

# 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](http://ashipunov.info/shipunov/school/biol_240)



A. Shipunov, and others.

*Visual statistics. Use R!*

Ongoing translation from Russian.

