

# **explanation\_data\_analysis**

```
library(tidyverse)
```

```
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr     1.1.4     v readr     2.1.6
v forcats   1.0.1     v stringr   1.6.0
v ggplot2   4.0.1     v tibble    3.3.0
v lubridate  1.9.4     v tidyr    1.3.1
v purrr    1.2.0
-- Conflicts -----
x dplyr::filter() masks stats::filter()
x dplyr::lag()    masks stats::lag()
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become non-conflicting
```

```
library(readr)
library(ggplot2)
library(dplyr)
library(brms)
```

```
Loading required package: Rcpp
Loading 'brms' package (version 2.23.0). Useful instructions
can be found by typing help('brms'). A more detailed introduction
to the package is available through vignette('brms_overview').
```

```
Attaching package: 'brms'
```

```
The following object is masked from 'package:stats':
```

```
ar
```

```
library(lme4)
```

Loading required package: Matrix

Attaching package: 'Matrix'

The following objects are masked from 'package:tidy়':

expand, pack, unpack

Attaching package: 'lme4'

The following object is masked from 'package:brms':

ngrps

```
library(lmerTest)
```

Attaching package: 'lmerTest'

The following object is masked from 'package:lme4':

lmer

The following object is masked from 'package:stats':

step

```
library(ggpubr)
```

```
df_full <- read_csv("/home/pulapura/Documents/MSc Speech and Language Processing/diss_data_e
```

Rows: 64 Columns: 103

-- Column specification -----

Delimiter: ","

chr (103): StartDate, EndDate, Status, IPAddress, Progress, Duration (in sec...)

i Use `spec()` to retrieve the full column specification for this data.

i Specify the column types or set `show\_col\_types = FALSE` to quiet this message.

```

#I manually coded to itemname-itemtype-condition-question format in excel
# so I renamed the headers where appropriate (qualtrics output was the full question text)
old_names <- names(df_full)
new_names <- df_full[2, ] |> as.character()
new_names[is.na(new_names) | new_names == ""] <- old_names[is.na(new_names) | new_names == ""]
names(df_full) <- new_names
df_clean <- df_full[-c(1:4), ]

```

```

#Create a new row per participant per item per question
#and replace likert labels with numbers

df_pivot <- df_clean %>%
  pivot_longer(
    cols = matches("^[A-Za-z]+-(exp|unexp)-(r|nr)-[1-3]$$"),
    names_to = c("item", "type", "condition", "question"),
    names_pattern = "([A-Za-z]+)-(exp|unexp)-(r|nr)-([1-3])",
    values_to = "rating"
  )

df_pivot <- df_pivot |>
  mutate(
    type = toupper(type),
    condition = toupper(condition),
    question = recode(question,"1" = "Q1","2" = "Q2", "3"="Q3")
  )

likert_map <- c(
  "Very unfavorable" = 1,
  "Unfavorable" = 2,
  "Somewhat unfavorable" = 3,
  "Neither unfavorable nor favorable" = 4,
  "Somewhat favorable" = 5,
  "Favorable" = 6,
  "Very favorable" = 7
)

df_numeric <- df_pivot %>%
  mutate(rating_num = likert_map[rating])

df_numeric$type <- factor(df_numeric$type, levels = c("UNEXP", "EXP"))

```

```
#quick summary stats

df_summary <- df_numeric %>%
  filter(question %in% c("Q1", "Q2")) %>%
  group_by(type, condition, question) %>%
  summarise(
    mean = mean(rating_num, na.rm = TRUE),
    sd = sd(rating_num, na.rm = TRUE),
  )
```

`summarise()` has grouped output by 'type', 'condition'. You can override using the `groups` argument.

```
df_summary
```

```
# A tibble: 8 x 5
# Groups:   type, condition [4]
  type  condition question  mean     sd
  <fct> <chr>     <chr>    <dbl>   <dbl>
1 UNEXP NR        Q1      4.95   1.38
2 UNEXP NR        Q2      5.08   1.42
3 UNEXP R         Q1      4.38   1.62
4 UNEXP R         Q2      5.09   1.39
5 EXP   NR        Q1      5.06   1.28
6 EXP   NR        Q2      5.18   1.22
7 EXP   R          Q1      5.05   1.60
8 EXP   R          Q2      5.60   1.26
```

```
#violin plots for Q1 (Did Alice expect Bob to react favorably?)

ggplot(df_numeric %>% filter(question == "Q1"),
       aes(x = condition, y = rating_num, fill = condition)) +
  geom_violin(alpha = 0.6) +
  geom_jitter(width = 0.1, size = 1.5, alpha = 0.15) +
  geom_boxplot(width = 0.1, fill = "white") +
  facet_grid(cols = vars(type)) +
  stat_compare_means( #will replace this with the proper test later
    method = "wilcox.test",
    paired = FALSE,
    label = "p.signif",
  ) +
```

```
labs(
  x = "Condition", y = "Rating", title = "RQ1: Ratings per Condition")
```

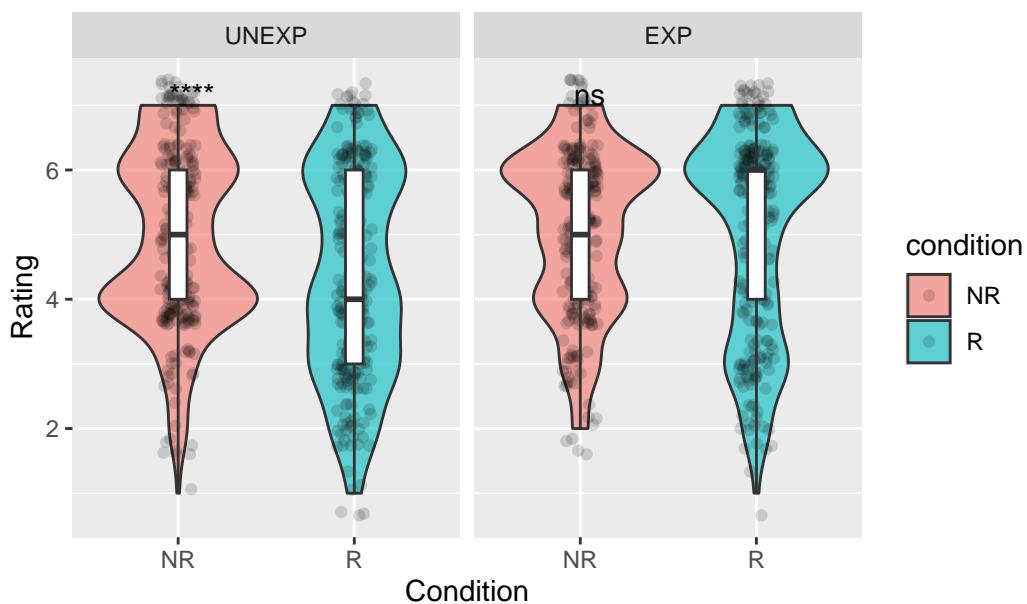
Warning: Removed 967 rows containing non-finite outside the scale range (`stat\_ydensity()`).

Warning: Removed 967 rows containing non-finite outside the scale range (`stat\_boxplot()`).

Warning: Removed 967 rows containing non-finite outside the scale range (`stat\_compare\_means()`).

Warning: Removed 967 rows containing missing values or values outside the scale range (`geom\_point()`).

### RQ1: Ratings per Condition



#Violin plots for Q2 ("How will Bob actually feel?")

```
ggplot(df_numeric %>% filter(question == "Q2"),
       aes(x = condition, y = rating_num, fill = condition)) +
  geom_violin(alpha = 0.6) +
```

```

geom_jitter(width = 0.1, size = 1.5, alpha = 0.15) +
geom_boxplot(width = 0.1, fill = "white") +
facet_grid(cols = vars(type)) +
stat_compare_means( #ditto Q1
  method = "wilcox.test",
  paired = FALSE,
  label = "p.signif",
) +
labs(x = "Condition", y = "Rating", title = "RQ2: Ratings per Condition")

```

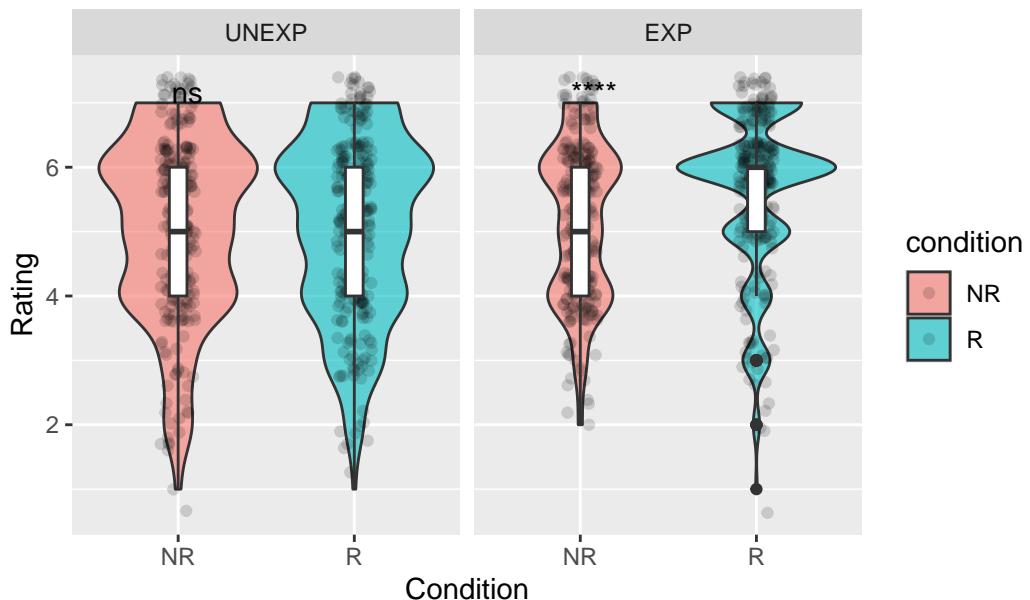
Warning: Removed 960 rows containing non-finite outside the scale range  
(`stat\_ydensity()`).

Warning: Removed 960 rows containing non-finite outside the scale range  
(`stat\_boxplot()`).

Warning: Removed 960 rows containing non-finite outside the scale range  
(`stat\_compare\_means()`).

Warning: Removed 960 rows containing missing values or values outside the scale range  
(`geom\_point()`).

## RQ2: Ratings per Condition



```

df_diff <- df_numeric %>%
  filter(question %in% c("Q1", "Q2")) %>%
  select(ResponseId, item, type, condition, question, rating_num) %>%
  pivot_wider(
    names_from = question,
    values_from = rating_num
  ) %>%
  mutate(diff = Q1 - Q2) #ended up not using this yet but I kept the df name sorry

```

```
#lme first cause im lazy
```

```

lme_Q1 <- lmer(Q1 ~ condition * type +
                 (1 | ResponseId) +
                 (1 | item),
                 data = df_diff)

```

```
summary(lme_Q1)
```

Linear mixed model fit by REML. t-tests use Satterthwaite's method [  
 lmerModLmerTest]

Formula: Q1 ~ condition \* type + (1 | ResponseId) + (1 | item)  
 Data: df\_diff

REML criterion at convergence: 3143.3

Scaled residuals:

Min	1Q	Median	3Q	Max
-3.2078	-0.6610	0.0688	0.6724	2.4549

Random effects:

Groups	Name	Variance	Std.Dev.
ResponseId	(Intercept)	0.3653	0.6044
item	(Intercept)	0.5495	0.7413
Residual		1.3499	1.1619

Number of obs: 953, groups: ResponseId, 60; item, 16

Fixed effects:

	Estimate	Std. Error	df	t value	Pr(> t )
(Intercept)	4.9270	0.2835	17.6528	17.376	1.52e-12 ***
conditionR	-0.5248	0.1061	876.0176	-4.945	9.12e-07 ***
typeEXP	0.1456	0.3858	15.1591	0.377	0.71118

```
conditionR:typeEXP    0.5103      0.1507 876.2277    3.386  0.00074 ***  
---  
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Correlation of Fixed Effects:  
          (Intr) cndtnR typEXP  
conditionR -0.187  
typeEXP     -0.679  0.138  
cndtnR:tEXP  0.132 -0.704 -0.197
```

```
lme_Q2 <- lmer(Q2 ~ condition * type +  
                 (1 | ResponseId) +  
                 (1 | item),  
                 data = df_diff)  
  
summary(lme_Q2)
```

```
Linear mixed model fit by REML. t-tests use Satterthwaite's method [  
lmerModLmerTest]  
Formula: Q2 ~ condition * type + (1 | ResponseId) + (1 | item)  
Data: df_diff
```

```
REML criterion at convergence: 2930.6
```

```
Scaled residuals:  
    Min     1Q   Median     3Q     Max  
-3.8830 -0.5956  0.1043  0.6490  2.8917
```

```
Random effects:  
Groups      Name        Variance Std.Dev.  
ResponseId (Intercept) 0.2176   0.4665  
item        (Intercept) 0.5373   0.7330  
Residual     1.0669   1.0329  
Number of obs: 960, groups: ResponseId, 60; item, 16
```

```
Fixed effects:  
            Estimate Std. Error       df t value Pr(>|t|)  
(Intercept)  5.05756   0.27429  16.44247 18.439 2.06e-12 ***  
conditionR    0.04737   0.09434 883.06257   0.502  0.61571  
typeEXP      0.10980   0.37844 14.91161   0.290  0.77571  
conditionR:typeEXP 0.40124   0.13342 883.06257   3.007  0.00271 **  
---
```

```
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

```
Correlation of Fixed Effects:  
            (Intr) cndtnR typEXP  
cndtnR    -0.172  
typEXP     -0.690  0.125  
cndtnR:tEXP  0.122 -0.707 -0.176
```

```
#bayesian ordinal model cause im paranoid  
  
fit_Q1 <- brm(Q1 ~ condition * type +  
                 (1|ResponseId) +  
                 (1|item),  
                 data = df_diff,  
                 family = cumulative())
```

```
Warning: Rows containing NAs were excluded from the model.
```

```
Compiling Stan program...
```

```
Start sampling
```

```
SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 1).  
Chain 1:  
Chain 1: Gradient evaluation took 0.000628 seconds  
Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 6.28 seconds.  
Chain 1: Adjust your expectations accordingly!  
Chain 1:  
Chain 1:  
Chain 1: Iteration:  1 / 2000 [  0%] (Warmup)  
Chain 1: Iteration:  200 / 2000 [ 10%] (Warmup)  
Chain 1: Iteration:  400 / 2000 [ 20%] (Warmup)  
Chain 1: Iteration:  600 / 2000 [ 30%] (Warmup)  
Chain 1: Iteration:  800 / 2000 [ 40%] (Warmup)  
Chain 1: Iteration: 1000 / 2000 [ 50%] (Warmup)  
Chain 1: Iteration: 1001 / 2000 [ 50%] (Sampling)  
Chain 1: Iteration: 1200 / 2000 [ 60%] (Sampling)  
Chain 1: Iteration: 1400 / 2000 [ 70%] (Sampling)  
Chain 1: Iteration: 1600 / 2000 [ 80%] (Sampling)  
Chain 1: Iteration: 1800 / 2000 [ 90%] (Sampling)
```

```
Chain 1: Iteration: 2000 / 2000 [100%]  (Sampling)
Chain 1:
Chain 1:   Elapsed Time: 33.263 seconds (Warm-up)
Chain 1:           31.442 seconds (Sampling)
Chain 1:           64.705 seconds (Total)
Chain 1:

SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 2).
Chain 2:
Chain 2: Gradient evaluation took 0.000566 seconds
Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 5.66 seconds.
Chain 2: Adjust your expectations accordingly!
Chain 2:
Chain 2:
Chain 2: Iteration:    1 / 2000 [  0%]  (Warmup)
Chain 2: Iteration:   200 / 2000 [ 10%]  (Warmup)
Chain 2: Iteration:   400 / 2000 [ 20%]  (Warmup)
Chain 2: Iteration:   600 / 2000 [ 30%]  (Warmup)
Chain 2: Iteration:   800 / 2000 [ 40%]  (Warmup)
Chain 2: Iteration:  1000 / 2000 [ 50%]  (Warmup)
Chain 2: Iteration: 1001 / 2000 [ 50%]  (Sampling)
Chain 2: Iteration: 1200 / 2000 [ 60%]  (Sampling)
Chain 2: Iteration: 1400 / 2000 [ 70%]  (Sampling)
Chain 2: Iteration: 1600 / 2000 [ 80%]  (Sampling)
Chain 2: Iteration: 1800 / 2000 [ 90%]  (Sampling)
Chain 2: Iteration: 2000 / 2000 [100%]  (Sampling)
Chain 2:
Chain 2:   Elapsed Time: 33.58 seconds (Warm-up)
Chain 2:           43.624 seconds (Sampling)
Chain 2:           77.204 seconds (Total)
Chain 2:

SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 3).
Chain 3:
Chain 3: Gradient evaluation took 0.000627 seconds
Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 6.27 seconds.
Chain 3: Adjust your expectations accordingly!
Chain 3:
Chain 3:
Chain 3: Iteration:    1 / 2000 [  0%]  (Warmup)
Chain 3: Iteration:   200 / 2000 [ 10%]  (Warmup)
Chain 3: Iteration:   400 / 2000 [ 20%]  (Warmup)
Chain 3: Iteration:   600 / 2000 [ 30%]  (Warmup)
```

```

Chain 3: Iteration: 800 / 2000 [ 40%] (Warmup)
Chain 3: Iteration: 1000 / 2000 [ 50%] (Warmup)
Chain 3: Iteration: 1001 / 2000 [ 50%] (Sampling)
Chain 3: Iteration: 1200 / 2000 [ 60%] (Sampling)
Chain 3: Iteration: 1400 / 2000 [ 70%] (Sampling)
Chain 3: Iteration: 1600 / 2000 [ 80%] (Sampling)
Chain 3: Iteration: 1800 / 2000 [ 90%] (Sampling)
Chain 3: Iteration: 2000 / 2000 [100%] (Sampling)
Chain 3:
Chain 3: Elapsed Time: 33.756 seconds (Warm-up)
Chain 3: 31.587 seconds (Sampling)
Chain 3: 65.343 seconds (Total)
Chain 3:

SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 4).
Chain 4:
Chain 4: Gradient evaluation took 0.000611 seconds
Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 6.11 seconds.
Chain 4: Adjust your expectations accordingly!
Chain 4:
Chain 4:
Chain 4: Iteration: 1 / 2000 [ 0%] (Warmup)
Chain 4: Iteration: 200 / 2000 [ 10%] (Warmup)
Chain 4: Iteration: 400 / 2000 [ 20%] (Warmup)
Chain 4: Iteration: 600 / 2000 [ 30%] (Warmup)
Chain 4: Iteration: 800 / 2000 [ 40%] (Warmup)
Chain 4: Iteration: 1000 / 2000 [ 50%] (Warmup)
Chain 4: Iteration: 1001 / 2000 [ 50%] (Sampling)
Chain 4: Iteration: 1200 / 2000 [ 60%] (Sampling)
Chain 4: Iteration: 1400 / 2000 [ 70%] (Sampling)
Chain 4: Iteration: 1600 / 2000 [ 80%] (Sampling)
Chain 4: Iteration: 1800 / 2000 [ 90%] (Sampling)
Chain 4: Iteration: 2000 / 2000 [100%] (Sampling)
Chain 4:
Chain 4: Elapsed Time: 33.437 seconds (Warm-up)
Chain 4: 28.017 seconds (Sampling)
Chain 4: 61.454 seconds (Total)
Chain 4:

```

```
summary(fit_Q1)
```

Family: cumulative

```

Links: mu = logit
Formula: Q1 ~ condition * type + (1 | ResponseId) + (1 | item)
Data: df_diff (Number of observations: 953)
Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
      total post-warmup draws = 4000

Multilevel Hyperparameters:
~item (Number of levels: 16)
Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
sd(Intercept) 1.25 0.27 0.83 1.89 1.00 1151 1838

~ResponseId (Number of levels: 60)
Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
sd(Intercept) 1.13 0.14 0.89 1.43 1.00 919 1761

Regression Coefficients:
Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
Intercept[1] -5.76 0.57 -6.90 -4.66 1.00 1152 1977
Intercept[2] -3.46 0.49 -4.39 -2.49 1.00 1011 1994
Intercept[3] -1.93 0.47 -2.84 -0.99 1.00 991 1910
Intercept[4] -0.43 0.47 -1.34 0.49 1.00 969 1787
Intercept[5] 0.34 0.47 -0.53 1.26 1.00 991 1759
Intercept[6] 2.69 0.48 1.78 3.67 1.00 1033 1995
conditionR -0.77 0.17 -1.09 -0.43 1.00 3489 2773
typeEXP 0.19 0.66 -1.12 1.52 1.00 883 1608
conditionR:typeEXP 0.83 0.24 0.35 1.30 1.00 3401 3085

```

#### Further Distributional Parameters:

	Estimate	Est.Error	l-95% CI	u-95% CI	Rhat	Bulk_ESS	Tail_ESS
disc	1.00	0.00	1.00	1.00	NA	NA	NA

Draws were sampled using sampling(NUTS). For each parameter, Bulk\_ESS and Tail\_ESS are effective sample size measures, and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat = 1).

```

fit_Q2 <- brm(Q2 ~ condition * type +
  (1|ResponseId) +
  (1|item),
  data = df_diff,
  family = cumulative())

```

Warning: Rows containing NAs were excluded from the model.

Compiling Stan program...

Start sampling

```
SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 1).
Chain 1:
Chain 1: Gradient evaluation took 0.000647 seconds
Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 6.47 seconds.
Chain 1: Adjust your expectations accordingly!
Chain 1:
Chain 1:
Chain 1: Iteration: 1 / 2000 [ 0%] (Warmup)
Chain 1: Iteration: 200 / 2000 [ 10%] (Warmup)
Chain 1: Iteration: 400 / 2000 [ 20%] (Warmup)
Chain 1: Iteration: 600 / 2000 [ 30%] (Warmup)
Chain 1: Iteration: 800 / 2000 [ 40%] (Warmup)
Chain 1: Iteration: 1000 / 2000 [ 50%] (Warmup)
Chain 1: Iteration: 1001 / 2000 [ 50%] (Sampling)
Chain 1: Iteration: 1200 / 2000 [ 60%] (Sampling)
Chain 1: Iteration: 1400 / 2000 [ 70%] (Sampling)
Chain 1: Iteration: 1600 / 2000 [ 80%] (Sampling)
Chain 1: Iteration: 1800 / 2000 [ 90%] (Sampling)
Chain 1: Iteration: 2000 / 2000 [100%] (Sampling)
Chain 1:
Chain 1: Elapsed Time: 46.089 seconds (Warm-up)
Chain 1:           43.231 seconds (Sampling)
Chain 1:           89.32 seconds (Total)
Chain 1:

SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 2).
Chain 2:
Chain 2: Gradient evaluation took 0.000649 seconds
Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 6.49 seconds.
Chain 2: Adjust your expectations accordingly!
Chain 2:
Chain 2:
Chain 2: Iteration: 1 / 2000 [ 0%] (Warmup)
Chain 2: Iteration: 200 / 2000 [ 10%] (Warmup)
Chain 2: Iteration: 400 / 2000 [ 20%] (Warmup)
Chain 2: Iteration: 600 / 2000 [ 30%] (Warmup)
Chain 2: Iteration: 800 / 2000 [ 40%] (Warmup)
```

```
Chain 2: Iteration: 1000 / 2000 [ 50%] (Warmup)
Chain 2: Iteration: 1001 / 2000 [ 50%] (Sampling)
Chain 2: Iteration: 1200 / 2000 [ 60%] (Sampling)
Chain 2: Iteration: 1400 / 2000 [ 70%] (Sampling)
Chain 2: Iteration: 1600 / 2000 [ 80%] (Sampling)
Chain 2: Iteration: 1800 / 2000 [ 90%] (Sampling)
Chain 2: Iteration: 2000 / 2000 [100%] (Sampling)
Chain 2:
Chain 2: Elapsed Time: 44.36 seconds (Warm-up)
Chain 2:           45.368 seconds (Sampling)
Chain 2:           89.728 seconds (Total)
Chain 2:
```

SAMPLING FOR MODEL 'anon\_model' NOW (CHAIN 3).

```
Chain 3:
Chain 3: Gradient evaluation took 0.000657 seconds
Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 6.57 seconds.
Chain 3: Adjust your expectations accordingly!
Chain 3:
Chain 3:
Chain 3: Iteration: 1 / 2000 [  0%] (Warmup)
Chain 3: Iteration: 200 / 2000 [ 10%] (Warmup)
Chain 3: Iteration: 400 / 2000 [ 20%] (Warmup)
Chain 3: Iteration: 600 / 2000 [ 30%] (Warmup)
Chain 3: Iteration: 800 / 2000 [ 40%] (Warmup)
Chain 3: Iteration: 1000 / 2000 [ 50%] (Warmup)
Chain 3: Iteration: 1001 / 2000 [ 50%] (Sampling)
Chain 3: Iteration: 1200 / 2000 [ 60%] (Sampling)
Chain 3: Iteration: 1400 / 2000 [ 70%] (Sampling)
Chain 3: Iteration: 1600 / 2000 [ 80%] (Sampling)
Chain 3: Iteration: 1800 / 2000 [ 90%] (Sampling)
Chain 3: Iteration: 2000 / 2000 [100%] (Sampling)
Chain 3:
Chain 3: Elapsed Time: 46.069 seconds (Warm-up)
Chain 3:           44.79 seconds (Sampling)
Chain 3:           90.859 seconds (Total)
Chain 3:
```

SAMPLING FOR MODEL 'anon\_model' NOW (CHAIN 4).

```
Chain 4:
Chain 4: Gradient evaluation took 0.000603 seconds
Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 6.03 seconds.
Chain 4: Adjust your expectations accordingly!
```

```

Chain 4:
Chain 4:
Chain 4: Iteration: 1 / 2000 [  0%] (Warmup)
Chain 4: Iteration: 200 / 2000 [ 10%] (Warmup)
Chain 4: Iteration: 400 / 2000 [ 20%] (Warmup)
Chain 4: Iteration: 600 / 2000 [ 30%] (Warmup)
Chain 4: Iteration: 800 / 2000 [ 40%] (Warmup)
Chain 4: Iteration: 1000 / 2000 [ 50%] (Warmup)
Chain 4: Iteration: 1001 / 2000 [ 50%] (Sampling)
Chain 4: Iteration: 1200 / 2000 [ 60%] (Sampling)
Chain 4: Iteration: 1400 / 2000 [ 70%] (Sampling)
Chain 4: Iteration: 1600 / 2000 [ 80%] (Sampling)
Chain 4: Iteration: 1800 / 2000 [ 90%] (Sampling)
Chain 4: Iteration: 2000 / 2000 [100%] (Sampling)
Chain 4:
Chain 4: Elapsed Time: 41.748 seconds (Warm-up)
Chain 4:           37.875 seconds (Sampling)
Chain 4:           79.623 seconds (Total)
Chain 4:

```

```
summary(fit_Q2)
```

```

Family: cumulative
Links: mu = logit
Formula: Q2 ~ condition * type + (1 | ResponseId) + (1 | item)
Data: df_diff (Number of observations: 960)
Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
      total post-warmup draws = 4000

Multilevel Hyperparameters:
~item (Number of levels: 16)
Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
sd(Intercept)     1.40      0.31      0.94      2.12 1.00      1325      2387

~ResponseId (Number of levels: 60)
Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
sd(Intercept)     1.03      0.13      0.80      1.32 1.00      1316      2262

Regression Coefficients:
Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
Intercept[1]       -6.23      0.74     -7.69     -4.80 1.00      1629      2298
Intercept[2]       -3.88      0.57     -4.94     -2.72 1.00      1212      1991

```

Intercept[3]	-2.40	0.54	-3.43	-1.31	1.00	1119	1675
Intercept[4]	-0.65	0.54	-1.68	0.43	1.00	1115	1899
Intercept[5]	0.50	0.54	-0.52	1.57	1.00	1111	1810
Intercept[6]	2.79	0.55	1.73	3.90	1.00	1135	1890
conditionR	0.15	0.17	-0.20	0.50	1.00	4484	3216
typeEXP	0.25	0.76	-1.25	1.79	1.00	947	1346
conditionR:typeEXP	0.75	0.24	0.28	1.22	1.00	4706	3127

Further Distributional Parameters:

	Estimate	Est.Error	1-95% CI	u-95% CI	Rhat	Bulk_ESS	Tail_ESS
disc	1.00	0.00	1.00	1.00	NA	NA	NA

Draws were sampled using sampling(NUTS). For each parameter, Bulk\_ESS and Tail\_ESS are effective sample size measures, and Rhat is the potential scale reduction factor on split chains (at convergence, Rhat = 1).

the end :)