

# ModBalsMassBoth.R

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```
library(ggplot2)
library(lme4)
```

```
## Loading required package: Matrix
```

```
library(nlme)
```

```
##
## Attaching package: 'nlme'

## The following object is masked from 'package:lme4':
##
##      lmList
```

```
library(lsmeans)
```

```
## Warning: package 'lsmeans' was built under R version 3.2.5

## Loading required package: estimability

## Warning: package 'estimability' was built under R version 3.2.5
```

```
library(lubridate)
```

```
## Warning: package 'lubridate' was built under R version 3.2.5

##
## Attaching package: 'lubridate'

## The following object is masked from 'package:base':
##
##      date
```

```
library(multcompView)
```

```
## Warning: package 'multcompView' was built under R version 3.2.5
```

```
library(car)
```

```
## Warning: package 'car' was built under R version 3.2.5
```

```
setwd("D:/Iowa State University/Debinski Lab/Nectar data/MAL")

balssug15 <- read.csv("nectar analysis/data files/balssugar15.csv", header = T)
balssug16 <- read.csv("nectar analysis/data files/balssugar16.csv", header = T)
balssugboth <- rbind(balssug15, balssug16)

balssugboth$year <- as.factor(year(balssugboth$date))

cellN <- with(balssugboth, table(treatment, year))
cellN
```

```
##           year
## treatment 2015 2016
##           C   30   80
##           H   56   71
```

```
cellMean <- with(balssugboth, tapply(mass, list(treatment, year), mean))
cellMean
```

```
##           2015           2016
## C 0.09528345 0.04865045
## H 0.10940649 0.05275608
```

```
modmass <- lmer(mass ~ treatment * year + (1|plot/plant), data = balssugboth)
mass.grid <- ref.grid(modmass)
```

```
## Loading required namespace: lmerTest
```

```
summary(mass.grid)
```

```
## treatment year prediction      SE    df
## C           2015 0.09847256 0.013422314 49.23
## H           2015 0.11243507 0.010582621 19.55
## C           2016 0.04738959 0.009105025 15.97
## H           2016 0.05223706 0.009519857 14.17
##
## Degrees-of-freedom method: satterthwaite
```

```
lsmeans(mass.grid, "treatment")
```

```
## NOTE: Results may be misleading due to involvement in interactions
```

```
## treatment    lsmean      SE    df  lower.CL  upper.CL
## C           0.07293107 0.009012363 14.76 0.05369497 0.09216718
## H           0.08233607 0.008366112  8.30 0.06316496 0.10150718
##
## Results are averaged over the levels of: year
## Degrees-of-freedom method: satterthwaite
## Confidence level used: 0.95
```

```
lsmeans(mass.grid, "year")
```

```
## NOTE: Results may be misleading due to involvement in interactions
```

```
##   year    lsmean      SE    df  lower.CL  upper.CL
## 2015 0.10545382 0.008546203 33.66 0.08807934 0.12282829
## 2016 0.04981332 0.006586524 15.03 0.03577693 0.06384972
##
## Results are averaged over the levels of: treatment
## Degrees-of-freedom method: satterthwaite
## Confidence level used: 0.95
```

```
mass.treat <- lsmeans(mass.grid, "treatment")
```

```
## NOTE: Results may be misleading due to involvement in interactions
```

```
pairs(mass.treat)
```

```
## contrast      estimate      SE    df t.ratio p.value
## C - H      -0.009404992 0.01229693 11.1  -0.765  0.4603
##
## Results are averaged over the levels of: year
```

```
pairs.treat <- pairs(mass.treat)
test(pairs.treat, joint = T)
```

```
## df1 df2    F p.value
##   1 11.1 0.585 0.4603
```

```
mass.year <- lsmeans(mass.grid, "year")
```

```
## NOTE: Results may be misleading due to involvement in interactions
```

```
pairs(mass.year)
```

```
## contrast      estimate      SE    df t.ratio p.value
## 2015 - 2016 0.05564049 0.00903467 154.43  6.159 <.0001
##
## Results are averaged over the levels of: treatment
```

```
pairs.year <- pairs(mass.year)
test(pairs.year, joint = T)
```

```
## df1 df2    F p.value
##   1 154.43 37.928 <.0001
```

```
int.mass <- pairs(mass.grid, by = "year")
int.mass
```

```
## year = 2015:
## contrast      estimate      SE    df t.ratio p.value
## C - H      -0.013962511 0.01709241 33.66  -0.817  0.4197
##
## year = 2016:
## contrast      estimate      SE    df t.ratio p.value
## C - H      -0.004847473 0.01317305 15.03  -0.368  0.7180
```

```
int.masstable <- update(int.mass, by = NULL)
int.masstable
```

```
## contrast year      estimate      SE    df t.ratio p.value
## C - H      2015 -0.013962511 0.01709241 33.66  -0.817  0.4197
## C - H      2016 -0.004847473 0.01317305 15.03  -0.368  0.7180
```

```
test(pairs(int.masstable), joint = T)
```

```
## df1    df2      F p.value
##    1 154.43 0.254  0.6147
```

```
Anova(modmass, type = 3)
```

```
## Analysis of Deviance Table (Type III Wald chisquare tests)
##
## Response: mass
##           Chisq Df Pr(>Chisq)
## (Intercept)  53.8240  1  2.193e-13 ***
## treatment      0.6673  1  0.4139948
## year          12.9676  1  0.0003169 ***
## treatment:year  0.2545  1  0.6139467
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```