

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

| 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<br>(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)<br>Caution! „arm” package required                                 | <code>discrete.histogram(data)</code>  |
| Histogram (for grouped data)  | <code>hist(data, main=title, xlab=label of x)</code>   |
| Automatic declaration of graphics titles e.g. using “for”<br>procedure (with space or without respectively) | <code>paste(„text”, name)</code><br><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<br>binomial<br>Poisson<br>exponential<br>normal<br>t-Student<br>chi-square<br>F Snedecor | <code>binom</code><br><code>pois</code><br><code>exp</code><br><code>norm</code><br><code>t</code><br><code>chisq</code><br><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 $\sigma$<br>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<br>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 $\sigma^2$<br>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 $\sigma$<br>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<br>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 $\sigma^2$<br>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)<br>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<br>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)<br>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<br>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!<br>data.frame(measurements, methods) |   |
| Bartlett test of homogeneity of variances  | bartlett.test(measurements~treatments)  |
| Analysis of variance   | anova(lm(measurements~treatments))  |
| TukeyTest HSD of homogeneous treatments  | TukeyHSD(aov(measurements~treatments),ordered=TRUE)   |
| Drawing simultaneous confidence intervals  | plot(TukeyHSD(aov(measurements~treatments),ordered=TRUE))   |
| REGRESSION ANALYSIS  |   |
| Covariance   | cov(data1, data2)   |
| Correlation  | cor(data1, data2)   |
| Point graph of bivariate relation  | plot(x, y)  |
| Regression line  | lm(y~x)   |
| Hypothesis about significance of regression  | anova(lm(y~x)) lub summary(lm(y~x))   |
| Point graph and regression line together   | plot(x, y); abline(regression line)   |
| Prediction of missing values   | predict(regression line, data.frame(x=c(x1, ..., xk)))  |
| CHI-SQUARE TESTS   |   |
| Goodness of fit test (qualitative data)  | chisq.test(observed frequencies, p=expected probabilities)  |
| Normality tests<br>Caution! "nortest" package required   | pearson.test(data, adjusted=T)    pearson.test(data, adjusted=F)<br>lillie.test(data)<br>shapiro.test(data) |
| Test of independence of two variables  | chisq.test(data.frame(data1, data2))  |