# Reanalysis of 24-Fujiwara

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01/06/2020

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### Reference

Fujiwara, E. (2018). Looking at the eyes interferes with facial emotion recognition in alexithymia. Journal of Abnormal Psychology, 127(6), 571–577. https://doi.org/10.1037/abn0000361

# Notes from reading methods section

This study assessed the two Alexithymic groups, high and low, with the individuals' ability to identify clear and ambiguous blends of emotions in faces and their underlying visual attention patterns. Using the Toronto Alexithymia Scale, students with high alexithymia (HA, n = 73) or low alexithymia (LA, n = 76) were enrolled in this study.

• Dependant variable: deviance score

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- Independent variables:
  - 2 levels ambiguity (clear vs. ambiguous emotional blends in the target face)
  - 6 levels of emotion (anger, disgust, fear, happiness, sadness, surprise is dominant emotion)
  - 2 groups (HA, LA; high and low Alexithymia score)
- Covariate:
  - DASS-21 (Depression, Anxiety and Stress total score)
  - Maybe log transformed response times were also used as covariate
- Design: 2 (w) x 6 (w) x 2 (b) mixed ANCOVA (w within; b between)

### Reading data

Data is loaded, reshaped into long form, and factors are specified.

```
##
          id
                                                                           TAS_group
                          dev
                                                               amb
##
    1
           :
              12
                    Min.
                            : 1.426
                                      happiness:298
                                                       ambigious:894
                                                                         Min.
                                                                                :1.00
                                                                         1st Qu.:1.00
##
    2
              12
                                                                 :894
                    1st Qu.: 8.822
                                      anger
                                                :298
                                                        clear
                                                :298
##
              12
                    Median :11.546
                                      disgust
                                                                         Median:1.00
##
              12
                    Mean
                            :13.562
                                      fear
                                                :298
                                                                         Mean
                                                                                :1.98
##
    5
              12
                    3rd Qu.:17.497
                                      surprise :298
                                                                         3rd Qu.:3.00
##
              12
                    Max.
                            :75.620
                                      sadness :298
                                                                         Max.
                                                                                :3.00
    (Other):1716
##
##
       DASS tot
                         log_lat
                                                     group
           : 2.00
                                       High Alexithymia:876
##
                      Min.
                              :3.343
    1st Qu.: 26.00
                      1st Qu.:3.626
                                       Low Alexithymia:912
  Median : 36.00
                      Median :3.735
##
           : 44.95
                              :3.730
##
    Mean
                      Mean
    3rd Qu.: 62.00
##
                      3rd Qu.:3.845
##
   Max.
            :120.00
                      Max.
                              :4.143
##
```

## Descriptives

#### Dependant variable

Number of samples and mean (SD) in levels of the independent variables. We reproduce Table 3 and Figure 2A of the study.

```
d.1 = aggregate(dev ~ DASS_tot + log_lat + id*group, data = data, FUN = mean)
d.2 = aggregate(dev ~ DASS_tot + log_lat + id*emo, data = data, FUN = mean)
d.3 = aggregate(dev ~ DASS_tot + log_lat + id*amb, data = data, FUN = mean)

p1 = ggplot(d.1, aes(y=dev, x=group, color=group)) +
    geom_boxplot() +
    geom_point(position = position_jitter(width = 0.15, height = 0)) +
    theme_minimal() + theme(axis.text.x = element_blank())

p2 = ggplot(d.2, aes(y=dev, x=emo, color=emo)) +
    geom_point(position = position_jitter(width = 0.15, height = 0)) +
    theme_minimal() + theme(axis.text.x = element_blank())
```

```
p3 = ggplot(d.3, aes(y=dev, x=amb, color=amb)) +
  geom_boxplot() +
  geom_point(position = position_jitter(width = 0.15, height = 0)) +
 theme_minimal() + theme(axis.text.x = element_blank())
plot_grid(p1, p2, p3, nrow = 1, ncol = 3)
                                                            happiness
 30
                                                                                               amb
                                                            anger
                                                                        30
≱ 20
                                                                       de√
                     High Alexithymia
                                   <u>§</u> 30
                                                            disgust
                                                                                               ambigious
                     Low Alexithymia
                                                            ear fear
                                                                                               clear
                                                            surprise
```

Averaging subjects across along the third factor to see the two-way interactions. Plots show means (95%-CI as error bar)

```
d.A = aggregate(dev ~ group*emo, data = data, FUN = mean)
d.B = aggregate(dev ~ group*amb, data = data, FUN = mean)
d.C = aggregate(dev ~ emo*amb, data = data, FUN = mean)
# get SD
d.A$sd = aggregate(dev ~ group*emo, data = data, FUN = sd)[,3]
d.A\$se = d.A\$sd/sqrt(rep(c(73, 76), 6))
d.B$sd = aggregate(dev ~ group*amb, data = data, FUN = sd)[,3]
d.B\$se = d.B\$sd/sqrt(rep(c(73, 76), 2))
d.C$sd = aggregate(dev ~ emo*amb, data = data, FUN = sd)[,3]
d.C\$se = d.C\$sd/sqrt(rep(N, 12))
pA = ggplot(d.A, aes(y=dev, x=emo, group=group, color=group)) +
  geom_errorbar(aes(ymin=dev-1.96*se, ymax=dev+1.96*se), width=.2) +
  geom_line() + geom_point() +
  theme_minimal() + theme(axis.text.x = element_text(angle = 20) )
pB = ggplot(d.B, aes(y=dev, x=amb, group=group, color=group)) +
  geom errorbar(aes(ymin=dev-1.96*se, ymax=dev+1.96*se), width=.2) +
  geom_line() + geom_point() +
  theme minimal()
pC = ggplot(d.C, aes(y=dev, x=emo, group=amb, color=amb)) +
  geom_errorbar(aes(ymin=dev-1.96*se, ymax=dev+1.96*se), width=.2) +
  geom_line() + geom_point() +
  theme_minimal() + theme(axis.text.x = element_text(angle = 20))
plot_grid(pA, pB, pC, nrow = 1, ncol = 3)
                                                                   20
 16
                                                                                        amb
                   group
                                                    group

    High Alexithymia

                                  14
                                                                                         ambigious

    High Alexithymia

                      Low Alexithymia
                                                                                           clear
                                  12
                                                       Low Alexithymia
                                  10
                                     ambigious
                                             clea
         emo
                                                                            emo
```

#### Covariate(s)

The two covariates are DASS\_tot scores and log of time.

```
p1 = ggplot(d.1, aes(y=DASS_tot, x=group, color=group)) +
  geom_boxplot() +
  geom_point(position = position_jitter(width = 0.15, height = 0)) +
theme minimal() + theme(axis.text.x = element blank())
p2 = ggplot(d.1, aes(y=log_lat, x=group, color=group)) +
  geom_boxplot() +
  geom_point(position = position_jitter(width = 0.15, height = 0)) +
theme_minimal() + theme(axis.text.x = element_blank())
plot_grid(p1, p2, nrow = 1, ncol = 2)
  125
  100
                                                   4.00
                               group
                                                                                group
   75
                                                 <u>a</u>
                                                 3.75
0
                               High Alexithymia
                                                                                 High Alexithymia
   50
                               Low Alexithymia
                                                                                 Low Alexithymia
   25
                                                   3.50
    0
              group
                                                                group
```

### Main analysis ANCOVA

Independent variable: TAS\_group (Between Group)

```
# Orthogonal contrasts
contrasts(data$group) = contr.helmert(2)
contrasts(data$amb) = contr.helmert(2)
contrasts(data$emo) = contr.helmert(6)
fit.ancova = aov(dev ~ log_lat + DASS_tot + (group*emo*amb) + Error(id/(emo*amb)) + group,
                 data = data)
result = summary(fit.ancova) # Type I
print(result)
##
## Error: id
              Df Sum Sq Mean Sq F value Pr(>F)
##
                                  0.010 0.9201
## log_lat
                      2
                            1.5
               1
                    437
                          437.0
                                  2.852 0.0934
## DASS tot
               1
                                  0.613 0.4349
## group
               1
                     94
                           93.9
## Residuals 145
                  22213
                          153.2
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Error: id:emo
##
              Df Sum Sq Mean Sq F value Pr(>F)
## emo
                   4506
                          901.3 45.282 <2e-16 ***
```

```
## group:emo
                           20.3
                                  1.019 0.405
             5
                    101
## Residuals 735 14629
                           19.9
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Error: id:amb
##
              Df Sum Sq Mean Sq F value Pr(>F)
## amb
                 14048
                          14048 130.571 <2e-16 ***
## group:amb
              1
                    424
                            424
                                  3.936 0.0491 *
                            108
## Residuals 147 15816
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Error: id:emo:amb
##
                  Df Sum Sq Mean Sq F value Pr(>F)
## emo:amb
                   5
                       2042
                              408.3 18.863 <2e-16 ***
                   5
                                      0.265 0.932
## group:emo:amb
                         29
                               5.7
## Residuals
                 735
                     15911
                               21.6
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
fit2.ancova = ezANOVA(dv = .(dev), wid = .(id), between = .(group), within = .(emo, amb),
                     within_covariates = .(DASS_tot, log_lat), type=3, detailed=TRUE, data = data)
## Warning: Data is unbalanced (unequal N per group). Make sure you specified a
## well-considered value for the type argument to ezANOVA().
## Warning: Implementation of ANCOVA in this version of ez is experimental and
## not yet fully validated. Also, note that ANCOVA is intended purely as a tool
## to increase statistical power; ANCOVA can not eliminate confounds in the data.
## Specifically, covariates should: (1) be uncorrelated with other predictors and
## (2) should have effects on the DV that are independent of other predictors.
## Failure to meet these conditions may dramatically increase the rate of false-
## positives.
## Warning: Covariate"DASS_tot" is numeric and will therefore be fit to a linear
## effect.
## Warning: Covariate"log_lat" is numeric and will therefore be fit to a linear
## effect.
print(fit2.ancova)
## $ANOVA
                                                                        p p<.05
##
            Effect DFn DFd
                                    SSn
                                             SSd
       (Intercept)
## 1
                     1 147 329109.61359 22453.25 2154.659410 1.045880e-89
## 2
            group
                     1 147
                              291.72820 22453.25
                                                    1.909926 1.690680e-01
## 3
                     5 735
                             4511.33271 14629.06
                                                   45.332085 7.777113e-41
               emo
## 5
              amb
                     1 147 13944.48409 15815.77 129.607304 6.306088e-22
## 4
                     5 735
                              101.37071 14629.06
                                                    1.018623 4.054881e-01
         group:emo
## 6
        group:amb
                    1 147
                              423.50614 15815.77
                                                    3.936287 4.911711e-02
## 7
                    5 735
                            2035.47814 15910.78
                                                   18.805826 1.283640e-17
          emo:amb
                     5 735
                               28.65974 15910.78
                                                   0.264788 9.322917e-01
## 8 group:emo:amb
##
             ges
## 1 0.827077994
## 2 0.004221790
## 3 0.061529198
```

```
## 5 0.168506591
## 4 0.001471055
## 6 0.006117170
## 7 0.028731698
## 8 0.000416339
##
## $`Mauchly's Test for Sphericity`
                                         p p<.05
##
            Effect
## 3
               emo 0.8032470 4.294652e-03
## 4
         group:emo 0.8032470 4.294652e-03
           emo:amb 0.5848216 7.154692e-11
## 8 group:emo:amb 0.5848216 7.154692e-11
## $`Sphericity Corrections`
##
                                    p[GG] p[GG]<.05
            Effect
                         GGe
                                                           HFe
                                                                       p[HF]
## 3
               emo 0.9180654 1.042569e-37
                                                   * 0.9512955 5.618448e-39
## 4
                                                     0.9512955 4.036846e-01
         group:emo 0.9180654 4.023647e-01
           emo:amb 0.8399564 3.577157e-15
                                                   * 0.8677406 1.344884e-15
## 8 group:emo:amb 0.8399564 9.080457e-01
                                                     0.8677406 9.128726e-01
    p[HF]<.05
## 3
## 4
## 7
## 8
```

### Comparing ANCOVA in original study with reanalysis

#### Independant variable

```
Main effect ambiguity
```

```
tab.IV = rbind(stats.orig.IV.amb, stats.rep.IV.amb, stats.rep2.IV.amb)
rownames(tab.IV) = c("original Study", "reanalysis type I SS", "reanalysis type III SS")
print(t(tab.IV))
          original Study reanalysis type I SS reanalysis type III SS
## Fvalue "4.67"
                          "130.57"
                                               "129.61"
          "1"
                          "1"
                                               "1"
## df1
          "145"
                         "147"
                                               "147"
## df2
                          "< 0.0001"
                                               "< 0.0001"
## pvalue "0.03"
Interaction ambiguity x emotion
tab.IV = rbind(stats.orig.IV.emoXamb, stats.rep.IV.emoXamb, stats.rep2.IV.emoXamb)
rownames(tab.IV) = c("original Study", "reanalysis type I SS", "reanalysis type III SS")
print(t(tab.IV))
          original Study reanalysis type I SS reanalysis type III SS
                          "18.86"
                                               "18.81"
## Fvalue "2.61"
## df1
          "4.21"
                          "5.00"
                                               "5.00"
## df2
          "610.65"
                          "735.00"
                                               "735.00"
## pvalue "0.03"
                          "< 0.0001"
                                               "< 0.0001"
```

### 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.

```
tapply(d.1$dev, d.1$group, sd)

## High Alexithymia Low Alexithymia
## 4.218945 2.803665

leveneTest(dev ~ group, data = d.1)

## Levene's Test for Homogeneity of Variance (center = median)
## Df F value Pr(>F)
## group 1 1.5935 0.2088
## 147
```

#### 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.

```
fit.cv1 = aov(DASS_tot ~ group, data = d.1)
summary(fit.cv1)
##
                Df Sum Sq Mean Sq F value
                                           Pr(>F)
                           14094
                                   27.13 6.31e-07 ***
## group
                   14094
## Residuals
               147
                   76366
                             519
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
fit.cv2 = aov(log_lat ~ group, data = d.1)
summary(fit.cv2)
##
                Df Sum Sq Mean Sq F value Pr(>F)
## group
                 1 0.007 0.007117
                                    0.303 0.583
## Residuals
              147 3.454 0.023496
```

#### 3. Homogeneity of regression slopes

```
fit.hrs = aov(dev ~ DASS_tot*group, data = d.1)
Anova(fit.hrs, type = "III")

## Anova Table (Type III tests)
##
```

```
## Response: dev
##
                  Sum Sq Df F value Pr(>F)
## (Intercept)
                 2149.18
                           1 168.3272 <2e-16 ***
                               0.9438 0.3329
## DASS_tot
                   12.05
## group
                    1.71
                           1
                               0.1339 0.7149
                    0.00
                               0.0002 0.9881
## DASS_tot:group
                           1
## Residuals
                 1851.34 145
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
ggplot(d.1, aes(y=dev, x=DASS_tot, color=group, shape=group)) +
  geom_point() +
 geom_smooth(formula = y ~ x,method=lm, se=FALSE, fullrange=TRUE) +
 theme minimal() +
 theme(legend.title = element_blank())
  30
                               High Alexithymia
  20
                                Low Alexithymia
  10
        25
           50 75 100 125
     0
          DASS tot
fit.hrs = aov(dev ~ DASS_tot*emo + Error(id/emo), data = d.2)
summary(fit.hrs)
##
## Error: id
             Df Sum Sq Mean Sq F value Pr(>F)
## DASS_tot
             1
                   218 218.20
                                 2.876 0.0921 .
## Residuals 147 11154
                        75.88
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Error: id:emo
##
                Df Sum Sq Mean Sq F value Pr(>F)
## emo
                 5
                     2253
                            450.6 45.235 <2e-16 ***
                 5
                              8.6
                                    0.864 0.505
## DASS_tot:emo
                       43
## Residuals
               735
                     7322
                             10.0
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
ggplot(d.2, aes(y=dev, x=DASS_tot, color=emo, shape=emo)) +
 geom_point() +
  geom_smooth(formula = y ~ x,method=lm, se=FALSE, fullrange=TRUE) +
 theme_minimal() +
 theme(legend.title = element_blank())
```

```
happiness

anger

disgust

fear

surprise

sadness

DASS_tot

happiness

happiness

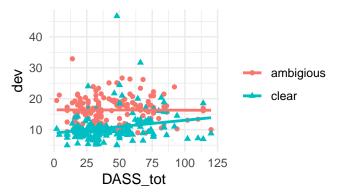
anger

disgust

Fear
```

```
fit.hrs = aov(dev ~ DASS_tot*amb + Error(id/amb), data = d.3)
summary(fit.hrs)
```

```
##
## Error: id
              Df Sum Sq Mean Sq F value Pr(>F)
##
## DASS_tot
                     73
                          72.73
                                  2.876 0.0921 .
## Residuals 147
                   3718
                          25.29
##
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
##
## Error: id:amb
##
                 Df Sum Sq Mean Sq F value Pr(>F)
                  1 2341.4 2341.4 131.132 <2e-16 ***
## amb
## DASS_tot:amb
                  1
                      81.8
                              81.8
                                     4.584 0.0339 *
## Residuals
                147 2624.7
                              17.9
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
ggplot(d.3, aes(y=dev, x=DASS_tot, color=amb, shape=amb)) +
 geom_point() +
  geom_smooth(formula = y ~ x,method=lm, se=FALSE, fullrange=TRUE) +
 theme_minimal() +
  theme(legend.title = element_blank())
```



### Notes

- Log transform of outcome could have been considered.
- Not all main effects and interactions were reported.

- Could not numerically reproduce the two reported results, but conclusions remain the same.
- Homogeneity of variances was met for groups.
- Independence between covariate DASS\_tot and the IV group was not met.
- Homogeneity of regression slopes was mostly met, i.e. for group and emotion, but not across levels of ambiguity (but only weak evidence of violation).

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