

FB_demo

Chi Zhang

2025-12-09

Packages

Data

This demo is used to replicate (and potentially optimise) the data analysis in Stengelin, R., Petrović, L., Thiele, M., Hepach, R., & Haun, D. B. (2024). Social reward predicts false belief understanding in Namibian Hai||om children. Social Development, 33(4), e12767. Lets' first get familiar with the dataset. The Independent Variable is FB (false belief), binary with 1 stands for passing the test and 0 fail. social touch from 0-8 represent how much time children touch on pad to see more adult/peer, as social rewards. We are trying to argue that social rewards (and what kind of social rewards) predicts children passing FB test.

```
data <- read.csv("data/sode12767-sup-0002-suppmat.csv",
  header = TRUE,
  dec = ",",
  sep = ";",
  stringsAsFactors = FALSE)
```

data

##	ID	Age	Sex	FB	SocialTouches_Adult	SocialTouches_Peer	Caregiver_Presence
## 1	1	2.97	M	1	4	4	1
## 2	9	2.31	M	1	3	4	1
## 3	13	3.33	M	0	3	3	1
## 4	25	5.19	F	1	4	4	0
## 5	17	4.2	F	0	4	4	1
## 6	4	3.14	M	0	3	4	1
## 7	20	2.28	F	0	4	4	1
## 8	16	3.68	M	1	4	4	1
## 9	8	4.59	M	0	4	4	0
## 10	12	4.71	M	1	5	2	0
## 11	28	3.35	F	0	4	5	1
## 12	11	4.49	M	0	4	4	1
## 13	3	4.22	M	1	4	4	1
## 14	15	2.64	M	0	4	3	1
## 15	23	4.3	F	1	4	4	1
## 16	2	5.86	M	0	4	4	1
## 17	6	4.51	M	0	0	5	1
## 18	30	4.39	F	1	4	4	0
## 19	18	2.9	F	0	4	6	1
## 20	26	3.82	F	1	8	6	1

## 21	29	4.9	F	1	5	3	1
## 22	14	4.61	M	1	3	4	0
## 23	21	5.56	F	1	4	5	1
## 24	5	5.86	M	0	4	4	0
## 25	19	5.79	F	0	4	4	0
## 26	27	5.36	F	1	4	4	1
## 27	100	5.24	M	0	4	4	0
## 28	180	7.37	F	1	4	4	0
## 29	260	7.24	F	1	4	3	0
## 30	200	6.97	M	1	4	2	0
## 31	3000	7.35	F	0	3	4	0
## 32	600	7.8	M	1	4	4	0
## 33	700	6.94	M	1	5	3	0
## 34	310	6.96	F	1	8	8	0
## 35	150	7.16	M	1	3	4	0
## 36	230	7.53	F	1	5	3	0
## 37	270	6.82	F	1	4	3	0
## 38	320	7.53	F	1	4	4	0
## 39	120	6.65	M	1	4	0	0
## 40	240	7.38	F	1	6	5	0
## 41	160	7.08	M	1	4	4	0
## 42	400	7.07	M	1	3	4	0
## 43	200	7.12	F	1	4	5	0
## 44	800	7.54	M	1	5	5	0
## 45	140	7.79	M	1	4	4	0
## 46	900	7.28	M	1	4	2	0
## 47	170	5.84	F	0	2	4	1
## 48	250	4.56	F	0	4	4	1
## 49	210	5.98	F	0	4	4	1
## 50	500	7.95	M	1	4	4	0
## 51	300	8.01	M	1	4	4	0
## 52	7	5.4	M	0	NA	4	0
## 53	190	3.84	F	0	3	7	0
## 54	24	3.88	F	1	4	4	0
## 55	280	4.53	F	1	4	4	0
## 56	32	5.67	F	0	5	3	0
## 57	111	3.62	M	0	3	5	0
## 58	133	5.83	M	1	5	2	0
## 59	220	4.62	F	1	4	4	0

In the original paper, the authors preprocessed the data in an unorthodox way: specifically, they standardized both sex and age. This is understandable from an engineering perspective, as it facilitates faster MCMC sampling. However, it renders the model output considerably harder to interpret. With standardized variables, one can no longer make statements such as “for each additional year of age, children are x% more likely to pass the test,” because the units of age and sex no longer carry natural meaning. Instead, one is limited to statements like “for one standard deviation increase in age...”—which is less intuitive. Here I name a `age.C` as age - mean age, that age variable is still centering, yet each unit of the variable is still naturally 1 year. And for sex, I make it a factor variable, but 0 point stands for average of male and female, which help make the intercept in the output understandable. I like the idea to make social touch as `mo()`, here I keep it as it is in original settings.

```
data$ID <- as.factor(data$ID)
data$Sex <- as.factor(data$Sex)
contrasts(data$Sex) <- contr.sum(2)
```

```

data$social.adult <- as.numeric(data$SocialTouches_Adult)
data$social.peer <- as.numeric(data$SocialTouches_Peer)
data$FB <- as.numeric(data$FB)
data$Age <- as.numeric(data$Age)
data$age.C <- data$Age - mean(data$Age)

```

Prior

I also re-assign reasonable prior. For intercept, now we are expecting “an average child around 5.5 years old with 0 score in touching task, would have x% chance to pass the FB test”. I agree to set it at (0, 1.5), this means the base probability is around 5%~95%. While for age prior, the original setting is normal(0, 0.375), this is less intuitive. In the original setting, after scaling, one SD of age is around 1.6 year, thus the OD for each year grow is around $\exp(0.375/1.6) = 1.3$. This is a very narrow prior, since we can expect children suddenly grow in one year (normally find in 3-5 years old as a milestone?), thus the original setting is about to say “I don’t quite believe that children can grow so fast in one year”. I expand a bit to (0,1) for age.C, means I am not surprised to see if the odds of passing rise 3 times for each year grow, which is more reasonable. For social rewards, which is specified as a monotonic effect, I set the prior as Normal(0, 1). The original prior of Normal(0, 1.5) is overly diffuse when considered in context. Taking two standard deviations as a reasonable upper bound (i.e., a coefficient of 3), this implies that children at the highest level of social reward (8) compared to those at the lowest level (0) could differ by a factor of approximately 20 in their odds of passing the task ($e^3 \approx 20$). Such an extreme effect size is implausible for a cognitive task of this nature. A tighter prior of Normal(0, 1) constrains the estimates to a more realistic range while still remaining weakly informative.

```

common_prior <- c(
  set_prior("normal(0, 1.5)", class = "Intercept"),
  set_prior("normal(0, 1)", class = "b", coef = "age.C"),
  set_prior("normal(0, 0.5)", class = "b", coef = "Sex1")
)

adult_prior <- common_prior +
  set_prior("normal(0, 1)", class = "b", coef = "mosocial.adult")

peer_prior <- common_prior +
  set_prior("normal(0, 1)", class = "b", coef = "mosocial.peer")

null_prior <- c(
  set_prior("normal(0, 1.5)", class = "Intercept"),
  set_prior("normal(0, 0.5)", class = "b", coef = "Sex1")
)

```

Multiple imputation

There was one missing value in the social touch (adult) variable. Ordinarily, I would perform multiple imputation to generate several completed datasets (e.g., five) and then fit the model using `brm_multiple()` to properly account for imputation uncertainty. However, given that this is a demonstration and the proportion of missing data is negligible, I opted to impute a single completed dataset for simplicity.

```
imp <- mice(data, m = 1, method = 'pmm', maxit = 5, seed = 123)

##
## iter imp variable
## 1 1 SocialTouches_Adult*
## 2 1 SocialTouches_Adult*
## 3 1 SocialTouches_Adult*
## 4 1 SocialTouches_Adult*
## 5 1 SocialTouches_Adult*

## Warning: Number of logged events: 14

data_imputed <- complete(imp)
data_imputed$social.adult <- as.numeric(data_imputed$SocialTouches_Adult)
```

Fitting model

The iteration number is reduced to 2000 for quicker sampling, but I find this is more than enough as all models converge well.

```
# Model 1: Adult Social Reward + Age + Sex
FB_model_adult_age <- brm(
  formula = FB ~ mo(social.adult) + age.C + Sex,
  data = data_imputed,
  prior = adult_prior,
  family = bernoulli(link = "logit"),
  # Demo Mode Settings
  cores = 4,
  warmup = 1000,
  iter = 2000,
  backend = "cmdstanr",
  control = list(adapt_delta = 0.99),
  sample_prior="yes",
  seed = 1234,
  file = "models/demo_FB_model_adult_age"
)

# Model 2: Peer Social Reward + Age + Sex
FB_model_peer_age <- brm(
  formula = FB ~ mo(social.peer) + age.C + Sex,
  data = data_imputed,
  prior = peer_prior,
  family = bernoulli(link = "logit"),
  cores = 4,
  warmup = 1000,
  iter = 2000,
  backend = "cmdstanr",
  control = list(adapt_delta = 0.99),
  seed = 1234,
  file = "models/demo_FB_model_peer_age"
)
```

```

# Model 3: Age Only (+ Sex control)
FB_model_age <- brm(
  formula = FB ~ age.C + Sex,
  data = data_imputed,
  prior = common_prior,
  family = bernoulli(link = "logit"),
  cores = 4,
  warmup = 1000,
  iter = 2000,
  backend = "cmdstanr",
  control = list(adapt_delta = 0.99),
  seed = 1234,
  file = "models/demo_FB_model_age"
)

# Model 4: Null Model (Only Sex control)
FB_model_null <- brm(
  formula = FB ~ Sex,
  data = data_imputed,
  prior = null_prior,
  family = bernoulli(link = "logit"),
  cores = 4,
  warmup = 1000,
  iter = 2000,
  backend = "cmdstanr",
  control = list(adapt_delta = 0.99),
  seed = 1234,
  file = "models/demo_FB_model_null"
)

```

Results

Now we can better interpret our results. General patterns are the same as the original paper: we find that age significantly predicts (or more faithful to Bayesian genre, say it is a non-zero positive effect) children FB, and social reward (adult) also serve as a positive effect, though this is quite marginal [0.04, 1.36]. Beyond that, we can make more intuitive statements (that is really good for public communication and advertisement!) to say:

For an average 5.5-year-old child who shows extreme indifference in the adult social reward task (i.e., zero button presses), the model predicts a probability of passing the false belief (FB) test of only average around 24% (Intercept = -1.16). This is well below the sample's average passing rate of 37/59, suggesting that at this age, passing the FB test is challenging for children who lack social motivation toward adults. With each additional year of age, the odds of passing the test are expected to increase by a factor of $e^{0.59} = 1.8$ on average. This is consistent with developmental psychology, which identifies ages 3 to 6 as a critical period for ToM development. Regarding adult social reward, the odds of passing the test are $e^{0.58} = 1.79$ times higher for the most socially motivated children (8 presses) compared to the least socially motivated (0 presses). On average, each additional press of the adult social button corresponds to an approximate 7.5% increase in the odds of passing the FB test.

```
summary (FB_model_adult_age)
```

```
## Family: bernoulli
## Links: mu = logit
## Formula: FB ~ mo(social.adult) + age.C + Sex
## Data: data_imputed (Number of observations: 59)
## Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup draws = 4000
##
## Regression Coefficients:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## Intercept      -1.16      1.00   -3.32    0.61 1.00     3165     2499
## age.C           0.59      0.21    0.20    1.02 1.00     4626     2624
## Sex1            -0.09      0.27   -0.63    0.42 1.00     4794     2821
## mosocial.adult  0.58      0.33    0.04    1.36 1.00     2905     2339
##
## Monotonic Simplex Parameters:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## mosocial.adult1[1]  0.10      0.09    0.00    0.33 1.00     3497     1752
## mosocial.adult1[2]  0.10      0.10    0.00    0.35 1.00     4211     2201
## mosocial.adult1[3]  0.12      0.10    0.00    0.39 1.00     3358     2005
## mosocial.adult1[4]  0.11      0.10    0.00    0.36 1.00     3540     1900
## mosocial.adult1[5]  0.14      0.11    0.00    0.42 1.00     3734     2113
## mosocial.adult1[6]  0.15      0.12    0.00    0.46 1.00     4246     2194
## mosocial.adult1[7]  0.14      0.12    0.00    0.45 1.00     4512     2623
## mosocial.adult1[8]  0.14      0.12    0.00    0.44 1.00     3898     2566
##
## Draws were sampled using sample(hmc). 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).
```

```
summary (FB_model_peer_age)
```

```
## Family: bernoulli
## Links: mu = logit
## Formula: FB ~ mo(social.peer) + age.C + Sex
## Data: data_imputed (Number of observations: 59)
## Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup draws = 4000
##
## Regression Coefficients:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## Intercept       0.99      0.90   -0.82    2.79 1.00     2317     2218
## age.C           0.60      0.20    0.22    1.02 1.00     3668     2506
## Sex1            0.02      0.27   -0.53    0.55 1.00     4793     2721
## mosocial.peer   -0.09      0.22   -0.53    0.37 1.00     2544     2278
##
## Monotonic Simplex Parameters:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## mosocial.peer1[1]  0.12      0.11    0.00    0.40 1.00     3655     1841
## mosocial.peer1[2]  0.12      0.11    0.00    0.41 1.00     3633     1973
## mosocial.peer1[3]  0.13      0.11    0.00    0.41 1.00     4066     2227
```

```
## mosocial.peer1[4]      0.12      0.11      0.00      0.38 1.00      3568      2053
## mosocial.peer1[5]      0.12      0.11      0.00      0.40 1.00      4349      2221
## mosocial.peer1[6]      0.13      0.11      0.00      0.41 1.00      3506      1979
## mosocial.peer1[7]      0.13      0.11      0.00      0.41 1.00      4478      2470
## mosocial.peer1[8]      0.13      0.12      0.00      0.44 1.00      4152      2602
##
## Draws were sampled using sample(hmc). 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).
```

```
summary(FB_model_age)
```

```
## Family: bernoulli
## Links: mu = logit
## Formula: FB ~ age.C + Sex
## Data: data_imputed (Number of observations: 59)
## Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup draws = 4000
##
## Regression Coefficients:
##      Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## Intercept      0.64      0.31    0.06    1.28 1.00      2552      2231
## age.C           0.61      0.20    0.24    1.02 1.00      2801      2408
## Sex1           -0.00      0.26   -0.50    0.52 1.00      2399      2026
##
## Draws were sampled using sample(hmc). 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).
```

```
summary(FB_model_null)
```

```
## Family: bernoulli
## Links: mu = logit
## Formula: FB ~ Sex
## Data: data_imputed (Number of observations: 59)
## Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup draws = 4000
##
## Regression Coefficients:
##      Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## Intercept      0.52      0.25    0.03    1.00 1.00      2344      2241
## Sex1           -0.02      0.24   -0.50    0.46 1.00      2135      1700
##
## Draws were sampled using sample(hmc). 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).
```

Prior sensitive analysis

I won't go through interpreting each model, and I really like the visualisation in the original paper, I won't go through these in detail. But one thing the original paper is missing: prior sensitive analysis. It seems ok

to not doing PSA as original paper finds the effect of social rewards adult is 0.67 [0.10, 1.59]. It's ok to think this range is safe. While in my model, it shows signal of danger (0.58 [0.04, 1.36]), it is very marginal as a non-zero effect. If being picky enough, people might ask whether this is because your prior providing extra information, questioning whether the pattern is data-oriented or prior-oriented. Here I give a tight prior at (0, 0.5) and wide prior (0, 1.5) (this is also the setting in the original paper.)

```
common_prior_fixed <- c(
  set_prior("normal(0, 1.5)", class = "Intercept"),
  set_prior("normal(0, 1)", class = "b", coef = "age.C"),
  set_prior("normal(0, 0.5)", class = "b", coef = "Sex1")
)

prior_tight <- common_prior_fixed +
  set_prior("normal(0, 0.5)", class = "b", coef = "mosocial.adult")

model_adult_tight <- brm(
  formula = FB ~ mo(social.adult) + age.C + Sex,
  data = data_imputed,
  prior = prior_tight,
  family = bernoulli(link = "logit"),
  cores = 4, warmup = 1000, iter = 2000, backend = "cmdstanr",
  seed = 1234,
  sample_prior="yes",
  file = "models/sens_model_adult_tight"
)

prior_wide <- common_prior_fixed +
  set_prior("normal(0, 1.5)", class = "b", coef = "mosocial.adult")

model_adult_wide <- brm(
  formula = FB ~ mo(social.adult) + age.C + Sex,
  data = data_imputed,
  prior = prior_wide,
  family = bernoulli(link = "logit"),
  cores = 4, warmup = 1000, iter = 2000, backend = "cmdstanr",
  seed = 1234,
  sample_prior="yes",
  file = "models/sens_model_adult_wide"
)
```

The results shows it passes the sensitive analysis, but still marginal in tight prior (0.44 [0, 0.97]), suggesting one should be more careful when interpreting this pattern.

```
summary(model_adult_tight)
```

```
## Family: bernoulli
## Links: mu = logit
## Formula: FB ~ mo(social.adult) + age.C + Sex
## Data: data_imputed (Number of observations: 59)
## Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup draws = 4000
##
```



```
## Regression Coefficients:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## Intercept      -0.83      0.91    -2.67     0.80 1.00     2228     2321
## age.C           0.58      0.21     0.21     1.01 1.00     3669     2633
## Sex1           -0.08      0.27    -0.62     0.45 1.00     4133     3074
## mosocial.adult   0.44      0.25    -0.00     0.97 1.00     2292     2577
##
## Monotonic Simplex Parameters:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## mosocial.adult1[1]  0.11      0.10     0.00     0.36 1.00     3522     1984
## mosocial.adult1[2]  0.11      0.10     0.00     0.37 1.00     3332     1676
## mosocial.adult1[3]  0.12      0.10     0.00     0.38 1.00     3520     1969
## mosocial.adult1[4]  0.12      0.10     0.00     0.37 1.00     2951     1691
## mosocial.adult1[5]  0.14      0.11     0.01     0.40 1.00     3237     2025
## mosocial.adult1[6]  0.14      0.11     0.00     0.42 1.00     3752     2116
## mosocial.adult1[7]  0.13      0.12     0.00     0.45 1.00     4077     2331
## mosocial.adult1[8]  0.13      0.11     0.00     0.42 1.00     3535     2604
##
## Draws were sampled using sample(hmc). 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).
```

```
summary (model_adult_wide)
```

```
## Family: bernoulli
## Links: mu = logit
## Formula: FB ~ mo(social.adult) + age.C + Sex
## Data: data_imputed (Number of observations: 59)
## Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
## total post-warmup draws = 4000
##
## Regression Coefficients:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## Intercept      -1.23      0.99    -3.31     0.50 1.00     2883     2546
## age.C           0.59      0.21     0.21     1.03 1.00     4588     2823
## Sex1           -0.09      0.27    -0.61     0.43 1.00     3795     2688
## mosocial.adult   0.64      0.39     0.06     1.58 1.00     2214     1465
##
## Monotonic Simplex Parameters:
##           Estimate Est.Error l-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## mosocial.adult1[1]  0.10      0.09     0.00     0.34 1.00     3733     2043
## mosocial.adult1[2]  0.10      0.09     0.00     0.33 1.00     3453     1783
## mosocial.adult1[3]  0.12      0.10     0.00     0.37 1.00     3105     2029
## mosocial.adult1[4]  0.11      0.09     0.00     0.34 1.00     2651     1447
## mosocial.adult1[5]  0.14      0.11     0.00     0.41 1.00     3555     2477
## mosocial.adult1[6]  0.15      0.13     0.00     0.47 1.00     3543     2356
## mosocial.adult1[7]  0.15      0.12     0.00     0.45 1.00     4295     2916
## mosocial.adult1[8]  0.14      0.13     0.00     0.46 1.00     3912     2116
##
## Draws were sampled using sample(hmc). 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).
```

Visualisation

Let's do a nicely plot that I saw in Winter et al (2025) (<https://www.sciencedirect.com/science/article/pii/S002244052400147X>).

```
extract_prior_post <- function(model) {
  draws <- as_draws_df(model)

  post_name <- "bsp_mosocial.adult"
  prior_name <- "prior_bsp_mosocial.adult"

  bind_rows(
    tibble(value = draws[[prior_name]], Type = "Prior"),
    tibble(value = draws[[post_name]], Type = "Posterior")
  )
}

d1 <- extract_prior_post(model_adult_tight)
d2 <- extract_prior_post(FB_model_adult_age)
d3 <- extract_prior_post(model_adult_wide)

plot_beautiful <- function(data, title_text) {

  post_vals <- data$value[data$Type == "Posterior"]

  ci_vals <- quantile(post_vals, probs = c(0.025, 0.975))

  sub_label <- sprintf("95% CI: [%.2f, %.2f]", ci_vals[1], ci_vals[2])

  ggplot(data, aes(x = value, fill = Type, color = Type, linetype = Type)) +
    geom_density(alpha = 0.4, size = 1) +
    scale_fill_manual(values = c("Posterior" = "gray80", "Prior" = "#E69F00")) +
    scale_color_manual(values = c("Posterior" = "gray40", "Prior" = "#E69F00")) +
    scale_linetype_manual(values = c("Posterior" = "solid", "Prior" = "dashed")) +
    geom_vline(xintercept = 0, linetype = "dotted", color = "black", alpha = 0.5) +
    scale_y_continuous(expand = c(0, 0)) +
    coord_cartesian(xlim = c(-2.5, 2.5)) +

    labs(title = title_text,
         subtitle = sub_label,
         x = NULL, y = NULL) +
    theme_classic() +
    theme(
      plot.title = element_text(face = "bold", size = 11),
      legend.position = "none",
      axis.line.y = element_blank(),
      axis.text.y = element_blank(),
      axis.ticks.y = element_blank()
    )
}
```

```
p1 <- plot_beautiful(d1, "1. Tight Prior (SD=0.5)")
```

```
## Warning: Using `size` aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use `linewidth` instead.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
```

```
p2 <- plot_beautiful(d2, "2. Medium Prior (SD=1)")
p3 <- plot_beautiful(d3, "3. Wide Prior (SD=1.5)")
```

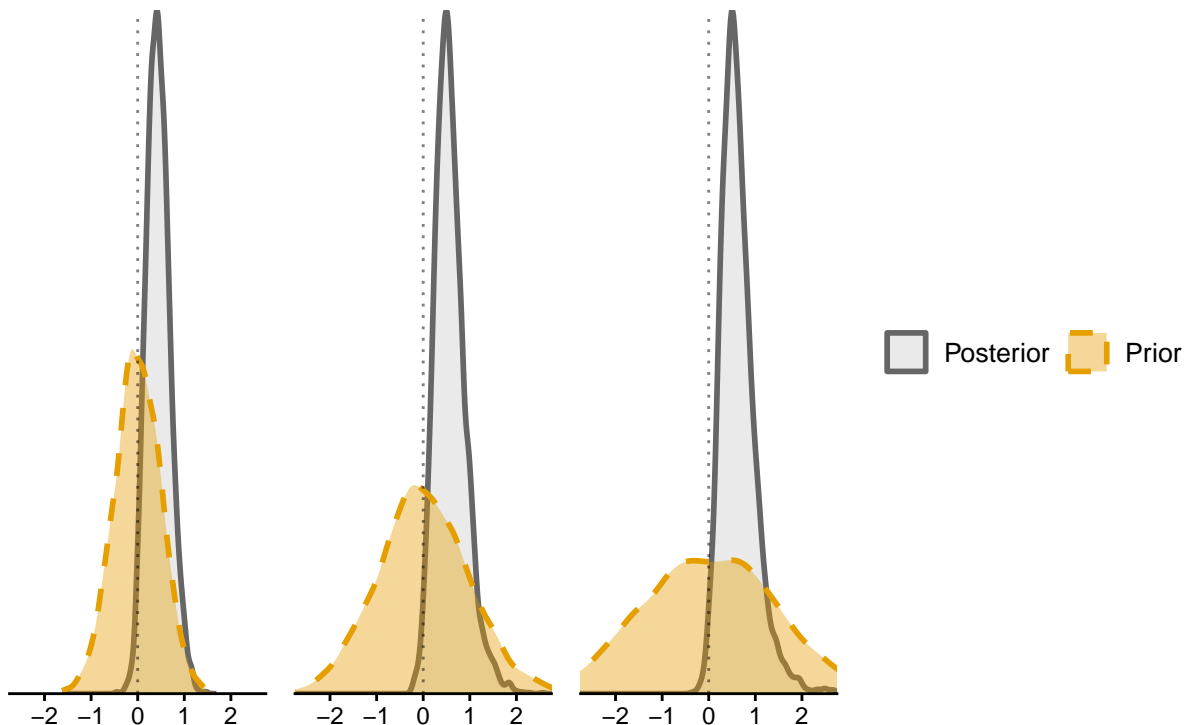
```
p_combined <- (p1 | p2 | p3) +
  plot_layout(guides = "collect") +
  theme(legend.position = "bottom",
        legend.title = element_blank(),

        legend.text = element_text(size = 10))
```

```
print(p_combined)
```

1. Tight Prior (SD=0.5) Medium Prior (SD=1) Wide Prior (SD=1.5)

95% CI: [-0.00, 0.97] 95% CI: [0.04, 1.36] 95% CI: [0.06, 1.58]



```
ggsave("slender_plot.png", p_combined, width = 12, height = 3.5, dpi = 300)
```

Model compare

Finally I'll do a brief model compare. we should be very careful when interpreting the loo compare and weight compare: that we can say FB_model_adult_age is the best model in weight compare, yet these four models provides similar information in terms of loo compare. This is a disappointing signal, suggesting that even the significant variable here is not strong enough to make difference. As a result we can also see Bayes R^2 is very low.

Nevertheless, given the small sample size of 59 for this logistic regression, such a pattern remains satisfactory. In this demo, I have attempted to make the entire analysis process more rigorous and comprehensive.

```
FB_model_adult_age <- add_criterion(FB_model_adult_age, "loo")
FB_model_peer_age  <- add_criterion(FB_model_peer_age, "loo")
FB_model_age       <- add_criterion(FB_model_age, "loo")
FB_model_null      <- add_criterion(FB_model_null, "loo")

loo_compare(FB_model_adult_age, FB_model_peer_age, FB_model_age, FB_model_null)
```

```
##               elpd_diff se_diff
## FB_model_adult_age  0.0         0.0
## FB_model_age       -1.8         1.5
## FB_model_peer_age -2.1         1.5
## FB_model_null     -6.0         3.4
```

```
model_weights(FB_model_adult_age, FB_model_peer_age, FB_model_age, FB_model_null)
```

```
## Warning: Some Pareto k diagnostic values are too high. See help('pareto-k-diagnostic') for details.
```

```
## FB_model_adult_age  FB_model_peer_age      FB_model_age      FB_model_null
##          9.719277e-01          1.454073e-08          8.988824e-07          2.807142e-02
```

```
bayes_R2(FB_model_adult_age)
```

```
##      Estimate Est.Error      Q2.5      Q97.5
## R2 0.2207915 0.06486884 0.08550987 0.3300632
```