

Examine dispersal values for males

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Goal

- Try to identify which males should share a common `disp_value` using the `'disp_flag == uniform_2'`.

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

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

## This data includes `motif_data_full`, which is unfiltered by males

load(file.path(input_dir, "data.processing_2023-04-05.Rda"),
      verbose = TRUE)

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

```
summary(motif_data_full)
```

```
##      male      round      trial_round      motif_count      motif_prop
## T247   :14   Min.    :1.00   Min.    :1.00   Min.    : 0   Min.    :0.000
## T229   :13   1st Qu.:1.00   1st Qu.:2.00   1st Qu.: 41   1st Qu.:0.043
## T231   :13   Median  :2.00   Median  :3.00   Median  : 84   Median  :0.081
## T234   :13   Mean    :2.18   Mean    :2.82   Mean    :107   Mean    :0.103
## T235   :13   3rd Qu.:3.00   3rd Qu.:4.00   3rd Qu.:167   3rd Qu.:0.126
## T236   :13   Max.    :3.00   Max.    :6.00   Max.    :425   Max.    :0.720
## (Other):67
##      temp_mean      humidity_mean      chamber      date      counter
## Min.    :25.7   Min.    :15.6   1:10   Length:146   Length:146
## 1st Qu.:34.3   1st Qu.:22.4   2:43   Class :character   Class :character
## Median  :38.9   Median  :24.2   3:10   Mode  :character   Mode  :character
## Mean    :37.7   Mean    :26.0   4:42
## 3rd Qu.:41.3   3rd Qu.:29.9   5: 5
## Max.    :45.8   Max.    :41.8   6:36
## NA's    :2     NA's    :46
##      test_order      temp_target      temp_median      humidity_mean      motif_rate
## Min.    :1.00   Min.    :27   Min.    :26.0   Min.    : 8.0   Min.    : 0.0
## 1st Qu.:2.00   1st Qu.:35   1st Qu.:34.5   1st Qu.:10.0   1st Qu.: 1.6
## Median  :3.00   Median  :40   Median  :40.0   Median  :11.3   Median  : 3.0
## Mean    :3.03   Mean    :38   Mean    :38.3   Mean    :12.7   Mean    : 3.6
## 3rd Qu.:4.00   3rd Qu.:42   3rd Qu.:42.0   3rd Qu.:16.2   3rd Qu.: 5.3
## Max.    :6.00   Max.    :44   Max.    :46.0   Max.    :19.5   Max.    :14.2
## NA's    :30     NA's    :2     NA's    :102   NA's    :75
##      mass      bill_length      bill_depth      bill_width      bill_date
## Min.    :15.0   Min.    :7.8   Min.    :7.2   Min.    :5.9   Length:146
## 1st Qu.:16.1   1st Qu.:8.2   1st Qu.:7.3   1st Qu.:6.4   Class :character
## Median  :17.8   Median  :8.6   Median  :7.5   Median  :6.6   Mode  :character
## Mean    :17.5   Mean    :8.5   Mean    :7.5   Mean    :6.6
## 3rd Qu.:18.9   3rd Qu.:8.7   3rd Qu.:7.7   3rd Qu.:6.8
## Max.    :20.0   Max.    :9.3   Max.    :8.0   Max.    :7.1
## NA's    :132   NA's    :132   NA's    :132   NA's    :132
##      n_obs_completed      motif_count_plus_1      log_motif_count_plus_1      temp
## Mode:logical   Min.    : 1   Min.    :0.00   Min.    :25.7
## TRUE:146       1st Qu.: 42   1st Qu.:3.74   1st Qu.:34.4
##               Median  : 84   Median  :4.44   Median  :39.0
##               Mean    :108   Mean    :4.06   Mean    :37.8
##               3rd Qu.:168   3rd Qu.:5.13   3rd Qu.:41.2
##               Max.    :426   Max.    :6.05   Max.    :45.8
##
##      count_total_round      n_obs_round      count_mean_round      count_sd_round
## Min.    : 0   Min.    :3.00   Min.    : 0.0   Min.    : 0.0
## 1st Qu.:253   1st Qu.:5.00   1st Qu.:58.8   1st Qu.:27.7
## Median  :498   Median  :5.00   Median :106.4   Median  :46.0
## Mean    :493   Mean    :4.63   Mean    :107.4   Mean    :55.1
## 3rd Qu.:651   3rd Qu.:5.00   3rd Qu.:140.1   3rd Qu.:71.0
## Max.    :1210   Max.    :6.00   Max.    :270.0   Max.    :159.0
##
##      count_cv_round      count_total      n_obs      count_mean      trial
## Min.    :0.072   Min.    : 25   Min.    : 5.0   Min.    : 3.6   Min.    : 1.00
## 1st Qu.:0.307   1st Qu.:601   1st Qu.: 8.0   1st Qu.:76.4   1st Qu.: 3.00
```

```
## Median :0.569   Median :1324   Median :13.0   Median :106.2   Median : 5.00
## Mean    :0.601   Mean    :1256   Mean    :11.2   Mean    :107.4   Mean    : 6.09
## 3rd Qu.:0.811   3rd Qu.:1861   3rd Qu.:13.0   3rd Qu.:143.2   3rd Qu.: 9.00
## Max.    :1.575   Max.    :2333   Max.    :14.0   Max.    :232.0   Max.    :14.00
## NA's    :3
## motif_prop_round  weights      svp      vpd      vpd_offset
## Min.    :0.000   Min.    :0.002   Min.    : 33.6   Min.    :20.1   Min.    : NA
## 1st Qu.:0.139   1st Qu.:0.006   1st Qu.: 55.9   1st Qu.:40.3   1st Qu.: NA
## Median :0.211   Median :0.012   Median : 72.4   Median :55.3   Median : NA
## Mean    :0.224   Mean    :0.112   Mean    : 70.1   Mean    :51.7   Mean    :NaN
## 3rd Qu.:0.299   3rd Qu.:0.024   3rd Qu.: 82.4   3rd Qu.:60.3   3rd Qu.: NA
## Max.    :0.842   Max.    :1.000   Max.    :105.7   Max.    :79.0   Max.    : NA
## NA's    :3      NA's    :2      NA's    :46      NA's    :146
```

```
data_working <- motif_data_full %>% relocate(male, temp, motif_count)
```

```
temp_stats_wide <- tibble(
  male = factor(0),
  mean = numeric(0),
  median = numeric(0),
  var = numeric(0),
  cv = numeric(0),
  disp = integer(0)
)
```

```
temp_stats_wide <- tibble()
```

```
for(threshold in 30:41) {
```

```
  data_tmp <- filter(data_working, temp <= threshold)
```

```
  stats_tmp <- data_tmp %>%
    group_by(male) %>%
    summarize(mean = mean(motif_count),
              median = median(motif_count),
              var = var(motif_count),
              cv = mean/sqrt(var),
              disp = var/mean,
              count = n()) %>%
```

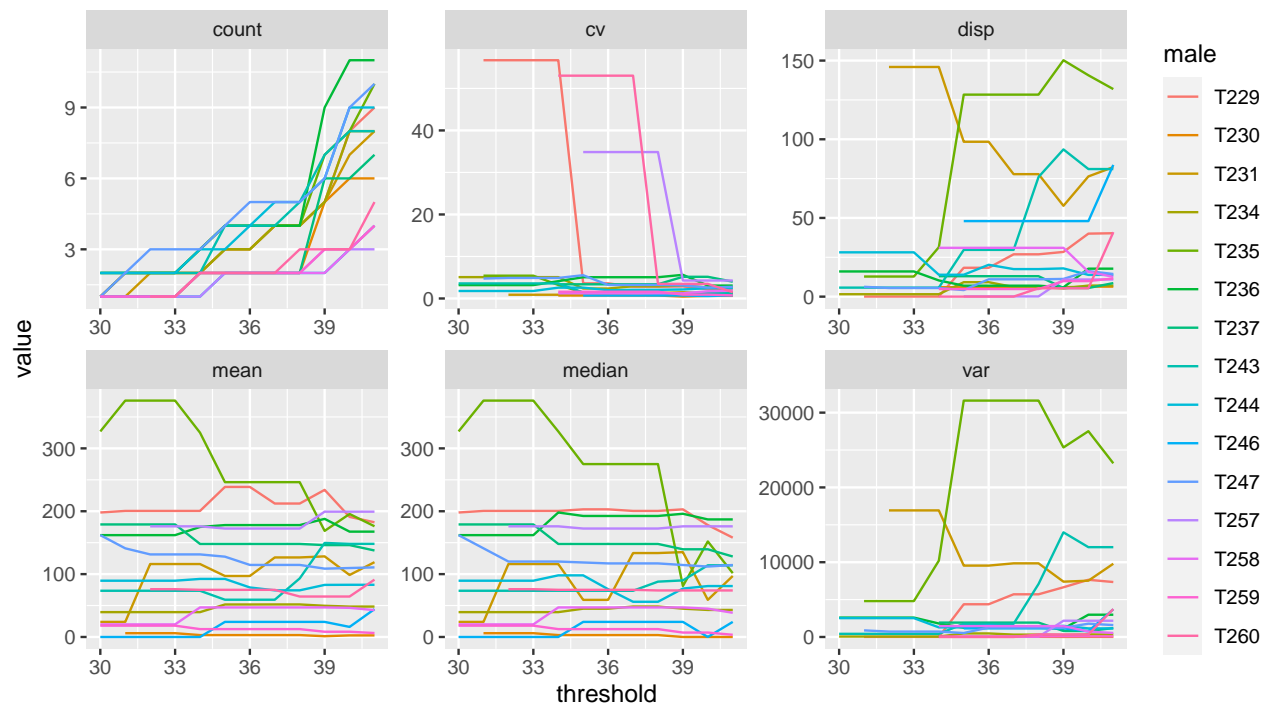
```
    mutate(threshold = threshold)
```

```
  temp_stats_wide <- bind_rows(temp_stats_wide, stats_tmp, .id = NULL)
```

```
}
```

```
temp_stats_long <- pivot_longer(temp_stats_wide,
                                cols = -c(male, threshold),
                                names_to = "statistic"
                                )
```

```
ggplot(data = temp_stats_long) +
  geom_line(aes(x = threshold, y = value, color = male)) +
  facet_wrap(facets = vars(statistic), scales = "free")
```



“ # Fit Models

See other folders created on this day.