

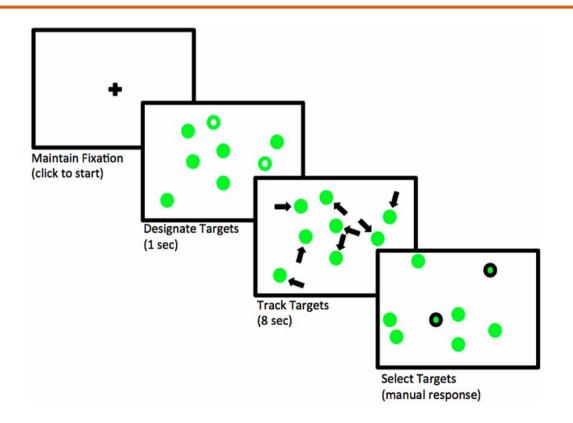
Example: Multiple object tracking

- The subject covertly tracks between zero and five objects among several randomly moving objects on a computer screen.
- First, several objects appear on the screen, and a subset of them are indicated as "targets" at the beginning.
- Then, the objects start moving randomly across the screen and become indistinguishable.
- After several seconds, the objects stop moving and the subject need to indicate which objects were the targets.

Our research goal is to examine how the attentional load affects pupil size.

Bayesian Data Analysis

Example: Multiple object tracking



Bayesian Data Analysis

Example: Multiple object tracking

A model for this experiment design:

$$p_size_n \sim Normal(\alpha + c_load_n \cdot \beta, \sigma)$$
 (1)

- \blacksquare n indicates the observation number with $n=1,\ldots,N$
- lacksquare c_load refers to centered load.
- Every data point is assumed to be independent (in frequentist terms: iid).

Bayesian Data Analysis

Example: Multiple object tracking: Prior specification

Some pilot data helps us work out priors:

```
data("df_pupil_pilot")
df_pupil_pilot$p_size %>% summary()

## Min. 1st Qu. Median Mean 3rd Qu. Max.
## 852 856 862 861 866 868
```

This suggests we can use the following regularizing prior for α :

```
\alpha \sim Normal(1000, 500)
```

(2)

What we are expressing with this prior:

20 1980

```
qnorm(c(.025, .975), mean = 1000, sd = 500)
```

Bayesian Data Analysis

Shravan Vasishth vasishth.github.io

Example: Multiple object tracking: Prior specification

For σ , we use an uninformative prior:

$$\sigma \sim Normal_{+}(0, 1000) \tag{3}$$

```
extraDistr::qtnorm(c(.025,0.975), mean = 0, sd = 1000, a = 0)
## [1] 31 2241
```

Bayesian Data Analysis

Example: Multiple object tracking: Prior specification

$$\beta \sim Normal(0, 100) \tag{4}$$

```
qnorm(c(.025, .975), mean = 0, sd = 100)
## [1] -196  196
```

Bayesian Data Analysis

Example: Multiple object tracking: Fit model

First, center the predictor:

```
data("df_pupil")
(df_pupil <- df_pupil %>%
  mutate(c load = load - mean(load)))
## # A tibble: 41 x 5
       subj trial load p_size c_load
##
      <int> <int> <int> <dbl> <dbl>
##
##
        701
                         1021. -0.439
##
        701
                          951. -1.44
                3
##
    3
        701
                         1064. 2.56
##
    4
        701
                          913. 1.56
    5
##
        701
                          603. -2.44
##
    6
        701
                          826. 0.561
##
        701
                          464. -2.44
```

Bayesian Data Analysis

Shravan Vasishth vasishth.github.io

Example: Multiple object tracking: Fit model

```
fit_pupil <- brm(p_size ~ 1 + c_load,
  data = df_pupil,
  family = gaussian(),
  prior = c(
    prior(normal(1000, 500), class = Intercept),
    prior(normal(0, 1000), class = sigma),
    prior(normal(0, 100), class = b, coef = c_load)
  )
)</pre>
```

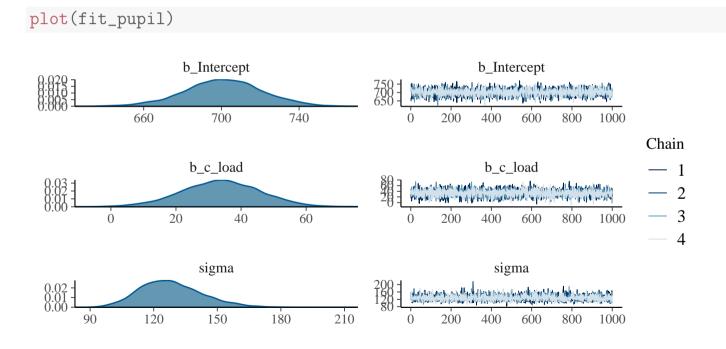
Bayesian Data Analysis

The next steps

Next, we will plot the posterior distributions of the parameters, and the posterior predictive distributions for the different load levels.

Bayesian Data Analysis

Example: Multiple object tracking: Summarize posteriors



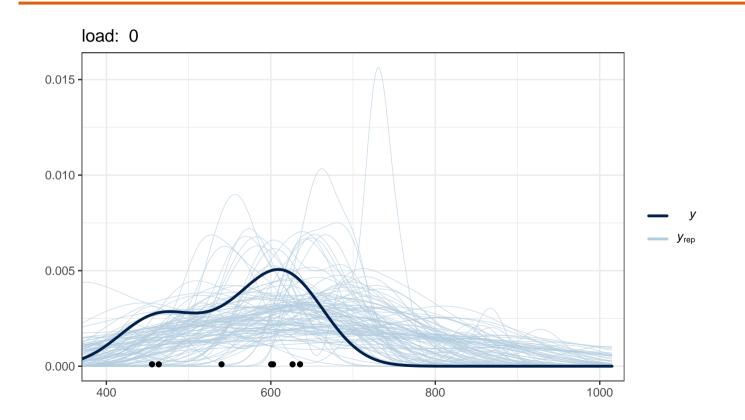
Bayesian Data Analysis

Example: Multiple object tracking: Summarize posteriors

```
## Note: short_summary is a function we wrote
short_summary(fit_pupil)
## ...
## Population-Level Effects:
           Estimate Est. Error 1-95% CI u-95% CI Rhat Bulk ESS Tail ESS
##
## Intercept 701.06
                    20.00 660.47 739.65 1.00
                                                      3761
                                                              2844
## c load 33.85 12.06 9.78 57.08 1.00
                                                              2859
                                                      3639
##
## Family Specific Parameters:
##
        Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
## sigma
          128.19 14.64 102.94 159.66 1.00
                                                  3751
                                                          2937
##
##
```

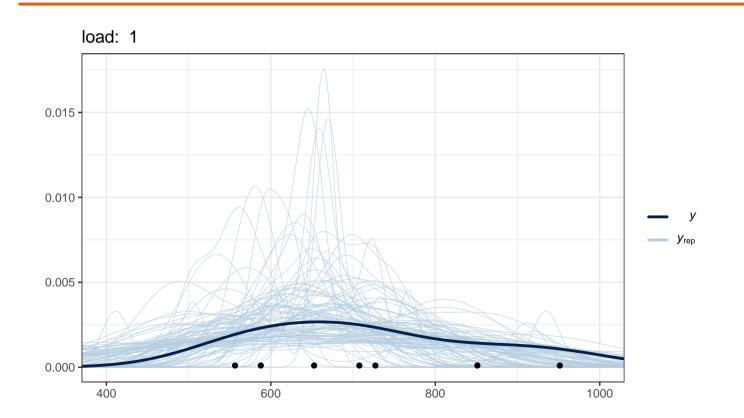
Bayesian Data Analysis

Shravan Vasishth vasishth.github.io



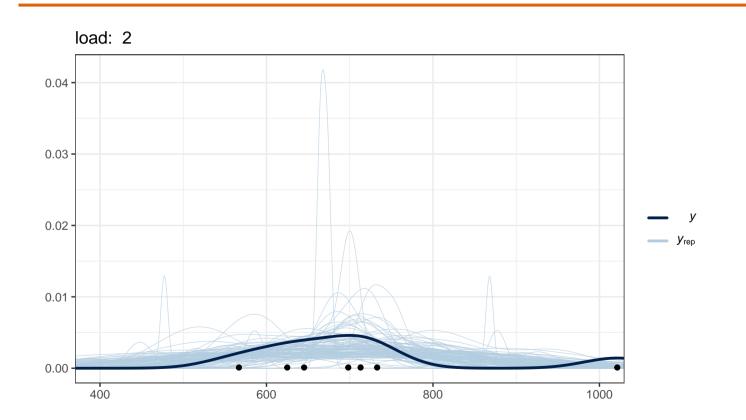
Bayesian Data Analysis

Shravan Vasishth vasishth.github.io



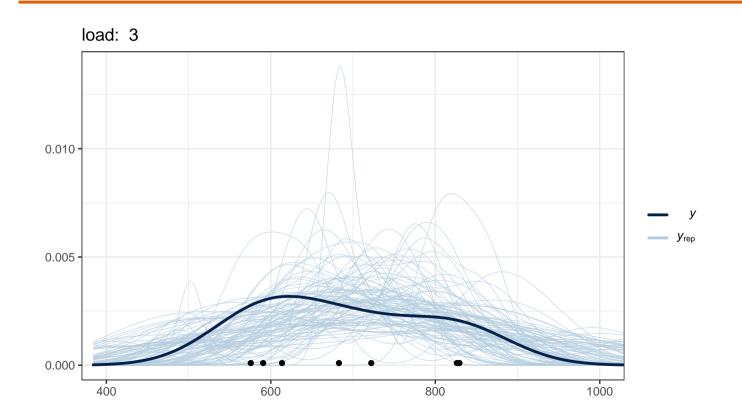
Bayesian Data Analysis

Shravan Vasishth vasishth.github.io



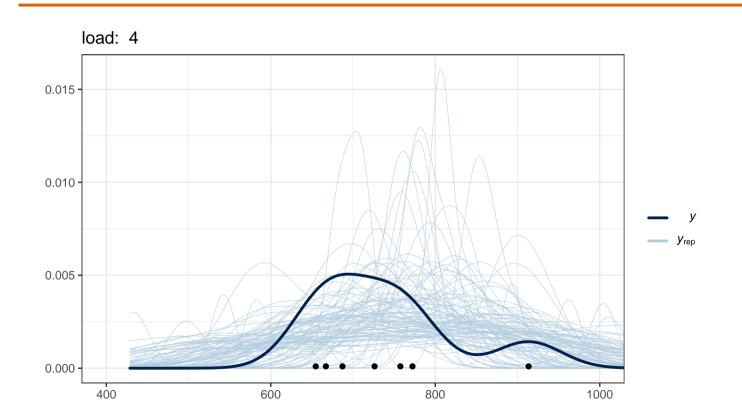
Bayesian Data Analysis

Shravan Vasishth vasishth.github.io



Bayesian Data Analysis

Shravan Vasishth vasishth.github.io



Bayesian Data Analysis

Shravan Vasishth vasishth.github.io

The next steps

Next, we will look at another example: the effect of trial id on button-pressing times. This time, we will use the log-normal likelihood.

Bayesian Data Analysis

```
df_spacebar <- df_spacebar %>%
    mutate(c_trial = trial - mean(trial))
```

Bayesian Data Analysis

If we assume that response times are log-normally distributed, we could proceed as follows:

$$rt_n \sim LogNormal(\alpha + c_trial_n \cdot \beta, \sigma)$$
 (5)

where

- lacksquare N is the total number of (independent!) data points
- $n=1,\ldots,N$, and
- \blacksquare rt is the dependent variable (response times in milliseconds).

Bayesian Data Analysis

The priors have to be defined on the log scale:

$$\alpha \sim Normal(6, 1.5)$$

$$\sigma \sim Normal_{+}(0, 1)$$
(6)

A new parameter, β , needs a prior specification:

$$\beta \sim Normal(0,1) \tag{7}$$

This prior on β is very uninformative.

Bayesian Data Analysis

Prior predictive distribution:

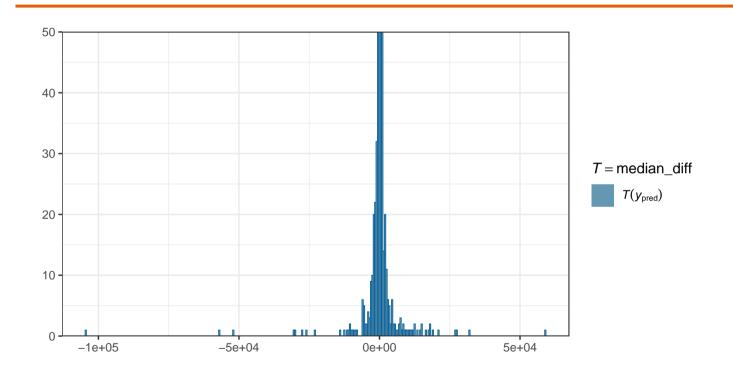
```
df_spacebar_ref <- df_spacebar %>%
  mutate(rt = rep(1, n()))
fit_prior_press_trial <- brm(t ~ 1 + c_trial,</pre>
  data = df_spacebar_ref,
  family = lognormal(),
  prior = c(
    prior(normal(6, 1.5), class = Intercept),
    prior(normal(0, 1), class = sigma),
    prior(normal(0, 1), class = b, coef = c_trial)
  sample_prior = "only",
  control = list(adapt_delta = .9)
```

Bayesian Data Analysis

Shravan Vasishth vasishth.github.io

```
median diff <- function(x) {</pre>
  median(x - lag(x), na.rm = TRUE)
pp_check(fit_prior_press_trial,
         type = "stat",
         stat = "median_diff",
  # show only prior predictive distributions
         prefix = "ppd",
  # each bin has a width of 500ms
         binwidth = 500) +
  # cut the top of the plot to improve its scale
  coord_cartesian(ylim = c(0, 50)) + theme_bw()
```

Bayesian Data Analysis



Bayesian Data Analysis

What would the prior predictive distribution look like if we set the following more informative prior on β ?

$$\beta \sim Normal(0, 0.01) \tag{8}$$

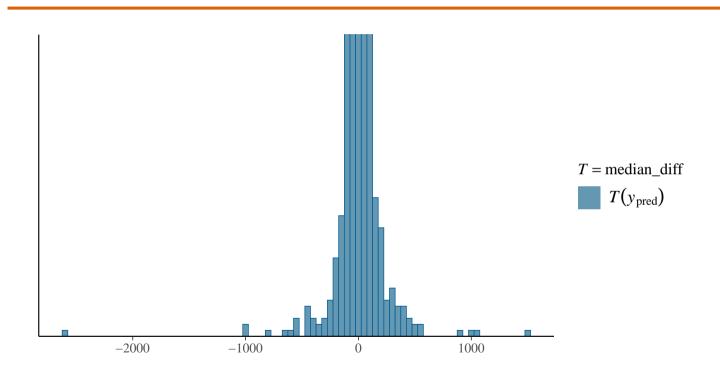
Bayesian Data Analysis

```
fit_prior_press_trial <- brm(t ~ 1 + c_trial,</pre>
  data = df_spacebar_ref,
  family = lognormal(),
  prior = c(
    prior(normal(6, 1.5), class = Intercept),
    prior(normal(0, 1), class = sigma),
    prior(normal(0, .01), class = b, coef = c_trial)
  sample_prior = "only",
  control = list(adapt_delta = .9)
```

Bayesian Data Analysis

```
fit_prior_press_trial <- brm(t ~ 1 + c_trial,</pre>
  data = df_spacebar_ref,
  family = lognormal(),
  prior = c(
    prior(normal(6, 1.5), class = Intercept),
    prior(normal(0, 1), class = sigma),
    prior(normal(0, .01), class = b, coef = c_trial)
  sample_prior = "only",
  control = list(adapt_delta = .9)
```

Bayesian Data Analysis



Bayesian Data Analysis

The next steps

Now that we have decided on our priors, we fit the model.

Bayesian Data Analysis

Fit the model:

```
fit_press_trial <- brm(t ~ 1 + c_trial,
  data = df_spacebar,
  family = lognormal(),
  prior = c(
    prior(normal(6, 1.5), class = Intercept),
    prior(normal(0, 1), class = sigma),
    prior(normal(0, .01), class = b, coef = c_trial)
  )
)</pre>
```

Bayesian Data Analysis

Summarize posteriors (graphically or in a table, or both):

plot(fit_press_trial) b_Intercept b_Intercept 5.1 5.1 5.1 1000 Chain b_c_trial b c trial 4e-04 5e-04 6e - 047e - 041000 3e-04sigma sigma 0.12 0.13 0.14 0.11 1000

Bayesian Data Analysis

Shravan Vasishth vasishth.github.io

Summarize results on the ms scale (the effect estimate from the middle of the expt to the preceding trial):

```
alpha_samples <- as_draws_df(fit_press_trial)$b_Intercept
beta_samples <- as_draws_df(fit_press_trial)$b_c_trial</pre>
beta_ms <- exp(alpha_samples) - exp(alpha_samples - beta_samples)
beta msmean <- round(mean(beta ms), 5)
beta_mslow <- round(quantile(beta_ms, prob = 0.025), 5)
beta_mshigh <- round(quantile(beta_ms, prob = 0.975), 5)
c(beta_msmean , beta_mslow, beta_mshigh)
## 2.5% 97.5%
## 0.087 0.066 0.108
```

Bayesian Data Analysis

Shravan Vasishth vasishth.github.io

The effect estimate at the first vs second trial:

```
first_trial <- min(df_spacebar$c_trial)</pre>
second_trial <- min(df_spacebar$c_trial) + 1</pre>
effect_beginning_ms <-
  exp(alpha_samples + second_trial * beta_samples) -
  exp(alpha_samples + first_trial * beta_samples)
## ms effect from first to second trial:
c(mean = mean(effect_beginning_ms),
  quantile(effect_beginning_ms, c(0.025, 0.975)))
  mean 2.5% 97.5%
## 0.079 0.062 0.096
```

Bayesian Data Analysis

Slowdown after 100 trials from the middle of the expt:

```
effect_100 <-
    exp(alpha_samples + 100 * beta_samples) -
    exp(alpha_samples)

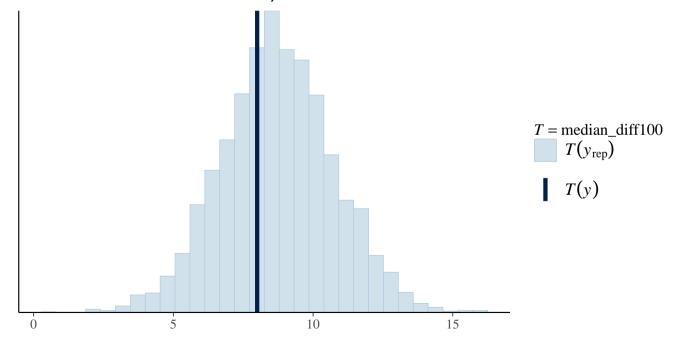
c(mean = mean(effect_100),
    quantile(effect_100, c(0.025, 0.975)))

## mean 2.5% 97.5%

## 9.0 6.7 11.2</pre>
```

Bayesian Data Analysis

The posterior predictive distribution (distribution of predicted median differences between the n and n-100th trial):

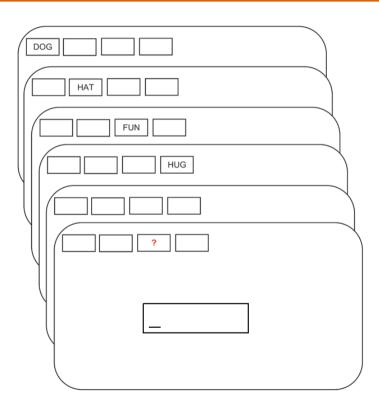


Bayesian Data Analysis

The next steps

Next: logistic regression.

Bayesian Data Analysis



Bayesian Data Analysis

```
data("df recall")
head(df_recall)
## # A tibble: 6 \times 7
##
     subj
           set_size correct trial session block tested
##
     <chr>
              <int>
                      <int> <int> <int> <int> <int>
## 1 10
                                 9
## 3 10
## 4 10
                                23
## 5 10
                                                       5
## 6 10
df_recall <- df_recall %>%
  mutate(c set size = set size - mean(set size))
```

Bayesian Data Analysis

Shravan Vasishth vasishth.github.io

38

```
# Set sizes in the data set:
df_recall$set_size %>%
  unique() %>% sort()
## [1] 2 4 6 8
```

Bayesian Data Analysis

```
# Trials by set size
df recall %>%
  group_by(set_size) %>%
  count()
## # A tibble: 4 \times 2
## # Groups: set_size [4]
##
     set_size
##
        <int> <int>
                  23
## 1
## 2
                  23
## 3
            6
                  23
                  23
## 4
```

Bayesian Data Analysis

$$correct_n \sim Bernoulli(\theta_n)$$
 (9)

$$\eta_n = g(\theta_n) = \log\left(\frac{\theta_n}{1 - \theta_n}\right)$$
(10)

Bayesian Data Analysis

```
x \leftarrow seq(0.001, 0.999, by = 0.001)
y < -\log(x / (1 - x))
logistic_dat <- data.frame(theta = x, eta = y)</pre>
p1 <- qplot(logistic_dat$theta, logistic_dat$eta, geom = "line") +
  xlab(expression(theta)) +
  ylab(expression(eta)) +
  ggtitle("The logit link") +
  annotate("text",
    x = 0.3, y = 4,
    label = expression(paste(eta, "=",
                              g(theta))), parse = TRUE,
    size = 8
```

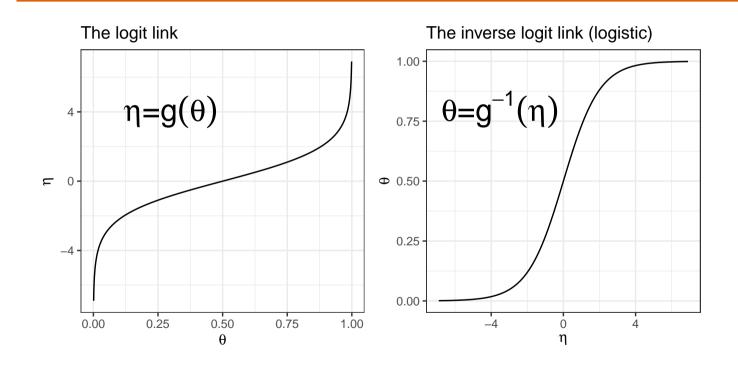
Bayesian Data Analysis

Shravan Vasishth vasishth.github.io

42

```
p2 <- qplot(logistic_dat$eta, logistic_dat$theta,
            geom = "line") + xlab(expression(eta)) +
 ylab(expression(theta)) +
  ggtitle("The inverse logit link (logistic)") +
  annotate("text",
 x = -3.5, y = 0.80,
 label = expression(paste(theta, "=", g^-1,
                           (eta))), parse = TRUE, size = 8
gridExtra::grid.arrange(p1, p2, ncol = 2)
```

Bayesian Data Analysis



Bayesian Data Analysis

The next steps

Next step: deciding on priors.

Bayesian Data Analysis

```
data("df recall")
head(df_recall)
## # A tibble: 6 \times 7
##
     subj
           set_size correct trial session block tested
##
     <chr>
              <int>
                      <int> <int> <int> <int> <int>
## 1 10
                                 9
## 3 10
## 4 10
                                23
## 5 10
                                                       5
## 6 10
df_recall <- df_recall %>%
  mutate(c set size = set size - mean(set size))
```

Bayesian Data Analysis

Shravan Vasishth vasishth.github.io

46

The linear model is now fit not to the 0,1 responses as the dependent variable, but to η_n , i.e., log-odds, as the dependent variable:

$$\eta_n = \log\left(\frac{\theta_n}{1 - \theta_n}\right) = \alpha + \beta \cdot c_set_size$$
(11)

Bayesian Data Analysis

- Unlike the linear models, the model is defined so that there is no residual error term (ε) in this model.
- Once η_n is estimated, one can solve the above equation for θ_n (in other words, we compute the inverse of the logit function and obtain the estimates on the probability scale).

This gives the above-mentioned logistic regression function:

$$\theta_n = g^{-1}(\eta_n) = \frac{\exp(\eta_n)}{1 + \exp(\eta_n)} = \frac{1}{1 + \exp(-\eta_n)}$$
(12)

Bayesian Data Analysis

In summary, the generalized linear model with the logit link fits the following Bernoulli likelihood:

$$correct_n \sim Bernoulli(\theta_n)$$
 (13)

- The model is fit on the log-odds scale, $\eta_n = \alpha + c_set_size_n \cdot \beta$.
- Once η_n has been estimated, the inverse logit or the logistic function is used to compute the probability estimates $\theta_n = \frac{\exp(\eta_n)}{1+\exp(\eta_n)}$.

Bayesian Data Analysis

There are two functions in R that implement the logit and inverse logit functions:

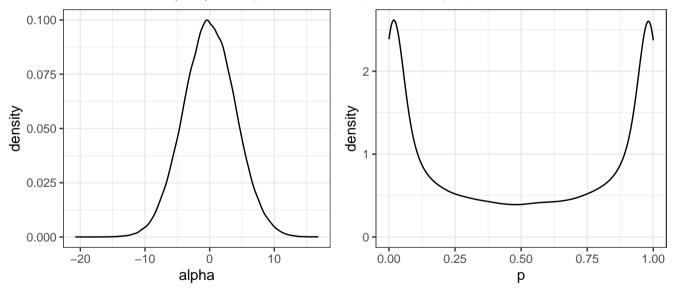
- qlogis(p) for the logit function and
- plogis(x) for the inverse logit or logistic function.

$$\alpha \sim Normal(0,4) \tag{14}$$

Let's plot this prior in log-odds and in probability scale by drawing random samples.

Bayesian Data Analysis

Prior for $\alpha \sim Normal(0,4)$ in log-odds and in probability space.



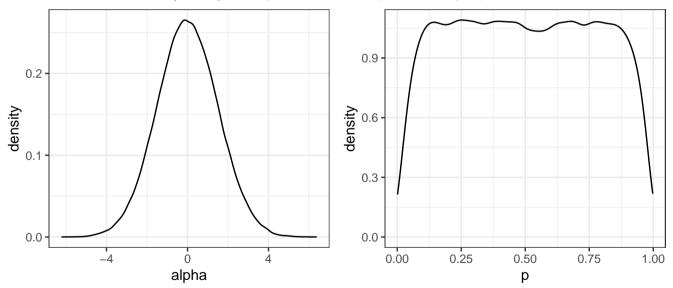
Bayesian Data Analysis

 $\alpha \sim Normal(0, 1.5)$

(15)

Bayesian Data Analysis

Prior for $\alpha \sim Normal(0, 1.5)$ in log-odds and in probability space.



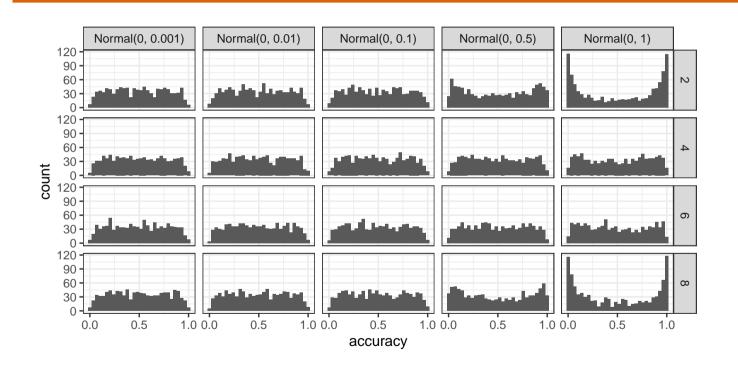
Bayesian Data Analysis

We can examine the consequences of each of the following prior specifications:

- 1. $\beta \sim Normal(0,1)$
- 2. $\beta \sim Normal(0, .5)$
- **3.** $\beta \sim Normal(0, .1)$
- **4.** $\beta \sim Normal(0, .01)$
- **5.** $\beta \sim Normal(0, .001)$

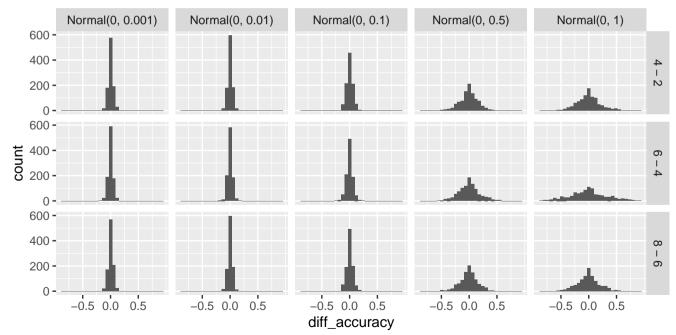
(The R code is in the textbook, chapter 4!)

Bayesian Data Analysis



Bayesian Data Analysis

It's usually more useful to look at the predicted differences in accuracy between set sizes.



Bayesian Data Analysis

These priors seem reasonable:

$$\alpha \sim Normal(0, 1.5)$$

$$\beta \sim Normal(0, 0.1)$$
(16)

Bayesian Data Analysis

The next steps

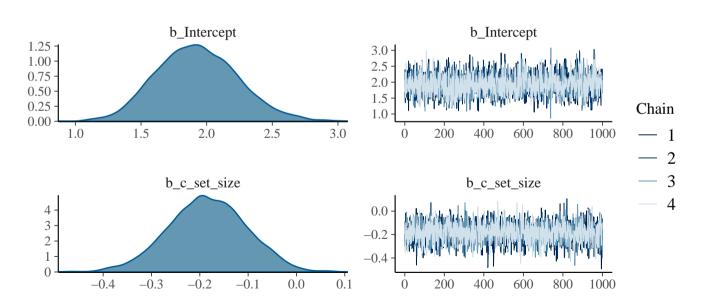
Next: fit the model and examine the posterior distributions of the parameters.

Bayesian Data Analysis

```
fit_recall <- brm(correct ~ 1 + c_set_size,
  data = df_recall,
  family = bernoulli(link = logit),
  prior = c(
    prior(normal(0, 1.5), class = Intercept),
    prior(normal(0, .1), class = b, coef = c_set_size)
posterior_summary(fit_recall,
                  variable = c("b_Intercept", "b_c_set_size"))
```

Bayesian Data Analysis

plot(fit_recall)



Bayesian Data Analysis

```
alpha_samples <- as_draws_df(fit_recall)$b_Intercept</pre>
beta_samples <- as_draws_df(fit_recall)$b_c_set_size</pre>
beta_mean <- round(mean(beta_samples), 5)</pre>
beta_low <- round(quantile(beta_samples, prob = 0.025), 5)
beta_high <- round(quantile(beta_samples, prob = 0.975), 5)
alpha_samples <- as_draws_df(fit_recall)$b_Intercept</pre>
av_accuracy <- plogis(alpha_samples)</pre>
c(mean = mean(av_accuracy), quantile(av_accuracy, c(0.025, 0.975)))
   mean 2.5% 97.5%
## 0.87 0.80 0.93
```

Bayesian Data Analysis

Find out the decrease in accuracy in proportions or probability scale:

```
beta_samples <- as_draws_df(fit_recall)$b_c_set_size
effect_middle <- plogis(alpha_samples) -
    plogis(alpha_samples - beta_samples)
c(mean = mean(effect_middle),
    quantile(effect_middle, c(0.025, 0.975)))
## mean 2.5% 97.5%
## -0.0188 -0.0370 -0.0033</pre>
```

Bayesian Data Analysis

```
four <- 4 - mean(df_recall$set_size)
two <- 2 - mean(df_recall$set_size)
effect_4m2 <-
    plogis(alpha_samples + four * beta_samples) -
    plogis(alpha_samples + two * beta_samples)
c(mean = mean(effect_4m2),
    quantile(effect_4m2, c(0.025, 0.975)))
## mean 2.5% 97.5%
## -0.0294 -0.0535 -0.0063</pre>
```

Bayesian Data Analysis

The next steps

Next: Hierarchical models.

Bayesian Data Analysis