

GENERAL COMMANDS	
Help (for the command)	?command
Comment	#
Summation, subtraction	+, -
Multiplication, ratio	*, /
Variable assignment	= or <-
Not equal	!=
Sinus, cosinus	sin(argument), cos(argument)
$\pi$	pi
Exponentiation (power)	^
Square root	sqrt(argument)
Root of n-th degree	argument^(1/n)
e	exp(1)
Logarithm	log(argument, base=base of logarithm)
Vector	c(elements divided by commas)
Indicating i-th element of a vector	vector[i]
Vector – sequence with the number of components equally spread one from the other	seq(from, to, length=number of classes)
Vector – sequence with steps	seq(from, to, by=step)
Vector of replicated subvector	rep(vector, times=number of replications)
Vector of replicated components of subvector	rep(vector, each=number of replications)
Sum of components in a vector	sum(vector)
Number of components in vector	length(vector)
Changing the order of vector components	rev(vector)
Removing the components from vector	vector[-c(indices of components to remove)]
Indicating components of vector smaller than "k"	vector[vector < k]
Indicating indices of vector components smaller than "k"	which(vector < k)
Indicating indices of minimal and maximal component	which.min(vector) which.max(vector)
Matrix from column vectors	cbind(x1, x2, ..., xm)
Matrix from row vectors	rbind(x1, x2, ..., xm)
Multiplication of matrices	%*%
Determinant	det(matrix)
Indicating (i,j)-th entry of a matrix	matrix[[i,j]]
Indicating i-th row / j-th column of a matrix	matrix[i,]      matrix[,j]
Transposition	t(matrix)
Diagonal of matrix	diag(matrix)
Dimension of matrix	dim(matrix)
Inverse of matrix	solve(matrix)
Number of rows and columns of matrix	nrow(matrix), ncol(matrix)
Percentage notation („scales” package)	percent(number)
Division of window with graphs	par(mfrow=c(n,m))
Simple graph representing points (x,y) or data	plot(x, y)      plot(data)
Graph representing a function of one variable	curve(function of x, x lower bound, x upper bound)
PACKAGES	
Package Installing	install.packages(„name”)
Package loading	library(name)
„FOR”, „IF”, FUCTIONS AND PROCEDURES	
„for”	for (variable in beginning:end){what to do} Caution! Variable can be also a vector!
„if”	if (condition) {what to do} else {what to do}
functions/procedures	name= function (arguments){ what to do return(output)}

DESCRIPTIVE STATISTICS	
Loading data	<code>read.csv(“name”, sep=“;”)</code>
Loading data in polish coding („;” used for decimal)	<code>read.csv(“name”, sep=“;”, dec=“.”)</code>
Loading data with labels	<code>read.csv(“name”, sep=“;”, head=TRUE)</code>
Creating data as a table of data (x1, x2,..., xm are column labels)	<code>data.frame(x1, x2,..., xm)</code>
Class of data	<code>class(data)</code>
Vector of labels	<code>names(data)</code>
Mean	<code>mean(data)</code>
Minimum and maximum	<code>min(data), max(data)</code>
Quartiles	<code>quantile(data)</code>
Quantiles (chosen)	<code>quantile(data, probs=vector of probabilities)</code>
Variance and standard deviation	<code>var(data), sd(data)</code>
Function from rows of matrix (table)	<code>apply(matrix, 1, function)</code>
Function from columns of matrix (table)	<code>apply(matrix, 2, function)</code>
Central tendency measures – all	<code>summary(data)</code>
Histogram for discrete data (line graph)	<code>discrete.histogram(data)</code>
Caution! „arm” package required	
Histogram (for grouped data)	<code>hist(data, main=title, xlab=label of x)</code>
Automatic declaration of graphics titles e.g. using “for” procedure (with space or without respectively)	<code>paste(“text”, name)</code> <code>paste0(“text”, name)</code>
Computing e.g. mean if the data have different lengths	<code>mean(na.omit(data))</code>
Grouping data into frequency table	<code>table(data)</code>
Frequency table with intervals	<code>cut(data, breaks = # of classes)</code>
Pie chart (of point frequency table)	<code>pie(table(data))</code>
Pie chart (of point frequency table)	<code>pie(table(cut(data, breaks = # of classes)))</code>
Box plot	<code>boxplot(data)</code>
RANDOM VARIABLES AND DISTRIBUTIONS	
Probability/density function (d – density)	<code>dname</code>
CDF (p – probability)	<code>pname</code>
Quantile (q – quantile)	<code>qname</code>
Random generating (r – random)	<code>rname</code>
Distribution names binomial Poisson exponential normal t-Student chi-square F Snedecor	<code>binom</code> <code>pois</code> <code>exp</code> <code>norm</code> <code>t</code> <code>chisq</code> <code>f</code>
Line graph of discrete distribution	<code>plot(x, dname(x, parameters))</code>
Drawing density function	<code>curve(dname(x, parameters))</code>

CONFIDENCE INTERVALS (CI) AND HYPOTHESES TESTING	
CI for $\mu$ under normality with known $\sigma$ Caution! „BSDA” package required	<code>z.test(data, sigma.x=<math>\sigma</math>, conf.level=1-<math>\alpha</math>)</code>
CI for $\mu$ – large sample Caution! „BSDA” package required	<code>zsum.test(sample mean, (sample) stand. dev., sample size, conf.level=1-<math>\alpha</math>)</code>
CI for $\mu$ under normality when $\sigma$ is unknown	<code>t.test(data, conf.level=1-<math>\alpha</math>)</code>
CI for $\sigma^2$ Caution! „TeachingDemos” package required	<code>sigma.test(data, conf.level=1-<math>\alpha</math>)</code>
CI for probability of success (proportion) $p$	<code>binom.test(no. of successes, sample size, conf.level=1-<math>\alpha</math>)</code>
Only CI as output	<code>NameOfTest\$conf.int</code>
Hypothesis about $\mu$ under normality with known $\sigma$ Caution! „BSDA” package required	<code>z.test(data, sigma.x=<math>\sigma</math>, alternative="two.sided", mu=tested mean)</code>
Hypothesis about $\mu$ when the sample is large Caution! „BSDA” package required	<code>zsum.test(sample mean, sample stand. dev., sample size, alternative="greater", mu=tested mean)</code>
Hypothesis about $\mu$ under normality with unknown $\sigma$	<code>t.test(data, alternative="less", mu=tested mean)</code>
Hypothesis about $\sigma^2$ Caution! „TeachingDemos” package required	<code>sigma.test(data, sigma=tested sigma, alternative="two.sided")</code>
Hypothesis about probability of success (proportion) $p$	<code>binom.test(no. of successes, sample size, p=tested probability, alternative="two.sided")</code>
Only p-value as an output	<code>NameOfTest\$p.value</code>

COMPARISON OF TWO POPULATIONS	
CI for difference of means under normality (equal population variances)	<code>t.test(data1, data2, var.equal=TRUE, conf.level=1-<math>\alpha</math>)</code>
CI for difference of means under normality (unequal population variances)	<code>t.test(data1, data2, var.equal=FALSE, conf.level=1-<math>\alpha</math>)</code>
CI for difference of means (large samples) Caution! “BSDA” package required	<code>zsum.test(sample mean 1, stand. dev. 1, sample size 1, sample mean 2, stand. dev. 2, sample size 2, conf.level=1-<math>\alpha</math>)</code>
CI for the ratio of variances Caution! „PairedData” package required	<code>var.test(data1, data2, conf.level=1-<math>\alpha</math>)</code>
CI for the difference of proportions	<code>prop.test(c(T1,T2), c(n1,n2), conf.level=1-<math>\alpha</math>)</code>
Hypothesis about difference of means (equal population variances)	<code>t.test(data1, data2, mu=tested difference of means, var.equal=TRUE, alternative="two.sided")</code>
Hypothesis about difference of means (unequal population variances)	<code>t.test(data1, data2, mu=tested difference of means, var.equal=FALSE, alternative="two.sided")</code>
Hypothesis about difference of means (large samples) Caution! “BSDA” package required	<code>zsum.test(sample mean 1, stand. dev. 1, sample size 1, sample mean 2, stand. dev. 2, sample size 2, mu=tested difference of means, alternative="two.sided")</code>
Hypothesis about ratio of variances Caution! „PairedData” package required	<code>var.test(data1, data2, ratio=tested ratio of variances, alternative="two.sided")</code>
Hypothesis about equality of proportions	<code>prop.test(c(T1,T2), c(n1,n2), alternative="two.sided")</code>

ANALYSIS OF VARIANCE (ANOVA)	
CAUTION! To perform ANOVA data has to be arrange in a proper way! data.frame(measurements, methods)	
Bartlett test of homogeneity of variances	<code>bartlett.test(measurements~treatments)</code>
Analysis of variance	<code>anova(lm(measurements~treatments))</code>
TukeyTest HSD of homogeneous treatments	<code>TukeyHSD(aov(measurements~treatments), ordered=TRUE)</code>
Drawing simultaneous confidence intervals	<code>plot(TukeyHSD(aov(measurements~treatments), ordered=TRUE))</code>
REGRESSION ANALYSIS	
Covariance	<code>cov(data1, data2)</code>
Correlation	<code>cor(data1, data2)</code>
Point graph of bivariate relation	<code>plot(x, y)</code>
Regression line	<code>lm(y~x)</code>
Hypothesis about significance of regression	<code>anova(lm(y~x)) lub summary(lm(y~x))</code>
Point graph and regression line together	<code>plot(x, y); abline(regression line)</code>
Prediction of missing values	<code>predict(regression line, data.frame(x=c(x1, ..., xk)))</code>
CHI-SQUARE TESTS	
Goodness of fit test (qualitative data)	<code>chisq.test(observed frequencies, p=expected probabilities)</code>
Normality tests	<code>pearson.test(data, adjusted=T)      pearson.test(data, adjusted=F)</code>
Caution! "nortest" package required	<code>lillie.test(data) shapiro.test(data)</code>
Test of independence of two variables	<code>chisq.test(data.frame(data1, data2))</code>