

Biometry. Lecture 12

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- 1 One-dimensional data
 - One-dimensional tests
 - Normality and R functions



```
> setwd("<working folder>")  
or  
"Change dir"  
in menu!
```

On Mac, be sure that startup option is working: `getwd()`
(`getwd()` checks if R is in working folder, `dir()` checks the folder
content)



One-dimensional data

One-dimensional tests



t-test and Wilcoxon test for one-dimensional data

- Statistical tests allow to check how well the general characteristic calculated from *sample* represents a *population*
- t-test (Student's) takes into account the normality of sample whereas Wilcoxon test do not consider the distribution, it is non-parametric
- Both give a *confidence interval*



t-test for one variable

```
> t.test(salary, mu=mean(salary))
One Sample t-test

data:  salary
t = 0, df = 6, p-value = 1
alternative hypothesis: true mean is not equal to 32.28571
95 percent confidence interval:
 3.468127 61.103302
sample estimates:
mean of x
32.28571
```



Wilcoxon test for one variable

```
> wilcox.test(salary, mu=median(salary), conf.int=TRUE)
```

Wilcoxon signed rank test with continuity correction

```
data: salary
```

```
V = 10, p-value = 0.5896
```

```
alternative hypothesis: true location is not equal to 21
```

```
80 percent confidence interval:
```

```
 17.99999 63.50002
```

```
sample estimates:
```

```
(pseudo)median
```

```
 24.99994
```

This will test median, not mean! Wilcoxon test is more universal but less traditional.



Understanding the test output: theory

- Alternative hypothesis (“something”) and null hypothesis (“nothing”)
- Type I error (false alarm), p-value (probability to issue the false alarm) and significance level (matter of agreement)
- In science, Type I error is more dangerous so we follow the “0.005 rule”: **if p-value is more than 0.05, we must accept the null hypothesis**



Understanding the test output: quick and dirty

- Which hypothesis is null?
- Does p-value less than 0.05?
 - 1 No: accept the null hypothesis—"sit and relax"
 - 2 Yes: reject the null hypothesis—"jump and call the police"



How to understand which test to use? Normality.

- Normality tests will check if we can accept the normal distribution of our sample
- It is widely accepted that the strict normality testing is not generally required, it is enough, for example, to test normality graphically



Shapiro test for normality

```
> shapiro.test(salary) # What is a null hypothesis?!  
> shapiro.test(rnorm(1000)) # Null is normality!
```



Quantile-quantile plot for normality

```
> qqnorm(salary); qqline(salary) # Bad!  
> # Good:  
> set.seed(1); qqnorm(rnorm(100)); qqline(rnorm(100))
```

`set.seed()` helps to maintain the same set of random numbers in the session.



One-dimensional data

Normality and R functions



- `shapiro.test()` is good but it is hard to apply if for data frames, and output is not very helpful.
- We will create the user function which run Shapiro-Wilks test with better output.



What is a function

```
> Sum <- function(a, b)
+ {
+ a + b
+ }
> Sum(2, 3)
```

Function is a piece of code which may run independently. All R commands are functions. Please note that every functions requires two parts: **arguments** in *round brackets* and **body** in *curly brackets*. It is too boring to enter functions line-by-line. Instead, it is better to copy function from external editor. If function contains mistake(s), one may use `fix()` command.



Let us create a simple useful function

R has no command for coefficient of variation, we will create it ourselves:

```
> CV <- function(x)
+ {
+ (sd(x) / mean(x)) * 100
+ }
> CV(trees[,3])
> CV(trees$Volume)
> sapply(trees, CV)
```

We can then run `fix(CV)` and add `round(..., 2)` function to make numbers more readable.



Finishing...

Save your commands!

(`savehistory(<todaysdate>.r)` or File -> Save as... on Mac)



Summary: most important commands

- `t.test()`— checks the reliability of mean (assuming that data distribution is normal)
- `wilcox.test()`— checks the reliability of median (non-parametric)
- `shapiro.test()`—Shapiro-Wilks test for normality
- `function(){}` —creates a function
- `for(){}` —the cycle
- `ifelse(check, yes, no)`—selector



For Further Reading



A. Shipunov.

Biometry [Electronic resource].

2012—onwards.

Mode of access:

http://ashipunov.info/shipunov/school/biol_240



A. Shipunov, and others.

Visual statistics. Use R!

Ongoing translation from Russian.

