

# Drug\_test\_project\_Assignment\_Chi

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## Introduction:

The data of this project assignment is the “placebo\_new\_drug” data which is about results from testing new drug. The sample is small with 5 objects. Hence, I will choose non-parametric test to analyze the data.

## The Analysis:

Firstly, I import the libraries and the data:

```
library(data.table)
library(BSDA)
```

```
## Loading required package: lattice
```

```
##
## Attaching package: 'BSDA'
```

```
## The following object is masked from 'package:datasets':
##
##      Orange
```

```
library(dplyr)
```

```
##
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:data.table':
##
##      between, first, last
```

```
## The following objects are masked from 'package:stats':
##
##      filter, lag
```

```
## The following objects are masked from 'package:base':
##
##      intersect, setdiff, setequal, union
```

```
setwd("/Users/kienguyen/Documents/DATA SCIENCE/MSDS/03. MSMS 660/08. Week 8/In class")
drugdt <- read.csv(file = 'placebo_new_drug.csv', sep=",", header=T)
```

Next, I calculate the median of the 2 groups:

```
median(drugdt$Placebo)
```

```
## [1] 6
```

```
median(drugdt$New.Drug)
```

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

The mean of “Placebo” variable is greater than the “New.Drug” variable, hence, I will use “greater” alternative for the below tests about the means.

## Test the difference between medians of 2 groups:

**SIGN test:**

```
SIGN.test(x = drugdt$Placebo, y = drugdt$New.Drug , alternative = 'greater')
```

```
##
## Dependent-samples Sign-Test
##
## data: drugdt$Placebo and drugdt$New.Drug
## S = 4, p-value = 0.1875
## alternative hypothesis: true median difference is greater than 0
## 95 percent confidence interval:
## -0.64 Inf
## sample estimates:
## median of x-y
## 2
##
## Achieved and Interpolated Confidence Intervals:
##
## Conf.Level L.E.pt U.E.pt
## Lower Achieved CI 0.8125 2.00 Inf
## Interpolated CI 0.9500 -0.64 Inf
## Upper Achieved CI 0.9688 -1.00 Inf
```

p-value = 0.1875 > 0.05. We fail to reject the null hypothesis and state that the medians are NOT significantly different.

## Wilcoxon Signed Rank test

```
wilcox.test(drugdt$Placebo, drugdt$New.Drug, alternative = 'greater')
```

```
## Warning in wilcox.test.default(drugdt$Placebo, drugdt$New.Drug, alternative =  
## "greater"): cannot compute exact p-value with ties
```

```
##  
## Wilcoxon rank sum test with continuity correction  
##  
## data: drugdt$Placebo and drugdt$New.Drug  
## W = 22, p-value = 0.02928  
## alternative hypothesis: true location shift is greater than 0
```

We received the red warning because of the small size sample. So, with the  $p\text{-value} = 0.02928 < 0.05$ , I think I will take one more test to check rather than trust this result.

### Kruskal-Wallis

```
kruskal.test(drugdt)
```

```
##  
## Kruskal-Wallis rank sum test  
##  
## data: drugdt  
## Kruskal-Wallis chi-squared = 3.9865, df = 1, p-value = 0.04587
```

The p-value 0.04587 which is approximately equal to 0.05, so that we fail to reject the null hypothesis and state that the medians are NOT significantly different.

### Check the distribution difference between 2 groups:

#### Kolmogorov-Smirnov:

```
ks.test(drugdt$New.Drug, drugdt$Placebo, alternative = 'greater')
```

```
##  
## Exact two-sample Kolmogorov-Smirnov test  
##  
## data: drugdt$New.Drug and drugdt$Placebo  
## D+ = 0.6, p-value = 0.1429  
## alternative hypothesis: the CDF of x lies above that of y
```

The p-value is 0.1429 bigger than 0.05, so we accept the null hypothesis which mean that the two dataset values are from the same continuous distribution.

## Test if variables are correlated

### Spearman test

Now, let's check if those 2 variables are correlated together or not with Spearman test:

```
library(pspearman)
spearman.test(drugdt$Placebo, drugdt$New.Drug)

##
## Spearman's rank correlation rho
##
## data: drugdt$Placebo and drugdt$New.Drug
## S = 28, p-value = 0.5167
## alternative hypothesis: true rho is not equal to 0
## sample estimates:
## rho
## -0.4
```

According to the Spearman test:

- The p-value is  $0.5167 > 0.05$  which indicates that there's a weak evidence to reject the null Hypothesis. We fail to reject the null hypothesis and there is NO relationship between 2 variables.

### Kendall's Tau

```
library(Kendall)
Kendall(drugdt$Placebo, drugdt$New.Drug)

## tau = -0.4, 2-sided pvalue =0.46243
```

The p-value of this test also bigger than 0.05 which means there's NO relationship between 2 variables.

## SUMMARY:

I firstly import the "placebo\_new\_drug" data which is about results from testing new drug and choose non-parametric test to analyze the data:

- Check the difference between medians of 2 groups with: SIGN test, Wilcoxon, and Kruskal test: the p-values of SIGN and Kruskal test are bigger or equal to 0.05, so the medians are NOT significantly different. There was a red warning when running the Wilcoxon Signed Rank test due to the small size sample, so that I don't trust this result when the p-value =  $0.02928 < 0.05$ .
- Check the distribution difference: Kolmogorov-Smirnov test results in the p-value is 0.1429 bigger than 0.05, so we can accept the null hypothesis which mean that the two data set values are from the same continuous distribution.
- Check the correlation between 2 variables: Spearman test and Kendall's Tau both result in the conclusion that there's NO relationship between 2 variables.