Reanalysis of 17-DePascalis

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Reference

De Pascalis, V., Fracasso, F., & Corr, P. J. (2017). Personality and Augmenting/Reducing (A/R) in auditory event-related potentials (ERPs) during emotional visual stimulation. Scientific Reports, 7, 41588. https://doi.org/10.1038/srep41588

Notes from reading methods section

• Dependant variable: "N1/P2 complex" (n = 39)

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- Independent variables:
 - Emotion, 3 levels (negative, neutral, positive pictures)
 - Auditory intensity, 5 levels (59, 70, 79, 88, 96 dB SPL)
 - Recording Site, 3 levels (frontal Fz, central Cz, parietal Pz)
- Covariate: BIS (Behavioural Inhibition System) with RST-PQ questionnaire
- Design: 3-way 3x5x3 ANCOVA (all factors within)

Reading data

Data is loaded, reshaped if necessary, and factors are specified.

```
PATH = file.path(path.expand("~"), "Data", "ancova") # ancova project folder

raw = read_excel(file.path(PATH, "dataPrimaryStudies", "17-DePascalis", "BIS_cov_N1-P2peak_amplitude_r.

raw = as.data.frame(raw)

# correcting typo

colnames(raw)[1] = "Subject"
```

Reshaping data into long

```
N = 39
data = data.frame(subject = as.factor(rep(raw$Subject, 3*5*3)))
data$amplitude = c(as.matrix(raw[,2:46]))
data$emotion = as.factor(c(rep("negative", 5*3*N), rep("neutral", 5*3*N), rep("positive", 5*3*N)))
data$intensity = as.factor(rep(c(rep("level 1", 3*N), rep("level 2", 3*N), rep("level 3", 3*N), rep("location 3", N)), 3*5
data$bis = rep(raw$RST_BIS, 3*5*3)
summary(data)
```

```
##
       subject
                     amplitude
                                         emotion
                                                       intensity
                                                                           site
                                     negative:585
##
    S01
          : 45
                   Min.
                           : 2.322
                                                    level 1:351
                                                                   location 1:585
##
    S02
                   1st Qu.:10.778
                                     neutral:585
              45
                                                    level 2:351
                                                                   location 2:585
   S03
##
              45
                   Median :15.112
                                     positive:585
                                                    level 3:351
                                                                   location 3:585
## S04
              45
                   Mean
                           :17.944
                                                     level 4:351
##
    S05
              45
                   3rd Qu.:22.039
                                                    level 5:351
    S06
##
              45
                   Max.
                           :68.823
##
   (Other):1485
##
         bis
## Min.
           :33.00
  1st Qu.:46.00
## Median :53.00
## Mean
           :54.79
##
   3rd Qu.:64.00
##
  Max.
           :81.00
##
```

Running an initial ANCOVA

Descriptives

Dependent variable

We show descriptives plots and no tables as the factorial designs is rather complex (3x5x3).

Main effects

Averaging subjects across the other two levels

```
d.1 = aggregate(amplitude ~ bis + emotion*subject, data = data, FUN = mean)
d.2 = aggregate(amplitude ~ bis + intensity*subject, data = data, FUN = mean)
d.3 = aggregate(amplitude ~ bis + site*subject, data = data, FUN = mean)
p1 = ggplot(d.1, aes(y=amplitude, x=emotion, color=emotion)) +
  geom_boxplot() +
  geom_point(position = position_jitter(width = 0.15, height = 0)) +
  theme minimal()
p2 = ggplot(d.2, aes(y=amplitude, x=intensity, color=intensity)) +
  geom_boxplot() +
  geom_point(position = position_jitter(width = 0.15, height = 0)) +
  theme_minimal()
p3 = ggplot(d.3, aes(y=amplitude, x=site, color=site)) +
  geom_point(position = position_jitter(width = 0.15, height = 0)) +
  theme_minimal()
plot_grid(p1, p2, p3, nrow = 1, ncol = 3)
                                                          intensity
                        emotion
                                                                                          site
                                  30
                                                                                          location 2
                                                          level 4
                                                                                          location 3
                                                          level 5
     negative
           neutral
                                                                       location 1
                                                                            location 2 location 3
          emotion
                                            intensity
                                                                              site
```

Interaction effects (two-way)

Averaging subjects across along the third factor to see the two-way interactions. Plots show means (SD as error bar)

```
d.A = aggregate(amplitude ~ emotion*intensity, data = data, FUN = mean)
d.B = aggregate(amplitude ~ emotion*site, data = data, FUN = mean)
d.C = aggregate(amplitude ~ intensity*site, data = data, FUN = mean)

# get SD
d.A$sd = aggregate(amplitude ~ emotion*intensity, data = data, FUN = sd)[,3]
d.B$sd = aggregate(amplitude ~ emotion*site, data = data, FUN = sd)[,3]
d.C$sd = aggregate(amplitude ~ intensity*site, data = data, FUN = sd)[,3]
```

```
pA = ggplot(d.A, aes(y=amplitude, x=intensity, group=emotion, color=emotion)) +
  geom_errorbar(aes(ymin=amplitude-sd, ymax=amplitude+sd), width=.1) +
  geom_line() + geom_point() +
  theme minimal()
pB = ggplot(d.B, aes(y=amplitude, x=site, group=emotion, color=emotion)) +
  geom_errorbar(aes(ymin=amplitude-sd, ymax=amplitude+sd), width=.1) +
  geom_line() + geom_point() +
  theme_minimal()
pC = ggplot(d.C, aes(y=amplitude, x=intensity, group=site, color=site)) +
  geom_errorbar(aes(ymin=amplitude-sd, ymax=amplitude+sd), width=.1) +
  geom_line() + geom_point() +
  theme_minimal()
plot_grid(pA, pB, pC, nrow = 1, ncol = 3)
 40
                          emotion
                                      30
                                                               emotion
                                                                                                   site
amplitude 30

    negative

                                                                           30

    location 1

                             negative
                                      20
                             positive
                                                                                                      location 3
  10
                                                                           10
                                                                              level 1 level 2 level 3 level 4 level 5
    level 1 level 2 level 3 level 4 level 5
                                          location 1 location 2 location 3
           intensity
                                                                                    intensity
```

Covariate(s)

```
Descriptives, see Table 1
sprintf("BIS: %0.1f (%0.1f)", mean(data$bis[1:N]), sd(data$bis[1:N]))
## [1] "BIS: 54.8 (11.8)"
```

Main analysis ANCOVA

Error: subject:emotion

```
# Orthogonal contrasts
# contrasts(data$group) = contr.helmert(2)
fit.ancova = aov(amplitude ~ bis + emotion*intensity*site + Error(subject/(emotion*intensity*site)), da
result = summary(fit.ancova) # Type I
#result = Anova(fit.ancova, type=3) # Type III
print(result)
##
## Error: subject
##
            Df Sum Sq Mean Sq F value Pr(>F)
                 9585
                          9585
                                6.816 0.013 *
## bis
## Residuals 37 52027
                          1406
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
```

```
Df Sum Sq Mean Sq F value Pr(>F)
                 423 211.74
                             4.533 0.0138 *
## emotion
                3550 46.71
## Residuals 76
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Error: subject:intensity
            Df Sum Sq Mean Sq F value Pr(>F)
## intensity 4 71397 17849
                              96.29 <2e-16 ***
## Residuals 152 28176
                         185
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Error: subject:site
           Df Sum Sq Mean Sq F value Pr(>F)
## site
            2
               6710
                      3355
                              68.75 <2e-16 ***
                3709
## Residuals 76
                         49
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Error: subject:emotion:intensity
                    Df Sum Sq Mean Sq F value Pr(>F)
## emotion:intensity 8 2252 281.51
                                      8.319 3.3e-10 ***
## Residuals
                   304 10288
                              33.84
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Error: subject:emotion:site
               Df Sum Sq Mean Sq F value Pr(>F)
## emotion:site 4 16.75 4.188
                                 2.682 0.0337 *
## Residuals
            152 237.31
                          1.561
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Error: subject:intensity:site
                 Df Sum Sq Mean Sq F value Pr(>F)
## intensity:site
                    1871 233.86 50.06 <2e-16 ***
## Residuals
                304
                    1420
                             4.67
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Error: subject:emotion:intensity:site
                         Df Sum Sq Mean Sq F value Pr(>F)
## emotion:intensity:site 16
                             22.8 1.426 1.022 0.431
## Residuals
                        608 848.8
                                    1.396
```

Comparing ANCOVA in original study with reanalysis

Independent variable

```
tab.IV = rbind(stats.orig.IV, stats.rep.IV)
rownames(tab.IV) = c("original Study", "reanalysis")
```

print(t(tab.IV)) original Study reanalysis ## Fvalue NA## df1 NA NA ## df2 NA NA## pvalue NA NA ## MD NANA ## lowerCI NA NA ## upperCI NA NA

Covariate

```
tab.CV = rbind(stats.orig.CV.bis, stats.rep.CV.bis)
rownames(tab.CV) = c("original Study", "reanalysis")
print(t(tab.CV))
##
           original Study reanalysis
           "6.82"
                           "6.82"
## Fvalue
           "1"
                           "1"
## df1
           "37"
                           "37"
## df2
           "0.013"
                           "0.013"
## pvalue
## MD
           NA
                           NA
## lowerCI NA
                           NA
## upperCI NA
                           NA
```

Assumptions

1. Homogeneity of variance

- ANOVA/ANCOVA is fairly robust in terms of the error rate when sample sizes are equal.
- When groups with larger sample sizes have larger variances than the groups with smaller sample sizes, the resulting F-ratio tends to be conservative. That is, it's more likely to produce a non-significant result when a genuine difference does exist in the population.
- Conversely, when the groups with larger sample sizes have smaller variances than the groups with smaller samples sizes, the resulting F-ratio tends to be liberal and can inflate the false positive rate.

```
tmp = tapply(d.1$amplitude, d.1$emotion, sd)
tab = sprintf("%.2f", tmp)
names(tab) = names(tmp)
stats = leveneTest(amplitude ~ emotion, data = d.1)
tab[length(tab)+1] = formatPval(stats$`Pr(>F)`[1])
names(tab)[length(tab)] = "Levene's p-value"
print(tab)
##
           negative
                             neutral
                                              positive Levene's p-value
             "5.61"
##
                              "5.47"
                                                "7.27"
                                                                  "0.12"
tmp = tapply(d.2$amplitude, d.2$intensity, sd)
tab = sprintf("%.2f", tmp)
names(tab) = names(tmp)
stats = leveneTest(amplitude ~ intensity, data = d.2)
```

```
tab[length(tab)+1] = formatPval(stats$`Pr(>F)`[1])
names(tab)[length(tab)] = "Levene's p-value"
print(tab)
##
            level 1
                             level 2
                                               level 3
                                                                 level 4
                                                "5.13"
                                                                  "9.26"
##
             "2.92"
                               "3.87"
##
            level 5 Levene's p-value
            "11.27"
                          "< 0.0001"
##
tmp = tapply(d.3$amplitude, d.3$site, sd)
tab = sprintf("%.2f", tmp)
names(tab) = names(tmp)
stats = leveneTest(amplitude ~ site, data = d.3)
tab[length(tab)+1] = formatPval(stats$`Pr(>F)`[1])
names(tab)[length(tab)] = "Levene's p-value"
print(tab)
##
         location 1
                          location 2
                                            location 3 Levene's p-value
##
             "5.63"
                               "7.40"
                                                "5.30"
                                                                  "0.19"
```

2. Independence between covariate and IV

When the covariate and the experimental effect (independent variable) are not independent the treatment effect is obscured, spurious treatment effects can arise and the interpretation of the ANCOVA is seriously compromised.

We test whether our groups differ on the CV. If the groups do not significantly differ then is appropriate to use the covariate.

Not applicable for a within design as the covariate does not change across levels. So this assumption does not need testing.

3. Homogeneity of regression slopes

##

Error: subject:emotion

• We test the interaction between the IV and the CV

Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1

```
Df Sum Sq Mean Sq F value Pr(>F)
               2 28.23 14.116
                                4.944 0.00965 **
## emotion
## bis:emotion 2 25.42 12.708
                                 4.451 0.01495 *
             74 211.27
## Residuals
                          2.855
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
fit.hrs = aov(amplitude ~ bis*intensity + Error(subject/intensity), data = d.2)
summary(fit.hrs)
##
## Error: subject
            Df Sum Sq Mean Sq F value Pr(>F)
                 1065 1065.0
                                6.816 0.013 *
## bis
             1
## Residuals 37
                 5781
                        156.2
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Error: subject:intensity
##
                 Df Sum Sq Mean Sq F value Pr(>F)
## intensity
                      7933 1983.2 118.878 < 2e-16 ***
## bis:intensity
                  4
                       662
                             165.4
                                   9.914 3.9e-07 ***
## Residuals
                148
                      2469
                              16.7
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
fit.hrs = aov(amplitude ~ bis*site + Error(subject/site), data = d.3)
summary(fit.hrs)
##
## Error: subject
            Df Sum Sq Mean Sq F value Pr(>F)
## bis
             1
                  639
                        639.0
                                6.816 0.013 *
                         93.7
## Residuals 37
                 3468
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Error: subject:site
            Df Sum Sq Mean Sq F value Pr(>F)
## site
             2 447.4 223.68
                                68.31 <2e-16 ***
## bis:site
            2
                  5.0
                         2.49
                                 0.76 0.471
## Residuals 74 242.3
                         3.27
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
p1 = ggplot(d.1, aes(y=amplitude, x=bis, color=emotion, shape=emotion)) +
  geom_point() +
  geom_smooth(formula = y ~ x,method=lm, se=FALSE, fullrange=TRUE) +
 theme_minimal() +
 theme(legend.title = element_blank())
p2 = ggplot(d.2, aes(y=amplitude, x=bis, color=intensity, shape=intensity)) +
  geom_point() +
  geom_smooth(formula = y ~ x,method=lm, se=FALSE, fullrange=TRUE) +
 theme_minimal() +
 theme(legend.title = element_blank())
```

Notes

- Amplitudes were not normal, log transform could have had helped.
- The first reported ANCOVA result was reproduced: effect of the covariate BIS
- BIS was used as continuous covariate which showed a statistically sign. effect. on amplitude. However, authors reported an effect of low vs. high BIS participants on amplitude (categorical), with lower amplitudes in high BIS participants based on Figure 2.
- Not clear how low vs. high BIS was determined (which threshold was used?).
- Only parts of ANCOVA results were reported. There were altogether three main effects and three two-way interactions that were not all reported.
- Generally, there is a mismatch between the ANCOVA model specified and the result reported (BIS treated continuous or categorical?).
- Assumptions of homogeneity of regression slope was not met for intensity and emotion: the relationship between BIS and amplitude differed across levels of intensity and levels of emotion. The authors reported both, so this is fine.
- Assumption of independence between COV and IV cannot be tested in within-only designs.

Data was analyzed according to recommendations by Field, Miles, & Field (2012).