

Prediction First Pass

lib

11/05/2020

Importing the RT data sets we will be working with.

Below we are importing each data set and modifying the participant numbers to apply to the appropriate person. As we have had to extract the data in eyedry in batches of ~20 per extraction the participant numbers are not related to the actual participants. Here we are fixing this as well as importing the data.

```
#import the data set batch 1
FP_ED_batch1_corr <- read_csv("Duncans-Grant-master/FP_ED/FP_ED_batch1_corr.csv")
```

```
## Parsed with column specification:
## cols(
##   seq = col_double(),
##   subj = col_double(),
##   item = col_double(),
##   cond = col_double(),
##   R1 = col_double(),
##   R2 = col_double(),
##   R3 = col_double(),
##   R4 = col_double(),
##   R5 = col_double(),
##   R6 = col_double()
## )
```

#Rename the participant numbers in the batches back to their original participant numbers.

```
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 20] <-"38"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 19] <-"36"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 18] <-"35"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 17] <-"34"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 16] <-"32"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 15] <-"30"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 14] <-"28"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 13] <-"26"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 12] <-"24"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 11] <-"22"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 10] <-"20"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 9] <-"18"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 8] <-"16"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 7] <-"14"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 6] <-"12"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 5] <-"10"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 4] <-"8"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 3] <-"6"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 2] <-"4"  
FP_ED_batch1_corr$subj[FP_ED_batch1_corr$subj == 1] <-"2"
```

#import the data set batch 2B

```
FP_ED_batch2B_corr <- read_csv("Duncans-Grant-master/FP_ED/FP_ED_batch2B_corr.csv"  
)
```

Parsed with column specification:

```
## cols(  
##   seq = col_double(),  
##   subj = col_double(),  
##   item = col_double(),  
##   cond = col_double(),  
##   R1 = col_double(),  
##   R2 = col_double(),  
##   R3 = col_double(),  
##   R4 = col_double(),  
##   R5 = col_double(),  
##   R6 = col_double()  
## )
```

#Rename the participant numbers in the batches back to their original participant numbers.

```
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 19] <-"71"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 18] <-"70"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 17] <-"69"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 16] <-"68"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 15] <-"66"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 14] <-"64"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 13] <-"62"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 12] <-"60"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 11] <-"58"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 10] <-"56"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 9] <-"54"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 8] <-"50"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 7] <-"48"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 6] <-"46"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 5] <-"44"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 4] <-"43"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 3] <-"42"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 2] <-"40"  
FP_ED_batch2B_corr$subj[FP_ED_batch2B_corr$subj == 1] <-"39"
```

#import the data set batch 3

```
FP_ED_batch3_corr <- read_csv("Duncans-Grant-master/FP_ED/FP_ED_batch3_corr.csv")
```

Parsed with column specification:

```
## cols(  
##   seq = col_double(),  
##   subj = col_double(),  
##   item = col_double(),  
##   cond = col_double(),  
##   R1 = col_double(),  
##   R2 = col_double(),  
##   R3 = col_double(),  
##   R4 = col_double(),  
##   R5 = col_double(),  
##   R6 = col_double()  
## )
```

```
#Rename the participant numbers in the batches back to their original participant numbers.
```

```
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 21] <-"99"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 20] <-"95"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 19] <-"92"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 18] <-"91"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 17] <-"89"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 16] <-"87"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 15] <-"86"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 14] <-"85"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 13] <-"84"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 12] <-"83"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 11] <-"82"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 10] <-"81"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 9] <-"80"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 8] <-"79"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 7] <-"78"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 6] <-"77"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 5] <-"76"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 4] <-"75"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 3] <-"74"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 2] <-"73"  
FP_ED_batch3_corr$subj[FP_ED_batch3_corr$subj == 1] <-"72"
```

```
#import the data set batch 4
```

```
FP_ED_batch4_error <- read_csv("Duncans-Grant-master/FP_ED/FP_ED_batch4_error.csv"  
)
```

```
## Parsed with column specification:
```

```
## cols(  
##   seq = col_double(),  
##   subj = col_double(),  
##   item = col_double(),  
##   cond = col_double(),  
##   R1 = col_double(),  
##   R2 = col_double(),  
##   R3 = col_double(),  
##   R4 = col_double(),  
##   R5 = col_double(),  
##   R6 = col_double()  
## )
```

#Rename the participant numbers in the batches back to their original participant numbers.

```
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 20] <-"41"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 19] <-"37"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 18] <-"35"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 17] <-"33"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 16] <-"31"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 15] <-"29"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 14] <-"27"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 13] <-"25"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 12] <-"23"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 11] <-"21"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 10] <-"19"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 9] <-"17"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 8] <-"15"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 7] <-"13"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 6] <-"11"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 5] <-"9"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 4] <-"7"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 3] <-"5"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 2] <-"3"  
FP_ED_batch4_error$subj[FP_ED_batch4_error$subj == 1] <-"1"
```

#import the data set batch 5

```
FP_ED_batch5_error <- read_csv("Duncans-Grant-master/FP_ED/FP_ED_batch5_error.csv"  
)
```

Parsed with column specification:

```
## cols(  
##   seq = col_double(),  
##   subj = col_double(),  
##   item = col_double(),  
##   cond = col_double(),  
##   R1 = col_double(),  
##   R2 = col_double(),  
##   R3 = col_double(),  
##   R4 = col_double(),  
##   R5 = col_double(),  
##   R6 = col_double()  
## )
```

```
#Rename the participant numbers in the batches back to their original participant numbers.
```

```
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 19] <-"98"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 18] <-"97"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 17] <-"96"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 16] <-"94"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 15] <-"93"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 14] <-"90"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 13] <-"88"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 12] <-"67"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 11] <-"65"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 10] <-"63"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 9] <-"61"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 8] <-"59"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 7] <-"57"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 6] <-"55"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 5] <-"53"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 4] <-"51"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 3] <-"49"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 2] <-"47"  
FP_ED_batch5_error$subj[FP_ED_batch5_error$subj == 1] <-"45"
```

Creating the RT data set we will be working with.

Below we are combining the 5 batches into 1 data set.

```
#Let's combine the data  
all_data <- rbind(FP_ED_batch1_corr, FP_ED_batch2B_corr, FP_ED_batch3_corr, FP_ED_  
batch4_error,  
                FP_ED_batch5_error)  
  
# creating subj as a factor so we can actually combine our individual difference m  
easures with this data set.  
  
all_data$subj <- as.factor(all_data$subj)
```

Importing the Individual difference (ID) data set we will be working with.

Below we are importing the ID data set that will be used to investigate our covariates. We are also renaming the participant column to subj so it can be successfully combined with our RT data set on the column subj.

```
#Import Individual difference measures  
ID_Measures <- read_csv("ID Measures.csv")
```

```
## Parsed with column specification:
## cols(
##   .default = col_double(),
##   SRS2_classification = col_character(),
##   WRMT_WI_RPI = col_character(),
##   WRMT_WA_RPI = col_logical(),
##   WRMT_WC_RPI = col_logical(),
##   WRMT_PC_RPI = col_logical(),
##   WRMT_LC_RPI = col_logical(),
##   WRMT_ORF_RPI = col_logical(),
##   WRMT_WI_stand = col_character(),
##   WRMT_WA_stand = col_character(),
##   WRMT_WC_stand = col_character(),
##   WRMT_PC_stand = col_character(),
##   WRMT_LC_stand = col_character(),
##   WRMT_ORF_stand = col_character(),
##   WRMT_total_reading_RPI = col_character()
## )
```

```
## See spec(...) for full column specifications.
```

```
#View(ID_Measures)
```

```
# Rename Participabt in ID_measures to subj to be the same as current data set
ID_Measures <- rename(ID_Measures, subj = Participant)
ID_Measures$subj <- as.factor(ID_Measures$subj)
```

Removing participants with missing data.

Below we are removing participants from the ID and RT data sets who have missing individual difference measures.

Additionally, this will ensure that when we next combined the two data sets they do not duplicate ID measures in the incorrect participant slot.

```
#Remove participants with missing data
ID_Measures_removed <- ID_Measures %>% filter(subj != 16)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 67)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 34)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 39)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 42)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 43)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 44)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 69)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 70)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 71)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 72)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 73)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 74)
ID_Measures_removed <- ID_Measures_removed %>% filter(subj != 75)
```


Next we will combine the two data sets together.

```
# Add the ID's to the data frame
all_data_join <- inner_join(all_data_removed, ID_Measures_removed, by = "subj")
```

```
## Warning: Column `subj` joining factors with different levels, coercing to
## character vector
```

Visualisation and analysis of *region 4* “the question”

Now that we’ve got a full data set ready to be analysed let’s first do some visualisation of the data to get a feel for how it is looking as well as the appropriate method. We might as well also relabel our conditions so they are easier to understand.

```
all_data_join$cond <- recode(all_data_join$cond, "1" = "facilitated", "2" = "unfacilitated")
```

```
all_data_join$cond <- as.factor(all_data_join$cond)
```

```
#Let's start with Region 4 "the question"
```

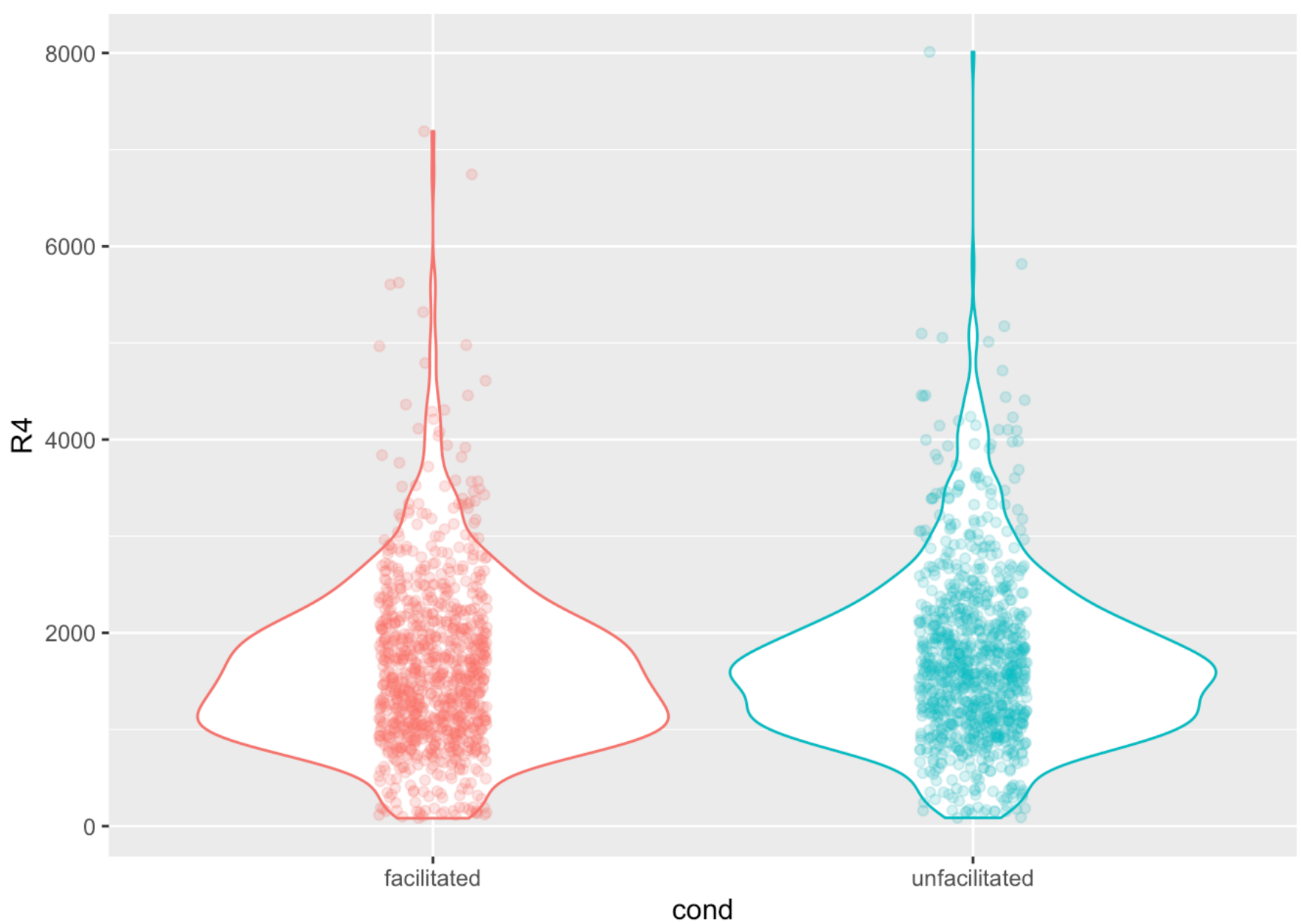
```
# Throw away zeroes
```

```
all_data_join <- all_data_join %>% filter(R4 != 0)
```

```
# Visualise
```

```
all_data_join %>%
  ggplot(aes(x = cond, y = R4, colour = cond)) +
  geom_violin() +
  geom_jitter(alpha = .2, width = .1) +
  stat_summary(fun.data = "mean_cl_boot", colour = "black") +
  guides(colour = FALSE)
```

```
## Warning: Computation failed in `stat_summary()`:
## Hmisc package required for this function
```



```
all_data_join %>%  
  group_by(cond) %>%  
  summarise(mean(R4), sd(R4))
```

```
## # A tibble: 2 x 3  
##   cond      `mean(R4)` `sd(R4)`  
##   <fct>      <dbl>    <dbl>  
## 1 facilitated    1631.    871.  
## 2 unfacilitated  1706.    882.
```

```
# Model assuming normality of residuals - singular fit error with more complex models  
model.null <- lmer(R4 ~ (1 | subj) + (1 + cond | item), all_data_join)  
model <- lmer(R4 ~ cond + (1 | subj) + (1 + cond | item), all_data_join)  
summary(model)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: R4 ~ cond + (1 | subj) + (1 + cond | item)
## Data: all_data_join
##
## REML criterion at convergence: 30563.5
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -3.5522 -0.5245 -0.0934  0.4131  8.1246
##
## Random effects:
## Groups      Name                Variance Std.Dev. Corr
## subj      (Intercept)          279730   528.9
## item      (Intercept)           49884   223.3
##           condunfacilitated    10564   102.8   -0.15
## Residual                        442335   665.1
## Number of obs: 1915, groups:  subj, 60; item, 32
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)    1629.88     81.75   83.10  19.937  <2e-16 ***
## condunfacilitated    76.39     35.43   30.09   2.156   0.0392 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr)
## condnfcltttd -0.198
```

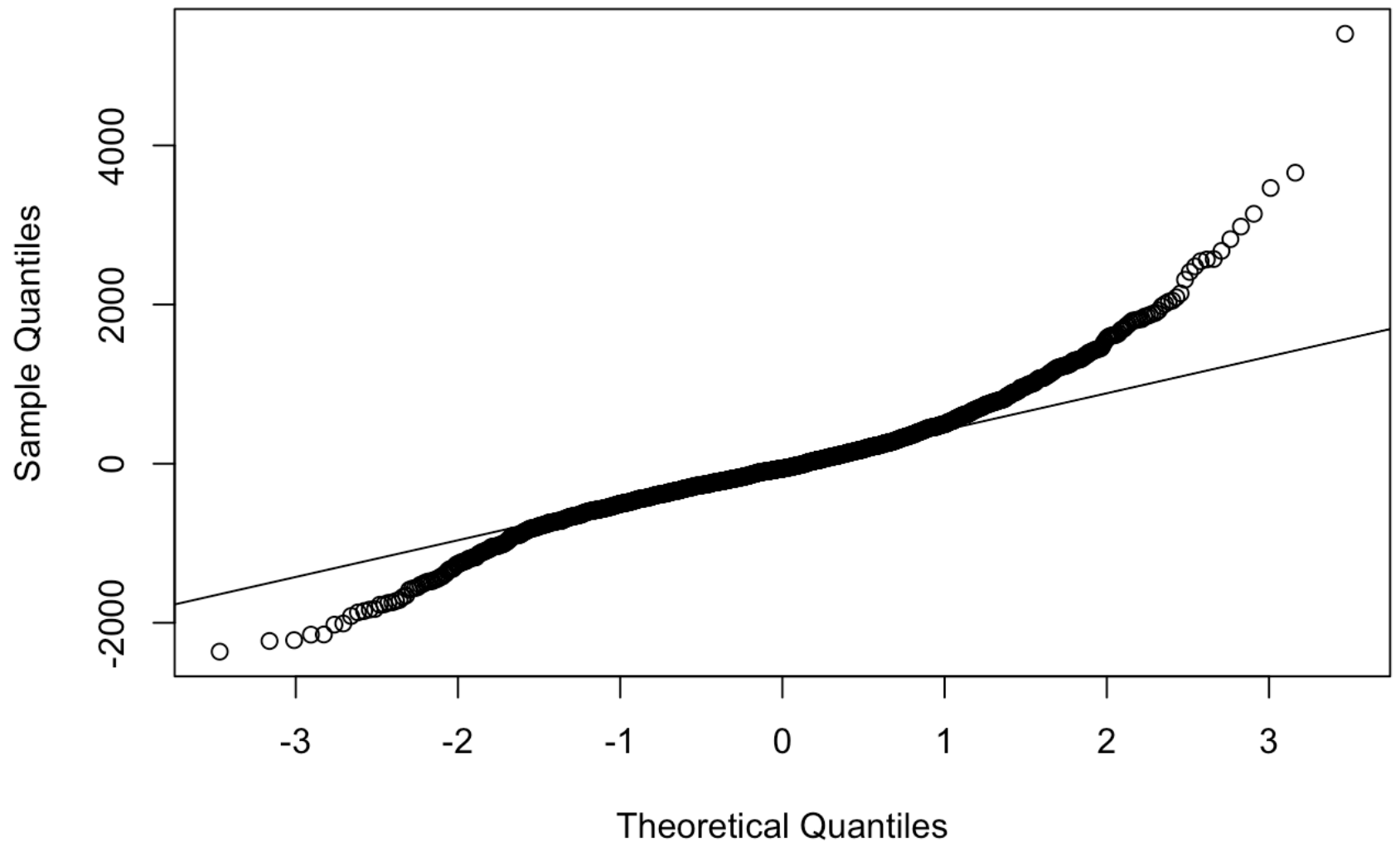
```
anova(model, model.null)
```

```
## refitting model(s) with ML (instead of REML)
```

```
## Data: all_data_join
## Models:
## model.null: R4 ~ (1 | subj) + (1 + cond | item)
## model: R4 ~ cond + (1 | subj) + (1 + cond | item)
##           Df    AIC    BIC logLik deviance  Chisq Chi Df Pr(>Chisq)
## model.null  6 30600 30633 -15294    30588
## model       7 30597 30636 -15292    30583 4.4661      1    0.03457 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

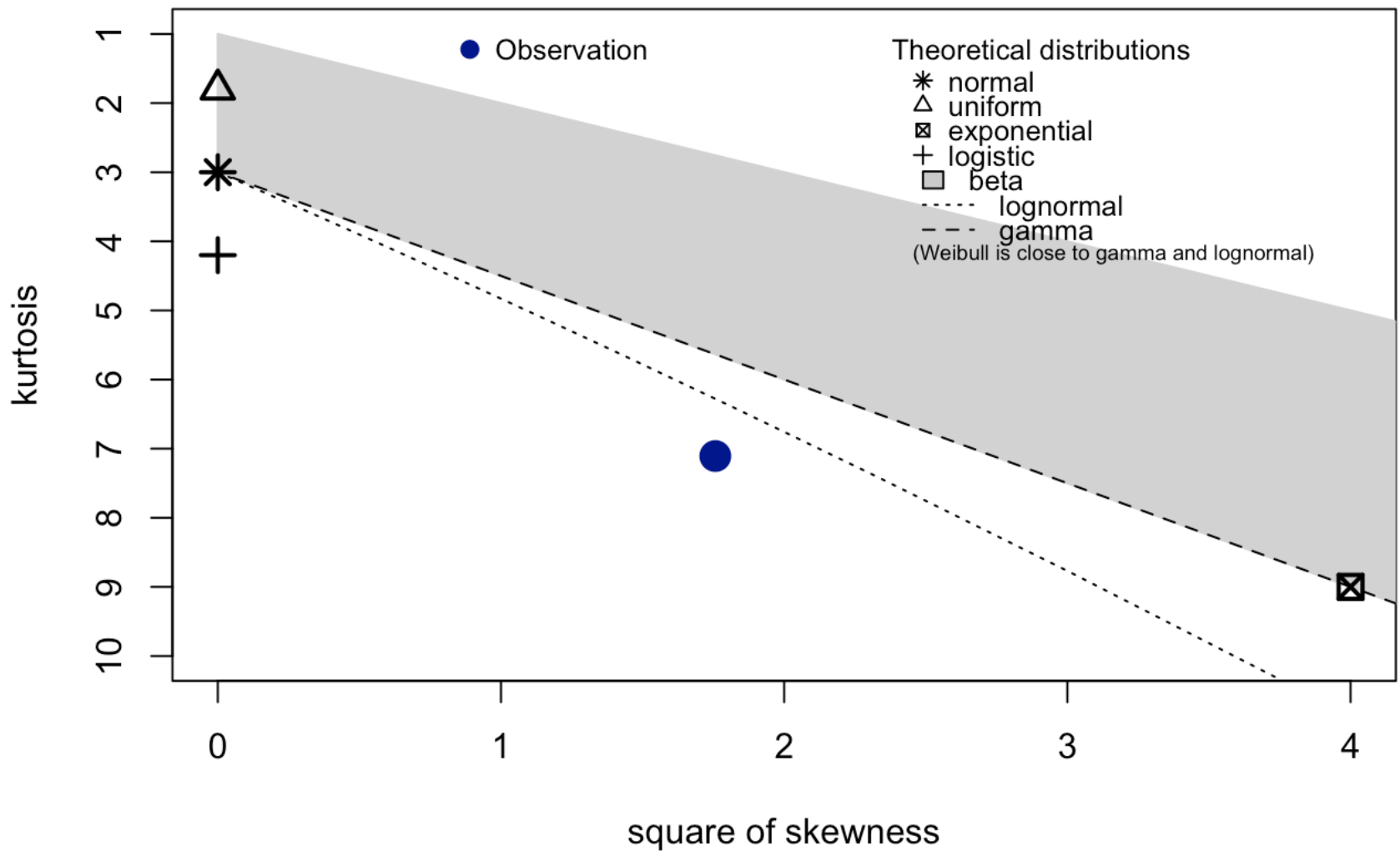
```
qqnorm(residuals(model))
qqline(residuals(model))
```

Normal Q-Q Plot



```
descdist(all_data_join$R4)
```

Cullen and Frey graph



```
## summary statistics
## -----
## min: 82    max: 8011
## median: 1561
## mean: 1668.422
## estimated sd: 876.8836
## estimated skewness: 1.325456
## estimated kurtosis: 7.106687
```

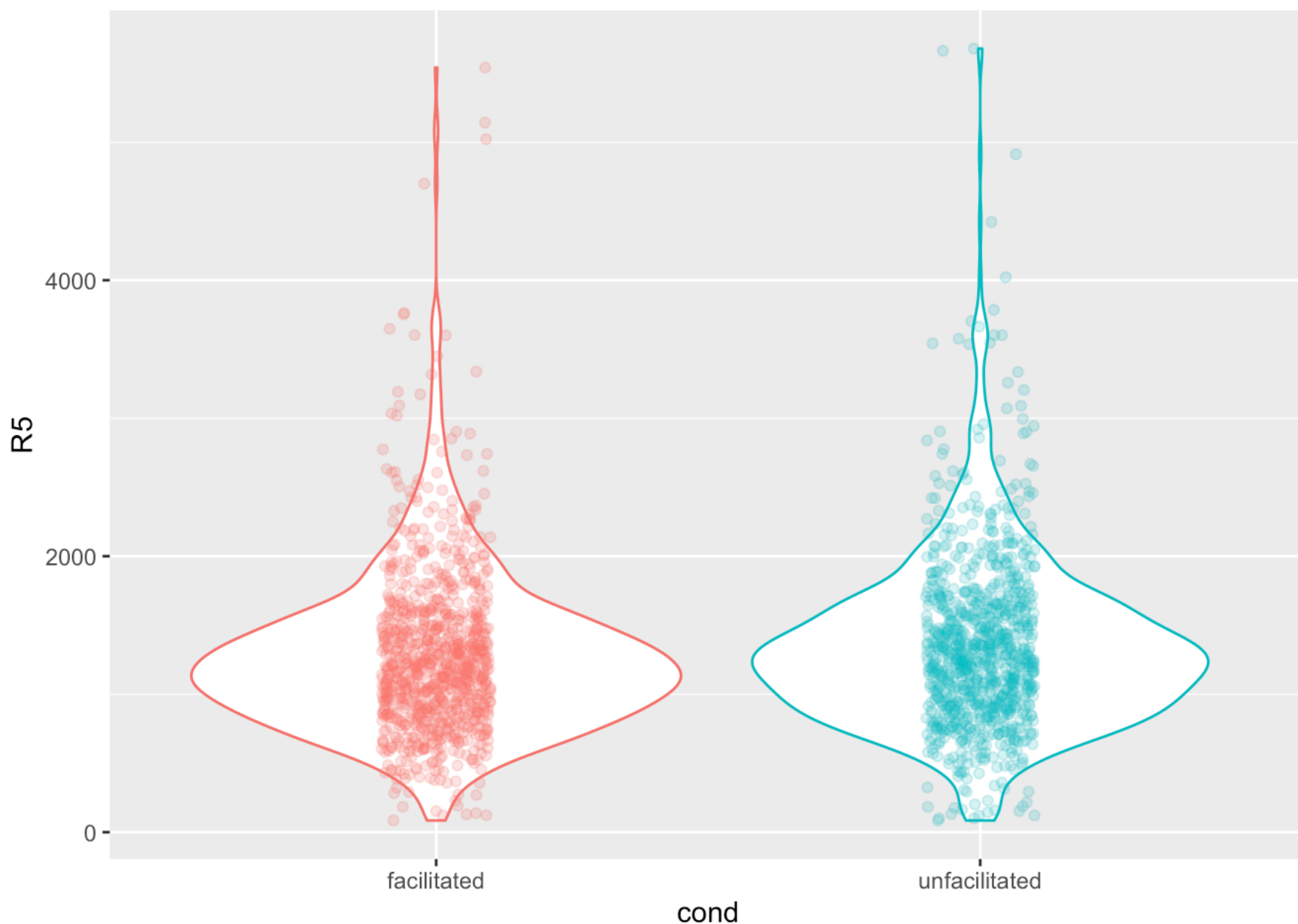
Visualisation and analysis of *region 5* “the reply”

What does region 5 look like?

```
all_data_join <- all_data_join %>% filter(R5 != 0)

# Visualise
all_data_join %>%
  ggplot(aes(x = cond, y = R5, colour = cond)) +
  geom_violin() +
  geom_jitter(alpha = .2, width = .1) +
  stat_summary(fun.data = "mean_cl_boot", colour = "black") +
  guides(colour = FALSE)
```

```
## Warning: Computation failed in `stat_summary()`:  
## Hmisc package required for this function
```



```
all_data_join %>%  
  group_by(cond) %>%  
  summarise(mean(R5), sd(R5))
```

```
## # A tibble: 2 x 3  
##   cond      `mean(R5)` `sd(R5)`  
##   <fct>      <dbl>    <dbl>  
## 1 facilitated    1286.    604.  
## 2 unfacilitated  1352.    634.
```

```
# Model assuming normality of residuals - singular fit error with more complex models  
model.null <- lmer(R5 ~ (1 | subj) + (1 + cond | item), all_data_join)  
model <- lmer(R5 ~ cond + (1 | subj) + (1 + cond | item), all_data_join)  
summary(model)
```

```
## Linear mixed model fit by REML. t-tests use Satterthwaite's method [
## lmerModLmerTest]
## Formula: R5 ~ cond + (1 | subj) + (1 + cond | item)
## Data: all_data_join
##
## REML criterion at convergence: 29029.7
##
## Scaled residuals:
##      Min       1Q   Median       3Q      Max
## -4.4843 -0.5294 -0.1056  0.3949  9.8484
##
## Random effects:
##   Groups      Name                Variance Std.Dev. Corr
##   subj      (Intercept)          144612   380.28
##   item      (Intercept)           34090   184.64
##              condunfacilitated    3681    60.67  0.51
## Residual                        201056   448.39
## Number of obs: 1912, groups:  subj, 60; item, 32
##
## Fixed effects:
##              Estimate Std. Error      df t value Pr(>|t|)
## (Intercept)      1286.45      60.71   84.61  21.189 < 2e-16 ***
## condunfacilitated    65.21      23.15   29.90   2.817  0.00851 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation of Fixed Effects:
##              (Intr)
## condnfclttd -0.022
```

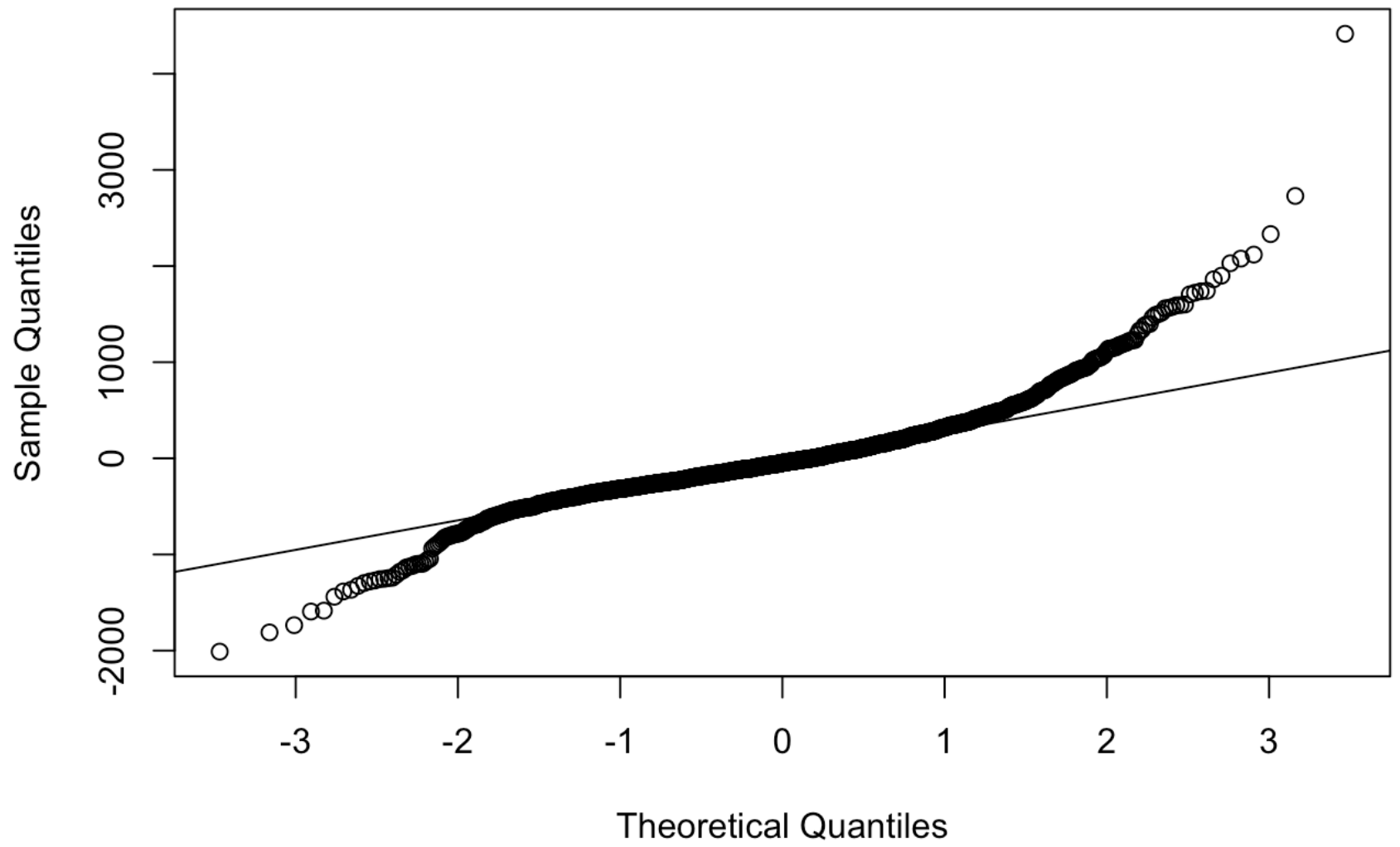
```
anova(model, model.null)
```

```
## refitting model(s) with ML (instead of REML)
```

```
## Data: all_data_join
## Models:
## model.null: R5 ~ (1 | subj) + (1 + cond | item)
## model: R5 ~ cond + (1 | subj) + (1 + cond | item)
##           Df   AIC   BIC logLik deviance  Chisq Chi Df Pr(>Chisq)
## model.null  6 29067 29100 -14528    29055
## model       7 29062 29101 -14524    29048 7.2758      1  0.006989 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

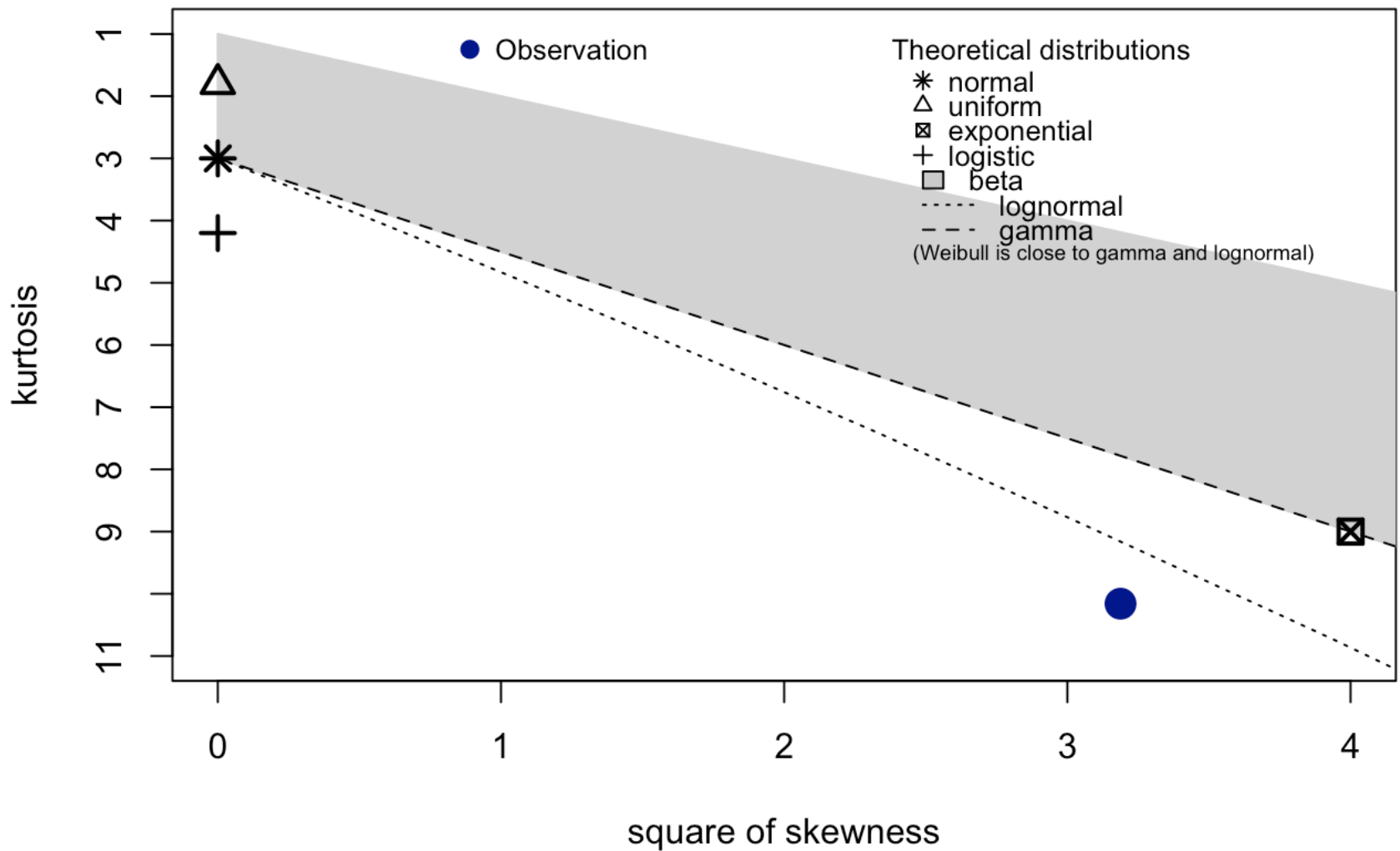
```
qqnorm(residuals(model))
qqline(residuals(model))
```

Normal Q-Q Plot



```
descdist(all_data_join$R5)
```


Cullen and Frey graph



```
## summary statistics
## -----
## min: 85    max: 5679
## median: 1230
## mean: 1319.181
## estimated sd: 620.0185
## estimated skewness: 1.785423
## estimated kurtosis: 10.15902
```

#Including covariates in the model. What does it look like when we start including our individual difference measures? First we scale our measures of interest.

```
#Let's include some covariates! Region 4
```

```
#Step 1: Scale the ID measures so as to be comparable.
```

```
all_data_join$SRS2_total_score_t <- scale(all_data_join$SRS2_total_score_t)
```

```
all_data_join$EQ <- scale(all_data_join$EQ)
```

```
all_data_join$WRMT_total_reading_score <- scale(all_data_join$WRMT_total_reading_score)
```

```
all_data_join$WRMT_WI_raw <- scale(all_data_join$WRMT_WI_raw)
```

```
# SINGULAR FIT
```

```
model_alldata <- lmer(R4 ~ cond + SRS2_total_score_t + EQ + WRMT_total_reading_score + WRMT_WI_raw +
```

```
      (1 |subj) + (0 + SRS2_total_score_t|cond) +
```

```
      (0 + EQ|cond) + (0 + WRMT_total_reading_score|cond) +
```

```
      (0 + WRMT_WI_raw|cond) , all_data_join, REML = TRUE)
```

```
## boundary (singular) fit: see ?isSingular
```

```
#Try again
```

```
#Simplified it loads and cannot get it to converge!
```