

Nested

First Example

```
#rm(list = ls()) # Clean the workspace
library(lsmmeans) # Call the library
library(lme4) # Call the library
library(car) # Call the library
# Read Data
mydata<-read.table(file = "C:/Users/toledo/Dropbox/UNIPD/Biostatistics Course R Spring 2018/corso STAT PhD 2018/
                    sep = "\t",header = TRUE,stringsAsFactors = TRUE)
mydata$Boar<-as.factor(mydata$Boar) # Set the variable as factor
str(mydata) # See the structure of my data
```

```
## 'data.frame': 20 obs. of 4 variables:
## $ Boar : Factor w/ 5 levels "1","2","3","4",...: 1 1 1 1 2 2 2 2 3 3 ...
## $ Sow : Factor w/ 10 levels "A","B","C","D",...: 1 1 2 2 3 3 4 4 5 5 ...
## $ Replicate: int 1 2 1 2 1 2 1 2 1 2 ...
## $ ADG : num 2.77 2.38 2.58 2.94 2.28 2.22 3.01 2.61 2.36 2.71 ...
```

```
contrasts(mydata$Boar)<-contr.SAS # Set the contrast as SAS
contrasts(mydata$Sow)<-contr.SAS # Set the contrast as SAS
table(mydata$Sow,mydata$Boar) # Frequencies for factors
```

```
##
## 1 2 3 4 5
## A 2 0 0 0 0
## B 2 0 0 0 0
## C 0 2 0 0 0
## D 0 2 0 0 0
## E 0 0 2 0 0
## F 0 0 2 0 0
## G 0 0 0 2 0
## H 0 0 0 2 0
## I 0 0 0 0 2
## L 0 0 0 0 2
```

```
mymodel<-lm(ADG ~ Boar + Sow%in%Boar, data = mydata ) # Fit the model with Nesting
summary(mymodel) # See the results
```

```
##
## Call:
## lm(formula = ADG ~ Boar + Sow %in% Boar, data = mydata)
##
## Residuals:
## Min 1Q Median 3Q Max
## -0.2050 -0.1113 0.0000 0.1113 0.2050
##
## Coefficients: (40 not defined because of singularities)
## Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.4900 0.1391 17.900 6.33e-09 ***
## Boar1 0.2700 0.1967 1.372 0.1999
## Boar2 0.3200 0.1967 1.627 0.1349
## Boar3 0.2400 0.1967 1.220 0.2505
## Boar4 -0.2150 0.1967 -1.093 0.3001
## Boar1:Sow1 -0.1850 0.1967 -0.940 0.3692
## Boar2:Sow1 NA NA NA NA
## Boar3:Sow1 NA NA NA NA
## Boar4:Sow1 NA NA NA NA
```

```
## Boar5:Sow1      NA      NA      NA      NA
## Boar1:Sow2      NA      NA      NA      NA
## Boar2:Sow2      NA      NA      NA      NA
## Boar3:Sow2      NA      NA      NA      NA
## Boar4:Sow2      NA      NA      NA      NA
## Boar5:Sow2      NA      NA      NA      NA
## Boar1:Sow3      NA      NA      NA      NA
## Boar2:Sow3     -0.5600    0.1967   -2.847    0.0173 *
## Boar3:Sow3      NA      NA      NA      NA
## Boar4:Sow3      NA      NA      NA      NA
## Boar5:Sow3      NA      NA      NA      NA
## Boar1:Sow4      NA      NA      NA      NA
## Boar2:Sow4      NA      NA      NA      NA
## Boar3:Sow4      NA      NA      NA      NA
## Boar4:Sow4      NA      NA      NA      NA
## Boar5:Sow4      NA      NA      NA      NA
## Boar1:Sow5      NA      NA      NA      NA
## Boar2:Sow5      NA      NA      NA      NA
## Boar3:Sow5     -0.1950    0.1967   -0.991    0.3449
## Boar4:Sow5      NA      NA      NA      NA
## Boar5:Sow5      NA      NA      NA      NA
## Boar1:Sow6      NA      NA      NA      NA
## Boar2:Sow6      NA      NA      NA      NA
## Boar3:Sow6      NA      NA      NA      NA
## Boar4:Sow6      NA      NA      NA      NA
## Boar5:Sow6      NA      NA      NA      NA
## Boar1:Sow7      NA      NA      NA      NA
## Boar2:Sow7      NA      NA      NA      NA
## Boar3:Sow7      NA      NA      NA      NA
## Boar4:Sow7      0.3900    0.1967    1.982    0.0756 .
## Boar5:Sow7      NA      NA      NA      NA
## Boar1:Sow8      NA      NA      NA      NA
## Boar2:Sow8      NA      NA      NA      NA
## Boar3:Sow8      NA      NA      NA      NA
## Boar4:Sow8      NA      NA      NA      NA
## Boar5:Sow8      NA      NA      NA      NA
## Boar1:Sow9      NA      NA      NA      NA
## Boar2:Sow9      NA      NA      NA      NA
## Boar3:Sow9      NA      NA      NA      NA
## Boar4:Sow9      NA      NA      NA      NA
## Boar5:Sow9      0.1600    0.1967    0.813    0.4350
```

```
## ---
```

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

```
##
```

```
## Residual standard error: 0.1967 on 10 degrees of freedom
```

```
## Multiple R-squared:  0.6315, Adjusted R-squared:  0.2999
```

```
## F-statistic: 1.904 on 9 and 10 DF,  p-value: 0.1649
```

```
anova(mymodel) # ANOVA table SS type III
```

```
## Analysis of Variance Table
```

```
##
```

```
## Response: ADG
```

```
##      Df Sum Sq Mean Sq F value Pr(>F)
```

```
## Boar    4 0.09973 0.024932  0.6443 0.64332
```

```
## Boar:Sow  5 0.56355 0.112710  2.9124 0.07067 .
```

```
## Residuals 10 0.38700 0.038700
```

```
## ---
```

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

```

#ref.grid(mymodel)
lsmeans(mymodel,"Boar")          # LSM for factor

## Boar lsmean          SE df lower.CL upper.CL
## 1      2.6675 0.09836158 10 2.448337 2.886663
## 2      2.5300 0.09836158 10 2.310837 2.749163
## 3      2.6325 0.09836158 10 2.413337 2.851663
## 4      2.4700 0.09836158 10 2.250837 2.689163
## 5      2.5700 0.09836158 10 2.350837 2.789163
##
## Results are averaged over the levels of: Sow
## Confidence level used: 0.95

lsmeans(mymodel,"Sow")          # LSM for factor

## Sow Boar lsmean          SE df lower.CL upper.CL
## A 1      2.575 0.1391043 10 2.265056 2.884944
## B 1      2.760 0.1391043 10 2.450056 3.069944
## C 2      2.250 0.1391043 10 1.940056 2.559944
## D 2      2.810 0.1391043 10 2.500056 3.119944
## E 3      2.535 0.1391043 10 2.225056 2.844944
## F 3      2.730 0.1391043 10 2.420056 3.039944
## G 4      2.665 0.1391043 10 2.355056 2.974944
## H 4      2.275 0.1391043 10 1.965056 2.584944
## I 5      2.650 0.1391043 10 2.340056 2.959944
## L 5      2.490 0.1391043 10 2.180056 2.799944
##
## Confidence level used: 0.95

mymodel.1<-aov(ADG ~ Boar + Error(Sow), data = mydata) # Set the factor as an error term
summary(mymodel.1)          # See the results

##
## Error: Sow
##           Df Sum Sq Mean Sq F value Pr(>F)
## Boar         4 0.0997 0.02493    0.221  0.916
## Residuals    5 0.5636 0.11271
##
## Error: Within
##           Df Sum Sq Mean Sq F value Pr(>F)
## Residuals  10 0.387  0.0387
#### Random Model

mymodel.3<-lmer(ADG ~ Boar + (1|Sow), data = mydata ) # Fit a model with Random effect
summary(mymodel.3)          # See the results

## Linear mixed model fit by REML ['lmerMod']
## Formula: ADG ~ Boar + (1 | Sow)
## Data: mydata
##
## REML criterion at convergence: 6.1
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -1.1527 -0.5563 -0.1764  0.6277  1.5054
##
## Random effects:
## Groups Name Variance Std.Dev.
## Sow (Intercept) 0.03701 0.1924
## Residual 0.03870 0.1967
## Number of obs: 20, groups: Sow, 10

```

```
##
## Fixed effects:
##           Estimate Std. Error t value
## (Intercept)  2.5700    0.1679  15.310
## Boar1        0.0975    0.2374   0.411
## Boar2       -0.0400    0.2374  -0.168
## Boar3        0.0625    0.2374   0.263
## Boar4       -0.1000    0.2374  -0.421
##
## Correlation of Fixed Effects:
##      (Intr) Boar1  Boar2  Boar3
## Boar1 -0.707
## Boar2 -0.707  0.500
## Boar3 -0.707  0.500  0.500
## Boar4 -0.707  0.500  0.500  0.500

Anova(mymodel.3,type = 3,test.statistic = "F") # Anova table SS Type III

## Analysis of Deviance Table (Type III Wald F tests with Kenward-Roger df)
##
## Response: ADG
##           F Df Df.res    Pr(>F)
## (Intercept) 234.4033  1      5 2.156e-05 ***
## Boar        0.2212  4      5  0.9155
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

lsmeans(mymodel.3,"Boar") # LSM for factor

## Boar lsmean      SE df lower.CL upper.CL
## 1      2.6675 0.1678616  5 2.235998 3.099002
## 2      2.5300 0.1678616  5 2.098498 2.961502
## 3      2.6325 0.1678616  5 2.200998 3.064002
## 4      2.4700 0.1678616  5 2.038498 2.901502
## 5      2.5700 0.1678616  5 2.138498 3.001502
##
## Degrees-of-freedom method: satterthwaite
## Confidence level used: 0.95
```

Second Example

```
#### Second Example ####

#rm(list = ls())      # Clean the workspace
library(lsmmeans)     # Call the library
library(lme4)
# Read Data
mydata<-read.table(file = "../Dropbox/UNIPD/Biostatistics Course R Spring 2018/curso STAT PhD 2018 ExpDesign/d
                      sep = "\t",header = TRUE,stringsAsFactors = TRUE)
mydata$Boar<-as.factor(mydata$Boar) # Set the variable as factor
str(mydata)                        # See the structure of my data
contrasts(mydata$Boar)<-contr.SAS   # Set the contrast as SAS
contrasts(mydata$Sow)<-contr.SAS    # Set the contrast as SAS
table(mydata$Sow,mydata$Boar)      # Frequencies for factors

#### Random Model in a for loop
# To store the results directly in a file use:
sink("../Dropbox/UNIPD/Biostatistics Course R Spring 2018/curso STAT PhD 2018 ExpDesign/data/results.txt",
      append = FALSE)
# For loop to do the analysis for several variables
for (i in 4:5){ # For the 4 and 5 column
  y <- mydata[,i] # Select the column i
  print(paste0("Results for ",colnames(mydata)[i])) # Print the number of the variable
  print("#####") #
  mymodel<-lmer(y ~ mydata$Boar + (1|mydata$Sow)) # Fit the random model
  print(summary(mymodel)) # See the results
  print(Anova(mymodel,type = 3,test.statistic = "F")) # Anova table SS type III
  print(lsmmeans(mymodel,"mydata$Boar")) # LSM for factor
  print("#####")
}
sink() # Return the result to the R console
```