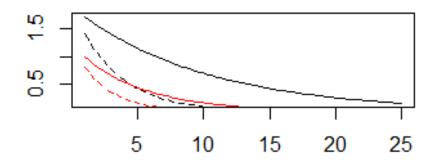
Heather Siart ECO-634 Environmental Data Analysis – Lab 10/27/2021 – LATE Partners: Jahiya Clark

Lab 5: Uncertainty, Samples, and Populations

```
Q1:
    Exp_fun = function (x, a, b)
    {
        return(a * exp ( -b * x ))
      }
      curve(
        exp_fun(x, 0.3, (1.15)), add = FALSE, from = 0, to = 50, ann = FALSE, axes = TRUE, ylab = "f(x)"); box()

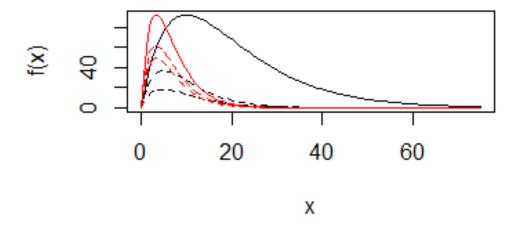
Q2:
```



Q3: When you change parameter "a" the line begins at a different height on the y axis.

Q4: When you change parameter "b" it causes a drastic curve in the line along the x axis.

Q5:



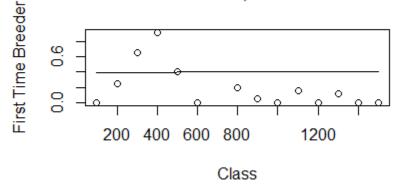
Q6: By changing parameter "a" you change the slope of the line. The red lines have a much more dramatic curve where the black lines are rounder, and softer.

Q7: Changing parameter "b" changes how high the peak of the curve is on the graph.

Q8: To figure out the values for my line I used locator(1) and tried to pick a location that visually looked balanced within the data. This gave me and x value of 503.032, and a Y value of 0.3993503.

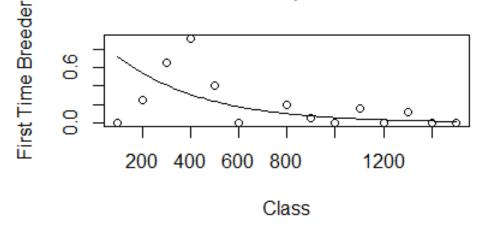
Q9:

Marbled Salamander, first time breeders



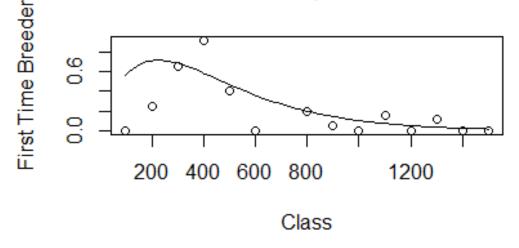
Q10: I used 0.97 for parameter "a" and 1/350 for parameter "b". I choice both of these values by changing them around and seeing visually what curve fit the data points best. **Q11:**

Marbled Salamander, first time breeders



Q12: I chose 0.0087 for parameter "a" and 1/225 for parameter "b". I used these values because they visually fit the data points best. My curve touches some data points but stays in the middle of them too. **Q13:**

Marbled Salamander, first time breeders



Q14:

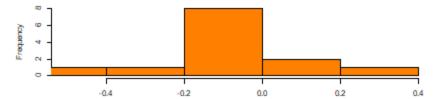
Locator(15)

dat_dispersal\$resids_ricker <- c(0.386, 0.386, 0.386, 0.386, 0.399, 0.399, 0.399, 0.412, 0.425, 0.425, 0.412, 0.412, 0.425)

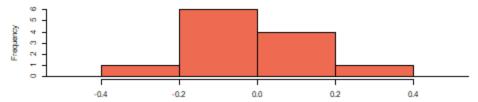
dat_dispersal\$resids_linear <- c(0.672, 0.646, 0.606, 0.550, 0.489, 0.454, 0.393, 0.332, 0.241, 0.196, 0.150, 0.0850, 0.0496, 0.00408, 0.00331)

```
dat_dispersal$resids_exp <- c(0.728, 0.531, 0.399, 0.307,
              0.241, 0.175, 0.135, 0.096,
              0.0961, 0.069, 0.056, 0.0301,
              0.0301, 0.0301, 0.043)
View(dat_dispersal)
resid linear <- c(dat dispersal$disp.rate.ftb - dat dispersal$resids linear)
resid_exp <- c(dat_dispersal$disp.rate.ftb - dat_dispersal$resids_exp)</pre>
resid_ricker <- c(dat_dispersal$disp.rate.ftb - dat_dispersal$resids_ricker)</pre>
disp_resids <- data.frame(dat_dispersal$resids_linear, dat_dispersal$resids_exp,
dat dispersal$resids ricker)
require(here)
png(
 filename = here("images", "lab_05_hist.png"))
par(mfrow = c(3, 1))
hist(resid_linear, main = "Histogram of Linear Resids", xlab = "", col = "darkorange1", xlim = c(-0.5, 0.5))
hist(resid_exp, main = "Histogram of Exponential Resids", xlab = "", col = "coral2", xlim = c(-0.5, 0.5))
hist(resid_ricker, main = "Histogram of Ricker Resids", xlab = "", col = "darkgoldenrod2", xlim = c(-0.7,
0.3))
dev.off()
Q15:
```

Histogram of Linear Resids



Histogram of Exponential Resids



Histogram of Ricker Resids

