

# Prepare data for BRMS fits focusing on x0

Michael Gilchrist

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## Goal

- Process data as in 2023-02-07\_brms\_two.piece\_focus.on.x0.fittings/ for other routines

## Recap

## Set up

### Install libraries

```
# install packages user might not have by replacing FALSE with TRUE

## load libraries
library(stats)
library(MASS) # provides negative binomial fitting: glm.nb
library(ggplot2)
library(tidyverse)
library(viridisLite)
```

### Load Data

```
resave_data <- FALSE # flag for saving data_stan and possibly others in future

input_dir <- file.path("input")
output_dir <- file.path("output")

load(file.path(input_dir, "data.processing_2022-12-15.Rda"),
      verbose = TRUE)
```

```
## Loading objects:
## motif_data
## motif_data_40C
## motif_stats
## motif_stats_40C
## bird_bill_data
```

```
motif_data
```

```
## # A tibble: 146 x 28
##   male round trial_round motif_count motif_prop temp_mean humidity_mean
##   <fct> <dbl>      <dbl>      <int>      <dbl>      <dbl>      <dbl>
## 1 T229     1         1         0         0         45.8         NA
## 2 T229     1         2         24        0.0131        42.3         NA
## 3 T229     1         3        114        0.0622        40.7         NA
## 4 T229     1         4        198        0.108         26.2         NA
## 5 T229     1         5        315        0.172         34.9         NA
## 6 T231     1         1         57        0.0431        42.8         NA
## 7 T231     1         2          7        0.00529        45.0         NA
## 8 T231     1         3         86        0.0650        41.1         NA
## 9 T231     1         4         24        0.0181        27.2         NA
## 10 T231    1         5        215        0.162         36.5         NA
## # i 136 more rows
## # i 21 more variables: chamber <fct>, date <chr>, counter <chr>,
## #   test_order <int>, temp_target <dbl>, temp_median <dbl>, humidity_mean <dbl>,
## #   motif_rate <dbl>, mass <dbl>, n_obs_completed <lgl>,
## #   motif_count_plus_1 <int>, log_motif_count_plus_1 <dbl>, temp <dbl>,
## #   n_obs_round <int>, n_obs <int>, trial <int>, motif_prop_round <dbl>,
## #   weights <dbl>, svp <dbl>, vpd <dbl>, vpd_offset <dbl>
```

## Process Data

### Create Working Dataset

```
filter_data <- TRUE

if(filter_data) {
  males_filtered_disp <- motif_stats_40C %>%
    filter(dispersion < 50) %>%
    pull(male)

  males_filtered_mean <- motif_stats %>%
    filter(mean > 10) %>% # changing from 10 to 40 removes previous male 7 (T258)
    pull(male)

  male_vector <- intersect(males_filtered_mean, males_filtered_disp)
} else {
  male_vector <- motif_data %>% select(male) %>% distinct()
}

data_ind <- motif_data %>%
  filter(male %in% male_vector) %>%
  mutate(male = droplevels(male)) %>%
  mutate(index = as.integer(male)) %>%
  mutate(male = as.character(male)) %>%
  arrange(index) %>%
  select(male, index, motif_count, temp_target, temp, round, trial_round, date, counter) %>%
```

```

## left_join(index_shape, by = "index") %>%
group_by(male) %>% mutate(y0_simple_est = mean(motif_count), phi_ind = var(motif_count)/y0_simple_est,
ungroup() %>%
mutate()

save(data_ind, file = file.path(output_dir, "data_ind.Rda"))
write_csv(data_ind, file = file.path(output_dir, "data_ind.csv"))

stats_ind <- motif_stats %>%
  filter(male %in% male_vector)

save(stats_ind, file = file.path(output_dir, "stats_ind.Rda"))
write_csv(stats_ind, file = file.path(output_dir, "stats_ind.csv"))

summary(data_ind)

```

```

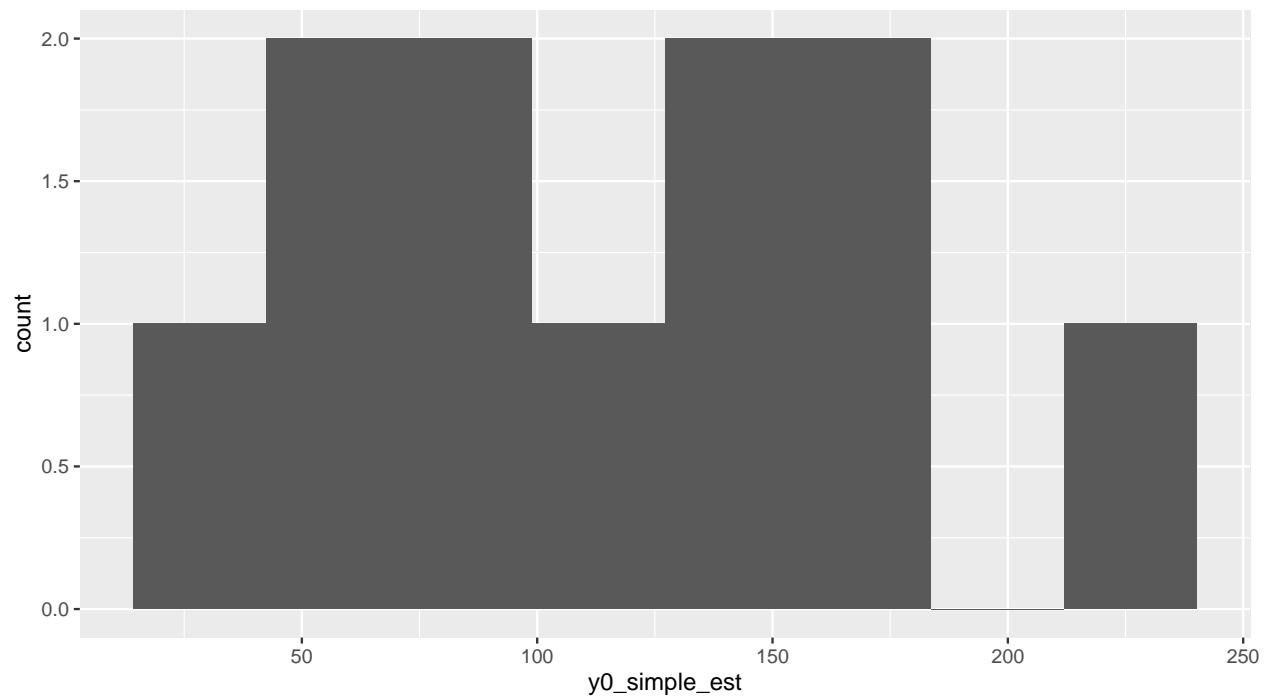
##      male           index      motif_count      temp_target      temp
## Length:107      Min.   : 1.00      Min.   : 0      Min.   :27      Min.   :25.7
## Class :character 1st Qu.: 3.00      1st Qu.: 56      1st Qu.:35      1st Qu.:34.3
## Mode  :character Median : 5.00      Median : 90      Median :40      Median :38.8
##              Mean  : 5.14      Mean  :116      Mean  :38      Mean  :37.6
##              3rd Qu.: 8.00      3rd Qu.:175      3rd Qu.:42      3rd Qu.:41.1
##              Max.   :11.00      Max.   :425      Max.   :44      Max.   :45.6
##      round      trial_round      date      counter
## Min.   :1.00      Min.   :1.00      Length:107      Length:107
## 1st Qu.:1.00      1st Qu.:2.00      Class :character  Class :character
## Median :2.00      Median :3.00      Mode  :character  Mode  :character
## Mean   :2.15      Mean   :2.83
## 3rd Qu.:3.00      3rd Qu.:4.00
## Max.   :3.00      Max.   :6.00
## y0_simple_est      phi_ind
## Min.   : 34.2      Min.   : 9.7
## 1st Qu.: 76.4      1st Qu.: 12.3
## Median :106.2      Median : 22.5
## Mean   :116.5      Mean   : 35.5
## 3rd Qu.:161.2      3rd Qu.: 58.9
## Max.   :232.0      Max.   :103.7

```

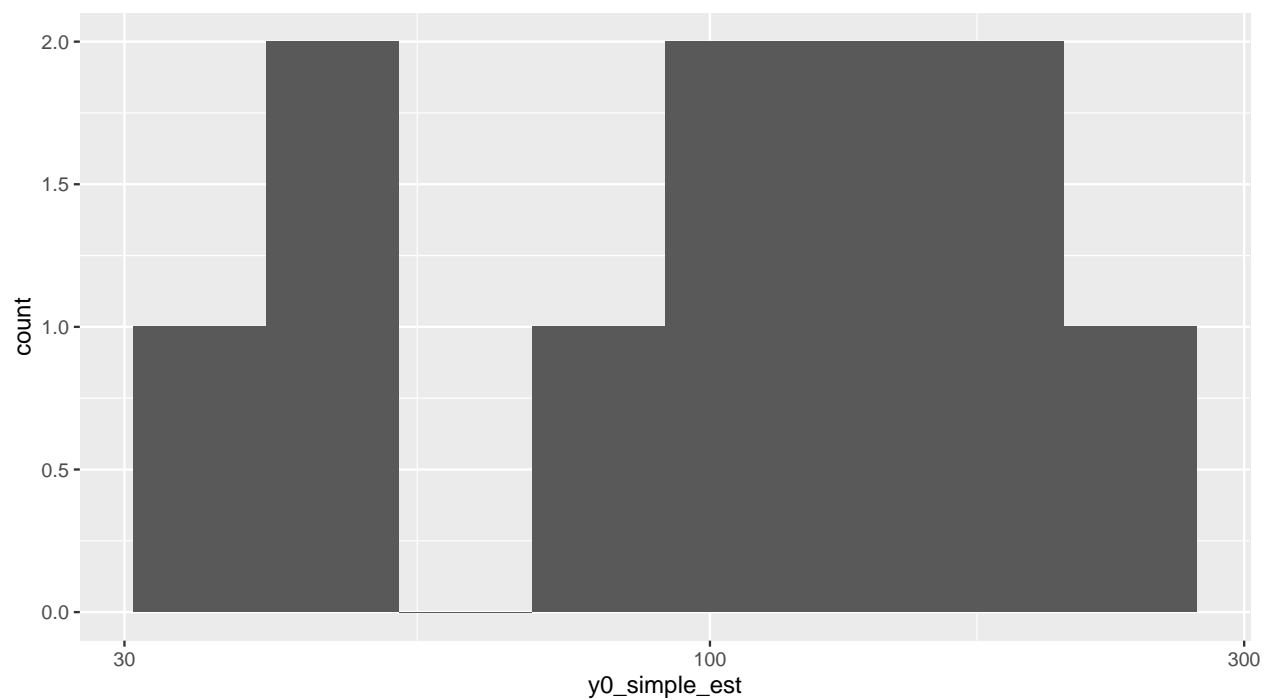
```

y0_hist <- data_ind %>%
  select(y0_simple_est) %>%
  unique() %>%
  ggplot(aes(y0_simple_est)) + geom_histogram(bins = 8)
print(y0_hist)

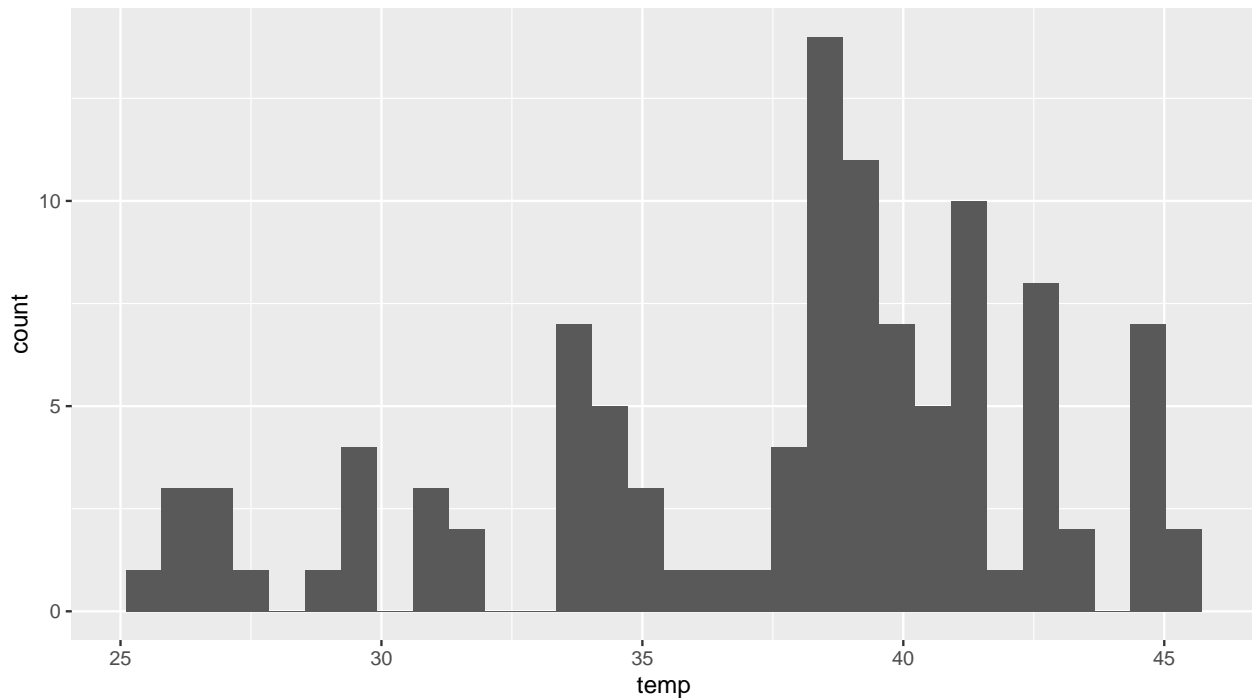
```



```
log_y0_hist <- y0_hist +
  scale_x_log10()
print(log_y0_hist)
```



```
temp_hist <- ggplot(data_ind, aes(temp)) + geom_histogram()
print(temp_hist)
```



```
n_male <- length(unique(data_ind$male))

## for calculation of 'theta_bar' (size in `rnbinom`) see note in `Set Up Simulate Data`
summary_stats <- data_ind %>%
  ungroup() %>%
  summarize(y0_bar = mean(y0_simple_est),
            y0_sd = sd(y0_simple_est),
            log_y0_bar = mean(log(y0_simple_est)),
            log_y0_sd = sd(log(y0_simple_est)),
            phi_bar = median(phi_ind), # qpoisson over dispersion
            theta_bar = y0_bar^2/(y0_sd^2 - y0_bar)) # nb shape parameter
comment(summary_stats) <- "summary stats for observed bird motifs"

save(summary_stats, file = file.path(output_dir, "obs_summary_stats.Rda"))
```

## Fit Models

See other folders created on this day.