

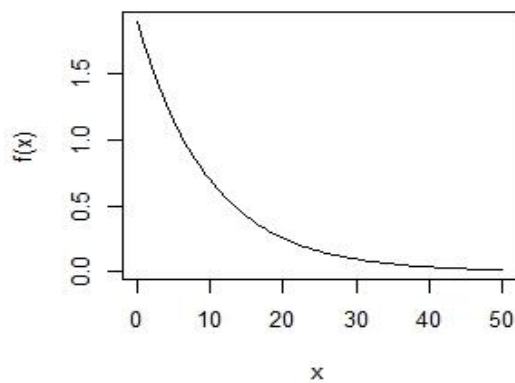
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10/12/22

1.

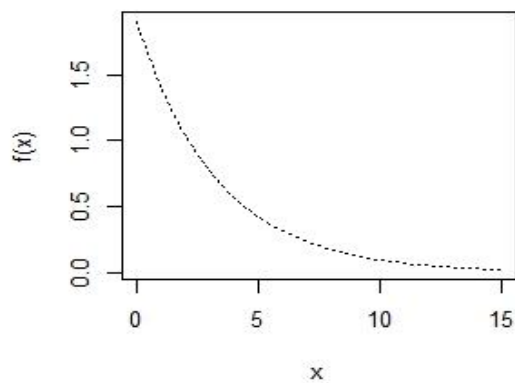
```
exp_fun <- function(a, b, x) {  
  return(a * exp(-b * x))  
}
```

2.

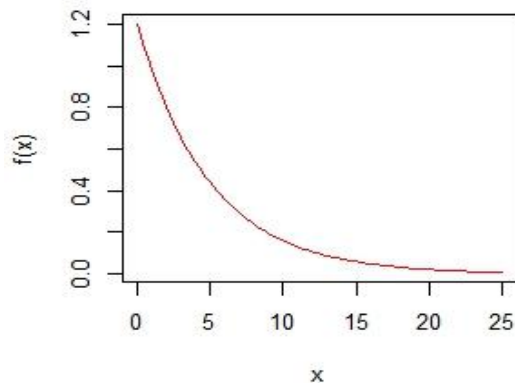
Exponential function (a = 1.9, b = 0.1)



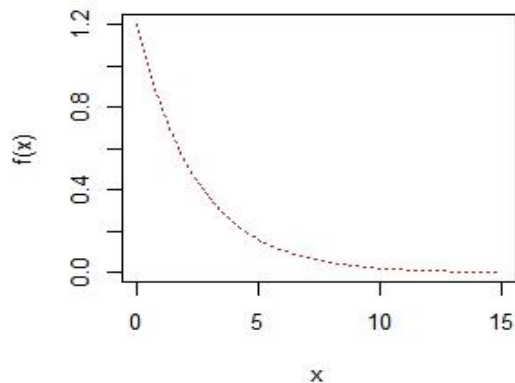
Exponential function (a = 1.9, b = 0.3)



Exponential function (a = 1.2, b = 0.2)



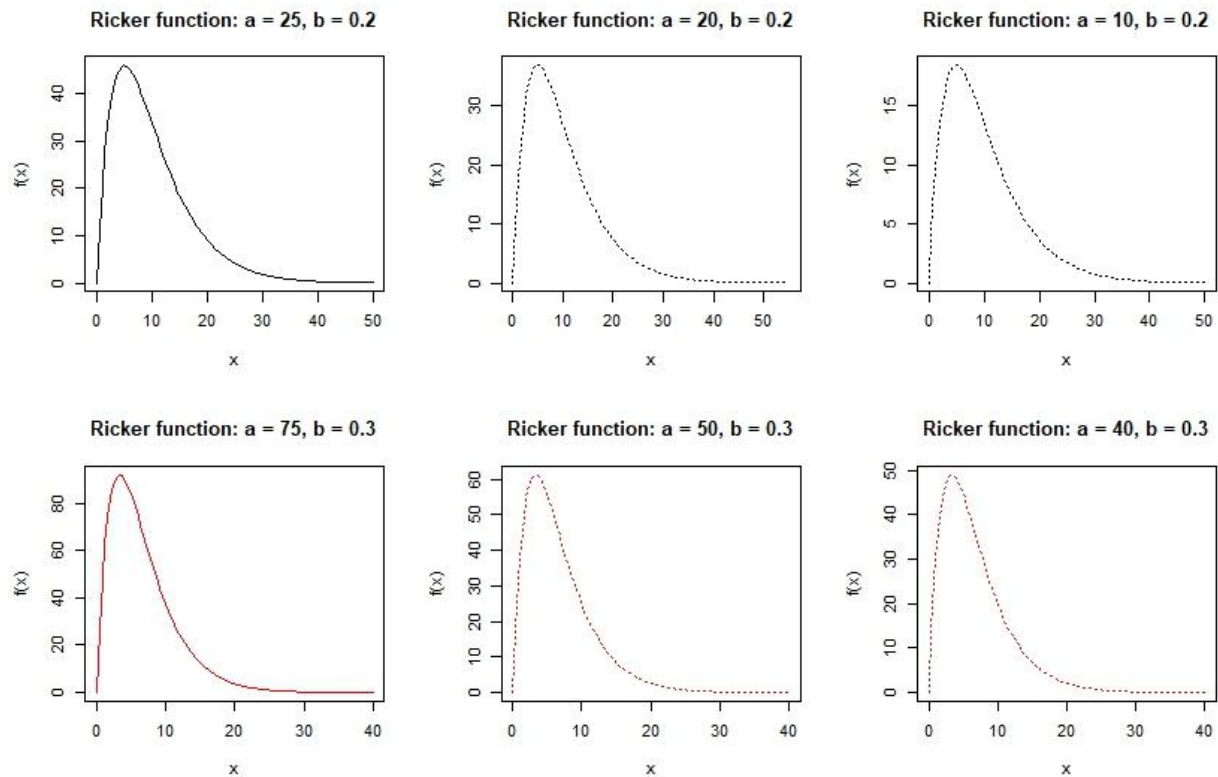
Exponential function (a = 1.2, b = 0.4)



3. As you vary parameter **a** in exp_fun, the starting height of the curve on the Y-axis is changed

4. As you vary parameter **b** in exp_fun, the rate of change in the curve is increased or decreased

5.

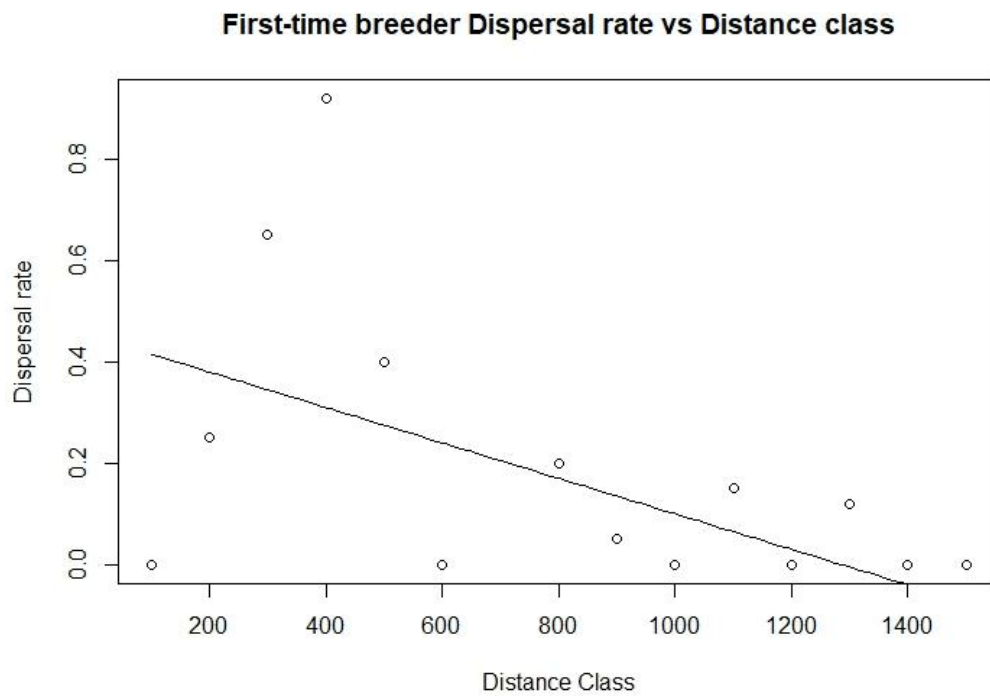


6. As you vary parameter **a** in `ricker_fun`, the max height of the initial curve on the y-axis is changed

7. As you vary parameter **b** in `ricker_fun`, the rate of change in the curve is increased or decreased

8. Slope = -0.00035, $x_1 = 0$, $y_1 = 0.45$. I chose the y_1 value by guessing where the majority of the plotted values were below. I chose the slope by taking y_1 and dividing it by an estimate of the largest x -value. x_1 is zero because that just looked right.

9.



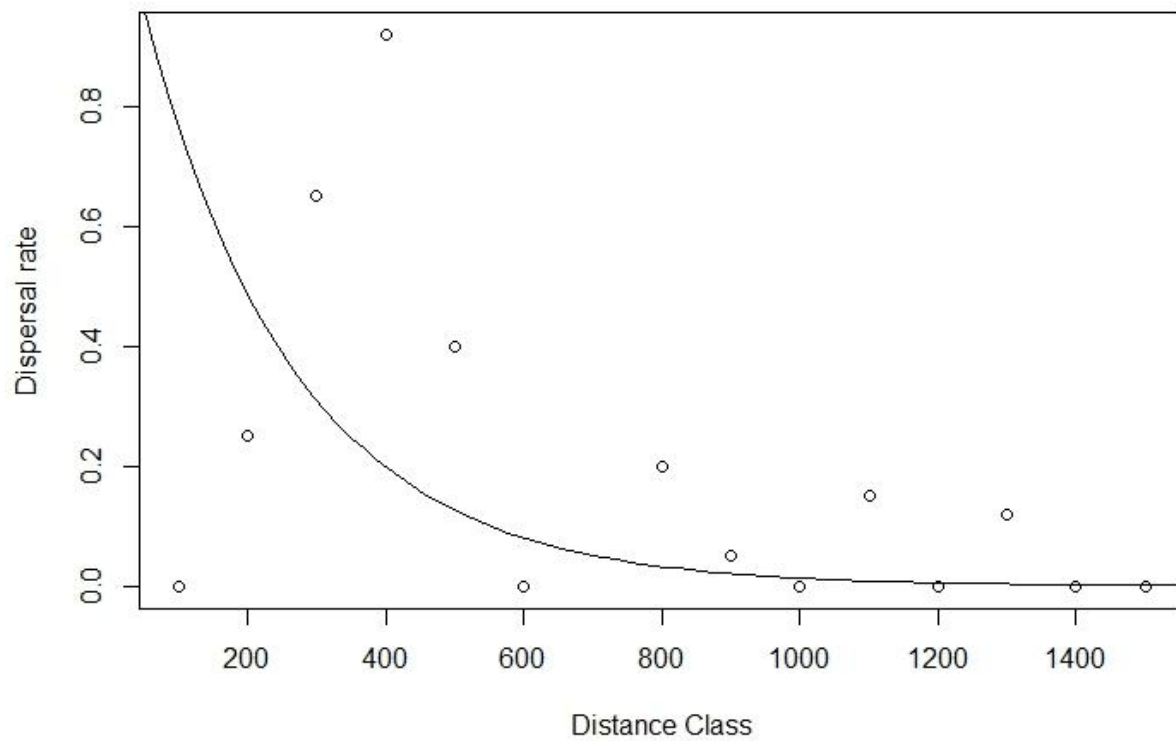
10.

$a = 1.2$, $b = 0.0045$

I chose these values because the value of **a** is inclusive of all the shown points and the line created from using **b** most closely resembles the rate of change in the given points

11.

First-time breeder Dispersal rate vs Distance class (exponential)



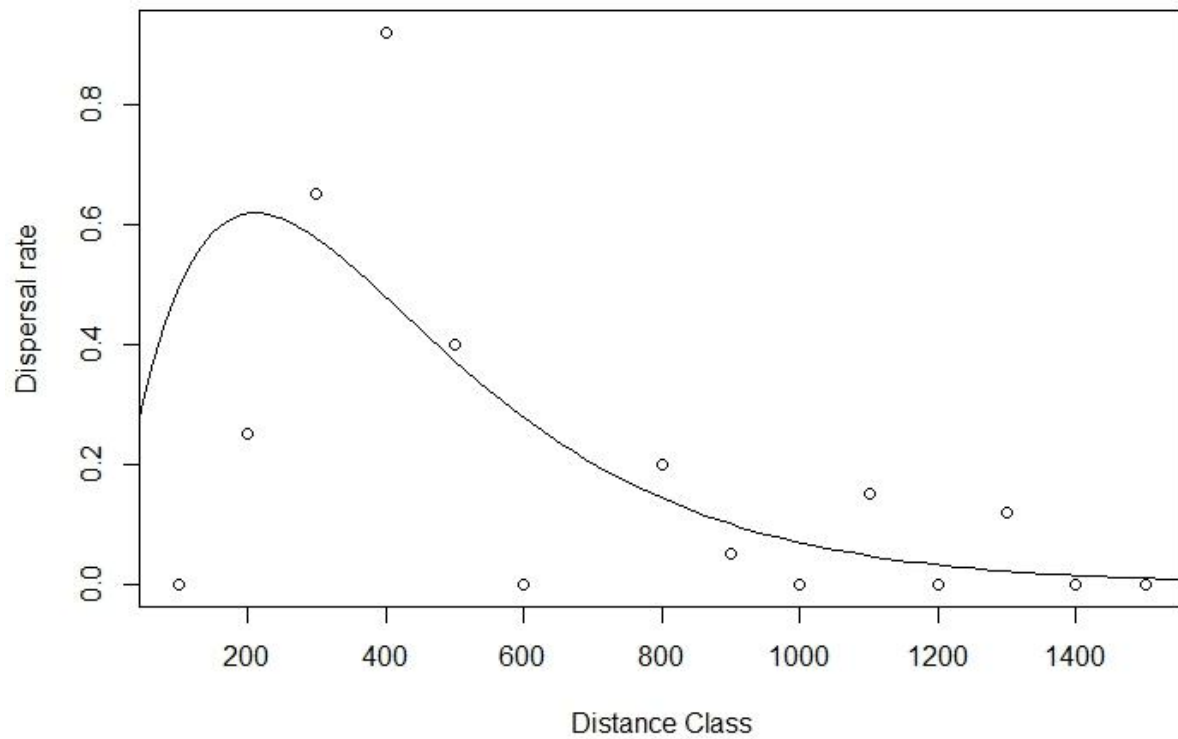
12.

$a = 0.008$, $b = 0.00475$

I chose these values because the initial slope given by **a** is sufficient to include the majority of the given points and the rate of change given by **b**

13.

First-time breeder Dispersal rate vs Distance class (Ricker)



14.

```
resids_linear <- data.frame(lm_obs - lm_exp)  
resids_exp <- data.frame(expo_obs - expo_exp)  
resids_ricker <- data.frame(rick_obs - rick_exp)
```

```
residual_model_df <- cbind(resids_linear, resids_exp, resids_ricker)
```

15.

