", width = 4, height = 4,
units = 'in', res = 300)
> plot(w, z)
> dev.off()
```



#### Advance Plot: qplot

- > library(ggplot2)
- > set.seed(1410)
- >dsmall <- diamonds[sample(nrow(diamonds), 100), ]</pre>

> qplot(carat, price, data = diamonds, colour = color)



#### Histogram

> z = rnorm(1000)
> w = rnorm(1000)

- >hist(z)
- > rug(z)
- > qplot(z, geom = "histogram")





#### Box Plot

```
> str(airquality)
> boxplot (airquality$Ozone)
> boxplot(airquality$Ozone,
main = "Mean ozone in parts
per billion at Roosevelt
Island",
xlab = "Parts Per Billion",
ylab = "Ozone",
col = "orange",
border = "brown",
horizontal = TRUE,
notch = TRUE
```

\$ Wind

\$ Temp

\$ Dav



#### Advance Boxplot

- > library(ggplot2)
- > diamonds

> qplot(color, price / carat, data = diamonds, geom = "boxplot", alpha=I(1/5))

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<pre>&gt; diamonds # A tibble: 53,940 x 10     carat cut    color clarity     <dbl> <ord> <or< td=""><td><pre>depth table price x y z <dbl> <dbl> <int> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> 61.5 55 326 3.95 3.98 2.43 59.8 61 326 3.89 3.84 2.31 56.9 65 327 4.05 4.07 2.31 62.4 58 334 4.2 4.23 2.63 63.3 58 335 4.34 4.35 2.75 62.8 57 336 3.94 3.96 2.48 62.3 57 336 3.95 3.98 2.47 61.9 55 337 4.07 4.11 2.53 65.1 61 337 3.87 3.78 2.49 59.4 61 338 4 4.05 2.39 ta = diamonds, geom = "boxplot",</dbl></dbl></dbl></dbl></dbl></dbl></dbl></dbl></dbl></int></dbl></dbl></pre></td><td></td><td></td></or<></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></ord></dbl></pre>	<pre>depth table price x y z <dbl> <dbl> <int> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> 61.5 55 326 3.95 3.98 2.43 59.8 61 326 3.89 3.84 2.31 56.9 65 327 4.05 4.07 2.31 62.4 58 334 4.2 4.23 2.63 63.3 58 335 4.34 4.35 2.75 62.8 57 336 3.94 3.96 2.48 62.3 57 336 3.95 3.98 2.47 61.9 55 337 4.07 4.11 2.53 65.1 61 337 3.87 3.78 2.49 59.4 61 338 4 4.05 2.39 ta = diamonds, geom = "boxplot",</dbl></dbl></dbl></dbl></dbl></dbl></dbl></dbl></dbl></int></dbl></dbl></pre>		
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### Descriptive Statistics

- It is aimed at summarizing, describing and presenting a series of values or a dataset.
- Two types:
  - 1. Location measures (mean, median, mode)
  - 2. Dispersion measures (variance, std deviation, quartile)

```
😨 R Console
                                                                - • ×
> dat <- iris
> head(dat)
  Sepal.Length Sepal.Width Petal.Length Petal.Width Species
            5.1
                        3.5
                                      1.4
                                                   0.2
                                                        setosa
           4.9
                        3.0
                                      1.4
                                                   0.2
                                                        setosa
3
            4.7
                        3.2
                                      1.3
                                                   0.2
                                                        setosa
            4.6
                        3.1
                                      1.5
                                                   0.2
                                                        setosa
            5.0
                        3.6
                                      1.4
                                                   0.2
                                                        setosa
           5.4
                        3.9
                                      1.7
                                                   0.4
                                                       setosa
> min(dat$Sepal.Length)
[1] 4.3
> rng <- range(dat$Sepal.Length)</p>
> mean(dat$Sepal.Length)
[1] 5.843333
> median(dat$Sepal.Length)
[1] 5.8
> quantile(dat$Sepal.Length, 0.25) # first quartile
25%
5.1
> sd(dat$Sepal.Length)
[1] 0.8280661
> var(dat$Sepal.Length)
[1] 0.6856935
> summary(dat)
  Sepal.Length
                   Sepal.Width
                                    Petal.Length
                                                     Petal.Width
        :4.300
                         :2.000
                                          :1.000
                                                           :0.100
 Min.
                  Min.
                                   Min.
                                                    Min.
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                  1st Qu.:2.800
                                   1st Qu.:1.600
                                                    1st Qu.:0.300
 Median :5.800
                  Median :3.000
                                   Median :4.350
                                                   Median :1.300
        :5.843
                         :3.057
                                          :3.758
                                                           :1.199
 Mean
                  Mean
                                   Mean
                                                    Mean
 3rd Ou.:6.400
                  3rd Ou.:3.300
                                   3rd Ou.:5.100
                                                    3rd Ou.:1.800
        :7.900
                  Max.
                         :4.400
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                                                           :2.500
 Max.
                                   Max.
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       Species
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```

versicolor:50 virginica :50

### Statistical Inference

- Statistical inference is the process of using data analysis to deduce properties of an underlying distribution of probability.
- ## Assess Normality
- > z = rnorm(1000)
- > qqnorm(z)
- > shapiro.test(z)
- The null hypothesis is that the data are normally distributed.
- Since p is quite high (> 0.05), we fail to reject the null hypothesis



#### Test of Mean Differences

- t-Tests for One Sample
- >iq <- c(105, 98, 110, 105, 95)
- >t.test(iq, mu = 100)

Note:

- The **null hypothesis** was that the sample was drawn from a population with mean equal to 100.
- Since p = 0.3892 is relatively large (certainly not smaller than some conventional level such as 0.05 or 0.01), we fail to reject the null hypothesis.

R Console
> iq <- c(105, 98, 110, 105, 95)
> iq [1] 105 68 110 105 65
> t.test(iq, mu = 100)
One Sample t-test
data: iq
t = 0.96495, df = 4, p-value = 0.3892
alternative hypothesis: true mean is not equal to 100
95 percent confidence interval:
95.11904 110.08096
sample estimates:
mean of x
102.6

 Two-Sample t-Test: A two-sample test is used to evaluate the null hypothesis that two population means are equal, or equivalently, that both samples were selected from the same population.

```
> grade.0 <- c(30, 25, 59, 42, 31)
> grade.1 <- c(140, 90, 95, 170, 120)
> t.test(grade.0, grade.1)
```

<pre>     R Console     grade.0 &lt;- c(30, 25, 59, 42, 31)     grade.1 &lt;- c(140, 90, 95, 170, 120)     t.test(grade.0, grade.1)     Welch Two Sample t-test  data: grade.0 and grade.1 t = -5.3515, df = 5.3094, p-value = 0.002549 alternative hypothesis: true difference in means is not equal to 0 95 percent confidence interval:     -126.00773 -45.19227 sample estimates: mean of x mean of y     27.4 122.0 </pre>		
<pre>&gt; grade.0 &lt;- c(30, 25, 59, 42, 31) &gt; grade.1 &lt;- c(140, 90, 95, 170, 120) &gt; t.test(grade.0, grade.1) Welch Two Sample t-test data: grade.0 and grade.1 t = -5.3515, df = 5.3094, p-value = 0.002549 alternative hypothesis: true difference in means is not equal to 0 95 percent confidence interval: -126.00773 -45.19227 sample estimates: mean of x mean of y 27.4 122.0</pre>	R Console	×
<pre>data: grade.0 and grade.1 t = -5.3515, df = 5.3094, p-value = 0.002549 alternative hypothesis: true difference in means is not equal to 0 95 percent confidence interval: -126.00773 -45.19227 sample estimates: mean of x mean of y 27.4 122.0</pre>	<pre>&gt; grade.0 &lt;- c(30, 25, 59, 42, 31) &gt; grade.1 &lt;- c(140, 90, 95, 170, 120) &gt; t.test(grade.0, grade.1) Welch Two Sample t-test</pre>	^
mean of x mean of y	<pre>data: grade.0 and grade.1 t = -5.3515, df = 5.3094, p-value = 0.002549 alternative hypothesis: true difference in means is not equal to 0 95 percent confidence interval: -126.00773 -45.19227 sample estimates:</pre>	
57.1 125.0 V	mean of x mean of y 37.4 123.0	~

Since p-value is equal to 0.002549, we have evidence to reject the null hypothesis

#### Analysis of Variance (ANOVA)

Ho:  $\mu_1 = \mu_2 = \mu_3 = \mu_4$ *H*<sub>1</sub>:  $\mu_1 \neq \mu_2, \mu_3 \neq \mu_4$ , etc

Teacher						
1	2	3	4			
70	69	85	95			
67	68	86	94			
65	70	85	89			
75	76	76	94			
76	77	75	93			
73	75	73	91			
M = 71.00	M = 72.5	M = 80.0	M = 92.67			

Table 6.1 Achievement as a function of teacher.



R Graphics: Device 2 (ACTIVE)

#### Achievement as a Function of Teacher

> achiev <- read.table("achiev.txt", header = T)</pre>

> attach (achiev)

```
> boxplot(ac ~ teach, data = achiev,
main="Achievement as a Function of Teacher",
xlab = "Teacher", ylab = "Achievement")
```

```
> shapiro.test(ac)
```

> library(FSA)

```
> f.teach <- factor(teach)</pre>
```

```
> hist(ac~f.teach, data = achiev)
```

```
> fligner.test(ac~f.teach, data = achiev)
```

> aggregate(ac ~ f.teach, FUN = var)

> aggregate(ac ~ f.teach, FUN = mean)

```
> anova.fit <- aov(ac ~ f.teach, data = achiev)
> summary(anova.fit)
```

> boxplot(ac ~ teach, data = achiev, main="Achievement as a Function o\$ > shapiro.test(ac)

Shapiro-Wilk normality test

```
data: ac
W = 0.90565, p-value = 0.02842
```

```
>
```

```
> library(FSA)
```

> f.teach <- factor(teach)</pre>

> hist(ac~f.teach, data = achiev)

> fligner.test(ac~f.teach, data = achiev)

Fligner-Killeen test of homogeneity of variances

data: ac by f.teach
Fligner-Killeen:med chi-squared = 10.813, df = 3, p-value =
0.01278

```
> aggregate(ac ~ f.teach, FUN = var)
 f.teach
                ac
       1 19,600000
        2 15.500000
       3 35.200000
        4 5.066667
 aggregate(ac ~ f.teach, FUN = mean)
 f.teach
               ac
       1 71.00000
       2 72.50000
       3 80.00000
       4 92.66667
> anova.fit <- aov(ac ~ f.teach, data = achiev)
> summary(anova.fit)
           Df Sum Sq Mean Sq F value Pr(>F)
            3 1764.1 588.0 31.21 9.68e-08 ***
f.teach
Residuals 20 376.8
                       18.8
Signif. codes: 0 `***' 0.001 `**' 0.01 `*' 0.05 `.' 0.1 ` ' 1
```

p-value equal to 9.68e-08, So We reject the null hypothesis that population achievement means are equal across teachers

## Wawasan: Penelitian dengan R

#### Research using R language: R Packages

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- 2. Riza, L. S., & Nugroho, E. P. (2018). MetaheuristicOpt: An R Package for Optimisation Based on Meta-Heuristics Algorithms. Pertanika Journal of Science & Technology, 26(3).
- **3. Riza, L. S.,** Bergmeir, C. N., Herrera Triguero, F., & Benítez Sánchez, J. M. (2015). **frbs**: Fuzzy rule-based systems for classification and regression in R. American Statistical Association.
- 4. Riza, L. S., Janusz, A., Bergmeir, C., Cornelis, C., Herrera, F., Śle, D., & Benítez, J. M. (2014). Implementing algorithms of rough set theory and fuzzy rough set theory in the R package "RoughSets". Information Sciences, 287, 68-89.

#### Research using R language: Implementations/Computational Models

- Mediayani, M., Wibisono, Y., Riza, L. S., & Pérez, A. R. (2019). Determining trending topics in twitter with a data-streaming method in R. Indonesian Journal of Science and Technology, 4(1), 148-157.
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- 3. Riza, L. S., Rachmat, A. B., Munir, T. H., & Nazir, S. (2019). Genomic Repeat Detection Using the Knuth-Morris-Pratt Algorithm on R High-Performance-Computing Package. Int. J. Advance Soft Compu. Appl, 11(1).
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- 6. Nazir, S., Shahzad, S., & **Riza, L. S.** (2017). **Birthmark-based software** classification using rough sets. Arabian Journal for Science and Engineering, 42(2), 859-871.

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Tattar, P. N., Ramaiah, S., & Manjunath, B. G. (2016). A Course in Statistics with R. John Wiley & Sons.

Wickham, H. (2016). ggplot2: elegant graphics for data analysis. springer.