

# E4 categorization

R. Cassano-Coleman

2025-09-30

This notebook analyzes categorization using Bayesian binomial generalized linear mixed effects models (GLMMs).

## Set up

```
set.seed(15000)
# other seeds for stability checks
#set.seed(20)
#set.seed(20000)

data <- read_csv('../data/E1-E2-E4/categorization.csv')

## Rows: 856 Columns: 6
## -- Column specification -----
## Delimiter: ","
## chr (3): response, scramble, Musician
## dbl (3): exp_subject_id, Trial_Nr, yrs_mus_exp
##
## 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.
```

Convert variables to factors.

```
data %>%
  mutate(scramble = ifelse(scramble == 'intact', 'Intact', scramble)) %>%
  mutate(exp_subject_id = as.factor(exp_subject_id),
        response = ifelse(response == 'Correct', TRUE, FALSE),
        scramble = factor(scramble, levels = c('Intact', '8B', '2B', '1B')),
        Musician = factor(Musician, levels = c('No', 'Yes'))) %>%
  filter(!is.na(response))
```

Set the contrast for condition.

```
contrasts(data$scramble) <- contr.treatment(4)
print(contrasts(data$scramble))
```

```
##      2 3 4
## Intact 0 0 0
## 8B     1 0 0
## 2B     0 1 0
## 1B     0 0 1
```

Set the musician/non-musician contrast.

```
contrasts(data$Musician) <- contr.sum(2)
print(contrasts(data$Musician))
```

```
##      [,1]
## No       1
## Yes     -1
```

# Main analysis

## Priors

Priors are expressed in log(odds) space.

**Intercept:** Given that chance is 25%, we assume that participants will perform somewhere between chance and ceiling. We expect the center of the distribution of accuracy to be somewhere around 60% or 65%. If we use a center of 65% and an SD of 1.5, 95% of the values fall between 8.46% and 97.4%.

```
prior_intercept <- set_prior('normal(log(0.65 / (1 - 0.65)), 1)', class = 'Intercept')
```

**Group:** We might expect musicians to do slightly better than non-musicians, on average.

In this range, a difference in 0.25 log odds gives us about a 5% decrease in accuracy.

```
prior_mus <- set_prior('normal(-0.25, 1)', coef = 'Musician1')
```

**Scramble:** We expect performance to improve as scramble level decreases. If we code 8B as reference level, then we expect 8B > 2B and 8B > 1B.

Since we're keeping the musician slope at SD = 1, we'll keep these (and the interactions) at SD = 1. This seems to be a pretty weak prior.

```
prior_scramble8B <- set_prior('normal(-0.1, 1)', coef = 'scramble2')
prior_scramble2B <- set_prior('normal(-0.2, 1)', coef = 'scramble3')
prior_scramble1B <- set_prior('normal(-0.3, 1)', coef = 'scramble4')
```

**Interaction:** We expect no interaction between group and scramble.

```
prior_int8B <- set_prior('normal(0, 1)', coef = 'Musician1:scramble2')
prior_int2B <- set_prior('normal(0, 1)', coef = 'Musician1:scramble3')
prior_int1B <- set_prior('normal(0, 1)', coef = 'Musician1:scramble4')
```

**Random slope for subjects:** *Leave this as default for now, may update.*

## Main model with group and condition

```
mus_scram <- brm(response ~ Musician + scramble + (1 | exp_subject_id), data = data,
                    family = bernoulli(),
                    prior = c(prior_intercept, prior_mus,
                              prior_scramble8B, prior_scramble2B, prior_scramble1B),
                    save_pars = save_pars(all = TRUE), iter = 20000,
                    file = 'models/E4_mus_scram')

## Compiling Stan program...

## Trying to compile a simple C file

## Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
## using C compiler: 'Apple clang version 16.0.0 (clang-1600.0.26.6)'
## using SDK: 'MacOSX15.2.sdk'
## clang -arch arm64 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I"/Library/Frameworks/R.framework/Resources/include"
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/StanHeaders/include
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include/Eigen/src/Core
## /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include/Eigen/src/Core
##   679 | #include <cmath>
##       |           ^~~~~~~
## 1 error generated.
## make: *** [foo.o] Error 1

## Start sampling

##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 8.7e-05 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.87 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration:    1 / 20000 [  0%] (Warmup)
## Chain 1: Iteration:  2000 / 20000 [ 10%] (Warmup)
## Chain 1: Iteration:  4000 / 20000 [ 20%] (Warmup)
## Chain 1: Iteration:  6000 / 20000 [ 30%] (Warmup)
## Chain 1: Iteration:  8000 / 20000 [ 40%] (Warmup)
## Chain 1: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 1: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 1: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 1: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 1: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 1: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 1: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 3.774 seconds (Warm-up)
## Chain 1:                 4.864 seconds (Sampling)
## Chain 1:                 8.638 seconds (Total)
## Chain 1:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 2).
```

```

## Chain 2:
## Chain 2: Gradient evaluation took 3.5e-05 seconds
## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.35 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:
## Chain 2: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 2: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 2: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 2: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 2: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 2: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 2: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 2: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 2: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 2: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 2: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 2: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 3.721 seconds (Warm-up)
## Chain 2:           4.859 seconds (Sampling)
## Chain 2:           8.58 seconds (Total)
## Chain 2:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 3.9e-05 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0.39 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 3: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 3: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 3: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 3: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 3: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 3: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 3: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 3: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 3: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 3: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 3: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 3.736 seconds (Warm-up)
## Chain 3:           4.832 seconds (Sampling)
## Chain 3:           8.568 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 3.2e-05 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0.32 seconds.
## Chain 4: Adjust your expectations accordingly!

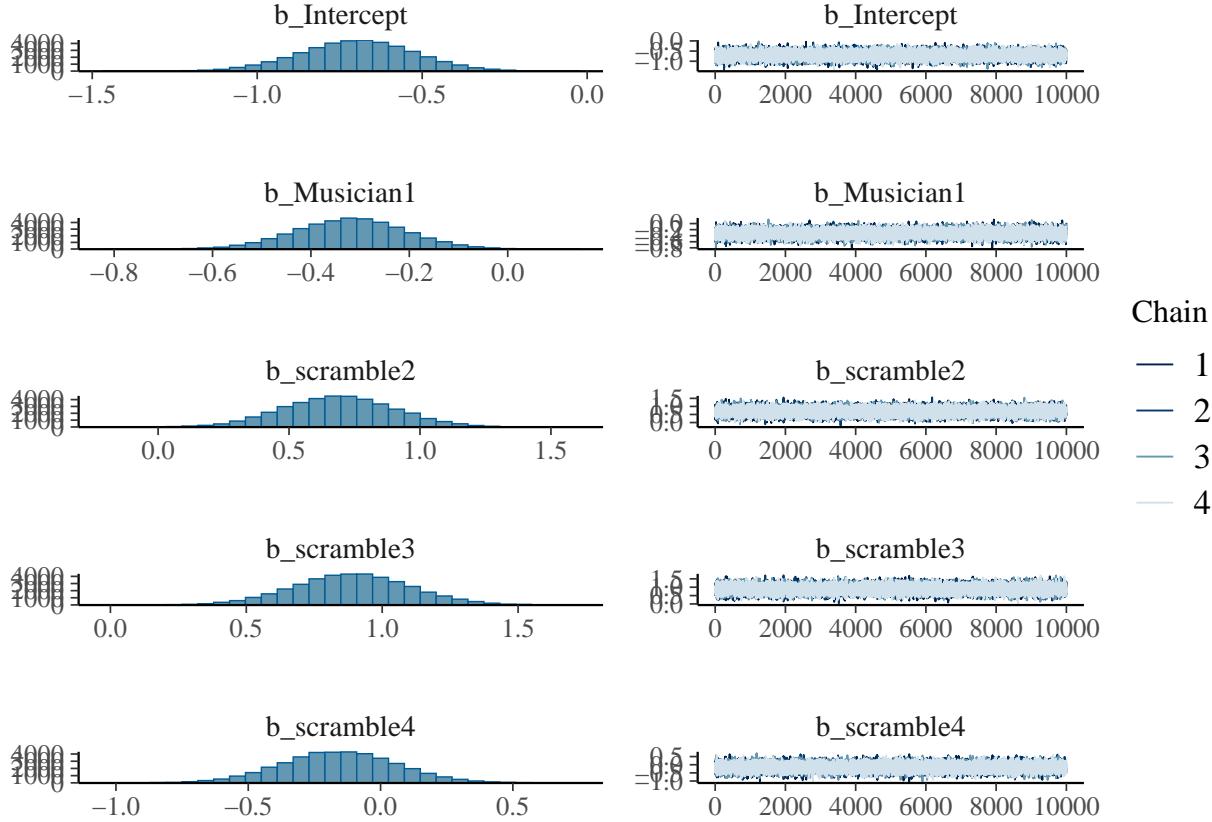
```

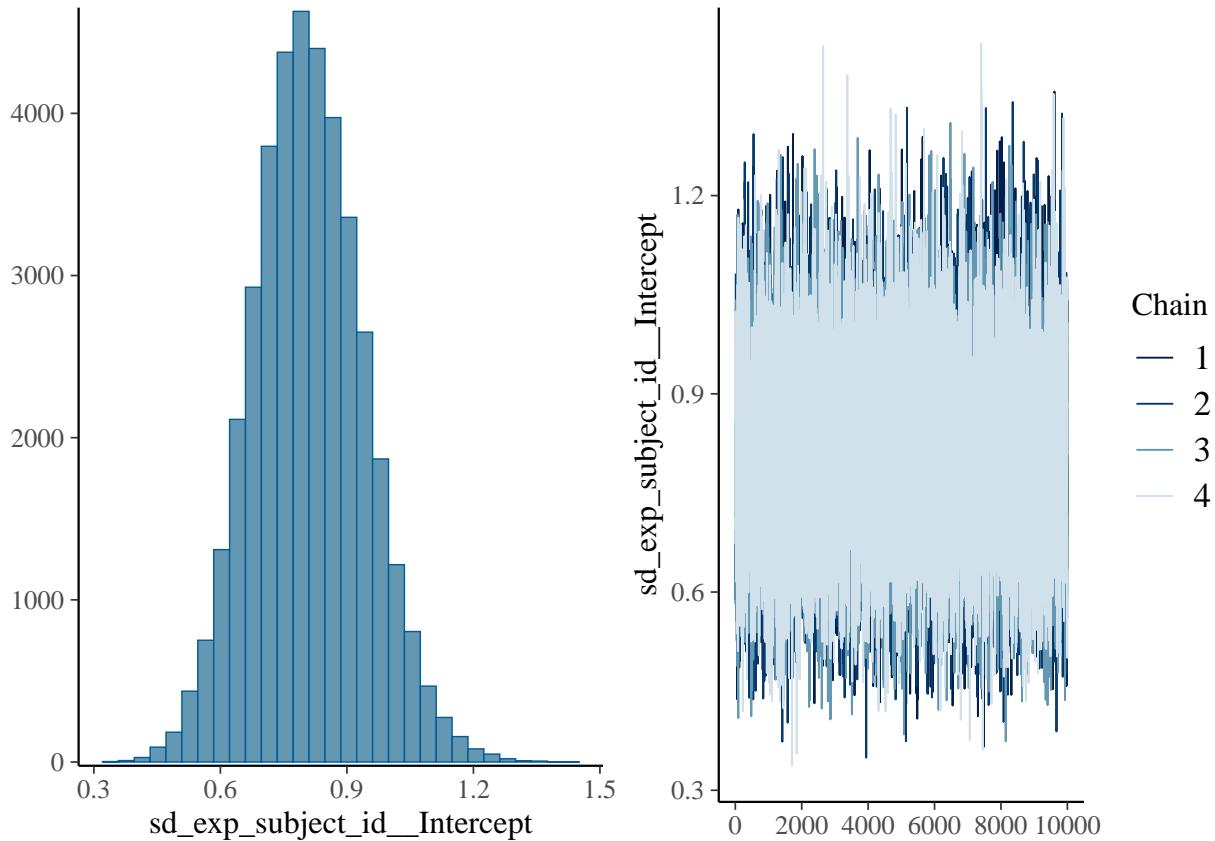
```

## Chain 4:
## Chain 4:
## Chain 4: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 4: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 4: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 4: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 4: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 4: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 4: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 4: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 4: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 4: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 4: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 4: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 3.739 seconds (Warm-up)
## Chain 4: 4.841 seconds (Sampling)
## Chain 4: 8.58 seconds (Total)
## Chain 4:

```

```
plot(mus_scram)
```





```

print(summary(mus_scram), digits = 4)

##  Family: bernoulli
##  Links: mu = logit
## Formula: response ~ Musician + scramble + (1 | exp_subject_id)
##  Data: data (Number of observations: 818)
##  Draws: 4 chains, each with iter = 20000; warmup = 10000; thin = 1;
##          total post-warmup draws = 40000
##
## Multilevel Hyperparameters:
## ~exp_subject_id (Number of levels: 106)
##             Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## sd(Intercept)  0.8094    0.1315   0.5621   1.0784 1.0000    15611    21521
## 
## Regression Coefficients:
##             Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## Intercept -0.6965    0.1712  -1.0364  -0.3652 1.0001    51118    29845
## Musician1 -0.3196    0.1118  -0.5391  -0.1033 1.0000    43053    31346
## scramble2  0.6903    0.2122   0.2813   1.1086 1.0002    54540    33660
## scramble3  0.8910    0.2131   0.4756   1.3106 1.0000    56685    34399
## scramble4 -0.1566    0.2169  -0.5805   0.2691 1.0001    60528    32682
## 
## 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).

```

```

emm_mus_scram_s <- emmeans(mus_scram, specs = "scramble")
summary(emm_mus_scram_s)

##   scramble   emmean lower.HPD upper.HPD
##   Intact    -0.69491   -1.043   -0.373
##   8B        -0.00553   -0.346    0.321
##   2B        0.19436   -0.136    0.532
##   1B        -0.85085   -1.199   -0.495
##
## Results are averaged over the levels of: Musician
## Point estimate displayed: median
## Results are given on the logit (not the response) scale.
## HPD interval probability: 0.95

contrast(emm_mus_scram_s, method = "pairwise")

##   contrast   estimate lower.HPD upper.HPD
##   Intact - 8B   -0.689   -1.104   -0.278
##   Intact - 2B   -0.890   -1.308   -0.474
##   Intact - 1B    0.157   -0.273    0.576
##   8B - 2B      -0.201   -0.613    0.220
##   8B - 1B      0.845    0.425    1.290
##   2B - 1B      1.048    0.607    1.474
##
## Results are averaged over the levels of: Musician
## Point estimate displayed: median
## Results are given on the log odds ratio (not the response) scale.
## HPD interval probability: 0.95

emm_mus_scram_ms <- emmeans(mus_scram, specs = c("Musician", "scramble"))
summary(emm_mus_scram_ms)

##   Musician scramble emmean lower.HPD upper.HPD
##   No       Intact   -1.013   -1.4179   -0.5957
##   Yes      Intact   -0.376   -0.7643    0.0198
##   No       8B       -0.326   -0.7282    0.0752
##   Yes      8B       0.313   -0.0825    0.7125
##   No       2B       -0.124   -0.5333    0.2684
##   Yes      2B       0.513    0.1238    0.9195
##   No       1B       -1.170   -1.5976   -0.7475
##   Yes      1B       -0.532   -0.9355   -0.1267
##
## Point estimate displayed: median
## Results are given on the logit (not the response) scale.
## HPD interval probability: 0.95

contrast(emm_mus_scram_ms, method = "pairwise")

##   contrast   estimate lower.HPD upper.HPD
##   No Intact - Yes Intact  -0.6369   -1.077   -0.206
##   No Intact - No 8B      -0.6892   -1.104   -0.278
##   No Intact - Yes 8B     -1.3258   -1.937   -0.716
##   No Intact - No 2B      -0.8905   -1.308   -0.474
##   No Intact - Yes 2B     -1.5252   -2.152   -0.921
##   No Intact - No 1B      0.1571   -0.273    0.576
##   No Intact - Yes 1B     -0.4829   -1.113    0.108

```

```

##  Yes Intact - No 8B    -0.0491   -0.650    0.536
##  Yes Intact - Yes 8B   -0.6892   -1.104   -0.278
##  Yes Intact - No 2B    -0.2514   -0.857    0.330
##  Yes Intact - Yes 2B   -0.8905   -1.308   -0.474
##  Yes Intact - No 1B     0.7962    0.180    1.403
##  Yes Intact - Yes 1B    0.1571   -0.273    0.576
##  No 8B - Yes 8B      -0.6369   -1.077   -0.206
##  No 8B - No 2B       -0.2009   -0.613    0.220
##  No 8B - Yes 2B      -0.8374   -1.455   -0.249
##  No 8B - No 1B       0.8452    0.425    1.290
##  No 8B - Yes 1B      0.2070   -0.405    0.805
##  Yes 8B - No 2B      0.4381   -0.175    1.039
##  Yes 8B - Yes 2B     -0.2009   -0.613    0.220
##  Yes 8B - No 1B      1.4832    0.860    2.110
##  Yes 8B - Yes 1B      0.8452    0.425    1.290
##  No 2B - Yes 2B      -0.6369   -1.077   -0.206
##  No 2B - No 1B       1.0478    0.607    1.474
##  No 2B - Yes 1B      0.4086   -0.199    1.012
##  Yes 2B - No 1B      1.6817    1.050    2.302
##  Yes 2B - Yes 1B      1.0478    0.607    1.474
##  No 1B - Yes 1B      -0.6369   -1.077   -0.206
##
## Point estimate displayed: median
## Results are given on the log odds ratio (not the response) scale.
## HPD interval probability: 0.95

```

## Main effects

```
main_BF <- describe_posterior(mus_scram,
                                estimate = "median", dispersion = TRUE,
                                ci = .95, ci_method = "HDI",
                                test = c("bayes_factor"))
print(main_BF, digits = 4)

## Summary of Posterior Distribution
##
## Parameter | Median | MAD | 95% CI | BF | Rhat | ESS
## -----
## (Intercept) | -0.6949 | 0.1712 | [-1.04, -0.37] | 798.50 | 1.000 | 50997.0000
## Musician1 | -0.3185 | 0.1117 | [-0.54, -0.10] | 7.04 | 1.000 | 42978.0000
## scramble2 | 0.6892 | 0.2139 | [ 0.28, 1.10] | 50.26 | 1.000 | 54437.0000
## scramble3 | 0.8905 | 0.2122 | [ 0.47, 1.31] | 1.14e+03 | 1.000 | 56586.0000
## scramble4 | -0.1571 | 0.2161 | [-0.58, 0.27] | 0.267 | 1.000 | 60477.0000
```

Moderate evidence for a main effect of group.

To get the main effect of scramble level, fit the “null” model with group only to compare.

```
mus_only <- brm(response ~ Musician + (1 | exp_subject_id), data = data,
                  family = bernoulli(),
                  prior = c(prior_intercept, prior_mus),
                  save_pars = save_pars(all = TRUE), iter = 20000,
                  file = 'models/E4_mus_only')

## Compiling Stan program...

## Trying to compile a simple C file

## Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
## using C compiler: 'Apple clang version 16.0.0 (clang-1600.0.26.6)'
## using SDK: 'MacOSX15.2.sdk'
## clang -arch arm64 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I"/Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/include"
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/StanHeaders/include.hpp:1
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include/RcppEigen.hpp:1
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include/Eigen/src/Core/Matrix.h:1
##   679 | #include <cmath>
##       |           ^
## 1 error generated.
## make: *** [foo.o] Error 1

## Start sampling

##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 9.1e-05 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.91 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 1: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 1: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 1: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 1: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 1: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 1: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 1: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 1: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 1: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 1: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 1: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 3.446 seconds (Warm-up)
## Chain 1:                 4.588 seconds (Sampling)
## Chain 1:                 8.034 seconds (Total)
## Chain 1:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 3.7e-05 seconds
```

```

## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.37 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:
## Chain 2: Iteration: 1 / 20000 [ 0%] (Warmup)
## Chain 2: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 2: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 2: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 2: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 2: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 2: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 2: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 2: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 2: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 2: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 2: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 3.447 seconds (Warm-up)
## Chain 2: 3.979 seconds (Sampling)
## Chain 2: 7.426 seconds (Total)
## Chain 2:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 3e-05 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0.3 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration: 1 / 20000 [ 0%] (Warmup)
## Chain 3: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 3: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 3: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 3: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 3: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 3: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 3: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 3: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 3: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 3: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 3: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 3.437 seconds (Warm-up)
## Chain 3: 4.626 seconds (Sampling)
## Chain 3: 8.063 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 3e-05 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0.3 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:

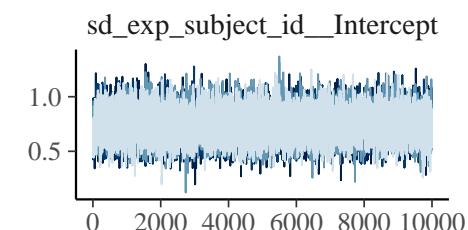
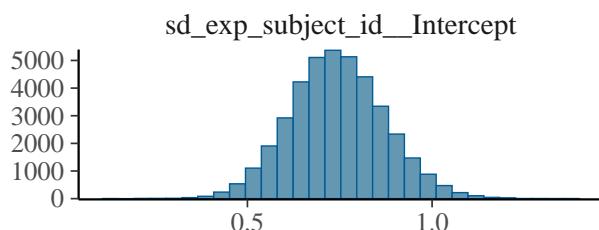
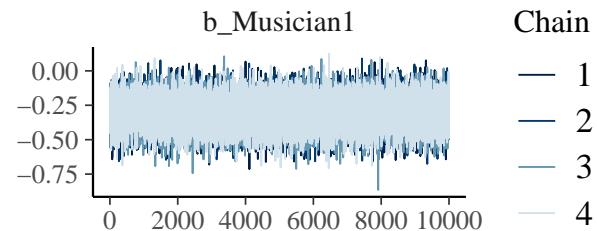
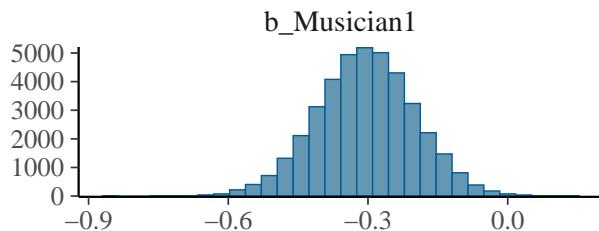
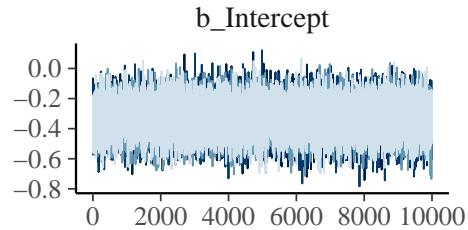
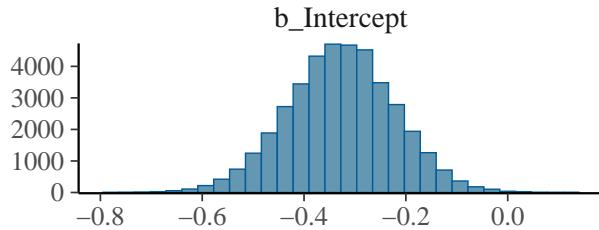
```

```

## Chain 4: Iteration:    1 / 20000 [  0%] (Warmup)
## Chain 4: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 4: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 4: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 4: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 4: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 4: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 4: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 4: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 4: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 4: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 4: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 3.441 seconds (Warm-up)
## Chain 4:                      4.601 seconds (Sampling)
## Chain 4:                      8.042 seconds (Total)
## Chain 4:

plot(mus_only)

```



```
print(summary(mus_only), digits = 4)
```

```

## Family: bernoulli
## Links: mu = logit
## Formula: response ~ Musician + (1 | exp_subject_id)
## Data: data (Number of observations: 818)
## Draws: 4 chains, each with iter = 20000; warmup = 10000; thin = 1;
##         total post-warmup draws = 40000
##

```

```

## Multilevel Hyperparameters:
## ~exp_subject_id (Number of levels: 106)
##           Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## sd(Intercept)  0.7434    0.1283   0.4996   1.0035 1.0008     14297    22399
##
## Regression Coefficients:
##           Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## Intercept  -0.3281    0.1055  -0.5376  -0.1240 1.0000     39158    30512
## Musician1 -0.3054    0.1053  -0.5148  -0.1002 1.0000     39893    31391
##
## 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).

BF_scramble <- bayes_factor(mus_scram, mus_only)

## Iteration: 1
## Iteration: 2
## Iteration: 3
## Iteration: 4
## Iteration: 5
## Iteration: 1
## Iteration: 2
## Iteration: 3
## Iteration: 4

print(BF_scramble)

## Estimated Bayes factor in favor of mus_scram over mus_only: 161997.86315

```

Very strong evidence for a main effect of scramble condition.

## Interaction between group and condition?

Add an interaction between group and condition, and compare the model with the interaction to the one without.

```
mus_scram_int <- brm(response ~ Musician*scramble + (1 | exp_subject_id), data = data,
                        family = bernoulli(),
                        prior = c(prior_intercept, prior_mus,
                                  prior_scramble8B, prior_scramble2B, prior_scramble1B,
                                  prior_int8B, prior_int2B, prior_int1B),
                        save_pars = save_pars(all = TRUE), iter = 20000,
                        file = 'models/E4_mus_scram_int')

## Compiling Stan program...

## Trying to compile a simple C file

## Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
## using C compiler: 'Apple clang version 16.0.0 (clang-1600.0.26.6)'
## using SDK: 'MacOSX15.2.sdk'
## clang -arch arm64 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I"/Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/StanHeaders/include"
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/StanHeaders/include
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include/Eigen/src/Core
##   679 | #include <cmath>
##       |           ^~~~~~~
## 1 error generated.
## make: *** [foo.o] Error 1

## Start sampling

##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 9e-05 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.9 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 1: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 1: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 1: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 1: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 1: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 1: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 1: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 1: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 1: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 1: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 1: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 4.363 seconds (Warm-up)
## Chain 1:                 5.078 seconds (Sampling)
## Chain 1:                 9.441 seconds (Total)
```

```

## Chain 1:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 3.3e-05 seconds
## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.33 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:
## Chain 2: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 2: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 2: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 2: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 2: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 2: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 2: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 2: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 2: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 2: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 2: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 2: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 4.386 seconds (Warm-up)
## Chain 2:           5.079 seconds (Sampling)
## Chain 2:           9.465 seconds (Total)
## Chain 2:
## 
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 3.3e-05 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0.33 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 3: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 3: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 3: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 3: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 3: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 3: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 3: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 3: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 3: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 3: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 3: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 4.39 seconds (Warm-up)
## Chain 3:           5.083 seconds (Sampling)
## Chain 3:           9.473 seconds (Total)
## Chain 3:
## 
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 4).
## Chain 4:

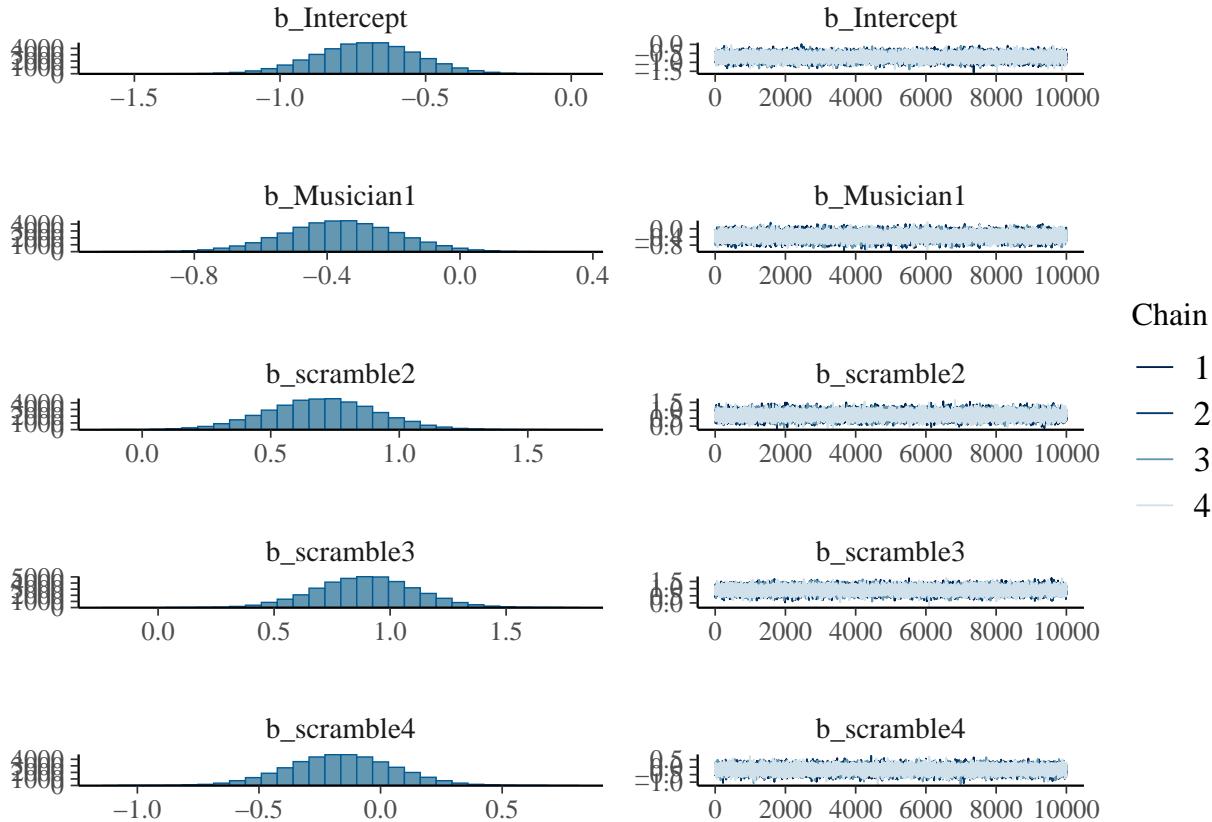
```

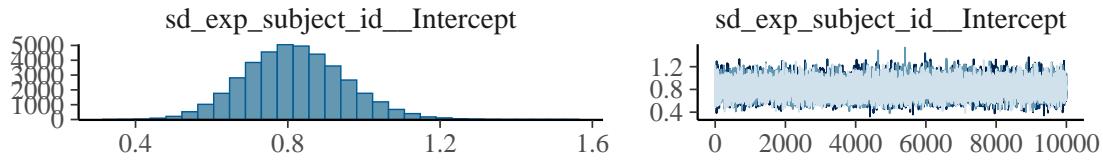
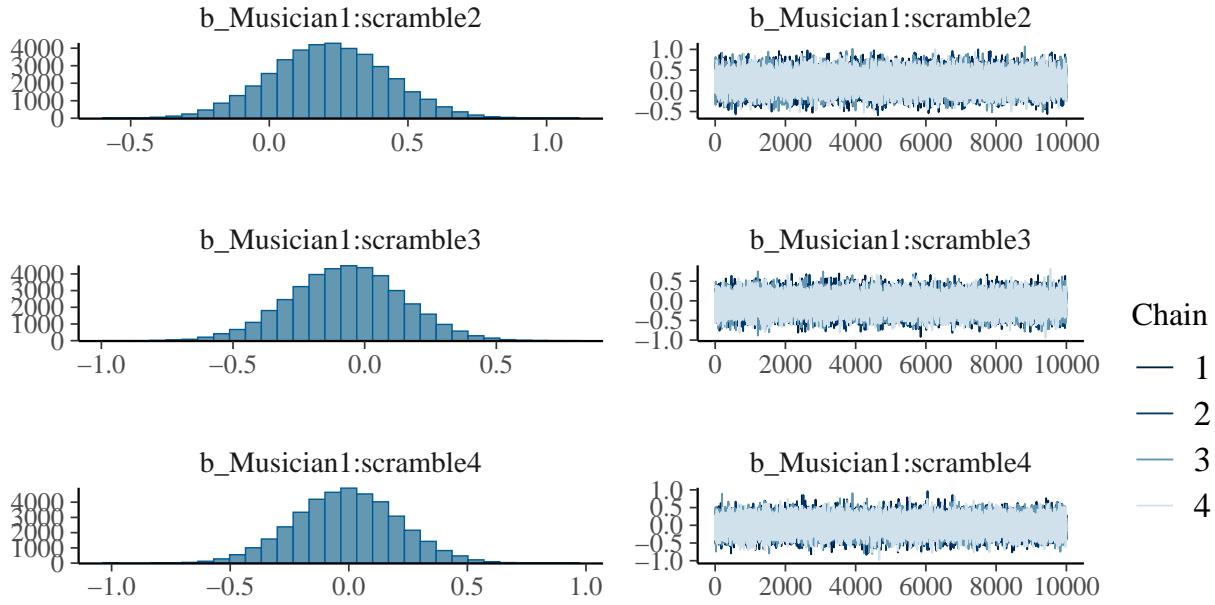
```

## Chain 4: Gradient evaluation took 3.7e-05 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0.37 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:
## Chain 4: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 4: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 4: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 4: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 4: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 4: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 4: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 4: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 4: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 4: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 4: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 4: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 4.373 seconds (Warm-up)
## Chain 4:           5.086 seconds (Sampling)
## Chain 4:           9.459 seconds (Total)
## Chain 4:

plot(mus_scram_int)

```





```
print(summary(mus_scram_int), digits = 4)
```

```
## Family: bernoulli
## Links: mu = logit
## Formula: response ~ Musician * scramble + (1 | exp_subject_id)
## Data: data (Number of observations: 818)
## Draws: 4 chains, each with iter = 20000; warmup = 10000; thin = 1;
##          total post-warmup draws = 40000
##
## Multilevel Hyperparameters:
## ~exp_subject_id (Number of levels: 106)
##                               Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## sd(Intercept)      0.8186     0.1322   0.5710   1.0895 1.0001    16177    22702
## 
## Regression Coefficients:
##                               Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS
## Intercept              -0.7028     0.1737  -1.0481  -0.3656 1.0001    45376
## Musician1                -0.3585     0.1699  -0.6943  -0.0256 1.0002    37265
## scramble2                 0.6984     0.2124   0.2828   1.1148 1.0002    54858
## scramble3                 0.8963     0.2144   0.4783   1.3192 1.0001    53643
## scramble4                 -0.1617     0.2168  -0.5851   0.2612 1.0000    56725
## Musician1:scramble2     0.2158     0.2114  -0.1949   0.6323 1.0000    44910
## Musician1:scramble3     -0.0698     0.2119  -0.4918   0.3445 1.0000    45835
## Musician1:scramble4     -0.0092     0.2181  -0.4375   0.4197 0.9999    45722
## 
##                               Tail_ESS
## Intercept                  29721
## Musician1                  31033
## scramble2                  33867
```

```

## scramble3          32999
## scramble4          34500
## Musician1:scramble2 32434
## Musician1:scramble3 33276
## Musician1:scramble4 33262
##
## 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).

BF_int <- bayes_factor(mus_scram_int, mus_scram)

## Iteration: 1
## Iteration: 2
## Iteration: 3
## Iteration: 4
## Iteration: 1
## Iteration: 2
## Iteration: 3
## Iteration: 4
## Iteration: 5

print(BF_int)

## Estimated Bayes factor in favor of mus_scram_int over mus_scram: 0.01949

```

Strong evidence against an interaction between group and condition.

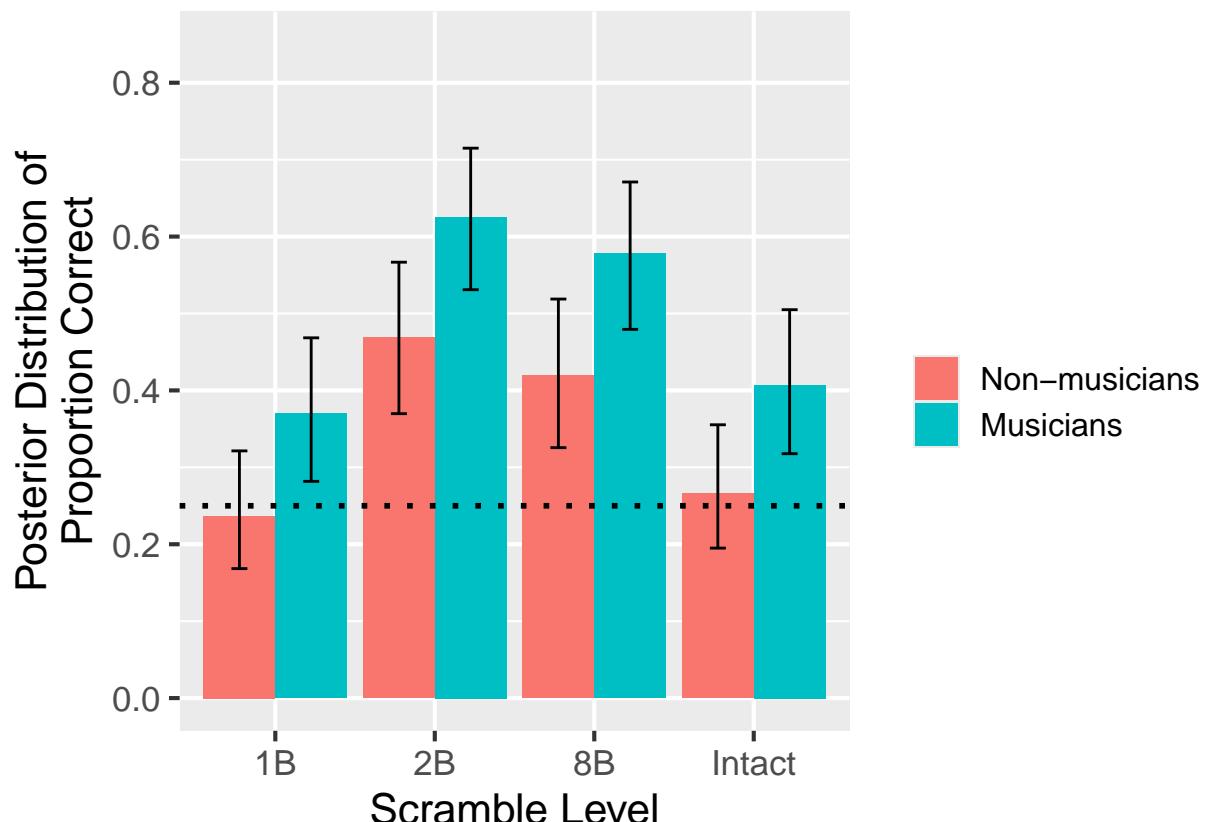
## Figure 5

Create a helper function for the conversion from log odds to probability.

```
calculate_prob_from_logodds <- function(logodds) {  
  return(exp(logodds) / (1 + exp(logodds)))  
}
```

Visualize with posterior estimates and 95% CrI on the scale of accuracy.

```
posterior_est <- as.data.frame(emm_mus_scram_ms)  
  
ggplot() +  
  geom_col(aes(x = scramble, y = calculate_prob_from_logodds(emmean), fill = Musician),  
           data = posterior_est,  
           position = "dodge") +  
  geom_errorbar(aes(x = scramble,  
                     ymin = calculate_prob_from_logodds(lower.HPD),  
                     ymax = calculate_prob_from_logodds(upper.HPD),  
                     fill = Musician),  
                data = posterior_est, position = position_dodge(width = 0.9), width = 0.2) +  
  geom_hline(yintercept = 0.25, linetype = "dotted", color = "black", linewidth = 1) +  
  theme_gray(base_size = 16) +  
  scale_x_discrete(limits = rev) +  
  ylim(0, 0.85) +  
  xlab('Scramble Level') +  
  ylab('Posterior Distribution of\nProportion Correct') +  
  scale_fill_discrete(name = "", labels = c('Non-musicians', 'Musicians')) +  
  theme(legend.text = element_text(size = 12))  
  
## Warning in geom_errorbar(aes(x = scramble, ymin =  
## calculate_prob_from_logodds(lower.HPD), : Ignoring unknown aesthetics: fill
```



## Years of experience

Keep only the subjects for which we have years of experience data and average accuracy per condition.

```
yrs_exp <- data %>%
  filter(!is.na(yrs_mus_exp)) %>%
  group_by(exp_subject_id, scramble, yrs_mus_exp) %>%
  summarize(count = n(),
            n_correct = sum(response),
            accuracy = n_correct / count)

## `summarise()` has grouped output by 'exp_subject_id', 'scramble'. You can
## override using the `.groups` argument.
```

## Priors

For this analysis, we're operating on the scale of accuracy. Because we don't see ceiling effects (i.e. participants aren't getting too close to perfect accuracy), a linear model is appropriate enough.

```
these_priors <- c(
  set_prior('normal(0.625, 0.1)', class = 'Intercept'),
  set_prior('normal(-0.1, 0.1)', coef = 'scramble2'),
  set_prior('normal(-0.2, 0.1)', coef = 'scramble3'),
  set_prior('normal(-0.3, 0.1)', coef = 'scramble4'),
  set_prior('normal(0, 0.1)', coef = 'yrs_mus_exp')
)
```

## Main model

```
years_mus_scram <- brm(accuracy ~ scramble + yrs_mus_exp + (1|exp_subject_id), data = yrs_exp,
                         prior = these_priors,
                         save_pars = save_pars(all = TRUE), iter = 20000,
                         file = 'models/E4_years')

## Compiling Stan program...

## Trying to compile a simple C file

## Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
## using C compiler: 'Apple clang version 16.0.0 (clang-1600.0.26.6)'
## using SDK: 'MacOSX15.2.sdk'
## clang -arch arm64 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I"/Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/StanHeaders/include"
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/StanHeaders/include
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include
## /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include/Eigen/src/Core
##   679 | #include <cmath>
##       |           ^
## 1 error generated.
## make: *** [foo.o] Error 1

## Start sampling

##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 7.9e-05 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.79 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration:    1 / 20000 [  0%] (Warmup)
## Chain 1: Iteration:  2000 / 20000 [ 10%] (Warmup)
## Chain 1: Iteration:  4000 / 20000 [ 20%] (Warmup)
## Chain 1: Iteration:  6000 / 20000 [ 30%] (Warmup)
## Chain 1: Iteration:  8000 / 20000 [ 40%] (Warmup)
## Chain 1: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 1: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 1: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 1: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 1: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 1: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 1: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 1.388 seconds (Warm-up)
## Chain 1:          0.939 seconds (Sampling)
## Chain 1:          2.327 seconds (Total)
## Chain 1:
## 
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 2).
## Chain 2:
## Chain 2: Gradient evaluation took 9e-06 seconds
```

```

## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.09 seconds.
## Chain 2: Adjust your expectations accordingly!
## Chain 2:
## Chain 2:
## Chain 2: Iteration: 1 / 20000 [ 0%] (Warmup)
## Chain 2: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 2: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 2: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 2: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 2: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 2: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 2: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 2: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 2: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 2: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 2: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 2:
## Chain 2: Elapsed Time: 1.247 seconds (Warm-up)
## Chain 2: 1.109 seconds (Sampling)
## Chain 2: 2.356 seconds (Total)
## Chain 2:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 3).
## Chain 3:
## Chain 3: Gradient evaluation took 8e-06 seconds
## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0.08 seconds.
## Chain 3: Adjust your expectations accordingly!
## Chain 3:
## Chain 3:
## Chain 3: Iteration: 1 / 20000 [ 0%] (Warmup)
## Chain 3: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 3: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 3: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 3: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 3: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 3: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 3: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 3: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 3: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 3: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 3: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 3:
## Chain 3: Elapsed Time: 1.153 seconds (Warm-up)
## Chain 3: 1.218 seconds (Sampling)
## Chain 3: 2.371 seconds (Total)
## Chain 3:
##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 4).
## Chain 4:
## Chain 4: Gradient evaluation took 8e-06 seconds
## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0.08 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:

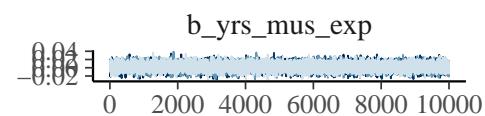
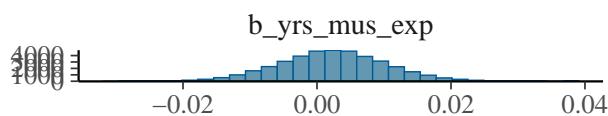
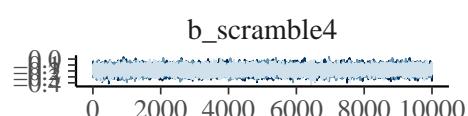
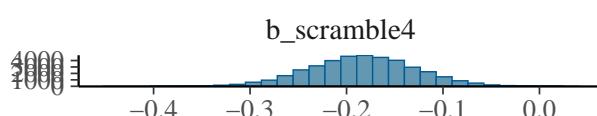
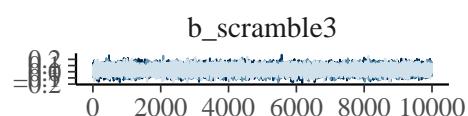
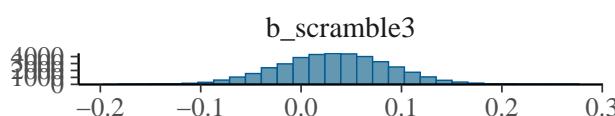
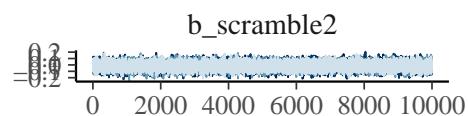
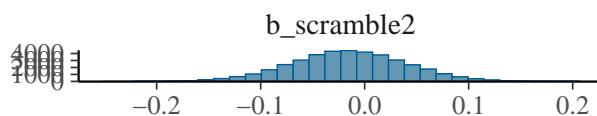
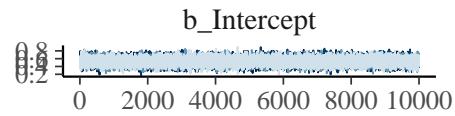
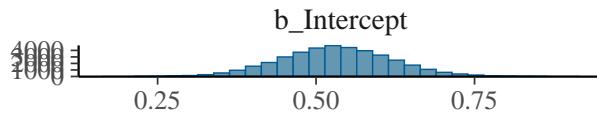
```

```

## Chain 4: Iteration:    1 / 20000 [  0%] (Warmup)
## Chain 4: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 4: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 4: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 4: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 4: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 4: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 4: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 4: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 4: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 4: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 4: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 1.274 seconds (Warm-up)
## Chain 4:                      1.225 seconds (Sampling)
## Chain 4:                      2.499 seconds (Total)
## Chain 4:

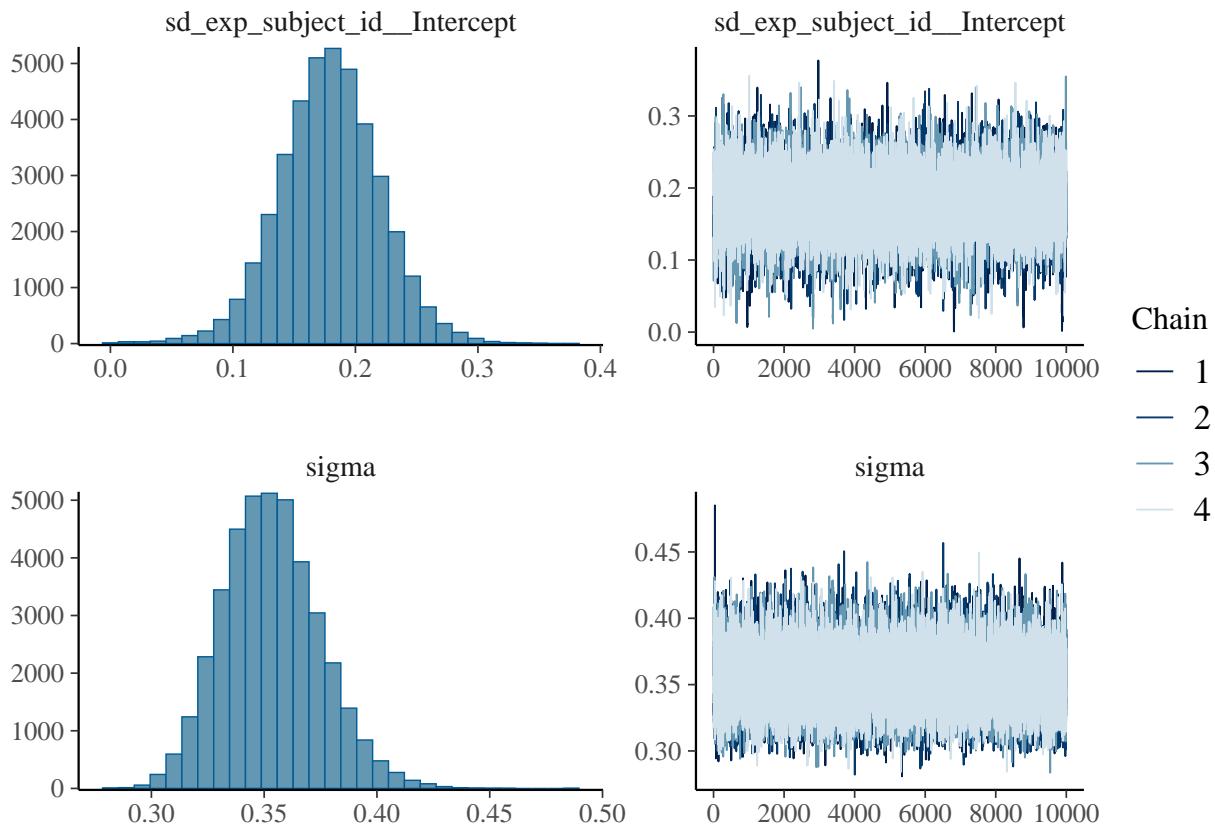
plot(years_mus_scram)

```



Chain

- 1
- 2
- 3
- 4



```

print(summary(years_mus_scram), digits = 4)

##  Family: gaussian
##  Links: mu = identity; sigma = identity
## Formula: accuracy ~ scramble + yrs_mus_exp + (1 | exp_subject_id)
##   Data: yrs_exp (Number of observations: 203)
##   Draws: 4 chains, each with iter = 20000; warmup = 10000; thin = 1;
##          total post-warmup draws = 40000
##
## Multilevel Hyperparameters:
## ~exp_subject_id (Number of levels: 51)
##             Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## sd(Intercept)  0.1790    0.0412   0.0970   0.2601 1.0002    11249    13637
## 
## Regression Coefficients:
##             Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## Intercept     0.5312    0.0875   0.3604   0.7044 1.0000    32698    28491
## scramble2    -0.0155    0.0551  -0.1250   0.0924 1.0002    49197    31529
## scramble3     0.0328    0.0553  -0.0760   0.1394 1.0000    49052    32666
## scramble4    -0.1836    0.0550  -0.2929  -0.0765 1.0000    49417    32925
## yrs_mus_exp   0.0023    0.0080  -0.0136   0.0181 1.0000    32125    29895
## 
## Further Distributional Parameters:
##             Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## sigma      0.3532    0.0215   0.3142   0.3983 1.0001    24441    27563
## 
## 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 for the estimate.

```

```
## scale reduction factor on split chains (at convergence, Rhat = 1).
```

## Null model (for plotting purposes)

```
years_mus <- brm(accuracy ~ yrs_mus_exp + (1|exp_subject_id), data = yrs_exp,
                   prior = c(
                     set_prior('normal(0.625, 0.1)', class = 'Intercept'),
                     set_prior('normal(0, 0.1)', coef = 'yrs_mus_exp')),
                   save_pars = save_pars(all = TRUE), iter = 20000,
                   file = 'models/E4_years_null')

## Compiling Stan program...

## Trying to compile a simple C file

## Running /Library/Frameworks/R.framework/Resources/bin/R CMD SHLIB foo.c
## using C compiler: 'Apple clang version 16.0.0 (clang-1600.0.26.6)'
## using SDK: 'MacOSX15.2.sdk'
## clang -arch arm64 -I"/Library/Frameworks/R.framework/Resources/include" -DNDEBUG -I"/Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/StanHeaders/include"
## In file included from <built-in>:1:
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/StanHeaders/include
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include
## In file included from /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include
## /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library/RcppEigen/include/Eigen/src/Core
##   679 | #include <cmath>
##       |           ^
## 1 error generated.
## make: *** [foo.o] Error 1

## Start sampling

##
## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 1).
## Chain 1:
## Chain 1: Gradient evaluation took 6.5e-05 seconds
## Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.65 seconds.
## Chain 1: Adjust your expectations accordingly!
## Chain 1:
## Chain 1:
## Chain 1: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 1: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 1: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 1: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 1: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 1: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 1: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 1: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 1: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 1: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 1: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 1: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 1:
## Chain 1: Elapsed Time: 1.333 seconds (Warm-up)
## Chain 1:                 1.225 seconds (Sampling)
## Chain 1:                 2.558 seconds (Total)
## Chain 1:
```

```

##  

## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 2).  

## Chain 2:  

## Chain 2: Gradient evaluation took 8e-06 seconds  

## Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.08 seconds.  

## Chain 2: Adjust your expectations accordingly!  

## Chain 2:  

## Chain 2:  

## Chain 2: Iteration: 1 / 20000 [ 0%] (Warmup)  

## Chain 2: Iteration: 2000 / 20000 [ 10%] (Warmup)  

## Chain 2: Iteration: 4000 / 20000 [ 20%] (Warmup)  

## Chain 2: Iteration: 6000 / 20000 [ 30%] (Warmup)  

## Chain 2: Iteration: 8000 / 20000 [ 40%] (Warmup)  

## Chain 2: Iteration: 10000 / 20000 [ 50%] (Warmup)  

## Chain 2: Iteration: 10001 / 20000 [ 50%] (Sampling)  

## Chain 2: Iteration: 12000 / 20000 [ 60%] (Sampling)  

## Chain 2: Iteration: 14000 / 20000 [ 70%] (Sampling)  

## Chain 2: Iteration: 16000 / 20000 [ 80%] (Sampling)  

## Chain 2: Iteration: 18000 / 20000 [ 90%] (Sampling)  

## Chain 2: Iteration: 20000 / 20000 [100%] (Sampling)  

## Chain 2:  

## Chain 2: Elapsed Time: 1.347 seconds (Warm-up)  

## Chain 2: 1.217 seconds (Sampling)  

## Chain 2: 2.564 seconds (Total)  

## Chain 2:  

##  

## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 3).  

## Chain 3:  

## Chain 3: Gradient evaluation took 8e-06 seconds  

## Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0.08 seconds.  

## Chain 3: Adjust your expectations accordingly!  

## Chain 3:  

## Chain 3:  

## Chain 3: Iteration: 1 / 20000 [ 0%] (Warmup)  

## Chain 3: Iteration: 2000 / 20000 [ 10%] (Warmup)  

## Chain 3: Iteration: 4000 / 20000 [ 20%] (Warmup)  

## Chain 3: Iteration: 6000 / 20000 [ 30%] (Warmup)  

## Chain 3: Iteration: 8000 / 20000 [ 40%] (Warmup)  

## Chain 3: Iteration: 10000 / 20000 [ 50%] (Warmup)  

## Chain 3: Iteration: 10001 / 20000 [ 50%] (Sampling)  

## Chain 3: Iteration: 12000 / 20000 [ 60%] (Sampling)  

## Chain 3: Iteration: 14000 / 20000 [ 70%] (Sampling)  

## Chain 3: Iteration: 16000 / 20000 [ 80%] (Sampling)  

## Chain 3: Iteration: 18000 / 20000 [ 90%] (Sampling)  

## Chain 3: Iteration: 20000 / 20000 [100%] (Sampling)  

## Chain 3:  

## Chain 3: Elapsed Time: 1.166 seconds (Warm-up)  

## Chain 3: 1.229 seconds (Sampling)  

## Chain 3: 2.395 seconds (Total)  

## Chain 3:  

##  

## SAMPLING FOR MODEL 'anon_model' NOW (CHAIN 4).  

## Chain 4:  

## Chain 4: Gradient evaluation took 8e-06 seconds

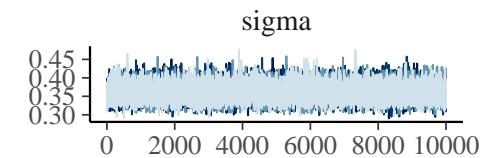
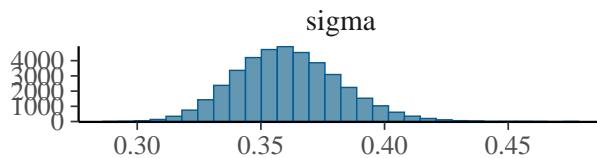
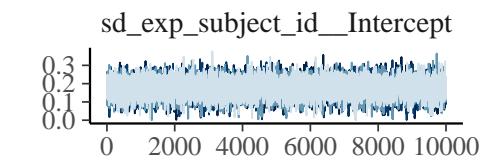
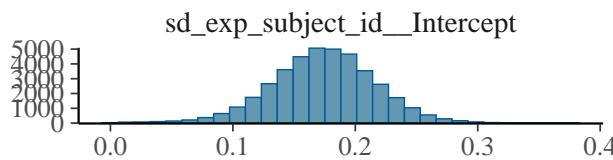
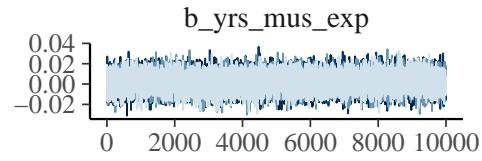
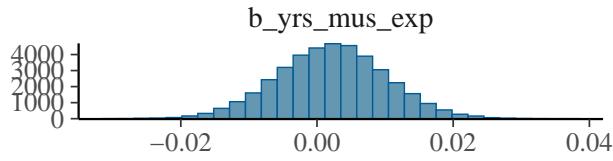
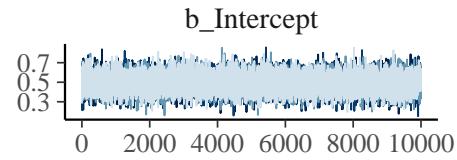
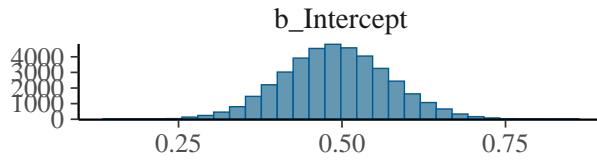
```

```

## Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0.08 seconds.
## Chain 4: Adjust your expectations accordingly!
## Chain 4:
## Chain 4:
## Chain 4: Iteration: 1 / 20000 [  0%] (Warmup)
## Chain 4: Iteration: 2000 / 20000 [ 10%] (Warmup)
## Chain 4: Iteration: 4000 / 20000 [ 20%] (Warmup)
## Chain 4: Iteration: 6000 / 20000 [ 30%] (Warmup)
## Chain 4: Iteration: 8000 / 20000 [ 40%] (Warmup)
## Chain 4: Iteration: 10000 / 20000 [ 50%] (Warmup)
## Chain 4: Iteration: 10001 / 20000 [ 50%] (Sampling)
## Chain 4: Iteration: 12000 / 20000 [ 60%] (Sampling)
## Chain 4: Iteration: 14000 / 20000 [ 70%] (Sampling)
## Chain 4: Iteration: 16000 / 20000 [ 80%] (Sampling)
## Chain 4: Iteration: 18000 / 20000 [ 90%] (Sampling)
## Chain 4: Iteration: 20000 / 20000 [100%] (Sampling)
## Chain 4:
## Chain 4: Elapsed Time: 1.236 seconds (Warm-up)
## Chain 4: 1.206 seconds (Sampling)
## Chain 4: 2.442 seconds (Total)
## Chain 4:

```

```
plot(years_mus)
```



```
print(summary(years_mus), digits = 4)
```

```

## Family: gaussian
## Links: mu = identity; sigma = identity
## Formula: accuracy ~ yrs_mus_exp + (1 | exp_subject_id)

```

```

##      Data: yrs_exp (Number of observations: 203)
##      Draws: 4 chains, each with iter = 20000; warmup = 10000; thin = 1;
##              total post-warmup draws = 40000
##
## Multilevel Hyperparameters:
## ~exp_subject_id (Number of levels: 51)
##           Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## sd(Intercept)  0.1735     0.0431   0.0845   0.2556 1.0002     10058    10953
##
## Regression Coefficients:
##           Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## Intercept     0.4906     0.0821   0.3311   0.6538 1.0000     41201    30923
## yrs_mus_exp   0.0021     0.0080  -0.0138   0.0179 1.0000     40497    31150
##
## Further Distributional Parameters:
##           Estimate Est.Error l-95% CI u-95% CI    Rhat Bulk_ESS Tail_ESS
## sigma       0.3610     0.0212   0.3224   0.4052 1.0000     25917    23933
##
## 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).

```

```

yrs_BF <- describe_posterior(years_mus_scram,
                               estimate = "median", dispersion = TRUE,
                               ci = .95, ci_method = "HDI",
                               test = c("bayes_factor"))
print(yrs_BF, digits = 4)

## Summary of Posterior Distribution
##
## Parameter | Median | MAD | 95% CI | BF | Rhat | ESS
## -----
## (Intercept) | 0.5308 | 0.0866 | [ 0.36, 0.70] | 7.70e+04 | 1.000 | 32738.0000
## scramble2 | -0.0156 | 0.0548 | [-0.12, 0.10] | 0.346 | 1.000 | 48817.0000
## scramble3 | 0.0333 | 0.0554 | [-0.08, 0.14] | 0.088 | 1.000 | 48990.0000
## scramble4 | -0.1834 | 0.0545 | [-0.29, -0.08] | 1.47 | 1.000 | 49511.0000
## yrs_mus_exp | 0.0023 | 0.0078 | [-0.01, 0.02] | 0.083 | 1.000 | 32026.0000

yrs_null_BF <- describe_posterior(years_mus,
                                   estimate = "median", dispersion = TRUE,
                                   ci = .95, ci_method = "HDI",
                                   test = c("bayes_factor"))
print(yrs_null_BF, digits = 4)

## Summary of Posterior Distribution
##
## Parameter | Median | MAD | 95% CI | BF | Rhat | ESS
## -----
## (Intercept) | 0.4895 | 0.0808 | [ 0.34, 0.66] | 9.93e+03 | 1.000 | 41339.0000
## yrs_mus_exp | 0.0022 | 0.0079 | [-0.01, 0.02] | 0.083 | 1.000 | 40379.0000

```

Strong evidence against an effect of years of musical experience.

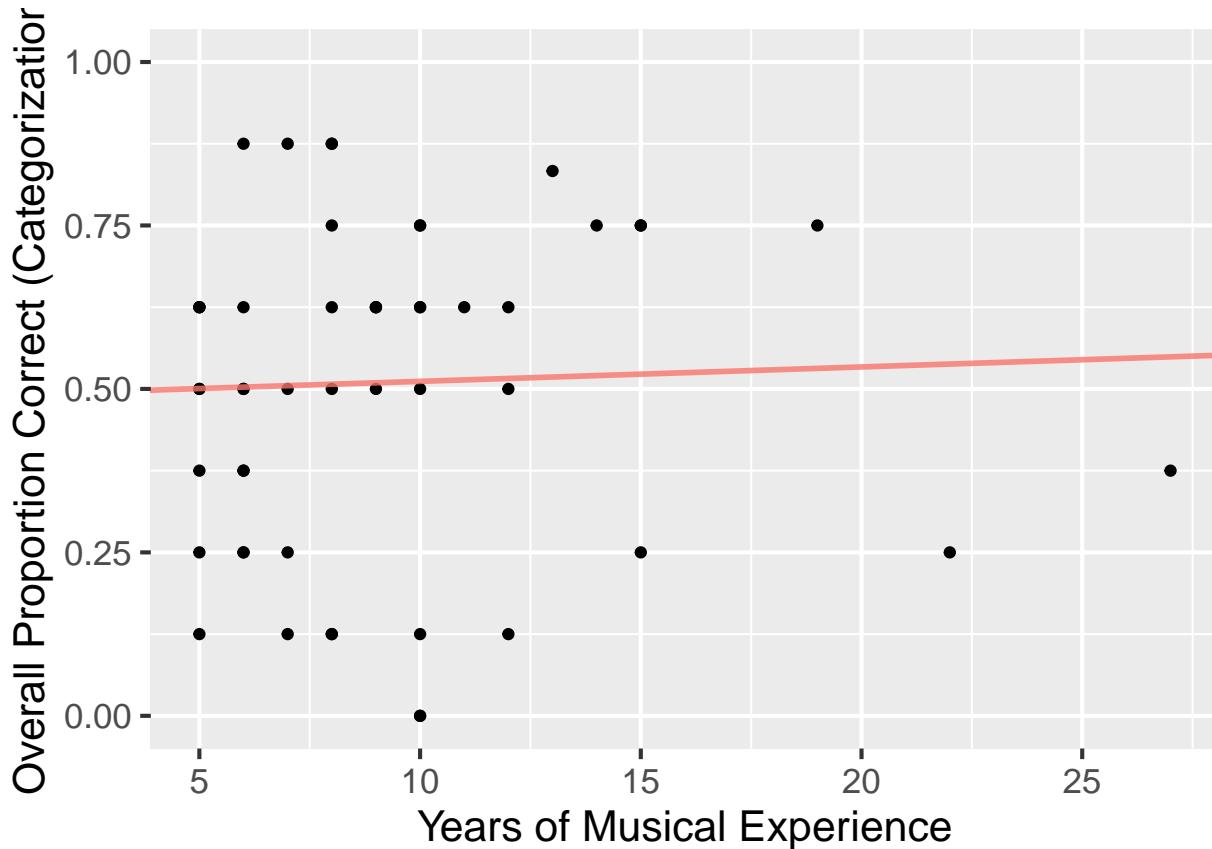
Figure S1C

```

yrs_exp %>%
  group_by(exp_subject_id, yrs_mus_exp) %>%
  summarize(mean_acc = mean(accuracy)) %>%
  ggplot(aes(yrs_mus_exp, mean_acc)) +
  geom_point() +
  geom_abline(intercept = yrs_null_BF$Median[1], slope = yrs_null_BF$Median[2],
              color = '#F8766D', linewidth = 1, alpha = 0.8) +
  xlab('Years of Musical Experience') +
  ylab('Overall Proportion Correct (Categorization)') +
  scale_x_continuous(breaks = seq(5,30,5)) +
  scale_y_continuous(breaks = seq(0, 1, 0.1)) +
  ylim(0,1) +
  theme_gray(base_size = 16)

## `summarise()` has grouped output by 'exp_subject_id'. You can override using
## the `.groups` argument.
## Scale for y is already present. Adding another scale for y, which will replace
## the existing scale.

```



```
ggsave('..../figures/FigS1C_categorization.png', width = 5, height = 5)
```