



Paired Samples Wilcoxon Test in R

Tools

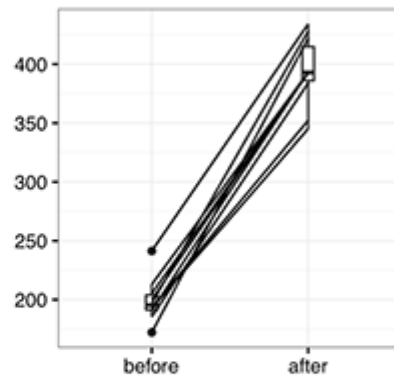
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The **paired samples Wilcoxon test** (also known as **Wilcoxon signed-rank test**) is a **non-parametric** alternative to paired t-test used to compare paired data. It's used when your data are not normally distributed. This tutorial describes how to compute paired samples Wilcoxon test in **R**.

Differences between paired samples should be distributed symmetrically around the median.

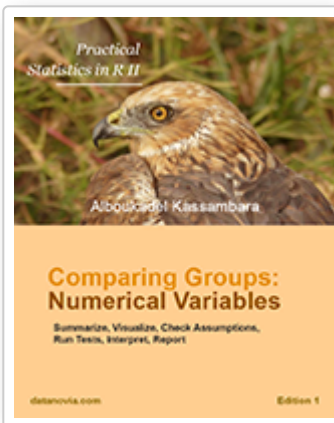
Paired Samples Wilcoxon Test in R

- + Definition
- + Research Questions & Statistics
- + Practical Examples in R
- + Interpretation



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Related Book:



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Visualize your data and compute paired samples Wilcoxon test in R

R function

The R function **wilcox.test()** can be used as follow:

```
wilcox.test(x, y, paired = TRUE, alternative = "two.sided")
```

- **x,y**: numeric vectors
- **paired**: a logical value specifying that we want to compute a paired Wilcoxon test
- **alternative**: the alternative hypothesis. Allowed value is one of "two.sided" (default), "greater" or "less".

Import your data into R

1. **Prepare your data** as specified here: [Best practices for preparing your data set for R](#)
2. **Save your data** in an external .txt tab or .csv files
3. **Import your data into R** as follow:

```
# If .txt tab file, use this
my_data <- read.delim(file.choose())
# Or, if .csv file, use this
my_data <- read.csv(file.choose())
```

Here, we'll use an example data set, which contains the weight of 10 mice before and after the treatment.

```
# Data in two numeric vectors
# ++++++
# Weight of the mice before treatment
before <-c(200.1, 190.9, 192.7, 213, 241.4, 196.9, 172.2, 185.5, 205.2, 193.7)
# Weight of the mice after treatment
after <-c(392.9, 393.2, 345.1, 393, 434, 427.9, 422, 383.9, 392.3, 352.2)
# Create a data frame
my_data <- data.frame(
  group = rep(c("before", "after"), each = 10),
  weight = c(before, after)
)
```

? We want to know, if there is any significant difference in the median weights before and after treatment?

Check your data

```
# Print all data
print(my_data)
```

```
   group weight
1  before  200.1
2  before  190.9
3  before  192.7
4  before  213.0
5  before  241.4
6  before  196.9
7  before  172.2
8  before  185.5
9  before  205.2
10 before  193.7
11 after   392.9
12 after   393.2
13 after   345.1
14 after   393.0
15 after   434.0
16 after   427.9
17 after   422.0
18 after   383.9
19 after   392.3
20 after   352.2
```

Compute summary statistics (median and inter-quartile range (IQR)) by groups using the dplyr package can be used.

- Install **dplyr** package:

```
install.packages("dplyr")
```

- Compute summary statistics by groups:

```
library("dplyr")
group_by(my_data, group) %>%
  summarise(
    count = n(),
    median = median(weight, na.rm = TRUE),
    IQR = IQR(weight, na.rm = TRUE)
  )
```

```
Source: local data frame [2 x 4]
  group count median  IQR
  (fctr) (int)  (dbl) (dbl)
```

```
1 after      10 392.95 28.800
2 before     10 195.30 12.575
```

Visualize your data using box plots

- To use R base graphs read this: [R base graphs](#). Here, we'll use the **ggpubr** R package for an easy ggplot2-based data visualization.
- Install the latest version of ggpubr from GitHub as follow (recommended):

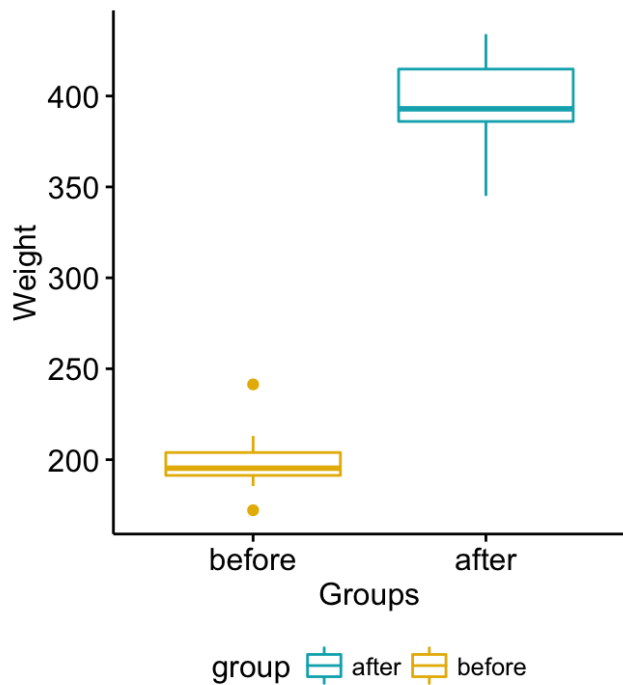
```
# Install
if(!require(devtools)) install.packages("devtools")
devtools::install_github("kassambara/ggpubr")
```

- Or, install from CRAN as follow:

```
install.packages("ggpubr")
```

- Visualize your data:

```
# Plot weight by group and color by group
library("ggpubr")
ggboxplot(my_data, x = "group", y = "weight",
           color = "group", palette = c("#00AFBB", "#E7B800"),
           order = c("before", "after"),
           ylab = "Weight", xlab = "Groups")
```



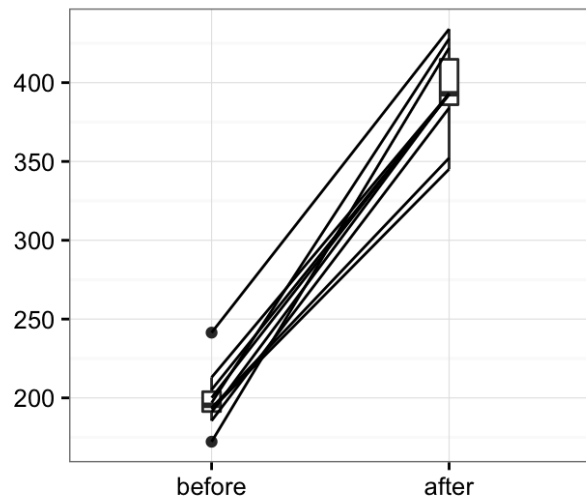
Box plots show you the increase, but lose the paired information. You can use the function **plot.paired()** [in **pairedData** package] to plot paired data ("before - after" plot).

- Install pairedData package:

```
install.packages("PairedData")
```

- Plot paired data:

```
# Subset weight data before treatment
before <- subset(my_data, group == "before", weight,
                 drop = TRUE)
# subset weight data after treatment
after <- subset(my_data, group == "after", weight,
               drop = TRUE)
# Plot paired data
library(PairedData)
pd <- paired(before, after)
plot(pd, type = "profile") + theme_bw()
```



Compute paired-sample Wilcoxon test

? Question : Is there any significant changes in the weights of mice before after treatment?

1) Compute paired Wilcoxon test - Method 1: The data are saved in two different numeric vectors.

```
res <- wilcox.test(before, after, paired = TRUE)
res
```

```
Wilcoxon signed rank test
data: before and after
V = 0, p-value = 0.001953
alternative hypothesis: true location shift is not equal to 0
```

2) Compute paired Wilcoxon-test - Method 2: The data are saved in a data frame.

```
# Compute t-test
res <- wilcox.test(weight ~ group, data = my_data, paired = TRUE)
res
```

```
Wilcoxon signed rank test
data: weight by group
V = 55, p-value = 0.001953
alternative hypothesis: true location shift is not equal to 0
```

```
# print only the p-value  
res$p.value
```

```
[1] 0.001953125
```

! As you can see, the two methods give the same results.

✓ The **p-value** of the test is 0.001953, which is less than the significance level $\alpha = 0.05$. We can conclude that the median weight of the mice before treatment is significantly different from the median weight after treatment with a **p-value** = 0.001953.

! Note that:

- if you want to test whether the median weight before treatment is less than the median weight after treatment, type this:

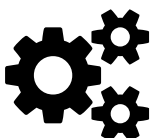
```
wilcox.test(weight ~ group, data = my_data, paired = TRUE,  
            alternative = "less")
```

- Or, if you want to test whether the median weight before treatment is greater than the median weight after treatment, type this

```
wilcox.test(weight ~ group, data = my_data, paired = TRUE,  
            alternative = "greater")
```

Online paired-sample Wilcoxon test calculator

You can perform **paired-sample Wilcoxon test, online**, without any installation by clicking the following link:



[Online paired-sample Wilcoxon test calculator](#)

See also

Paired Samples T-test (parametric)

Infos



This analysis has been performed using **R software** (ver. 3.2.4).



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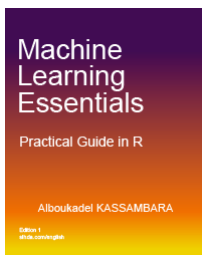
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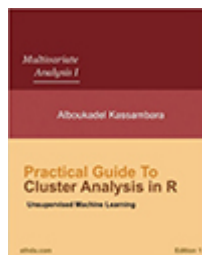
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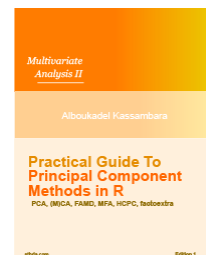
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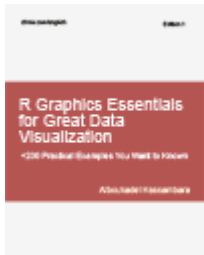
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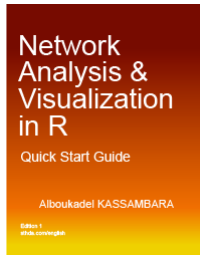
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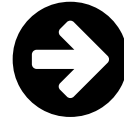
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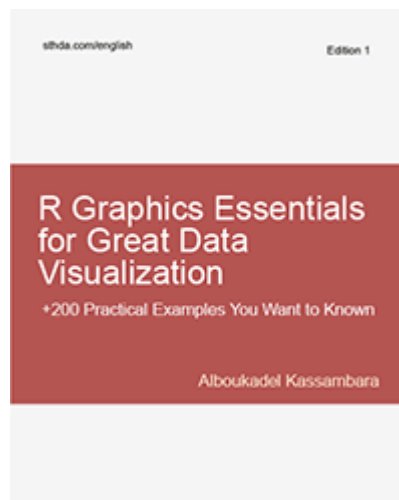
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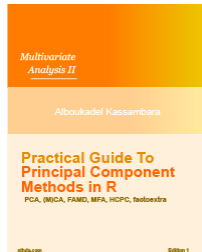


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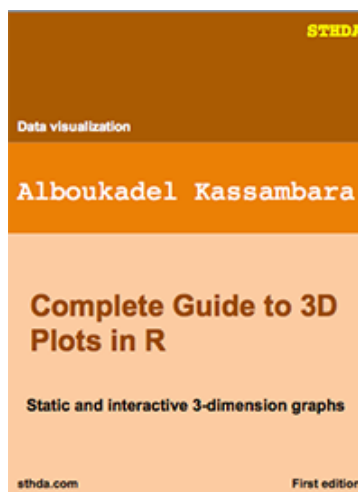
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