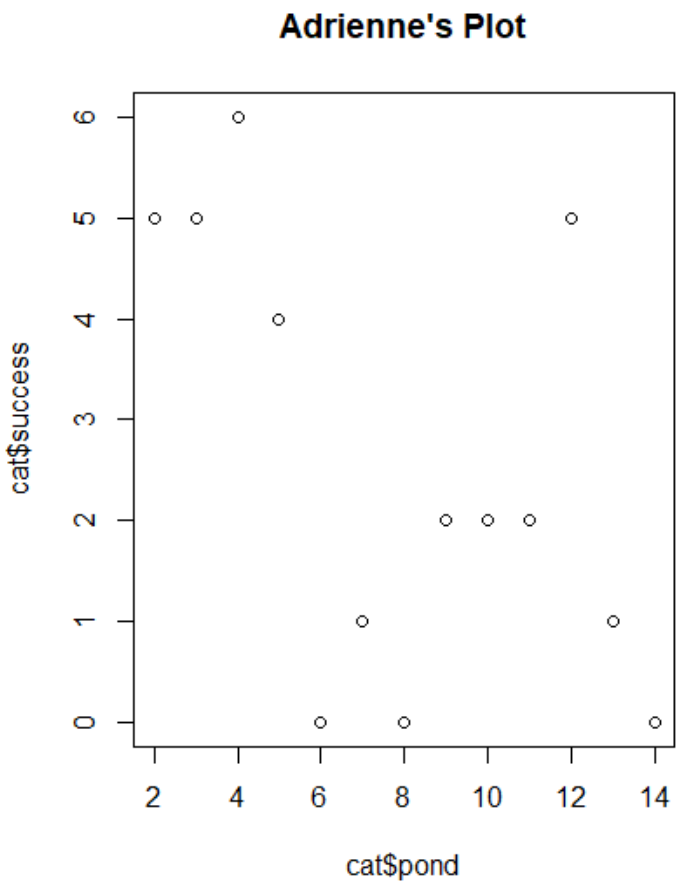
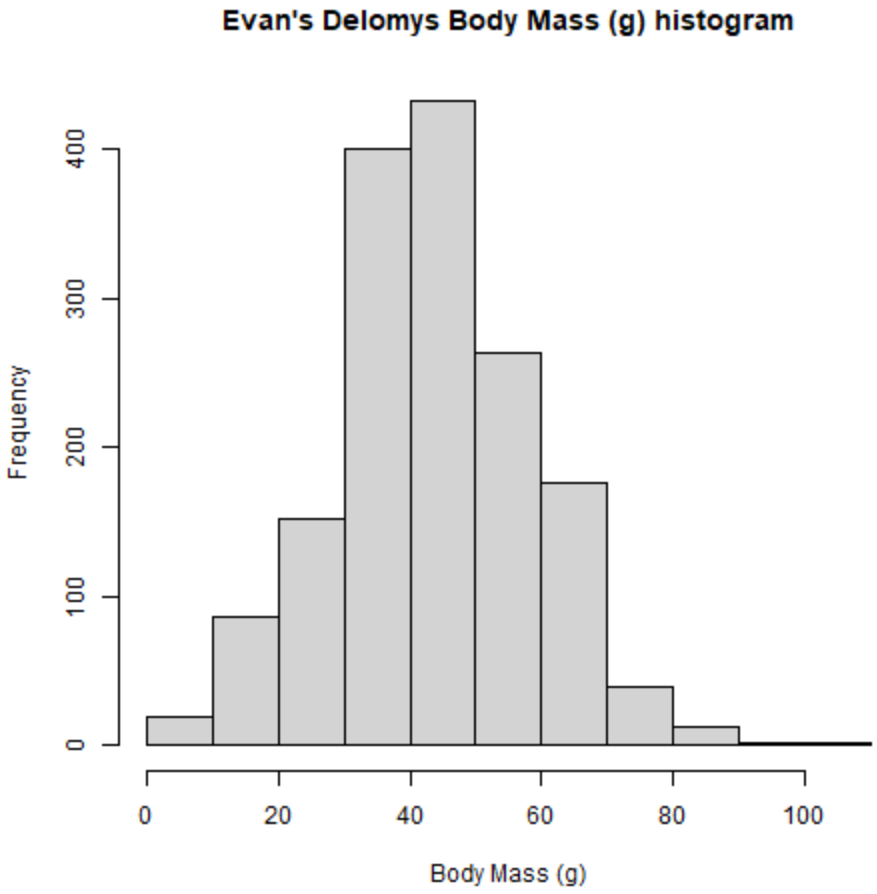
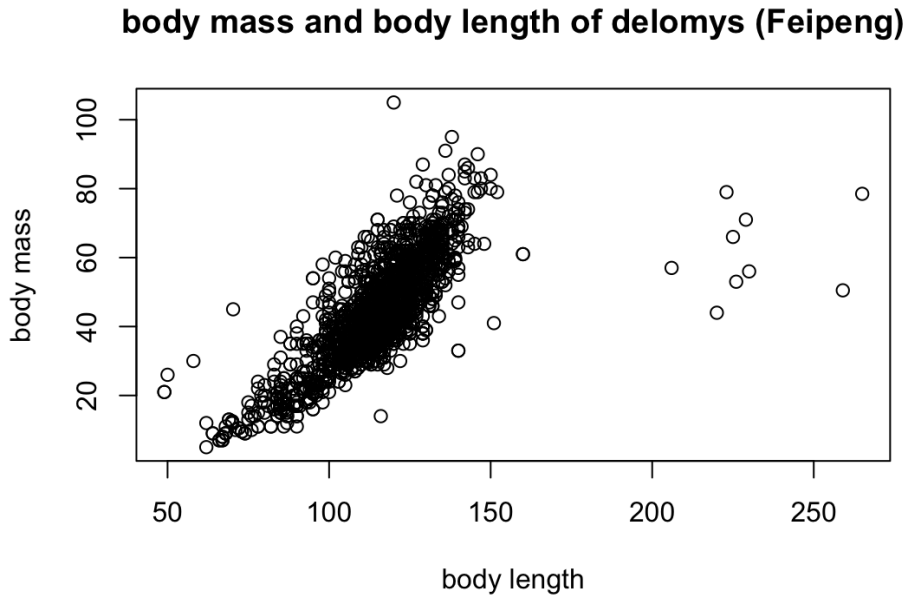
R Code:

```
> library(here)
here() starts at C:/Users/adrienne.dunk/Documents/environmental_data
> dat_catrate <- data.frame(read.csv(here("data", "catrate.csv")))
> head(dat_catrate)
> dat_delomys <- read.csv(here("data", "delomys.csv"))
> rope_dat <- read.csv(here("data", "rope.csv"))
plot(rope_dat, main = "Adrienne's Plot")
```

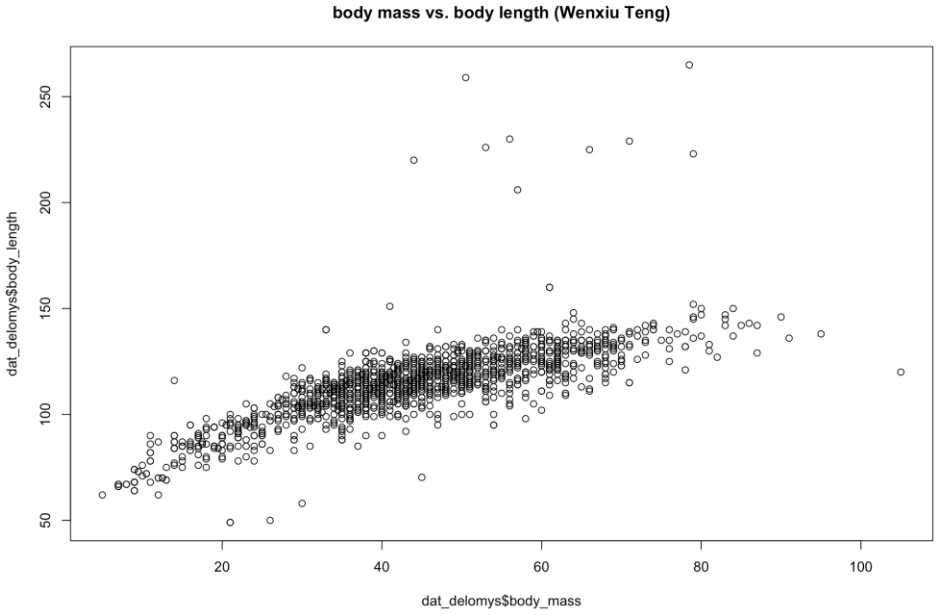


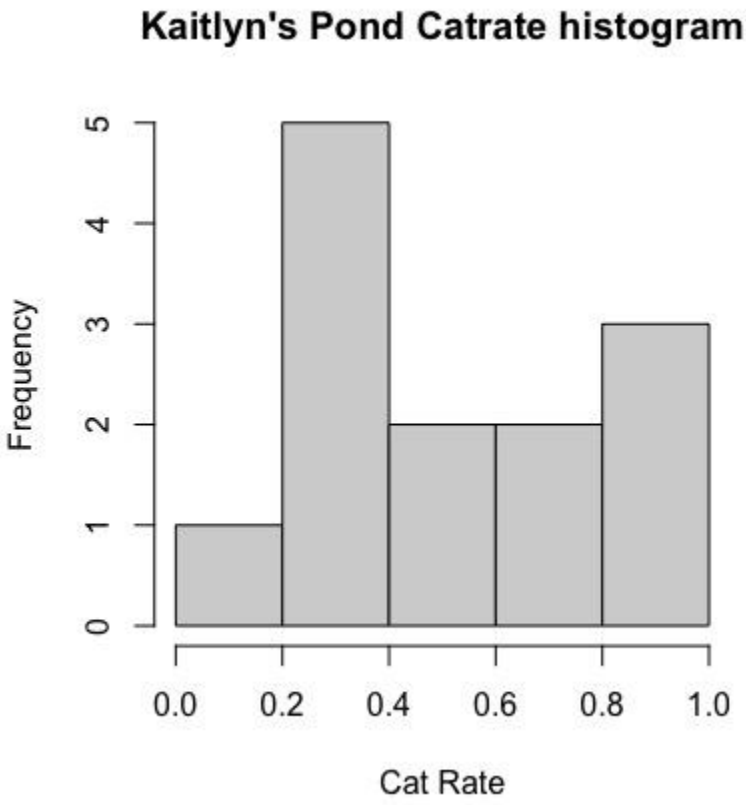

Evan's Delomys Body Mass (g) histogram
Frequency
0
100
200
300
400
0
20
40
60
80
100
Body Mass (g)


body mass and body length of delomys (Feipeng)
body mass
20
40
60
80
100
50
100
150
200
250
body length

```
dat_catrate = read.csv(here("data", "catrate.csv"))

dat_delomys = read.csv(here("data", "delomys.csv"))

dat_rope = read.csv(here("data", "rope.csv"))
```

```
require(here)

dat_delomys = read.csv(

  here("data", "delomys.csv")

)

head(dat_delomys)

plot(dat_delomys$body_mass,dat_delomys$body_length,main = "body mass vs. body length (Wenxiu Teng)")
```





We had trouble naming the correct axes when creating a histogram without an error. Instead of using x=, we figured out that the format “data$column” worked successfully each time.