

Introduction to parametric tests

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February 20, 2014

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1 Introduction

This is a detailed explanations of an R-session carried out in the context of an internal workshop at the [Crop Science Research Group](#) at the University of Bonn (Germany). In this session we will have an introductory exercise on Analysis of Variance.

You may have installed in your computer at least an actual version of [R](#) and of the editor [RStudio](#). Additionally a version of the package [SWEApack](#) will be provided by the tutor.

The data set used in this session belongs to the results of the SWEA project ([Agricultural Use and Vulnerability of Small Wetlands in East Africa](#)). To load the data set in your [R Workspace](#) you may type the following command lines.

```
> library(SWEApack)
> data(Africa)
```

Two data frames are downloaded in your session. We will work only with the table `Africa.head`. This table contains observations of 36 plots in four localities (two localities from Kenya and two localities from Tanzania). Plots are classified into three land uses ([Figure 1](#)), namely “unused fields” (semi-natural vegetation), “grazing lands” and “fallows” (long- and short-term abandoned crop fields).

Additional to country, locality and land use membership, `Africa.head` contains soil chemical parameters for each plot, which are organic carbon content (in g kg^{-1}), total nitrogen content (in g kg^{-1}), plant available phosphorous (in mg kg^{-1}), exchangeable potassium (in cmol kg^{-1}), electric conductivity (in dS m^{-1}) and pH ([Kamiri , 2010](#)).

2 Analysis of variance

The analysis of variance (ANOVA) assumes that the variability of measurements is influenced by determined factors. Additional effects by unknown factors are considered as random (non-systematic) and called “experimental errors”. For an easy explanation we will consider the effect of just one factor (independent variable) on a dependent variable, the so called one-way ANOVA. Just to contextualize this example in a scientific problematic, our question can be “*Is the electric conductivity of the soils depending on their country of origin?*” or in other words “*Is the country of origin influencing the electric conductivity of the soils?*” I do not really know, how interesting or logic this question can be but it works as a nice



Figure 1: Classification of land uses in the context of the SWEA project.

example. Then we may formulate a null hypothesis H_0 : “electric conductivity of soils is not depending on their country of origin”. The alternative hypothesis will be H_1 : “electric conductivity of is depending on country of origin and therefore different between countries”.

To carry out the analysis of variance we will use the function `aov`, which is internally calling a linear model (function `lm`). Alternatively you can use the function `anova`, but then you may previously fit the linear model.

```
> ANOVA <- aov(EC~Country, data=Africa.head)
> summary(ANOVA)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Country	1	11.447	11.447	39.94	3.32e-07 ***
Residuals	34	9.745	0.287		

```
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

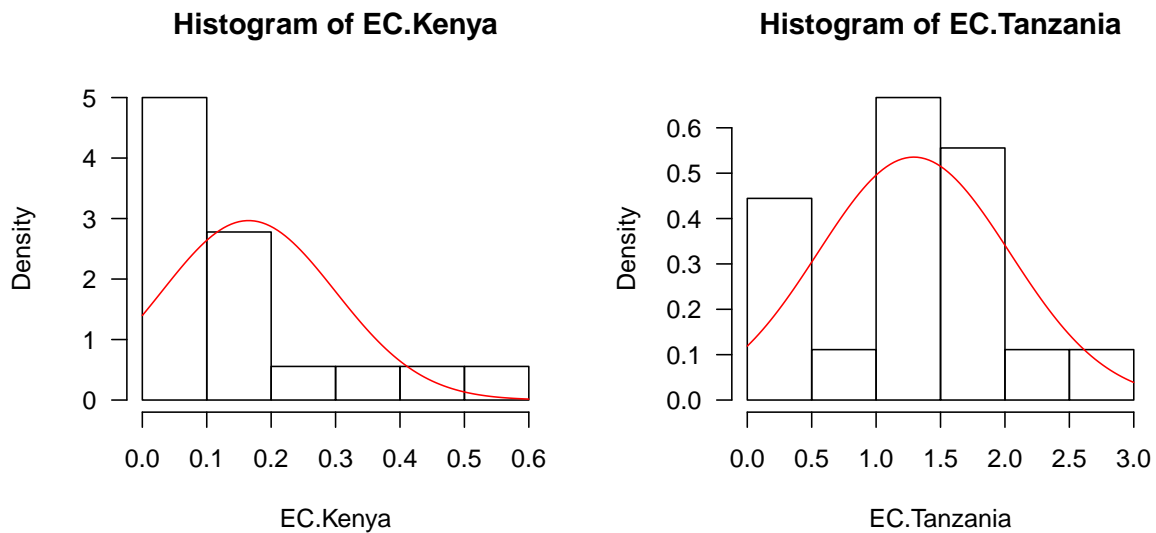
According to the result of `aov` there is a significant difference in the soil electric conductivity between countries. But is this result good enough?

The analysis of variance is a parametric test, that is to say, the values may be normal distributed within populations (in this case different countries). Additionally, the variances between compared populations may be homogeneously distributed (homoscedasticity).

2.1 Tests of normality

Previous to start a test of normality, we look a graphic overview of the distribution of values for the variable electric conductivity. For it, we first split the variable by country of origin and then we check their histograms.

```
> ## Electric conductivity in Kenyan soils
> EC.Kenya <- subset(Africa.head, Country=="Kenya")$EC
> ## Electric conductivity in Tanzanian soils
> EC.Tanzania <- subset(Africa.head, Country=="Tanzania")$EC
> ## Histograms
> par(mfrow=c(1,2), las=1)
> ## Electric conductivity in Kenyan soils
> hist(EC.Kenya, freq=FALSE)
> curve(dnorm(x, mean=mean(EC.Kenya), sd=sd(EC.Kenya)), col="red", add=TRUE)
> ## Electric conductivity in Tanzanian soils
> hist(EC.Tanzania, freq=FALSE)
> curve(dnorm(x, mean=mean(EC.Tanzania), sd=sd(EC.Tanzania)), col="red", add=TRUE)
```



To make life easy, we can alternatively check the distribution for the residuals in the models. Here, we will use an additional graphic option showing the deviance of observed distribution to a theoretical normal distribution, namely the normal quantile-quantile-plot (Q-Q-plot).

```
> ## Extraction of residuals
> Residuals <- resid(ANOVA)
> par(mfrow=c(1,2), las=1)
> ## Plotting histogram of residuals
> hist(Residuals, freq=FALSE, col="grey")
> curve(dnorm(x, mean=mean(Residuals), sd=sd(Residuals)), col="red", add=TRUE)
> ## Plotting q-q plots
> qqnorm(Residuals, asp=1)
> qqline(Residuals, lty=2)
```

So, what do you think? Is the distribution of residuals a normal one? While some authors recommend a visual assessment as a honest alternative to decide, if the distribution is normal, other authors consider that visual assessment cannot substitute a test.

To check normal distribution of values we have two alternatives, among others, the Kolmogorov-Smirnov test (function `ks.test`) and the Shapiro-Wilk test (function `shapiro.test`). In this session we will use the second option but you will find some advices how to use `ks.test` by [Dormann & Kühn \(2011\)](#).

```
> shapiro.test(Residuals)
```

Shapiro-Wilk normality test

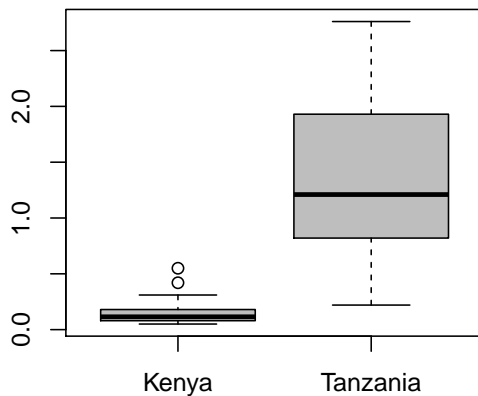
```
data: Residuals
W = 0.9026, p-value = 0.004017
```

In the Shapiro-Wilk test our null hypothesis is H_0 : “the distribution of residuals is not different from a normal one”. Considering a confidence interval of 95%, we will find significant differences when the p-value is lower or equal to 5% ($P < 0.05$), which is the case in this example.

2.2 Tests of homoscedasticity

The second condition for an ANOVA is that the variances between compared populations are homogeneous (homoscedastic). As in the previous example, we will first get a graphic overview, in this case a boxplot of the soil electric conductivity by country.

```
> boxplot(EC ~ Country, data=Africa.head, col="grey")
```



Obviously the variance in the Tanzanian samples is much bigger than the variance for the Kenyan samples. Herewith we may have also more than one alternative but we will just use the Bartlett test (function `bartlett.test`).

```
> bartlett.test(EC ~ Country, data=Africa.head)
```

Bartlett test of homogeneity of variances

data: EC by Country

Bartlett's K-squared = 34.7121, df = 1, p-value = 3.823e-09

Now our null hypothesis was H_0 : “the variances of soil electric conductivity compared between countries are homogeneous”. Once again, a p-value lower than 5% will indicate a significant difference between variances, which is the case here.

In conclusion, the variable electric conductivity does not fulfill the requirements for a parametric test and the results of the first ANOVA are not valid. What to do now?

2.3 Data transformations

Is in not everything lost for the moment. The salvation can be provided by data transformation. The most popular methods are relatively simple (see [Table 1](#)). Such transformations change not only the scale of the variables but they also modify the distribution of the values. Numerical variables are frequently transformed using the logarithmic, reciprocal and squared root transformations. The arcsine-squared root transformation is used to transform percentages and proportions.

Table 1: Formulas and functions used for data transformation.

Function or formula in R	description
<code>log(x+c, base=exp(1))</code>	logarithmic transformation
<code>1/(x+c)</code>	reciprocal transformation
<code>sqrt(x+c)</code>	squared root transformation
<code>asin(sqrt(x/c))*180/pi</code>	arcsine-squared root transformation
<code>(x-mean(x)/sd(x)), scale(x)</code>	standardisation

If the mentioned transformations did not succeed, it remains the alternative of a Box-Cox transformation may be carried out. For it the function `boxcox()` is available in the package [MASS](#).

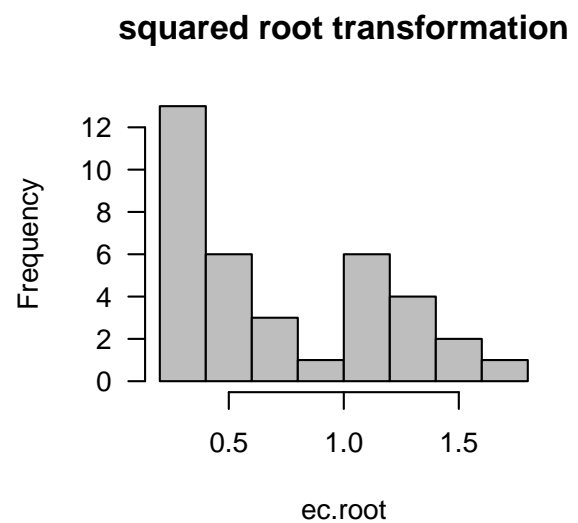
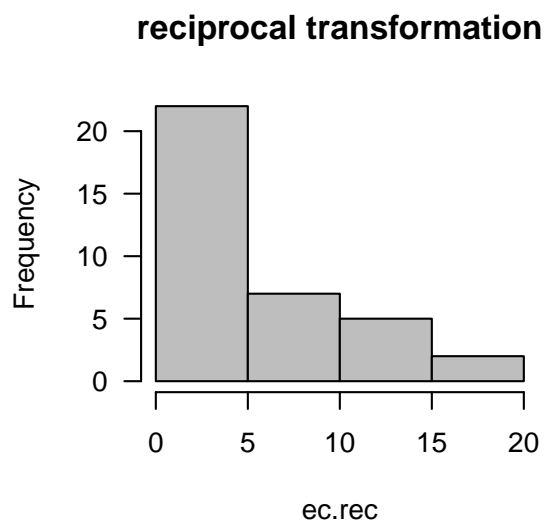
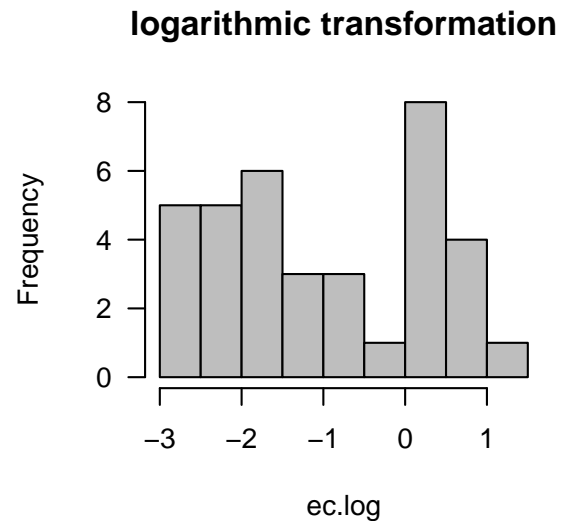
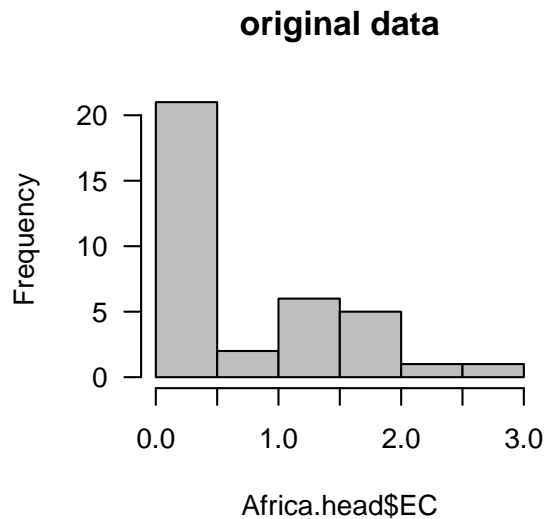
Some special cases are the dummy-transformation, applied for the analysis of nominal variables, and the standardization required mainly by multivariate statistics.

```
> # logarithmic transformation
> ec.log <- log(Africa.head$EC, base=exp(1))
> # reciprocal transformation
> ec.rec <- 1/Africa.head$EC
```

```
> # squared root transformation
> ec.root <- sqrt(Africa.head$EC)
```

Let us take a look on the effects of the previous data transformations in the distribution of values.

```
> par(mfrow=c(2,2), las=1)
> hist(Africa.head$EC, col="grey", main="original data")
> hist(ec.log, col="grey", main="logarithmic transformation")
> hist(ec.rec, col="grey", main="reciprocal transformation")
> hist(ec.root, col="grey", main="squared root transformation")
```



We test first the normality of residuals and homoscedasticity of variances in the logarithmic transformation of soil electric conductivity.

```
> # Test to normality
> shapiro.test(resid(aov(ec.log~Country, data=Africa.head)))
```

Shapiro-Wilk normality test

```
data: resid(aov(ec.log ~ Country, data = Africa.head))
W = 0.9831, p-value = 0.8442
```

```
> # Test to homoscedasticity
> bartlett.test(ec.log~Country, data=Africa.head)
```

Bartlett test of homogeneity of variances

```
data: ec.log by Country
Bartlett's K-squared = 0.5208, df = 1, p-value = 0.4705
```

Here we obtain for the logarithmic transformation of soil electric conductivity p-values higher than 0.05 in both, the Shapiro-Wilk test and the Bartlett test. That means, the residuals are normally distributed and the variances are homogeneous. But, what about the other transformation alternatives?

```
> # Tests for reciprocal transformation
> shapiro.test(resid(aov(ec.rec~Country, data=Africa.head)))
```

Shapiro-Wilk normality test

```
data: resid(aov(ec.rec ~ Country, data = Africa.head))
W = 0.9368, p-value = 0.04055
```

```
> bartlett.test(ec.rec~Country, data=Africa.head)
```

Bartlett test of homogeneity of variances

```
data: ec.rec by Country
Bartlett's K-squared = 22.5029, df = 1, p-value = 2.098e-06
```

```
> # Tests for squared root transformation
> shapiro.test(resid(aov(ec.root~Country, data=Africa.head)))
```

Shapiro-Wilk normality test

```
data: resid(aov(ec.root ~ Country, data = Africa.head))
W = 0.9436, p-value = 0.06584
```

```
> bartlett.test(ec.root~Country, data=Africa.head)
```

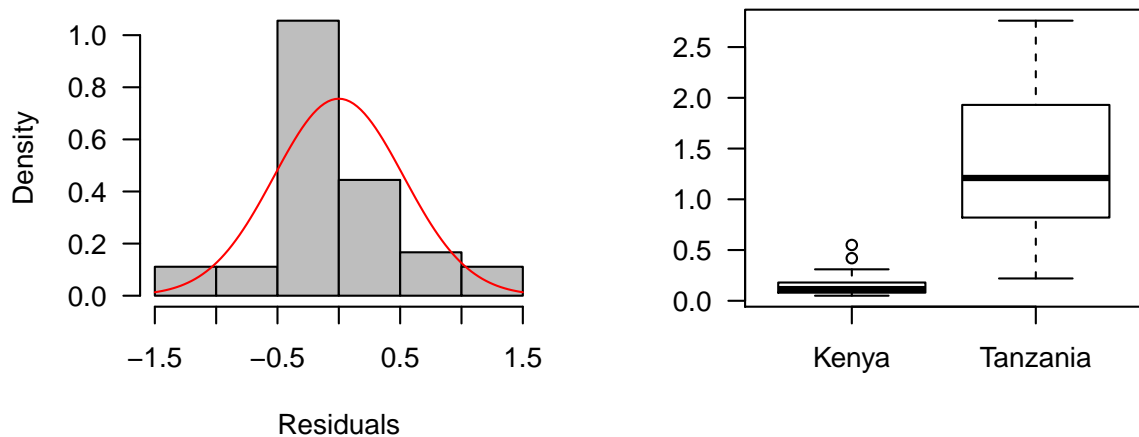
Bartlett test of homogeneity of variances

```
data: ec.root by Country
Bartlett's K-squared = 12.7333, df = 1, p-value = 0.0003592
```

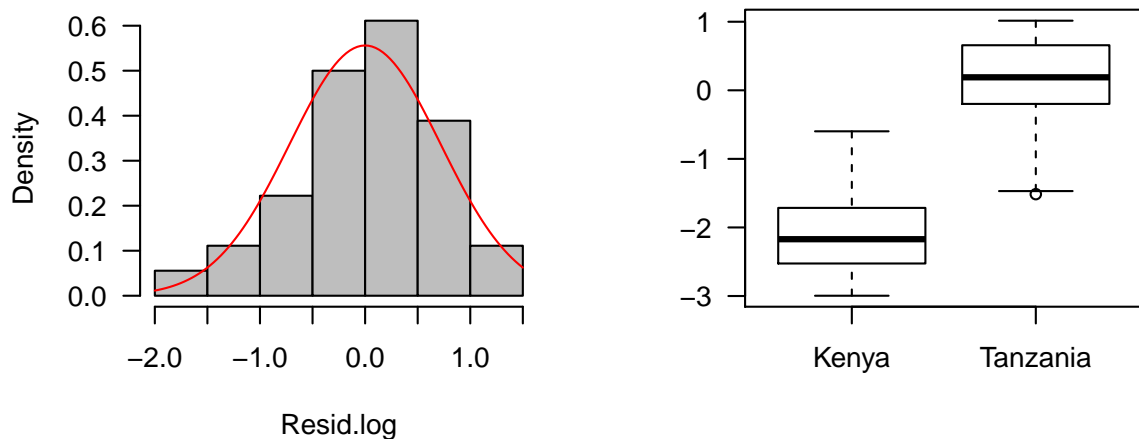
For the reciprocal transformation none of the requirements are fulfilled (considering the same significance interval of 95%), while for the squared root transformation only the normality of residuals can be achieved. Therefore, the logarithmic transformation is the only one achieving the requirements for a parametric test. We can compare original and transformed data in following graphics.

```
> # Calculation of residuals for logarithmic transformation
> Resid.log <- resid(aov(ec.log ~ Country, data=Africa.head))
> # Plotting distributions
> par(mfrow=c(2,2), las=1)
> hist(Residuals, freq=FALSE, col="grey", main="residuals from original data")
> curve(dnorm(x, mean=mean(Residuals), sd=sd(Residuals)), col="red", add=TRUE)
> boxplot(EC ~ Country, data=Africa.head)
> hist(Resid.log, freq=FALSE, col="grey", main="logarithmic transformation")
> curve(dnorm(x, mean=mean(Resid.log), sd=sd(Resid.log)), col="red", add=TRUE)
> boxplot(ec.log ~ Country, data=Africa.head)
```

residuals from original data



logarithmic transformation



You see? After logarithmic transformation the histogram seems to be better adjusted to a theoretical normal distribution and the boxplots are more similar between them.

2.4 ANOVA by the proper way

Once we find a proper transformation of the data, we can carry out the ANOVA and assure correct results.

```
> ANOVA <- aov(ec.log ~ Country, data=Africa.head)
> summary(ANOVA)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Country	1	38.21	38.21	72.11	6.44e-10 ***
Residuals	34	18.02	0.53		

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

After transformation there is still a highly significant difference between countries in the soil electric conductivity. And we can publish it.

3 Post-hoc contrasts

Previous to finish the parametric test, some advices regarding ANOVA. In the example we applied the ANOVA to a factor with just two levels (Kenya and Tanzania). In such case may be enough the use of

a paired two-sample t-test (function `pairwise.t.test`). We can apply the one-way ANOVA to factors with more than two levels instead, for example comparing soil reaction in different localities.

```
> ANOVA <- aov(pH ~ Locality, data=Africa.head)
> summary(ANOVA)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Locality	3	17.17	5.724	14.15	4.82e-06 ***
Residuals	32	12.94	0.404		

```
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

In some cases we may require the addition of a new explanatory (independent) variable, for example the land use type.

```
> ANOVA <- aov(pH ~ Locality + LandUse, data=Africa.head)
> summary(ANOVA)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Locality	3	17.172	5.724	14.410	5.4e-06 ***
LandUse	2	1.024	0.512	1.289	0.29
Residuals	30	11.916	0.397		

```
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Or even more, we may also need to know the interaction between factors.

```
> ANOVA <- aov(pH ~ Locality * LandUse, data=Africa.head)
> summary(ANOVA)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Locality	3	17.172	5.724	14.865	1.12e-05 ***
LandUse	2	1.024	0.512	1.329	0.283
Locality:LandUse	6	2.675	0.446	1.158	0.361
Residuals	24	9.242	0.385		

```
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

We come back to the first example and we test the effect of locality in the soil reaction (measured as pH). Of course, you may check previously the normality of residuals and the homoscedasticity of variances, but this is just one example... Here the ANOVA tell us “there is a difference in the soil reaction depending on sampling locality” but we do not know, which localities have soil pH values differetn to which ones. To know it, we require a *post-hoc* contrast. In this session we will use the Tukey’s honest significant difference (function `TukeyHSD`).

```
> ANOVA <- aov(pH ~ Locality, data=Africa.head)
> summary(ANOVA)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Locality	3	17.17	5.724	14.15	4.82e-06 ***
Residuals	32	12.94	0.404		

```
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

> Tukey <- TukeyHSD(ANOVA)
> Tukey
```

```
Tukey multiple comparisons of means
 95% family-wise confidence level
```

```
Fit: aov(formula = pH ~ Locality, data = Africa.head)
```

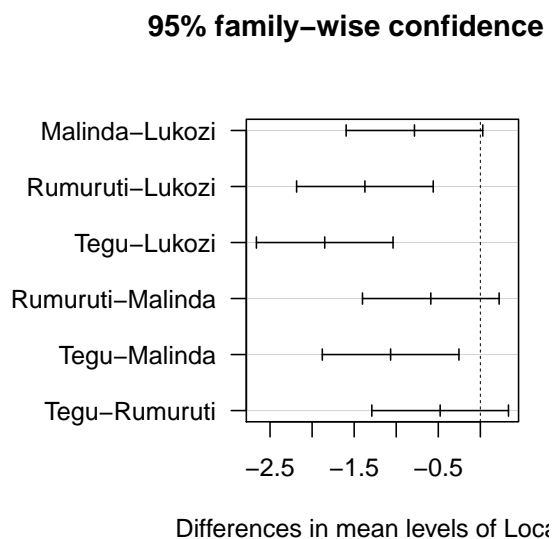
```
$Locality
```

	diff	lwr	upr	p adj
--	------	-----	-----	-------

Malinda-Lukozi	-0.7833333	-1.595514	0.02884776	0.0619263
Rumuruti-Lukozi	-1.3722222	-2.184403	-0.56004113	0.0003764
Tegu-Lukozi	-1.8500000	-2.662181	-1.03781891	0.0000038
Rumuruti-Malinda	-0.5888889	-1.401070	0.22329220	0.2225102
Tegu-Malinda	-1.0666667	-1.878848	-0.25448558	0.0062137
Tegu-Rumuruti	-0.4777778	-1.289959	0.33440331	0.3963421

In the output of `TukeyHSD` you may look for those pairwise comparisons where both, the upper and the lower limit of the difference are either negative or positive. If the mentioned values have different symbols, the difference is not significant. That can be also checked in the column showing the p-values (`p adj`). There is also a graphic option to show the results of `TukeyHSD`.

```
> # Plotting Tukey output
> par(las=1, mar=c(5,10,5,1))
> plot(Tukey)
```



4 Disclaimer

This handout is a preliminary version done by a non-statistician, therefore use carefully and at own risk. Since this text will be improved for further R-courses, any comment, correction and suggestion is most welcome (send them to malvarez@uni-bonn.de).

The content of this handout is based on German references (Ligges , 2008; Köhler et al. , 2002; Dormann & Kühn , 2011).

Some English references can be Zar (2009) and the tutorials by King (2014), Schumacher (2007) and Everitt & Hothorn (2005).

5 Exercises

- You ask “is there a difference in the soil pH between sampling localities?”, but this time you will check first the normal distribution of residuals and the homogeneity of variances between localities.
- Is there differences between applied insecticides in the amount of insects (survivors) found in experimental plots? Use the data set `InsectSprays`.
- Apply a factorial ANOVA to the data set `ToothGrowth`.
- Now try the same with your own data set.

6 Bibliography

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