Introduction to Statistics with R Session R05: ANOVA

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The example data set

This example data set contains scores on a sustainability scale for three groups, each with a distinct diet: meat-eating (meat), vegetarian (vegetarian) and plant-based (vegan). Furthermore, each participant's self-reported gender is denoted.

Tip: Use the parameter stringsAsFactors to interpret character strings as factors automatically:

```
df = read.csv("R05_notes_dataset.csv", stringsAsFactors = TRUE)
str(df)

## 'data.frame': 150 obs. of 4 variables:
## $ ID : int 1 2 3 4 5 6 7 8 9 10 ...
## $ diet : Factor w/ 3 levels "meat", "vegan",..: 1 1 1 1 1 1 1 1 1 1 1 ...
## $ gender : Factor w/ 2 levels "female", "male": 2 2 2 2 2 2 2 2 2 2 2 2 2 ...
## $ usutainability: int 18 NA 12 14 NA 9 19 9 22 10 ...
```

We can change the **ordering** of the **factor levels**:

```
df$diet = factor(df$diet, levels=c("vegan", "vegetarian", "meat"))
```

Missing data

In the R language, missing data are denoted as NA (not available). That's a dedicated symbol. It's not to be confused with NaN (not a number) that describes impossible data (e.g. from divison by 0).

head(df)

```
## 1 D diet gender sustainability
## 1 1 meat male 18
## 2 2 meat male NA
## 3 3 meat male 12
## 4 4 meat male 14
## 5 5 meat male NA
## 6 6 meat male 9
```

Some functions do not work properly when NAs are present:

mean(df\$sustainability)

[1] NA

In many cases, these functions implement the argument na.rm=TRUE (= remove NA values):

mean(df\$sustainability, na.rm=TRUE)

[1] 15.6338

There are different strategies for missing values:

- Delete the data
 - Delete the entire observation (row)
 - Exclude the observation only from those analyses where the missing value would be required
- Substitute the missing value with a *typical* value, like . . .
 - the mean \bar{x}
 - the median
 - the mode
 - **.** . . .
- Copy the last observation
- Estimate the missing value
 - Regression
 - Multiple imputation
 -

Introductory reading: https://doi.org/10.4097/kjae.2013.64.5.402

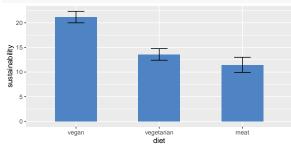
We can delete all rows that contain **any** missing data cell with na.omit(). That's simple but wasteful in applications where data are valuable.

df = na.omit(df)

Visualization: one-way ANOVA

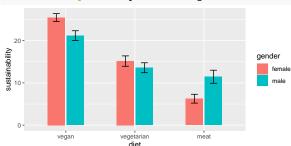
For a one-way ANOVA (one grouping variable), we can simply use a bar plot with error bars:

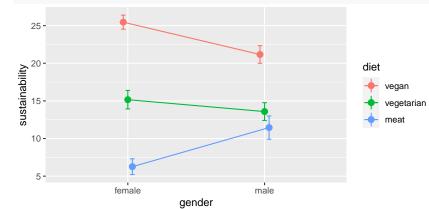
```
df %>% filter(gender=="male") %>%
  ggplot(., aes(y=sustainability, x = diet)) +
  stat_summary(fun = mean, geom = "bar", width=0.5, fill="#4E84C4") +
  stat_summary(fun.data = mean_se, geom = "errorbar", width=0.2)
```



Visualization: Two-way ANOVA

For two factors, we can use a bar plot with different fills. The bars and error bar need to be *dodged* a little:





Numerical ANOVA computation

The afex package implements convenient functions to compute ANOVAs. We will use the function aov_ez():

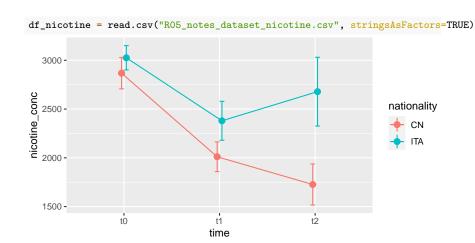
One-way ANOVA: Effect of gender

One-way ANOVA: Effect of diet

Two-way ANOVA: Effect of gender and diet

diet

```
aov_ez(id = "ID", dv = "sustainability",
          between = c("gender", "diet"), data = df) %>% summary()
  ## Anova Table (Type 3 tests)
  ##
  ## Response: sustainability
  ##
             num Df den Df MSE F
                                            ges
                                                  Pr(>F)
  ## gender
               1 136 33.591 0.0534 0.00039 0.8176617
  ## diet 2 136 33.591 74.0283 0.52122 < 2.2e-16 ***
  ## gender:diet 2 136 33.591 8.1932 0.10753 0.0004368 ***
  ## ---
  ## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
    25 -
sustainability
1
                                                       aender
                                                          female
    10 -
     5 -
                          vegetarian
             vegan
                                          meat
```



Within-subject ANOVA for factor time:

```
aov ez(id = "ID", dv = "nicotine conc",
        within = "time", data = df nicotine) %>% summary()
##
## Univariate Type III Repeated-Measures ANOVA Assuming Sphericity
##
##
                Sum Sq num Df Error SS den Df F value Pr(>F)
## (Intercept) 683527262 1 38774274 39 687.506 < 2.2e-16 ***
             17628370 2 64973788 78 10.581 8.593e-05 ***
## time
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
##
## Mauchly Tests for Sphericity
##
     Test statistic p-value
##
## time 0.72909 0.0024707
##
##
## Greenhouse-Geisser and Huvnh-Feldt Corrections
## for Departure from Sphericity
##
    GG eps Pr(>F[GG])
## time 0.78684 0.0003502 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
##
        HF eps Pr(>F[HF])
## time 0.8142253 0.0002922006
```

```
aov ez(id = "ID", dv = "nicotine conc", between="nationality",
        within = "time", data = df nicotine) %>% summary()
##
## Univariate Type III Repeated-Measures ANOVA Assuming Sphericity
##
##
                    Sum Sq num Df Error SS den Df F value Pr(>F)
## (Intercept) 674309848 1 31945156 38 802.1177 < 2.2e-16 ***
## nationality 6829118 1 31945156 38 8.1235 0.0070220 **
            13995826 2 61796338 76 8.6064 0.0004274 ***
## time
## nationality:time 3177450 2 61796338 76 1.9539 0.1487760
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##
## Mauchly Tests for Sphericity
##
##
               Test statistic p-value
## time
                   0.75397 0.0053836
## nationality:time 0.75397 0.0053836
##
##
## Greenhouse-Geisser and Huynh-Feldt Corrections
## for Departure from Sphericity
##
##
                 GG eps Pr(>F[GG])
                0.80255 0.001178 **
## time
## nationality:time 0.80255 0.158440
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

D-- (SE FIRE)