t-Test in R

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Contents

| Required Packages | 1 |
|--|---|
| in stall.packages ("reshape 2") | 1 |
| library(reshape 2) | 1 |
| One sample t-test | 2 |
| Manual Calculation | 2 |
| using function——— | 2 |
| Using Function | 4 |
| Two-samples t-test | 7 |
| Paired samples t-test | 9 |
| Required Packages | |
| install.packages("reshape2") | |
| library(reshape 2) | |
| <pre>library(ggplot2) library(reshape2) library(car) library(ggpubr)</pre> | |

One sample t-test

Manual Calculation

```
set.seed(123)
sample_data <- rnorm(30, mean = 50, sd = 10)</pre>
mean(sample_data)
## [1] 49.52896
sd(sample_data)
## [1] 9.810307
#mu_0=55
SE <- sd(sample_data)/sqrt(length(sample_data))</pre>
t_cal <- (mean(sample_data) - 55)/SE
print(t_cal)
## [1] -3.054553
t_crit <- qt(0.05/2, df = length(sample_data)-1, lower.tail=TRUE)
t_crit
## [1] -2.04523
t_cal <- t_crit
using function-
t.test(sample_data, mu=55)
##
    One Sample t-test
##
##
## data: sample_data
## t = -3.0546, df = 29, p-value = 0.004797
## alternative hypothesis: true mean is not equal to 55
## 95 percent confidence interval:
## 45.86573 53.19219
## sample estimates:
## mean of x
## 49.52896
```

```
t.test(sample_data, mu=55, conf.level=0.99)
##
##
    One Sample t-test
## data: sample_data
## t = -3.0546, df = 29, p-value = 0.004797
## alternative hypothesis: true mean is not equal to 55
## 99 percent confidence interval:
## 44.59198 54.46595
## sample estimates:
## mean of x
## 49.52896
t.test(sample_data, mu=55, conf.level=0.99)
##
##
    One Sample t-test
##
## data: sample_data
## t = -3.0546, df = 29, p-value = 0.004797
## alternative hypothesis: true mean is not equal to 55
## 99 percent confidence interval:
## 44.59198 54.46595
## sample estimates:
## mean of x
## 49.52896
Hypothesis:
H_0: \mu = \mu_0
H_1: \mu \neq \mu_0
# mu_0 = 55
SE <- sd(sample_data)/sqrt(length(sample_data))</pre>
t_cal <- (mean(sample_data) - 55)/SE
print(t_cal)
## [1] -3.054553
t_crit <- qt(0.05/2, df = length(sample_data)-1, lower.tail = TRUE)
t_crit
## [1] -2.04523
t_cal <= t_crit # decision: reject null</pre>
## [1] TRUE
```

```
pt(t_cal, df = 29, lower.tail = TRUE) + pt(-t_cal, df = 29, lower.tail = FALSE)

## [1] 0.0047971

mean(sample_data)+c(-1, 1)* abs(t_crit)*SE

## [1] 45.86573 53.19219

Calculating confidence interval:

mean(sample_data) + c(-1, 1) * abs(t_crit) * SE
```

```
## [1] 45.86573 53.19219
```

Using Function

Generate sample data:

```
set.seed(123)
sample_data <- rnorm(30, mean = 50, sd = 10)</pre>
```

Testing normality using Shapiro-Wilk test:

```
shapiro.test(sample_data)
```

```
##
## Shapiro-Wilk normality test
##
## data: sample_data
## W = 0.97894, p-value = 0.7966
```

 H_0 : Data follows normal distribution.

 H_1 : Data does not follow normal distribution.

Since the p-value is greater than the level of significance ($\alpha = 0.05$), we do not have enough statistical evidence to reject the null hypothesis.

One-sample t-test:

```
Hypothesis: H_0: \mu = 50

H_1: \mu \neq 50
```

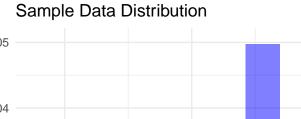
```
t.test(sample_data, mu = 50)
```

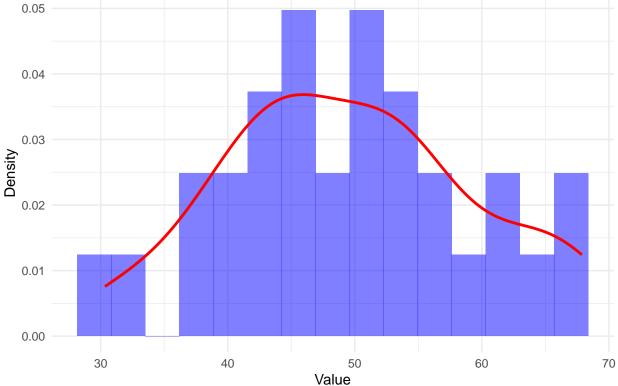
```
##
## One Sample t-test
##
## data: sample_data
```

```
## t = -0.26299, df = 29, p-value = 0.7944
## alternative hypothesis: true mean is not equal to 50
## 95 percent confidence interval:
## 45.86573 53.19219
## sample estimates:
## mean of x
## 49.52896
Using function, perform the two tailed t-test:
t.test(sample_data, mu = 53, conf.level = 0.95)
   One Sample t-test
##
##
## data: sample data
## t = -1.9379, df = 29, p-value = 0.06242
## alternative hypothesis: true mean is not equal to 53
## 95 percent confidence interval:
## 45.86573 53.19219
## sample estimates:
## mean of x
## 49.52896
If p-value < 0.05, then reject null. Decision is not rejected.
t.test(sample_data, mu = 53, conf.level = 0.99)
##
##
    One Sample t-test
##
## data: sample_data
## t = -1.9379, df = 29, p-value = 0.06242
## alternative hypothesis: true mean is not equal to 53
## 99 percent confidence interval:
## 44.59198 54.46595
## sample estimates:
## mean of x
## 49.52896
If p-value < 0.01, then reject null. Decision is not rejected.
t.test(sample_data, mu = 53, conf.level = 0.90)
##
##
   One Sample t-test
## data: sample_data
## t = -1.9379, df = 29, p-value = 0.06242
## alternative hypothesis: true mean is not equal to 53
```

90 percent confidence interval:

```
## 46.48564 52.57228
## sample estimates:
## mean of x
## 49.52896
If p-value < 0.10, then reject null. Decision is rejected.
Hypothesis:
H_0: \mu <= 40
H_1: \mu > 40
t.test(sample_data, mu = 40, alternative = "greater")
##
##
   One Sample t-test
##
## data: sample_data
## t = 5.3201, df = 29, p-value = 5.21e-06
## alternative hypothesis: true mean is greater than 40
## 95 percent confidence interval:
## 46.48564
                  Inf
## sample estimates:
## mean of x
## 49.52896
Hypothesis:
H_0: \mu >= 58
H_1: \mu < 58
t.test(sample_data, mu = 58, alternative = "less")
##
##
    One Sample t-test
## data: sample_data
## t = -4.7295, df = 29, p-value = 2.689e-05
## alternative hypothesis: true mean is less than 58
## 95 percent confidence interval:
##
        -Inf 52.57228
## sample estimates:
## mean of x
## 49.52896
ggplot(data.frame(Value = sample_data), aes(x = Value)) +
  geom_histogram(aes(y = after_stat(density)), bins = 15, fill = "blue", alpha = 0.5) +
  geom_density(color = "red", linewidth = 1) +
  labs(title = "Sample Data Distribution", x = "Value", y = "Density") +
  theme_minimal()
```





Two-samples t-test

Use built-in dataset: mtcars (comparing mpg for automatic vs manual cars):

```
data(mtcars)
```

Split into two groups based on transmission type":

```
auto_mpg <- mtcars$mpg[mtcars$am == 0]</pre>
                                                # Automatic
manual_mpg <- mtcars$mpg[mtcars$am == 1]</pre>
```

 H_0 : Automatic cars and manual cars have equal average mpg. H_1 : Automatic cars and manual cars have unequal average mpg.

```
mean(auto_mpg)
```

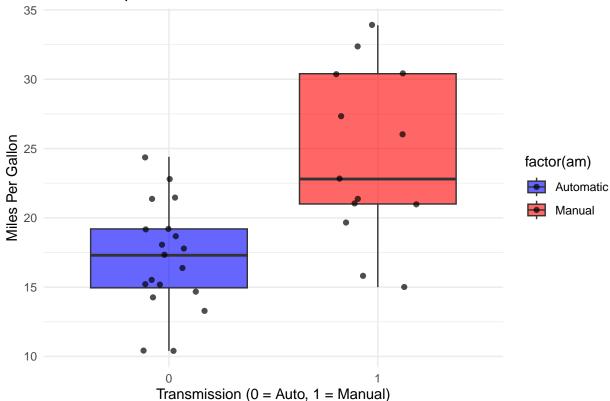
[1] 17.14737

```
mean(manual_mpg)
```

[1] 24.39231

```
var(auto_mpg)
## [1] 14.6993
var(manual_mpg)
## [1] 38.02577
Normality test for both groups:
shapiro.test(auto_mpg)
##
## Shapiro-Wilk normality test
## data: auto_mpg
## W = 0.97677, p-value = 0.8987
shapiro.test(manual_mpg)
##
## Shapiro-Wilk normality test
##
## data: manual_mpg
## W = 0.9458, p-value = 0.5363
Check variance homogeneity (Levene's test):
leveneTest(mpg ~ factor(am), data = mtcars, center = "mean")
## Levene's Test for Homogeneity of Variance (center = "mean")
       Df F value Pr(>F)
             5.921 0.02113 *
## group 1
##
        30
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
Perform two-sample t-test:
t.test(auto_mpg, manual_mpg, var.equal = FALSE)
##
## Welch Two Sample t-test
##
## data: auto_mpg and manual_mpg
## t = -3.7671, df = 18.332, p-value = 0.001374
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -11.280194 -3.209684
## sample estimates:
## mean of x mean of y
## 17.14737 24.39231
```

MPG Comparison: Automatic vs Manual



Paired samples t-test

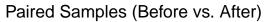
Generate 30 observations:

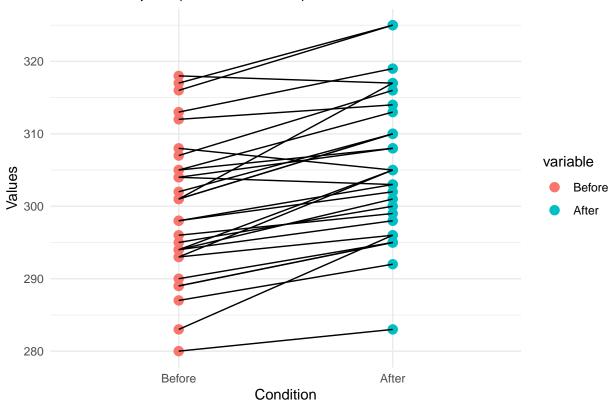
```
set.seed(123)
before <- round(rnorm(30, mean = 300, sd = 10), 0)
summary(before)

## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 280.0 293.2 299.5 299.6 305.0 318.0</pre>
```

```
# after a course
after <- before + round(rnorm(30, mean = 5, sd = 5), 0) # Simulating a increase
summary(after)
##
      Min. 1st Qu. Median Mean 3rd Qu.
                                                Max.
##
            298.2 305.0 305.5
                                      312.2
df <- data.frame(</pre>
  ID = 1:30,
  Before = before,
  After = after
mean(after) - mean(before)
## [1] 5.933333
H_0: Before and after the course the true GRE average score of the students stays the same.
H_1: Before and after the course the true GRE average score of the students does not remain the same.
Perform Paired t-test
t.test(x = after, y = before, paired = TRUE, alternative = "two.sided")
##
##
   Paired t-test
## data: after and before
## t = 7.7811, df = 29, p-value = 1.398e-08
## alternative hypothesis: true mean difference is not equal to 0
## 95 percent confidence interval:
## 4.373779 7.492888
## sample estimates:
## mean difference
          5.933333
# Visualize the differences
df <- data.frame(</pre>
  ID = 1:30,
 Before = before,
  After = after
df_long <- melt(df, id.vars = "ID")</pre>
ggplot(df_long, aes(x = variable, y = value, group = ID)) +
  geom_point(aes(color = variable), size = 3) +
  geom_line() +
  labs(title = "Paired Samples (Before vs. After)",
       y = "Values", x = "Condition") +
```

theme_minimal()





```
ggpaired(df_long,
    x = "variable",
    y = "value",
    color = "variable",
    line.color = "gray",
    line.size = 0.4,
    palette = "jco") +
    stat_compare_means(paired = TRUE, method = "t.test")
```



