

Preliminary Data Analysis

Project Description:

The data analysis will be conducted to evaluate the effect of zucchini planting date on the need for pollinator supplementation with managed bee hives. The data being used is unpublished and represents proportion analysis of bee visits over time. Dr. Ashley Leach, as co-PI, has been conducting experiments established across three field sites (Wooster, Willard, and Piketon). For this preliminary analysis, data from the Wooster site is being used, with additional site data to be incorporated in future analyses as it becomes available.

Predictor (Categorical) Variables:

1. Planting Date (June 1 vs July 1) - proxy for bloom time
2. Pollination Treatment (With managed bumblebees vs Without supplementation)

Response (Continuous) Variables:

1. Average number of fruit per plant (yield measure)
2. Average number of overall visits per flower per minute (pollination measure)

Loading the dataset into R:

```
rm(list=ls()) #clearing R's Brain
```

```
library(tidyverse)
```

```
— Attaching core tidyverse packages ————— tidyverse 2.0.0 —
✓ dplyr     1.1.4    ✓ readr     2.1.5
✓ forcats   1.0.0    ✓ stringr   1.5.2
✓ ggplot2   4.0.0    ✓ tibble    3.3.0
✓ lubridate 1.9.4    ✓ tidyr    1.3.1
✓ purrr    1.1.0
— Conflicts ————— tidyverse_conflicts() —
✖ dplyr::filter() masks stats::filter()
✖ dplyr::lag()    masks stats::lag()
ℹ Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become
errors
```

```
library(dplyr)
library(readxl)
library(ggplot2)
```

```
squash_beans_df <- read_excel("C:/Users/Rosita/OneDrive - The Ohio State University/Documents/AU
                                2025/Ent Tech&DataAnalysis/R_dataset1/Preliminary_data/pollinator_records.xlsx",
                                sheet="Pollinator.Records", "A1:Q97" )
```

```
summary(squash_beans_df)
```

date	treatment	planting	
Min. :2025-07-12 00:00:00	Length:96	Min. :1.00	
1st Qu.:2025-07-14 06:00:00	Class :character	1st Qu.:1.00	
Median :2025-07-18 12:00:00	Mode :character	Median :1.00	
Mean :2025-07-21 12:00:00		Mean :1.25	
3rd Qu.:2025-07-25 18:00:00		3rd Qu.:1.25	
Max. :2025-08-06 00:00:00		Max. :2.00	
site	observation	sampling	squash.beans
Length:96	Min. : 1.00	Min. :1.00	Min. : 0.000
Class :character	1st Qu.: 3.75	1st Qu.:1.00	1st Qu.: 3.000
Mode :character	Median : 6.50	Median :1.50	Median : 6.000
	Mean : 6.50	Mean :1.75	Mean : 6.177
	3rd Qu.: 9.25	3rd Qu.:2.25	3rd Qu.: 9.000
	Max. :12.00	Max. :3.00	Max. :16.000
bumblebees	honeybees	sweat.bees	other
Min. :0.0000	Min. :0.0000	Min. :0.0000	Min. :0
1st Qu.:0.0000	1st Qu.:0.0000	1st Qu.:0.0000	1st Qu.:0
Median :0.0000	Median :0.0000	Median :0.0000	Median :0
Mean :0.4688	Mean :0.4375	Mean :0.0625	Mean :0
3rd Qu.:0.0000	3rd Qu.:1.0000	3rd Qu.:0.0000	3rd Qu.:0
Max. :7.0000	Max. :3.0000	Max. :1.0000	Max. :0
total	flowers	male	female
Min. : 0.000	Min. :1.0	Min. :0.000	Min. :0.0000
1st Qu.: 4.000	1st Qu.:1.0	1st Qu.:1.000	1st Qu.:0.0000
Median : 6.500	Median :1.0	Median :1.000	Median :0.0000
Mean : 7.146	Mean :1.5	Mean :1.062	Mean :0.4375
3rd Qu.:11.000	3rd Qu.:2.0	3rd Qu.:1.000	3rd Qu.:1.0000
Max. :20.000	Max. :4.0	Max. :3.000	Max. :2.0000
visits.flower	visits.flower.min		
Min. : 0.000	Min. :0.0000		
1st Qu.: 2.500	1st Qu.:0.8333		
Median : 5.000	Median :1.6667		
Mean : 5.507	Mean :1.8356		
3rd Qu.: 8.000	3rd Qu.:2.6667		
Max. :15.000	Max. :5.0000		

```
head(squash_beans_df)
```

# A tibble: 6 × 17	date	treatment	planting	site	observation	sampling	squash.beans	<dttm>	<chr>	<dbl>	<chr>	<dbl>	<dbl>	<dbl>	
1	2025-07-12 00:00:00	Supplemental	1	woos...	1	1	5								
2	2025-07-12 00:00:00	Supplemental	1	woos...	2	1	1								
3	2025-07-12 00:00:00	Supplemental	1	woos...	3	1	15								
4	2025-07-12 00:00:00	Supplemental	1	woos...	4	1	5								
5	2025-07-12 00:00:00	Supplemental	1	woos...	5	1	2								
6	2025-07-12 00:00:00	Supplemental	1	woos...	6	1	0								

```
# i 10 more variables: bumblebees <dbl>, honeybees <dbl>, sweat.bees <dbl>,
#   other <dbl>, total <dbl>, flowers <dbl>, male <dbl>, female <dbl>,
#   visits.flower <dbl>, visits.flower.min <dbl>
```

```
num_data <- nrow(squash_beans_df)
print(num_data)
```

[1] 96

```
squash_beans_df$treatment <- as.factor(squash_beans_df$treatment)
str(squash_beans_df)
```

```
tibble [96 x 17] (S3:tbl_df/tbl/data.frame)
$ date           : POSIXct[1:96], format: "2025-07-12" "2025-07-12" ...
$ treatment      : Factor w/ 2 levels "Not supplemented",...: 2 2 2 2 2 2 2 2 2 ...
$ planting       : num [1:96] 1 1 1 1 1 1 1 1 1 ...
$ site           : chr [1:96] "wooster" "wooster" "wooster" "wooster" ...
$ observation    : num [1:96] 1 2 3 4 5 6 7 8 9 10 ...
$ sampling        : num [1:96] 1 1 1 1 1 1 1 1 1 ...
$ squash.beans   : num [1:96] 5 1 15 5 2 0 4 8 1 0 ...
$ bumblebees     : num [1:96] 0 0 0 0 0 0 0 0 0 ...
$ honeybees       : num [1:96] 0 0 0 0 0 0 0 1 0 0 ...
$ sweat.beans    : num [1:96] 0 0 0 0 0 0 0 0 0 ...
$ other           : num [1:96] 0 0 0 0 0 0 0 0 0 ...
$ total           : num [1:96] 5 1 15 5 2 0 4 9 1 0 ...
$ flowers          : num [1:96] 1 2 2 2 2 3 2 2 1 1 ...
$ male             : num [1:96] 1 1 2 1 2 2 1 2 1 1 ...
$ female            : num [1:96] 0 1 0 1 0 1 1 0 0 0 ...
$ visits.flower   : num [1:96] 5 0.5 7.5 2.5 1 0 2 4.5 1 0 ...
$ visits.flower.min: num [1:96] 1.667 0.167 2.5 0.833 0.333 ...
```

```
yield_df <- read_excel("C:/Users/Rosita/OneDrive - The Ohio State University/Documents/AU
2025/Ent Tech&DataAnalysis/R_dataset1/Preliminary_data/yield_prelim.xlsx",
sheet="yield_prelim", "A1:Q26")
```

```
summary(yield_df)
```

year	site	treatment	planting
Min. :2025	Length:25	Length:25	Min. :1.00
1st Qu.:2025	Class :character	Class :character	1st Qu.:1.00
Median :2025	Mode :character	Mode :character	Median :2.00
Mean :2025			Mean :1.56
3rd Qu.:2025			3rd Qu.:2.00
Max. :2025			Max. :2.00
harvest	num_harvest	damaged add	weight_harvest
Min. :1.00	Min. : 6.00	Min. : 0.00	Length:25
1st Qu.:2.00	1st Qu.: 15.00	1st Qu.: 0.00	Class :character
Median :2.00	Median : 45.00	Median : 0.00	Mode :character
Mean :2.12	Mean : 57.88	Mean : 7.16	

```

3rd Qu.:3.00  3rd Qu.: 61.00   3rd Qu.: 0.00
Max.    :3.00  Max.    :218.00  Max.    :61.00
  Cull_poll      Cull_poll_weight      Cull_other      Cull_other_weight
Min.    : 0.00  Length:25          Min.    :0.00  Length:25
  1st Qu.: 0.00  Class :character  1st Qu.:0.00  Class :character
  Median : 1.00  Mode   :character  Median :0.00  Mode   :character
  Mean   : 2.44                    Mean   :0.44
  3rd Qu.: 3.00                    3rd Qu.:0.00
  Max.   :21.00                    Max.   :3.00
  stand      functional stand  yield.plant      yield.plant_weight
Min.    :32.00  Min.    :20.00    Min.    :0.1034  Length:25
  1st Qu.:58.00 1st Qu.:33.00  1st Qu.:0.3333  Class :character
  Median :60.00  Median :53.00  Median :0.8491  Mode   :character
  Mean   :56.64  Mean   :49.24  Mean   :1.1292
  3rd Qu.:68.00 3rd Qu.:68.00  3rd Qu.:1.6500
  Max.   :76.00  Max.   :76.00  Max.   :3.7586

yield.plant_cull
Min.    :0.00000
  1st Qu.:0.00000
  Median :0.01887
  Mean   :0.05404
  3rd Qu.:0.09091
  Max.   :0.36207

```

```
head(yield_df)
```

```

# A tibble: 6 × 17
  year site treatment      planting harvest num_harvest `damaged` add` 
  <dbl> <chr>  <chr>           <dbl>   <dbl>     <dbl>       <dbl>
1 2025 wooster not supplemented      1       1        61        61
2 2025 wooster supplemented         1       2        42        42
3 2025 wooster supplemented         1       1        54        54
4 2025 wooster not supplemented     1       2        11        13
5 2025 wooster supplemented         1       2         9         9
6 2025 wooster not supplemented     1       2         8         0

# i 10 more variables: weight_harvest <chr>, Cull_poll <dbl>,
#   Cull_poll_weight <chr>, Cull_other <dbl>, Cull_other_weight <chr>,
#   stand <dbl>, `functional stand` <dbl>, yield.plant <dbl>,
#   yield.plant_weight <chr>, yield.plant_cull <dbl>

```

```

yield_df$treatment <- as.factor(yield_df$treatment)
str(yield_df)

```

```

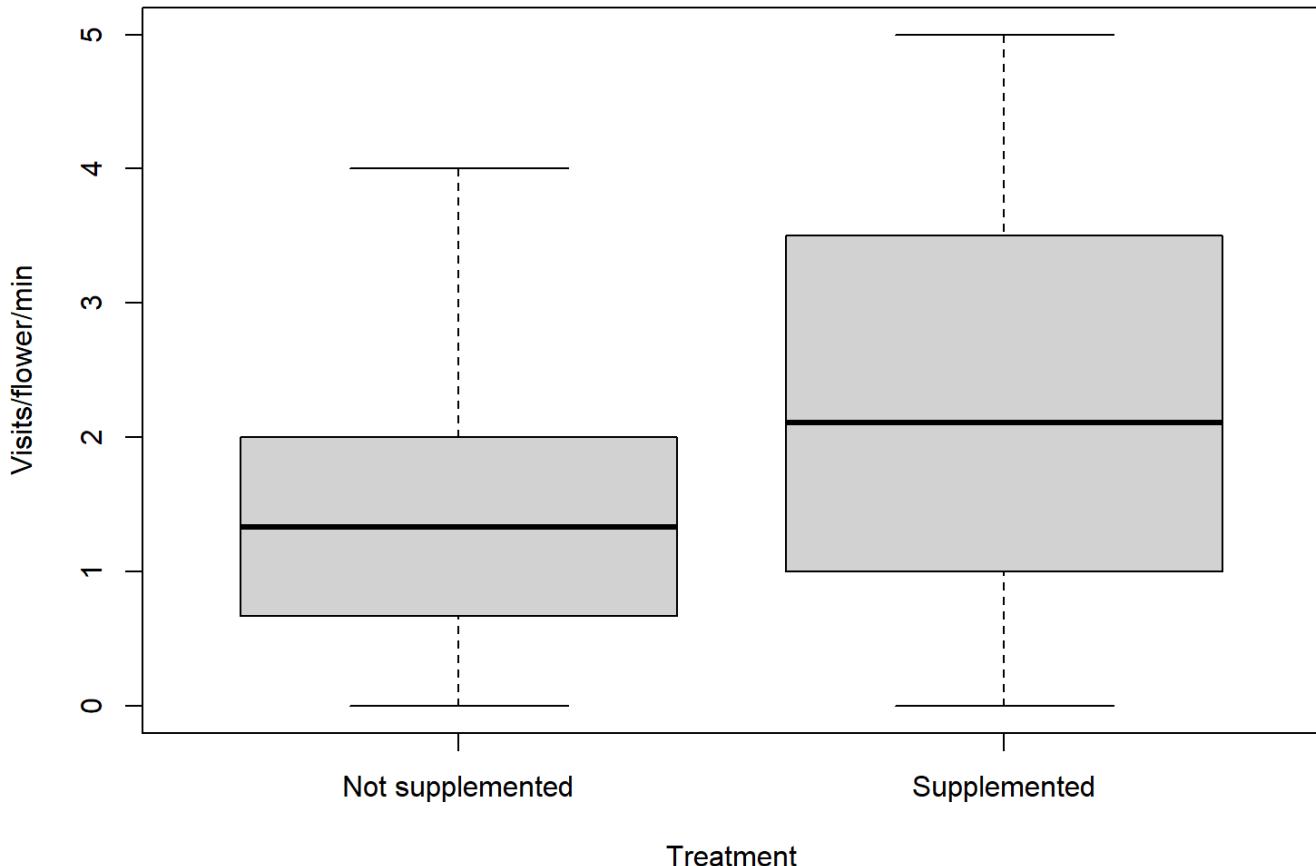
tibble [25 × 17] (S3: tbl_df/tbl/data.frame)
$ year           : num [1:25] 2025 2025 2025 2025 2025 ...
$ site            : chr [1:25] "wooster" "wooster" "wooster" "wooster" ...
$ treatment       : Factor w/ 2 levels "not supplemented",...: 1 2 2 1 2 1 2 1 2 1 ...
$ planting        : num [1:25] 1 1 1 1 1 1 1 1 1 ...
$ harvest         : num [1:25] 1 2 1 2 2 2 3 3 3 3 ...

```

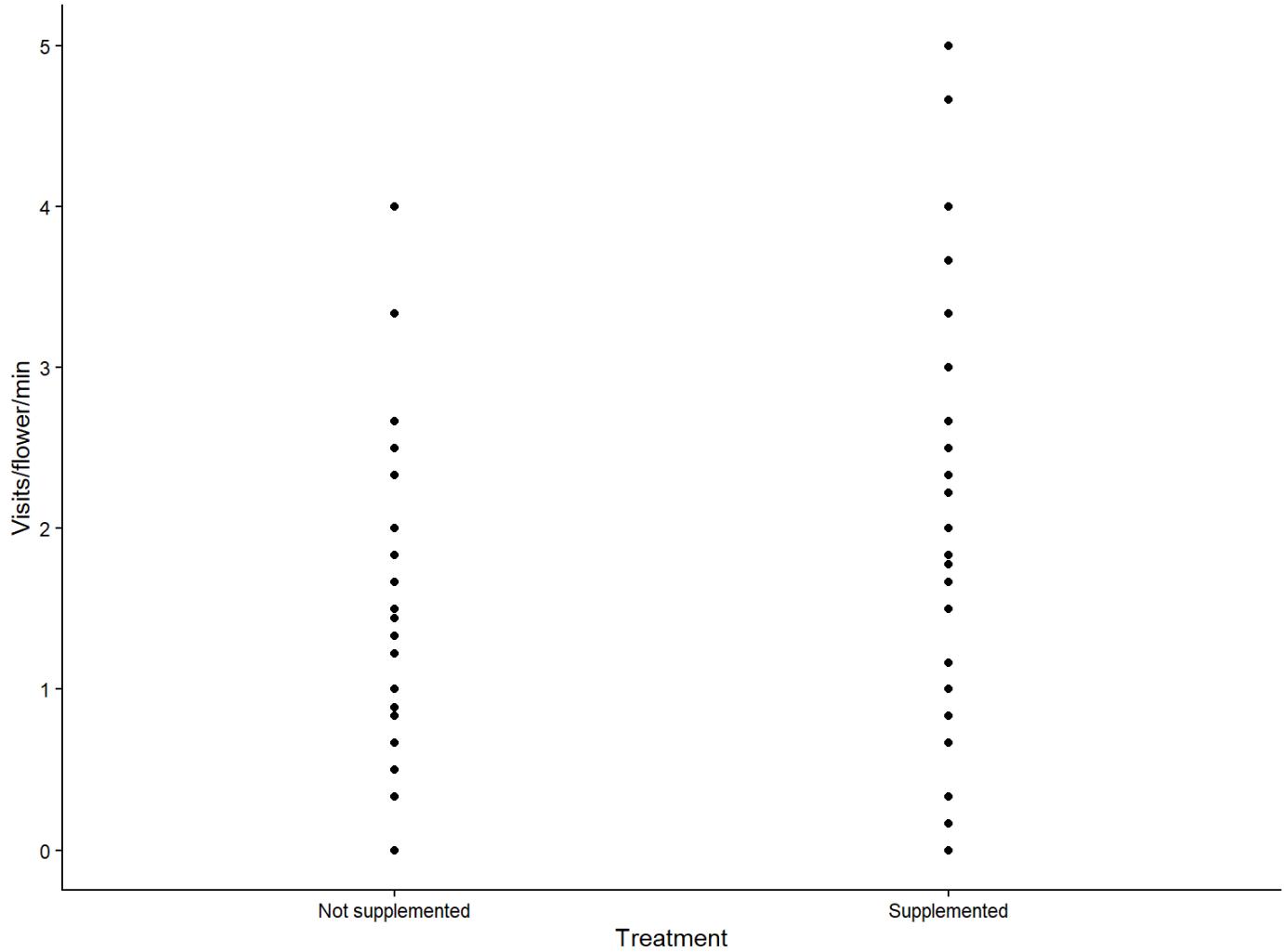
```
$ num_harvest      : num [1:25] 61 42 54 11 9 8 9 7 15 33 ...
$ damaged.add     : num [1:25] 61 42 54 13 9 0 0 0 0 0 ...
$ weight.harvest   : chr [1:25] "53.32900000000001" "42.616" "29.36" "8.08280000000007" ...
$ Cull.poll       : num [1:25] 8 2 0 3 0 0 0 3 3 2 ...
$ Cull.poll.weight : chr [1:25] "1.6990000000000001" "0.3420000000000003" "0"
"0.5500000000000004" ...
$ Cull.other       : num [1:25] 2 0 0 0 0 0 0 0 0 0 ...
$ Cull.other.weight: chr [1:25] "0.2800000000000003" "0" "0" "0" ...
$ stand            : num [1:25] 60 34 32 60 32 60 32 60 32 60 ...
$ functional.stand : num [1:25] 33 34 32 33 32 33 32 33 32 20 ...
$ yield.plant      : num [1:25] 1.848 1.235 1.688 0.333 0.281 ...
$ yield.plant.weight: chr [1:25] "1.6160303030000001" "1.2534117650000001" "0.9174999999999998"
"0.244933333" ...
$ yield.plant.cull : num [1:25] 0.2424 0.0588 0 0.0909 0 ...
```

Plot of visits/flower/min as a function of treatment:

```
plot(visits.flower.min~treatment, data= squash_beans_df, xlab= "Treatment",
      ylab= "Visits/flower/min")
```

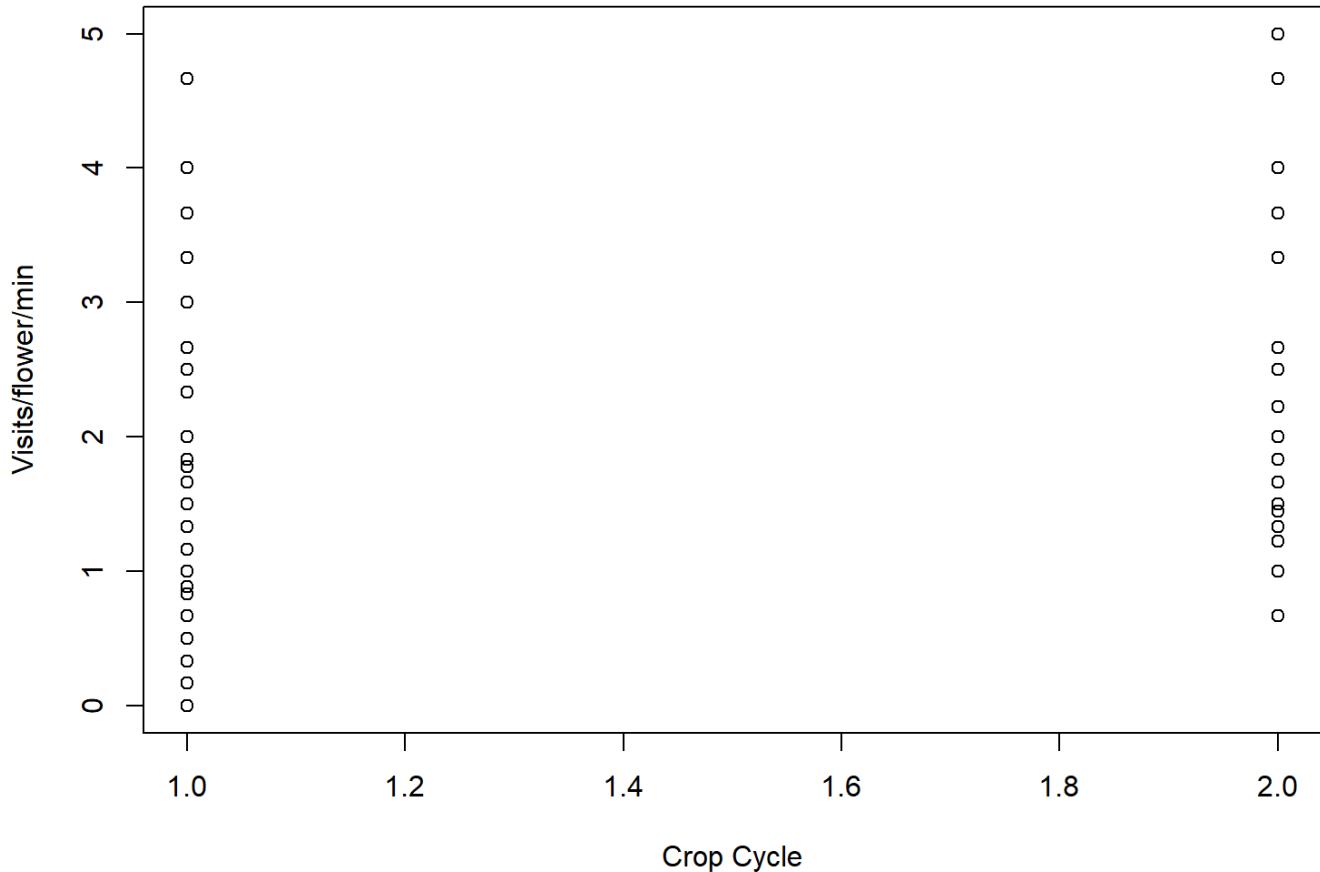


```
ggplot(data=squash_bees_df, mapping=aes(x=treatment , y=visits.flower.min)) +  
geom_point() + theme_classic() +  
xlab("Treatment") +  
ylab("Visits/flower/min")
```

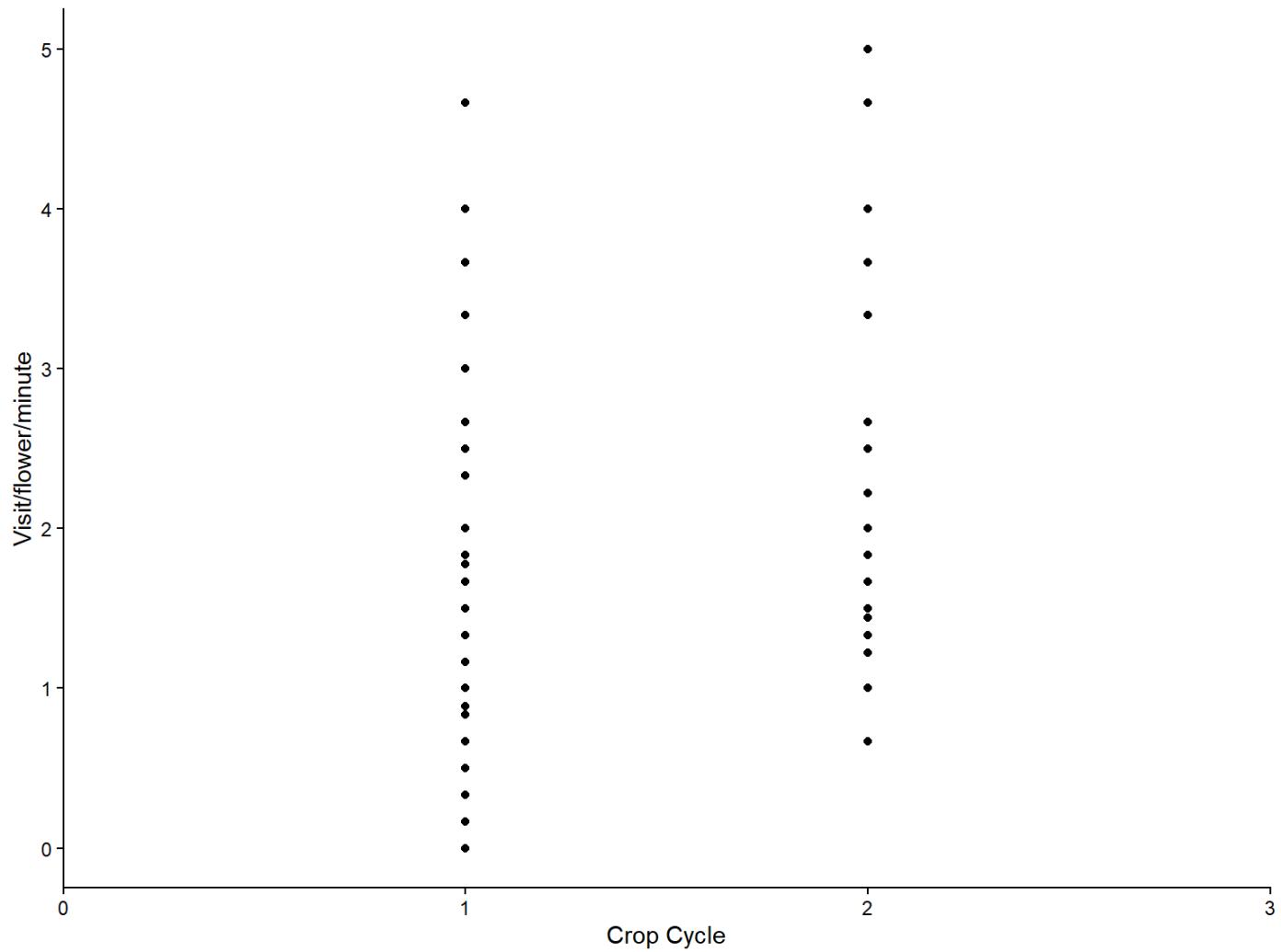


Plot of visits/flower/min as a function of planting date (~proxy for bloom date):

```
plot(visits.flower.min ~ planting, data= squash_bees_df, xlab="Crop Cycle", ylab="Visits/flower/min")
```

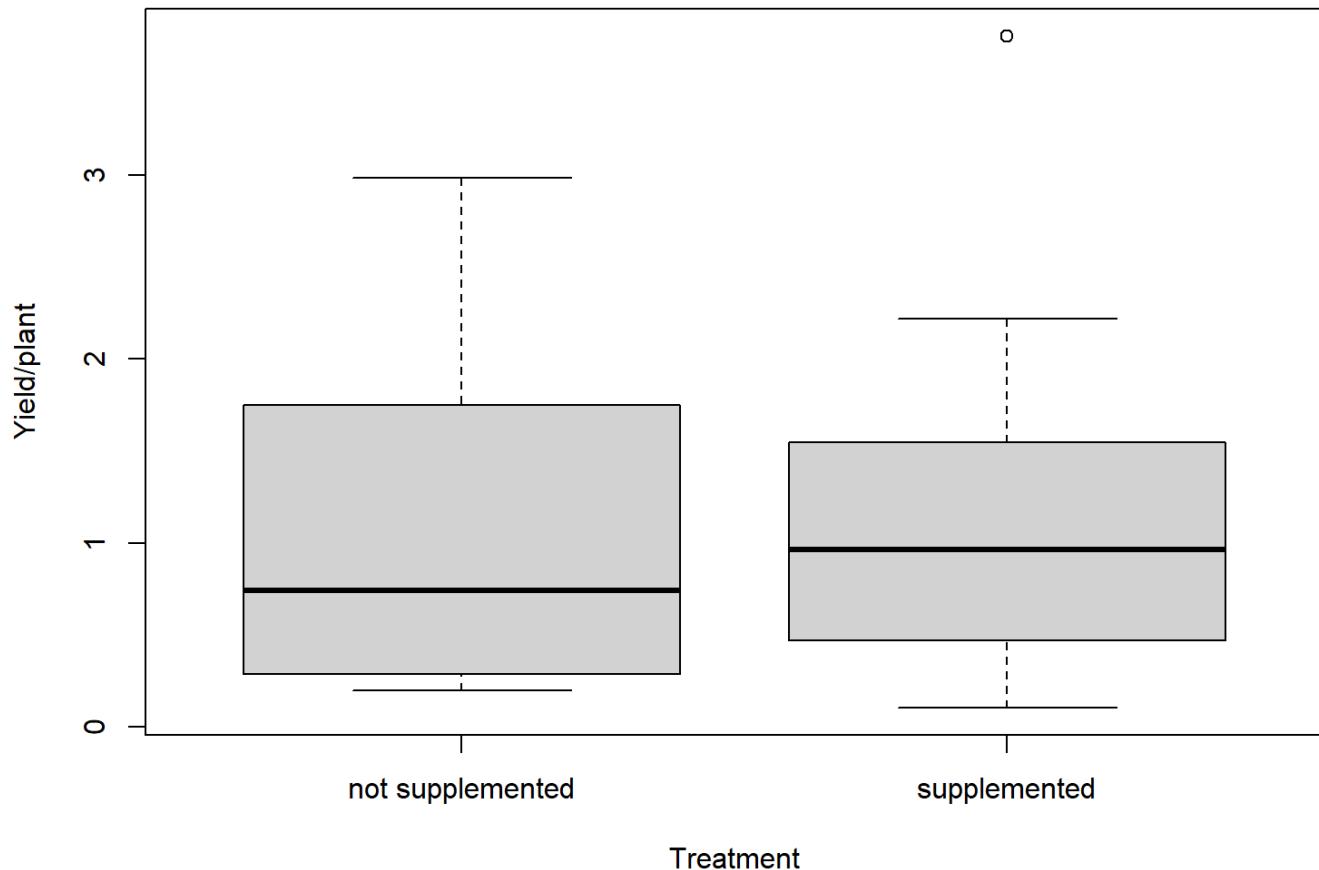


```
ggplot(data=squash_beans_df, mapping=aes(x=planting, y=visits.flower.minute)) +  
  geom_point() + theme_classic() +  
  scale_y_continuous(limits = c(0, 5), breaks = seq(0, 5, by = 1)) +  
  ylab("Visit/flower/minute") +  
  scale_x_continuous(limits = c(0, 3), breaks = seq(0, 3, by = 1),  
  expand=c(0,0)) +  
  xlab("Crop Cycle")
```

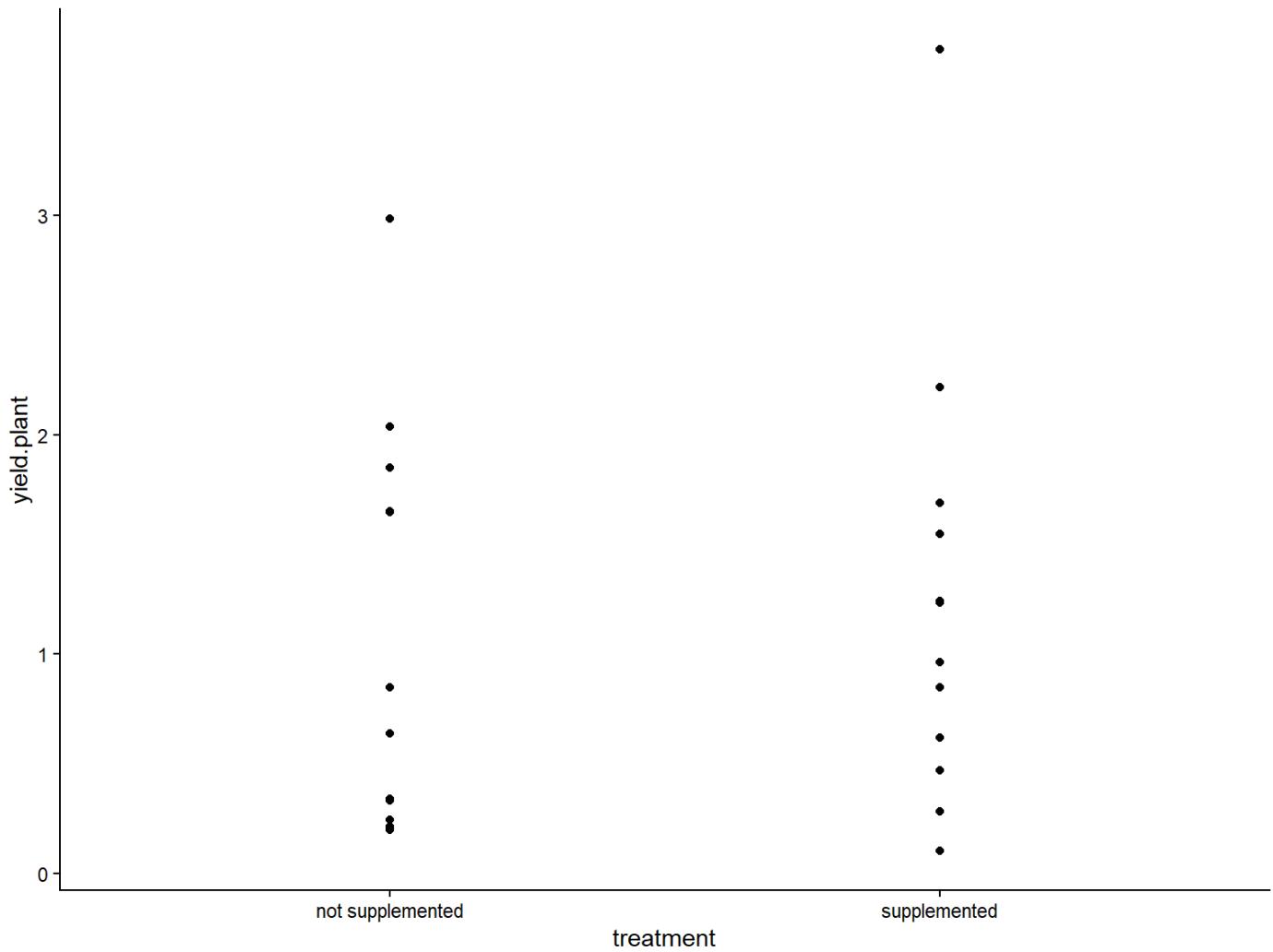


Plot of yield/plant as a function of treatment:

```
plot(yield.plant~treatment, data= yield_df, xlab= "Treatment", ylab= "Yield/plant")
```

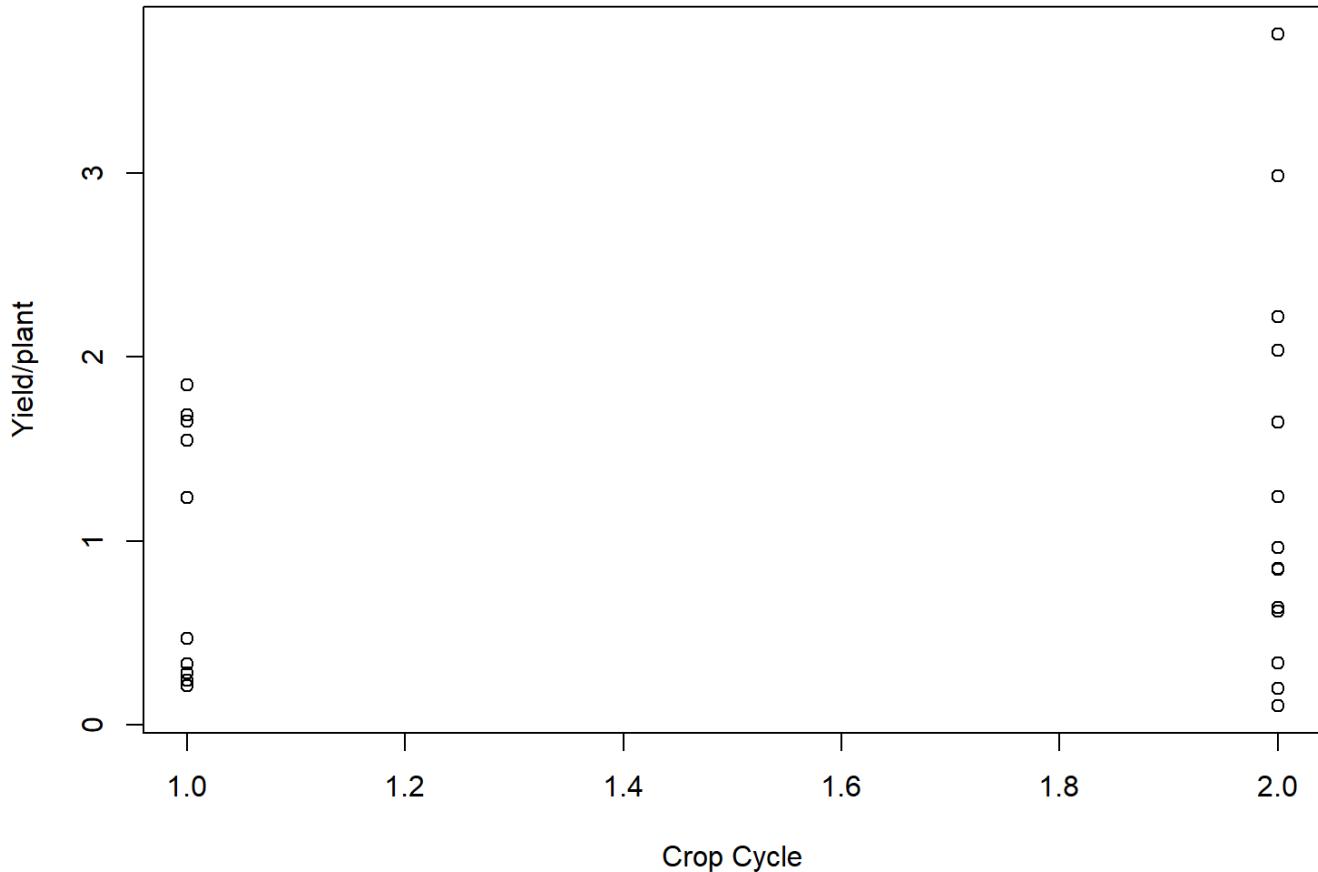


```
ggplot(data=yield_df, mapping=aes(x=treatment, y=yield.plant)) +  
  geom_point() + theme_classic()
```



Plot of yield/plant as a function of planting date (~proxy for bloom date):

```
plot(yield.plant~planting, data= yield_df, xlab= "Crop Cycle", ylab= "Yield/plant")
```



```
ggplot(data=yield_df, mapping=aes(x=planting, y=yield.plant)) +  
  geom_point() + theme_classic() +  
  scale_y_continuous(limits = c(0, 4), breaks = seq(0, 4, by = 0.25)) +  
  ylab("Yield/plant") +  
  scale_x_continuous(limits = c(0, 3), breaks = seq(0, 3, by = 1),  
  expand=c(0,0)) +  
  xlab("Crop Cycle")
```

