E1 memory

R. Cassano-Coleman

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

Set up

```
set.seed(15000)
data <- read_csv('../data/memory.csv')</pre>
## Rows: 3210 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(exp_subject_id = as.factor(exp_subject_id),
         response = ifelse(response == 'Correct', TRUE, FALSE),
         scramble = factor(scramble, levels = c('8B', '2B', '1B')),
         Musician = factor(Musician, levels = c('Yes', 'No'))) %>%
  filter(!is.na(response))
Set the contrast for condition.
contrasts(data$scramble) <- contr.treatment(3)</pre>
print(contrasts(data$scramble))
      2.3
## 8B 0 0
## 2B 1 0
## 1B 0 1
```

Main analysis

Priors

Priors are expressed in log(odds) space.

Intercept: Given that chance is 50%, we assume that participants will perform somewhere between chance and ceiling. We expect the center of the distribution of accuracy to be somewhere around 75% or 80%. If we use a center of 80% and an SD of 1, 95% of the values fall between 35.1% and 96.7%.

```
prior_intercept <- set_prior('normal(log(0.8 / (1 - 0.8)), 1)', class = 'Intercept')</pre>
```

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')</pre>
```

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_scramble2B <- set_prior('normal(-0.1, 1)', coef = 'scramble2')
prior_scramble1B <- set_prior('normal(-0.2, 1)', coef = 'scramble3')</pre>
```

Interaction: We expect no interaction between group and scramble.

```
prior_int2B <- set_prior('normal(0, 1)', coef = 'Musician1:scramble2')
prior_int1B <- set_prior('normal(0, 1)', coef = 'Musician1:scramble3')</pre>
```

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,</pre>
              family = bernoulli(),
              prior = c(prior_intercept, prior_mus,
                     prior_scramble2B, prior_scramble1B),
              save_pars = save_pars(all = TRUE), iter = 5000,
              file = '../models/E1_mus_scram')
plot(mus_scram)
              1.4 1.6 0 500 1000 1500 2000 2500
                           0.0 = 1000 1000 1500 2000 2500
                                                                    Chain
              _ 4
print(summary(mus scram), digits = 4)
## Warning: There were 1 divergent transitions after warmup. Increasing
## adapt_delta above 0.8 may help. See
## http://mc-stan.org/misc/warnings.html#divergent-transitions-after-warmup
## Family: bernoulli
   Links: mu = logit
## Formula: response ~ Musician + scramble + (1 | exp_subject_id)
     Data: data (Number of observations: 3153)
##
    Draws: 4 chains, each with iter = 5000; warmup = 2500; thin = 1;
          total post-warmup draws = 10000
##
##
## Multilevel Hyperparameters:
## ~exp_subject_id (Number of levels: 102)
             Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sd(Intercept) 0.1226 0.0706 0.0069 0.2655 1.0007
                                                       2427
                                                               4004
##
```

```
## Regression Coefficients:
##
            Estimate Est.Error 1-95% CI u-95% CI
                                                  Rhat Bulk ESS Tail ESS
                        0.0790 1.2742
                                          1.5821 1.0001
## Intercept 1.4241
                                                             8603
                        0.0429 -0.2182 -0.0500 1.0000
                                                            10909
                                                                      6935
## Musician1 -0.1336
## scramble2 -0.6391
                        0.1007 -0.8379 -0.4433 1.0004
                                                             9841
                                                                      7805
## scramble3 -0.9246
                        0.0987 -1.1192 -0.7340 1.0006
                                                            10169
                                                                      7911
## 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")</pre>
summary(emm_mus_scram_s)
   scramble emmean lower.HPD upper.HPD
## 8B
              1.423
                        1.277
                                  1.584
## 2B
              0.785
                        0.658
                                  0.927
## 1B
              0.500
                        0.376
                                  0.628
## 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
## 8B - 2B
               0.640
                         0.442
                                    0.836
## 8B - 1B
                0.925
                          0.732
                                    1.116
                                    0.466
## 2B - 1B
                0.285
                          0.110
## 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"))</pre>
summary(emm_mus_scram_ms)
## Musician scramble emmean lower.HPD upper.HPD
## Yes
            8B
                     1.557
                                 1.388
                                           1.746
## No
            8B
                       1.289
                                 1.121
                                           1.467
## Yes
            2B
                       0.919
                                 0.766
                                           1.085
## No
            2B
                       0.652
                                 0.488
                                           0.804
                                 0.480
## Yes
            1B
                       0.633
                                           0.783
## No
                       0.365
                                 0.208
                                           0.512
            1B
## 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
## Yes 8B - No 8B
                     0.2662
                                 0.101
                                           0.437
## Yes 8B - Yes 2B
                                           0.836
                     0.6402
                                 0.442
## Yes 8B - No 2B
                     0.9051
                                0.655
                                           1.181
```

```
## Yes 8B - Yes 1B
                                0.732
                                          1.116
                     0.9247
  Yes 8B - No 1B
                                          1.460
##
                     1.1920
                                0.940
  No 8B - Yes 2B
                     0.3705
                                          0.627
                                0.117
##
  No 8B - No 2B
                     0.6402
                                0.442
                                          0.836
  No 8B - Yes 1B
##
                     0.6575
                                0.416
                                          0.919
## No 8B - No 1B
                     0.9247
                                0.732
                                          1.116
## Yes 2B - No 2B
                     0.2662
                                0.101
                                          0.437
## Yes 2B - Yes 1B
                     0.2845
                                0.110
                                          0.466
##
   Yes 2B - No 1B
                     0.5528
                                0.315
                                          0.807
## No 2B - Yes 1B
                     0.0201
                               -0.221
                                          0.265
## No 2B - No 1B
                     0.2845
                                0.110
                                          0.466
## Yes 1B - No 1B
                     0.2662
                                0.101
                                          0.437
##
```

^{##} Point estimate displayed: median

^{##} Results are given on the log odds ratio (not the response) scale.

^{##} HPD interval probability: 0.95

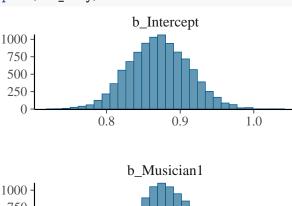
Main effects

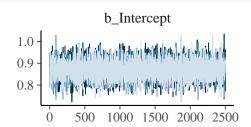
```
main_BF <- describe_posterior(mus_scram,</pre>
                             estimate = "median", dispersion = TRUE,
                             ci = .95, ci_method = "HDI",
                             test = c("bayes_factor"))
## Warning: Bayes factors might not be precise.
##
     For precise Bayes factors, sampling at least 40,000 posterior samples is
    recommended.
print(main_BF, digits = 4)
## Summary of Posterior Distribution
##
## Parameter | Median |
                            MAD |
                                     95% CI | BF | Rhat |
                                                                            ESS
## (Intercept) | 1.4226 | 0.0794 | [ 1.28, 1.58] | 1.41e+19 | 1.000 | 8576.0000
## Musician1 | -0.1331 | 0.0428 | [-0.22, -0.05] | 4.49 | 1.000 | 10880.0000
             | -0.6402 | 0.1010 | [-0.84, -0.44] | 3.74e+05 | 1.000 | 9833.0000
## scramble2
             | -0.9247 | 0.0999 | [-1.12, -0.73] | 9.22e+07 | 1.000 | 10152.0000
## scramble3
```

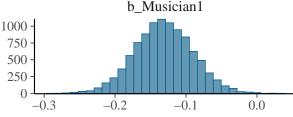
Moderate evidence for a main effect of group.

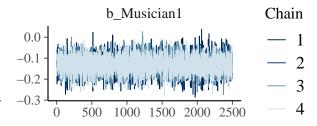
```
To get the main effect of scramble level, fit the "null" model with group only to compare.
```

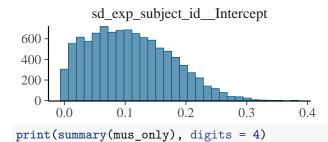
plot(mus_only)

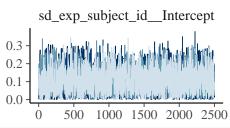












```
## Warning: There were 2 divergent transitions after warmup. Increasing
## adapt_delta above 0.8 may help. See
## http://mc-stan.org/misc/warnings.html#divergent-transitions-after-warmup
##
   Family: bernoulli
##
     Links: mu = logit
## Formula: response ~ Musician + (1 | exp_subject_id)
##
      Data: data (Number of observations: 3153)
##
     Draws: 4 chains, each with iter = 5000; warmup = 2500; thin = 1;
            total post-warmup draws = 10000
##
##
## Multilevel Hyperparameters:
  ~exp_subject_id (Number of levels: 102)
##
                 Estimate Est.Error 1-95% CI u-95% CI
                                                         Rhat Bulk_ESS Tail_ESS
## sd(Intercept)
                   0.1084
                             0.0668
                                      0.0057
                                               0.2485 1.0006
                                                                  2728
                                                                            5074
##
## Regression Coefficients:
```

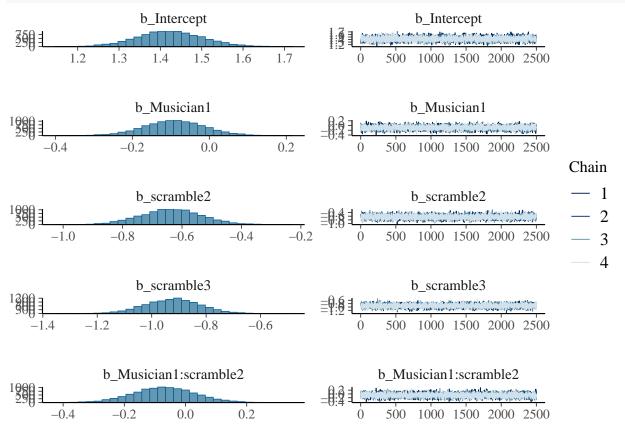
```
Estimate Est.Error 1-95% CI u-95% CI
                                                   Rhat Bulk_ESS Tail_ESS
## Intercept 0.8708
                        0.0412 0.7918
                                          0.9521 1.0006
                                                            12542
                                                                      6331
## Musician1 -0.1314
                        0.0412 -0.2125 -0.0505 1.0008
                                                            12688
                                                                      6331
##
## 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)</pre>
## Iteration: 1
## Iteration: 2
## Iteration: 3
## Iteration: 4
## Iteration: 5
## Iteration: 6
## Iteration: 1
## Iteration: 2
## Iteration: 3
## Iteration: 4
## Iteration: 5
print(BF_scramble)
```

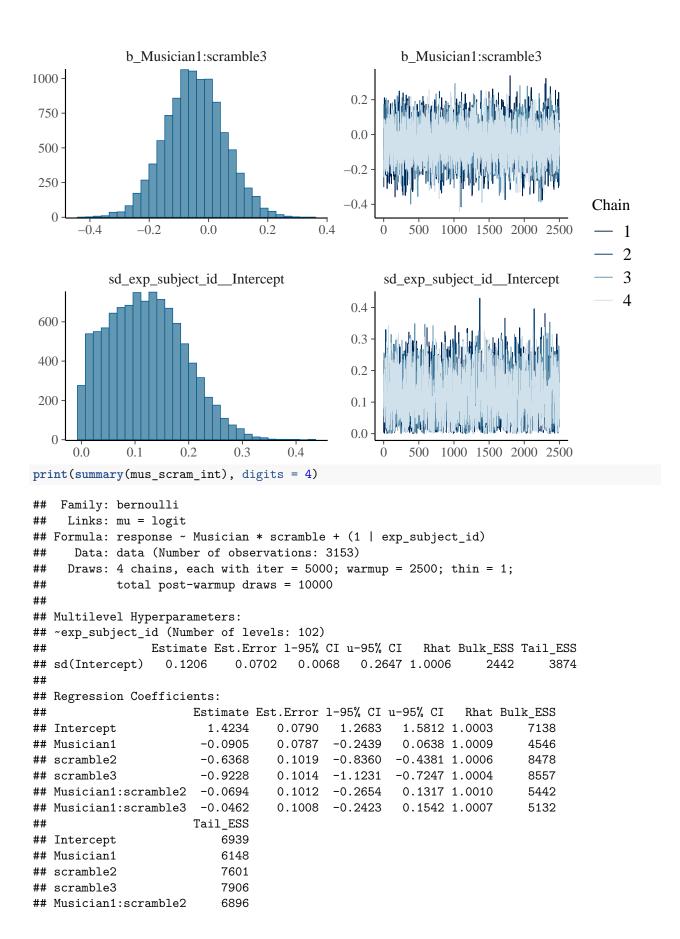
Estimated Bayes factor in favor of mus_scram over mus_only: 477225714046893824.00000 Very strong evidence for a main effect of scramble condition.

Interaction between group and condition?

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

plot(mus_scram_int)





```
## Musician1:scramble3
                           6924
##
## 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)</pre>
## Iteration: 1
## Iteration: 2
## Iteration: 3
## Iteration: 4
## Iteration: 5
## Iteration: 1
## Iteration: 2
## Iteration: 3
## Iteration: 4
## Iteration: 5
## Iteration: 6
print(BF_int)
```

Estimated Bayes factor in favor of mus_scram_int over mus_scram: 0.01095 Moderate evidence against an interaction between group and condition.

Figure 2A

generated.

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

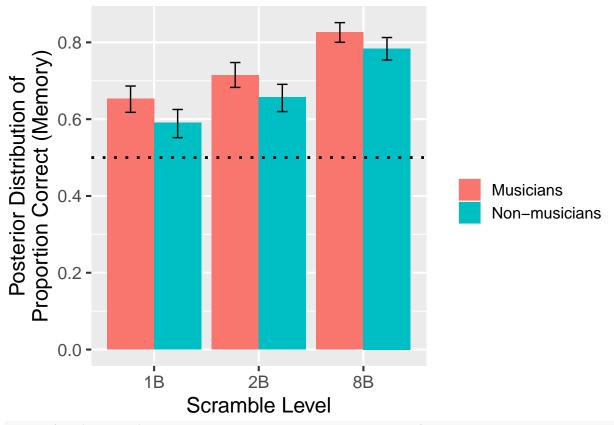
```
calculate_prob_from_logodds <- function(logodds) {
  return(exp(logodds) / (1 + exp(logodds)))
}</pre>
```

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

This warning is displayed once every 8 hours.

```
posterior_est <- as.data.frame(emm_mus_scram_ms)</pre>
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 = 0.5, linetype = "dotted", color = "black", size = 1) +
  theme_gray(base_size = 16) +
  scale_x_discrete(limits = rev) +
  xlab('Scramble Level') +
  ylab('Posterior Distribution of\nProportion Correct (Memory)') +
  scale_fill_discrete(name="", labels=c('Musicians', 'Non-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
## Warning: Using `size` aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use `linewidth` instead.
```

Call `lifecycle::last_lifecycle_warnings()` to see where this warning was



ggsave('../figures/Fig2A_memory.png', width = 7, height = 5)

Years of experience

Priors

Figure S1A