PSYC 7720 Lab

Lab 9 Activity

Andrew Graves, Department of Psychology, University of Virginia

Directions:

- A. Download the mccarthyData.csv. This is hypothetical data from 12 children who took the McCarthy Scales of Children's Abilites (MSCA) test at 30, 36, 42, and 48 months. The DV is the age-normed general cognitive score from the MSCA.
- B. Answer the following questions and save the code you used in an R script.
- C. You have until the end of lab to complete.

```
library(tidyverse)
library(ez)
library(emmeans)
options(contrasts = c("contr.sum","contr.poly"))
theme_set(theme_classic())
```

Questions:

1. Convert the mccarthyData from wide format to long format. Name the first column *id*, the second column *month*, the third column *mcsa*, and the dataframe *long_data*. Print the head and the dimensions of the long data.

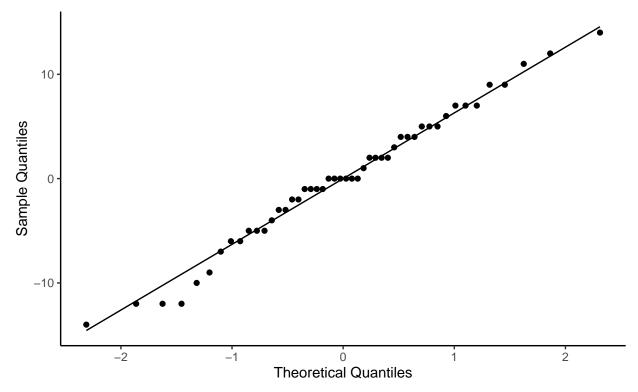
```
long data <- read csv("mccarthyData.csv") %>%
  rownames_to_column(var = "id") %>%
  pivot_longer(-id, names_to = "month", values_to = "mcsa") %>%
  mutate(id = factor(id),
         month = factor(month, ordered = TRUE))
head(long_data)
## # A tibble: 6 x 3
##
     id
           month
                     mcsa
##
     <fct> <ord>
                     <dbl>
## 1 1
           MONTHS30
                       108
## 2 1
           MONTHS36
                       96
## 3 1
           MONTHS42
                       110
## 4 1
           MONTHS48
                       122
## 5 2
           MONTHS30
                       103
## 6 2
           MONTHS36
                       117
dim(long_data)
```

[1] 48 3

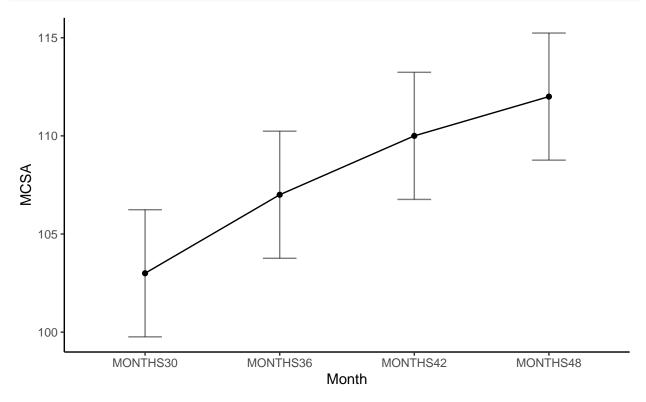
2. Run and interpret the results of an RM ANOVA using aov.

```
aov_mod <- aov(mcsa ~ month + Error(id/month), data = long_data)</pre>
summary(aov_mod)
##
## Error: id
##
            Df Sum Sq Mean Sq F value Pr(>F)
## Residuals 11 6624 602.2
##
## Error: id:month
            Df Sum Sq Mean Sq F value Pr(>F)
                 552 184.00
            3
                                3.027 0.0432 *
## Residuals 33
                2006 60.79
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
  • Month is a significant predictor of mcsa.
  3. Run and interpret the results of an RM ANOVA using ezANOVA.
ez_mod <- ezANOVA(data = long_data,</pre>
                 dv = mcsa,
                 within = month,
                 wid = id,
                 type = 3,
                 return_aov = TRUE,
                 detailed = TRUE
ez_mod
## $ANOVA
         Effect DFn DFd
                           SSn SSd
                                             F
                                                          p p<.05
## 1 (Intercept) 1 11 559872 6624 929.739130 5.586316e-12
                                                              * 0.98481975
          month 3 33
                           552 2006 3.026919 4.321863e-02
## 2
                                                              * 0.06011762
##
## $`Mauchly's Test for Sphericity`
##
   Effect
                   W
## 2 month 0.2426472 0.01771762
##
## $`Sphericity Corrections`
                         p[GG] p[GG]<.05
## Effect
                 GGe
                                            HFe
                                                        p[HF] p[HF]<.05
## 2 month 0.6095445 0.0747874
                                       0.7248502 0.06353773
##
## $aov
##
## Call:
## aov(formula = formula(aov_formula), data = data)
##
## Grand Mean: 108
##
## Stratum 1: id
##
## Terms:
##
                  Residuals
## Sum of Squares
                       6624
## Deg. of Freedom
                        11
```

```
##
## Residual standard error: 24.53939
##
## Stratum 2: id:month
##
## Terms:
##
                   month Residuals
## Sum of Squares
                     552
                               2006
## Deg. of Freedom
                        3
                                 33
##
## Residual standard error: 7.796658
## Estimated effects are balanced
# Check the residuals for normality
resid <- proj(ez_mod$aov)[[3]][, "Residuals"]</pre>
resid %>%
  tibble() %>%
  ggplot(aes(sample = `.`)) +
  geom_qq() +
  stat_qq_line() +
  labs(x = "Theoretical Quantiles",
       y = "Sample Quantiles")
```



```
adjust = "bonferroni"
  )
follow_up
##
    contrast estimate
                          SE df t.ratio p.value
##
    linear
                    30 10.1 33 2.980 0.0161
##
    quadratic
                    -2 4.5 33 -0.444
                                       1.0000
##
    cubic
                     0 10.1 33 0.000 1.0000
##
## P value adjustment: bonferroni method for 3 tests
ez_plot <- ezPlot(</pre>
  data = long_data,
  dv = mcsa,
  wid = id,
  within = month,
  x = month,
  x_lab = "Month",
  y_lab = "MCSA"
ez_plot
```



• After Greenhouse-Geisser correction due to violations of sphericity, *month* is not a significant predictor of *mcsa*. However, if we had planned a linear contrast across months, we could haved specifically tested the polynomial contrasts (or just the linear contrast if we had strong theory). In this case, after Bonferonni correction, there was a positive significant linear trend on *mcsa* across *months*.