

```

> data <- read_excel("R/daemen/324quiz7data1.xlsx")
> view(data)
> aov(Cholesterol ~ Group, data=data)
Call:
  aov(formula = Cholesterol ~ Group, data = data)

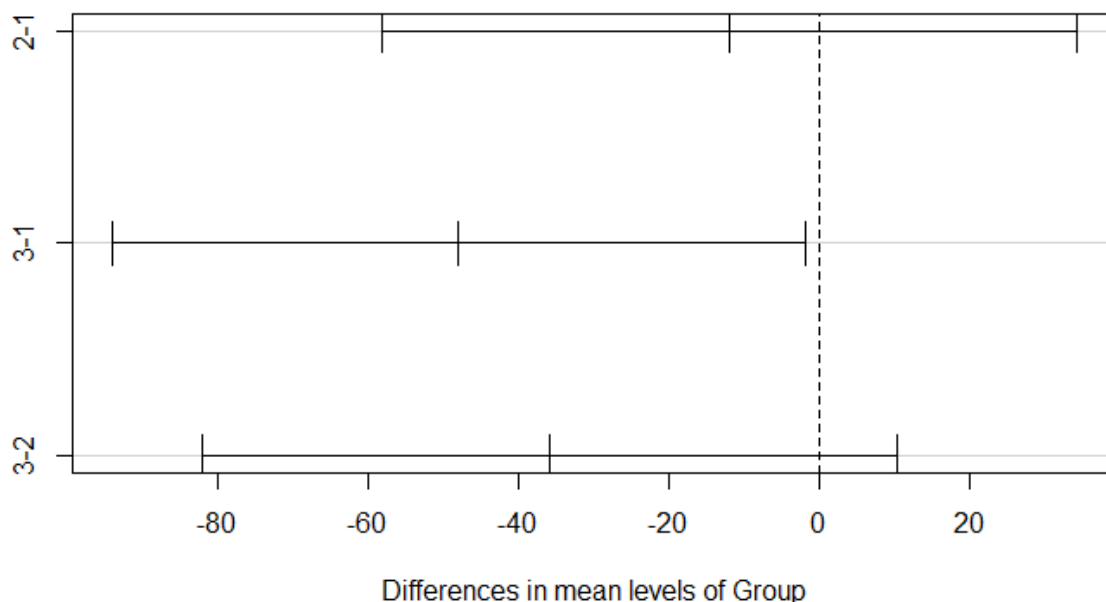
Terms:
              Group Residuals
Sum of Squares  5760      9480
Deg. of Freedom    1        13

Residual standard error: 27.00427
Estimated effects may be unbalanced
> summary(aov(Cholesterol ~ Group, data=data))
      Df Sum Sq Mean Sq F value Pr(>F)
Group   1  5760    5760   7.899 0.0147 *
Residuals 13  9480     729
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> |

> data$Group<-factor(data$Group)
> one_way<-aov(Cholesterol ~ Group, data=data)
> library(multcompview)
> tukey_one<-TukeyHSD(one_way)
> plot(tukey_one)
> |

```

### 95% family-wise confidence level

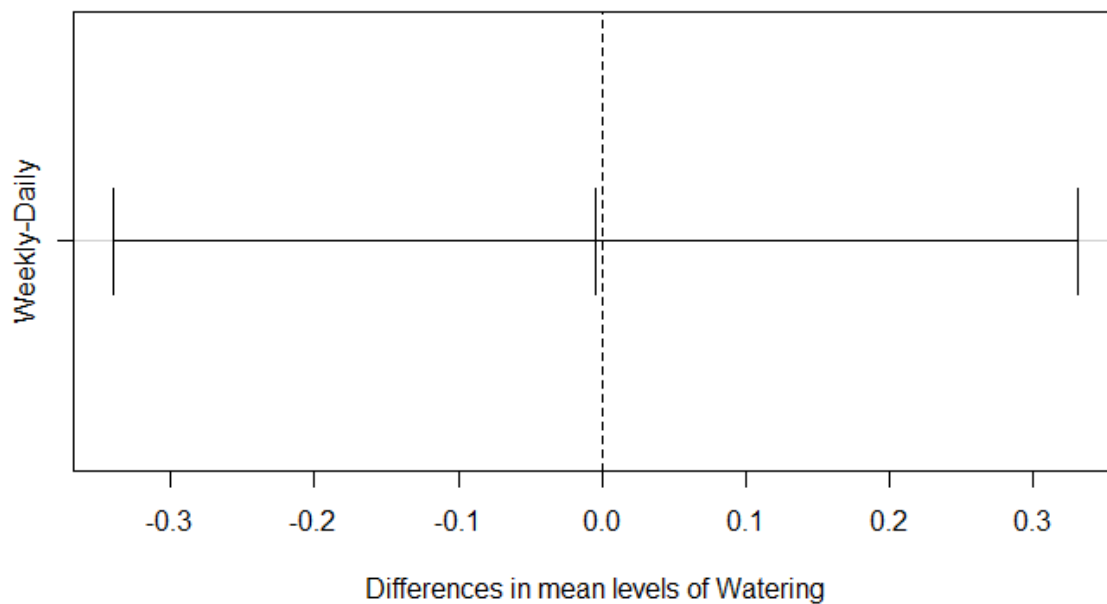


```

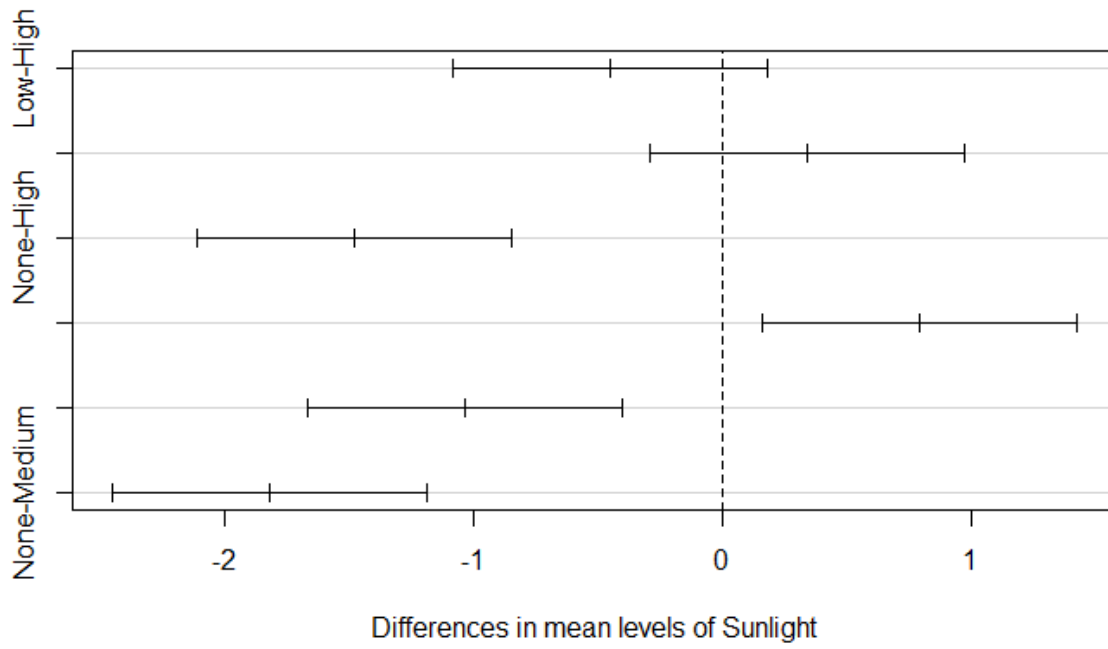
> data <- read_excel("R/daemen/324quiz7data2.xlsx")
> two_way<-aov(Height ~ watering*Sunlight, data=data)
> summary(two_way)
      Df Sum Sq Mean Sq F value Pr(>F)
watering  1  0.000   0.000   0.001  0.976
Sunlight  3 18.765   6.255  23.049 3.9e-08 ***
watering:sunlight 3  1.011   0.337   1.242  0.311
Residuals 32  8.684   0.271
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> tukey_two<-TukeyHSD(two_way)
> plot(tukey_two)
>

```

### 95% family-wise confidence level



### 95% family-wise confidence level



### 95% family-wise confidence level

