Day6 exercise solutions

Ali Movasati

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```
# Set global code chunk options
knitr::opts_chunk$set(warning = FALSE)

# load required libraries
library(skimr)
library(ggplot2)
library(ggpubr)
library(magrittr)
library(dplyr)
library(tibble)
library(lme4)
# define functions
`%notin%` <- Negate(`%in%`)
```

Problem 1

```
# read in the data
hearing <- read.table(file = "/Users/alimos313/Documents/studies/phd/university/courses/stat-modelling/</pre>
```

1.A)

```
# print descriptive
skim(hearing)
```

Table 1: Data summary

| Name | hearing |
|------------------------|---------|
| Number of rows | 96 |
| Number of columns | 3 |
| Column type frequency: | |
| character | 1 |
| numeric | 2 |
| Group variables | None |

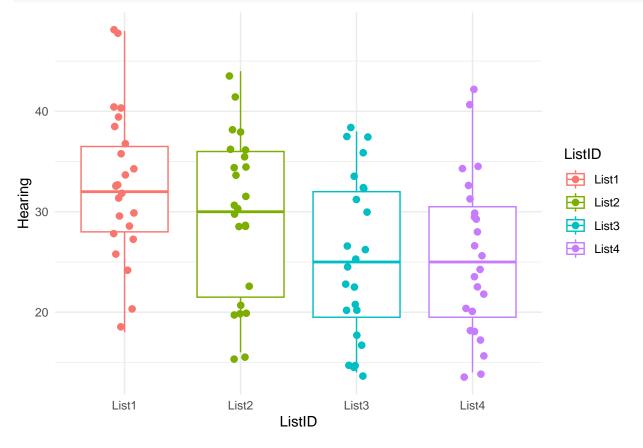
Variable type: character

| skim_variable | n_missing | $complete_rate$ | min | max | empty | n_unique | whitespace |
|---------------|-----------|------------------|-----|-----|-------|----------|------------|
| ListID | 0 | 1 | 5 | 5 | 0 | 4 | 0 |

Variable type: numeric

| skim_variable | n_missing | $complete_rate$ | mean | sd | p0 | p25 | p50 | p75 | p100 | hist |
|---------------|-----------|------------------|-------|---------------------|----|-------|------|-------|------|------|
| SubjectID | 0 | 1 | 12.50 | 6.96 | 1 | 6.75 | 12.5 | 18.25 | 24 | |
| Hearing | 0 | 1 | 28.31 | 8.37 | 14 | 20.00 | 30.0 | 34.00 | 48 | |

table(hearing\$ListID)



1.B)

```
lm_simple <- lm(Hearing ~ 1 + ListID, data = hearing)</pre>
sum_model_simple <- summary(lm_simple)</pre>
print(sum_model_simple)
##
## Call:
## lm(formula = Hearing ~ 1 + ListID, data = hearing)
## Residuals:
##
       Min
                 1Q
                      Median
                                   3Q
## -14.7500 -5.5833 -0.2083
                               6.3333 16.4167
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                32.750
                            1.612 20.315 < 2e-16 ***
## ListIDList2 -3.083
                            2.280 -1.352 0.17955
                -7.500
                            2.280 -3.290 0.00142 **
## ListIDList3
## ListIDList4 -7.167
                            2.280 -3.144 0.00225 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 7.898 on 92 degrees of freedom
## Multiple R-squared: 0.1382, Adjusted R-squared: 0.1101
## F-statistic: 4.919 on 3 and 92 DF, p-value: 0.00325
« comments »
```

Only 11.01% of variability in hearing measures are explained by different lists

We have enough evidence to state that the mean hearing score for List 3 and 4 are different than list 1, while for list 2 we cannot state that!

1.C)

```
# fit the mixed model
lm_mixed <- lmer(Hearing ~ 1 + ListID + (1|SubjectID), data = hearing)</pre>
sum_model_mixed <- summary(lm_mixed)</pre>
print(sum_model_mixed)
## Linear mixed model fit by REML ['lmerMod']
## Formula: Hearing ~ 1 + ListID + (1 | SubjectID)
##
      Data: hearing
##
## REML criterion at convergence: 635.4
## Scaled residuals:
        Min
                       Median
                                     3Q
                                              Max
                  1Q
## -1.86533 -0.56158 -0.01092 0.63222 2.69167
##
```

```
## Random effects:
                         Variance Std.Dev.
## Groups
             Name
                                  5.103
## SubjectID (Intercept) 26.04
## Residual
                         36.33
                                  6.027
## Number of obs: 96, groups: SubjectID, 24
##
## Fixed effects:
              Estimate Std. Error t value
##
                         1.612 20.315
## (Intercept) 32.750
## ListIDList2 -3.083
                            1.740 -1.772
## ListIDList3 -7.500
                            1.740 -4.311
## ListIDList4 -7.167
                            1.740 -4.119
## Correlation of Fixed Effects:
##
              (Intr) LsIDL2 LsIDL3
## ListIDList2 -0.540
## ListIDList3 -0.540 0.500
## ListIDList4 -0.540 0.500 0.500
# fit the model without ListID (null model)
lm_mixed_null <- lmer(Hearing ~ (1 | SubjectID), data = hearing)</pre>
sum_model_mixed_null <- summary(lm_mixed_null)</pre>
# Likelihood ratio test
anova(lm mixed null, lm mixed)
## refitting model(s) with ML (instead of REML)
## Data: hearing
## Models:
## lm_mixed_null: Hearing ~ (1 | SubjectID)
## lm_mixed: Hearing ~ 1 + ListID + (1 | SubjectID)
                               BIC logLik deviance Chisq Df Pr(>Chisq)
                      AIC
                npar
                   3 674.22 681.91 -334.11 668.22
## lm mixed null
## lm_mixed
                   6 657.70 673.09 -322.85 645.70 22.52 3 5.083e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
1.D)
```

Problem 2

```
# read in the data

termites <- read.table(file = "/Users/alimos313/Documents/studies/phd/university/courses/stat-modelling</pre>
```

- 2.A)
- 2.B)
- 2.C)
- 2.D)
- 2.E)
- 2.F)