

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 ----- tidyverse_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
```

```
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:tidyr':

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",
  ) +
```

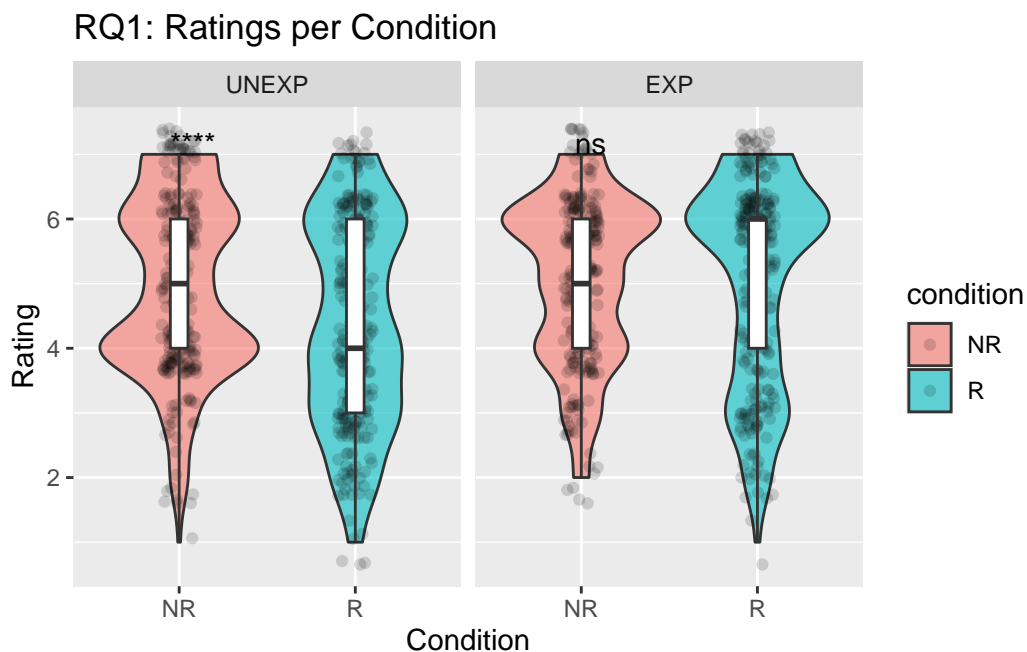
```
labs(
  x = "Condition", y = "Rating", title = "RQ1: Ratings per Condition")
```

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()``).



```
#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) +
```

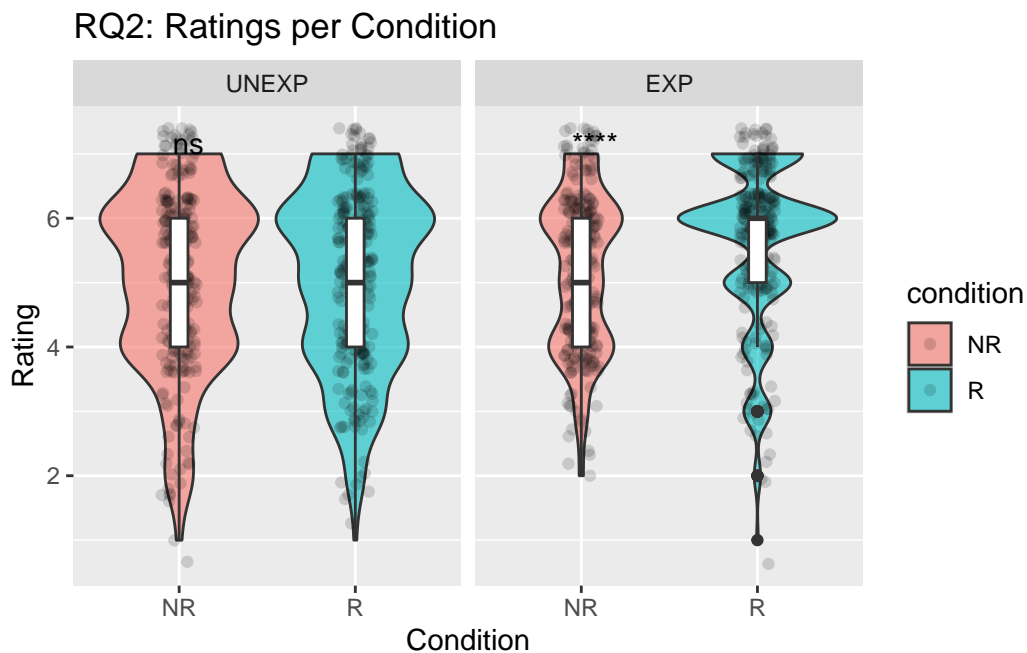
```
geom_jitter(width = 0.1, size = 1.5, alpha = 0.15) +
geom_boxplot(width = 0.1, fill = "white") +
facet_grid(cols = vars(type)) +
stat_compare_means( #ditto Q1
  method = "wilcox.test",
  paired = FALSE,
  label = "p.signif",
) +
labs(x = "Condition", y = "Rating", title = "RQ2: Ratings per Condition")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

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()`).



```
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  
conditionR -0.172  
typeEXP    -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 1-95% 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 1-95% 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	1-95% 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	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).

```

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 1-95% 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 1-95% 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 1-95% 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	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).

the end :)