# Modelling\_Group\_Project\_R

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November 27, 2017

# Group Project Question 1 - Lotka-Volterra Model

#### **Equations**

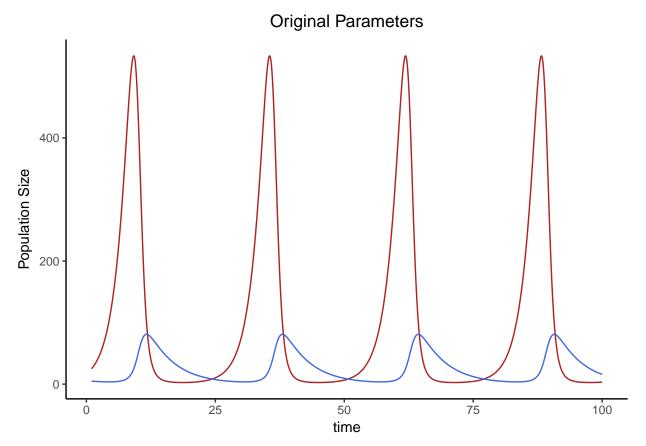
```
Prey dh/dt = b * H - a * P * H
Predator dP/dt = e * a * P * H - s * P
```

#### Initial Model

Below we load the required modules, set initial parameters, and use the function ode() to run our model.

```
# Load the deSolve package required for solving differential equations in R and ggplot2 package for gra
library(deSolve)
library(ggplot2)
# Create a custom function for simulating Lotka-Volterra Competition Model
predPreySim<-function(t,y,p){</pre>
 herbivore=y[1]
  carnivore=y[2]
 h_birthrate=p[1]
  c_attackrate=p[2]
  h_c_efficiency=p[3]
  c_{death=p[4]}
  dHerb_dt=(h_birthrate*herbivore)-(c_attackrate*carnivore*herbivore)
  dCarn_dt=(h_c_efficiency*c_attackrate*carnivore*herbivore)-(c_death*carnivore)
  return(list(c(dHerb_dt,dCarn_dt)))
}
# Set parameters, initial state variables, and simulation timepoints
y = c(25,5)
p = c(0.5, 0.02, 0.1, 0.2)
times=seq(from=1,to=100,by=0.1)
# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"</pre>
colnames(modelSim)[3] <- "predator"</pre>
# Convert to datafrome for plotting purposes
modelSim2 <- as.data.frame(modelSim)</pre>
```

```
# plot simulation
ggplot(data=modelSim2,aes(x=time)) + geom_line(aes(y=herbivore), colour="firebrick") + geom_line(aes(y=herbivore))
```



Initial summary of output below. How the parameters change will be determined by comparing results to above graph and summary below.

```
#show result summary of the data frame
summary(modelSim)
```

```
##
           herbivore
                        predator
## Min.
            2.642454
                        3.676633
                       5.386827
## 1st Qu.
            4.364783
## Median
            21.810086 15.707261
           105.735381 26.002543
## Mean
## 3rd Qu. 141.380539
                      41.663795
## Max.
           533.347325 80.982968
## N
           991.000000 991.000000
## sd
           155.689454 24.543401
```

#### Changing the number of prey

Adjust initial parameters. Change prey from 25 to 125.

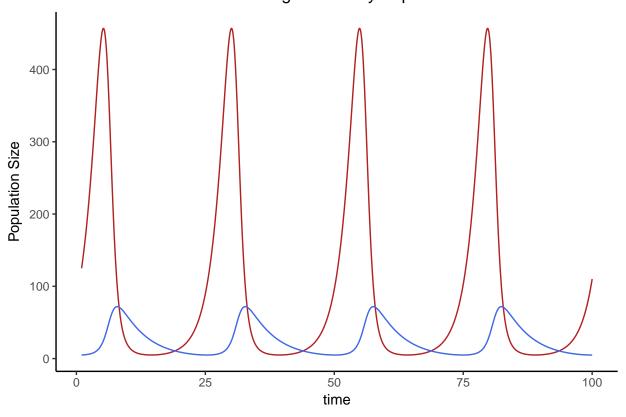
```
# simulate dynamics by resetting parameters and intitial conditions
y = c(125,5)
p = c(0.5,0.02,0.1,0.2)
times=seq(from=1,to=100,by=0.1)
```

```
# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)

# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=prediction))</pre>
```

# **Increasing Initial Prey Population**



# #show result summary of the data frame summary(modelSim)

```
##
            herbivore
                        predator
## Min.
             4.975246
                        4.933426
## 1st Qu.
             7.771253
                        7.059210
            26.336462
## Median
                       15.703783
## Mean
            99.961990 25.043921
## 3rd Qu. 142.406059
                       39.363238
## Max.
           456.995062 71.922218
## N
           991.000000 991.000000
## sd
           135.983485 21.480734
```

Adjust initial parameters again. Change prey from 25 to 5.

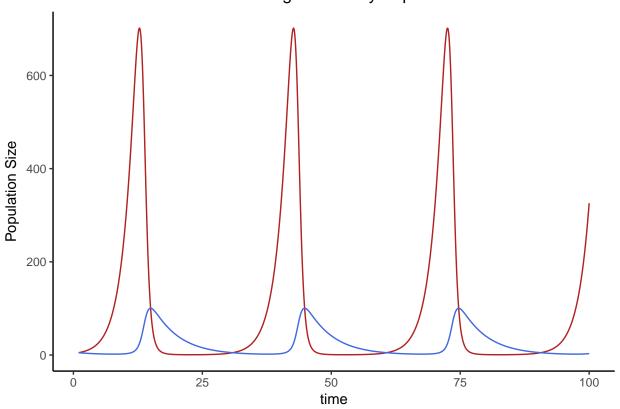
```
# simulate dynamics by resetting parameters and intitial conditions
y = c(5,5)
p = c(0.5,0.02,0.1,0.2)
times=seq(from=1,to=100,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)

# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=predator))</pre>
```

# **Decreasing Initial Prey Population**



# $\begin{tabular}{ll} \#show\ result\ summary\ of\ the\ data\ frame \\ summary\ (modelSim) \end{tabular}$

```
predator
##
             herbivore
## Min.
            0.6326619
                         1.952785
                         2.905456
## 1st Qu.
            1.4337351
## Median
            10.1132523
                         8.216084
## Mean
            97.7414364 22.871279
## 3rd Qu. 94.6878219 32.804991
           701.7623894 100.467088
## Max.
```

```
## N 991.000000 991.000000
## sd 177.0075387 28.422647
```

The above two models changed the number of prey from 25 to 125 and 5, respectively. Interestingly, changing the prey number does not change much of the overall model. There are some slight changes in overall min/max of predator/prey, however the frequency does not seem to change at all. Interestingly, decreasing the inital number of prey increased the maximum number of prey and predators. The relative numbers between the predator and prey does not change much.

#### Changing the number of predators

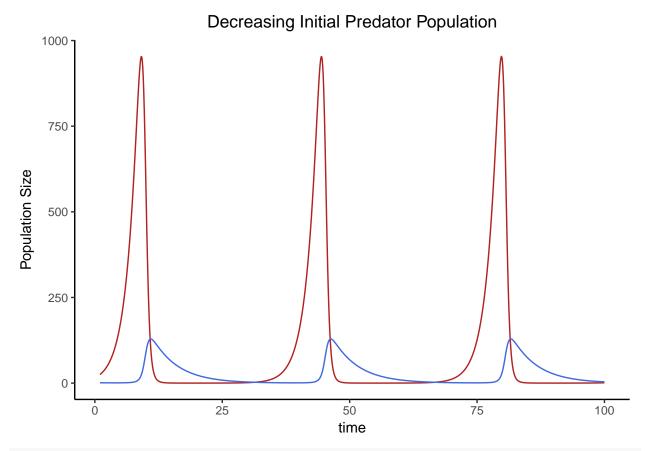
Now let's do the same thing for the pedators as we did for the prey above. Change predator initial values from 5 to 1.

```
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,1)
p = c(0.5,0.02,0.1,0.2)
times=seq(from=1,to=100,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)

# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=prediction))</pre>
```



# #show result summary of the data frame summary(modelSim)

```
##
              herbivore
                          predator
             0.06872585
                          0.7681323
## Min.
## 1st Qu.
             0.15660691
                          1.7932943
## Median
             1.50851963
                          8.6631537
## Mean
           106.43758841
                         26.5960674
## 3rd Qu. 63.43020431
                         37.5351994
## Max.
           953.84907677 128.8241713
## N
           991.00000000 991.0000000
           227.67255575 35.6732609
## sd
```

Change predator from 5 to 25

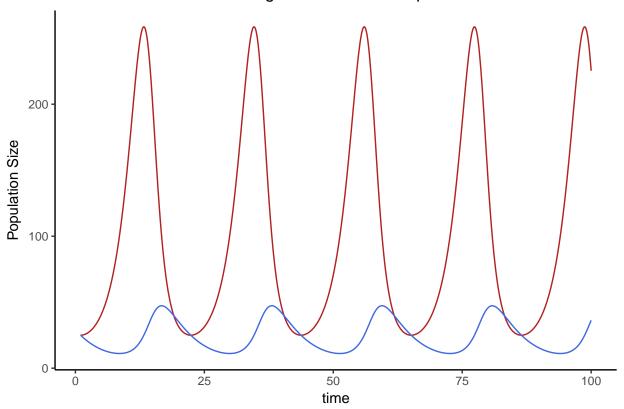
```
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,25)
p = c(0.5,0.02,0.1,0.2)
times=seq(from=1,to=100,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)</pre>
```

```
# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=pro
```





#show result summary of the data frame
summary(modelSim)

```
##
           herbivore predator
## Min.
            25.00000
                     11.12827
## 1st Qu.
            32.82907
                      13.07874
                      19.72100
## Median
            67.63201
           101.89641
## Mean
                      23.89643
## 3rd Qu. 165.16061
                      33.56128
           258.66607
                      47.30765
## Max.
## N
           991.00000 991.00000
## sd
            79.45680 12.04776
```

Unlike changing the intial number of prey, changing the inital number of predators dramatically changes the model output. For starters, the frequency for both prey/predator are decreased/increased when the inital predator count is decreased/increased. Additionally, the inital number of predators seems to have a bigger impact on both max/min numbers of both predator/prey. The relative numbers between the predator/prey does not change much.

## Changing the prey birthrate

Change prey birthrate from 0.5 to 0.1 (and time to 200)

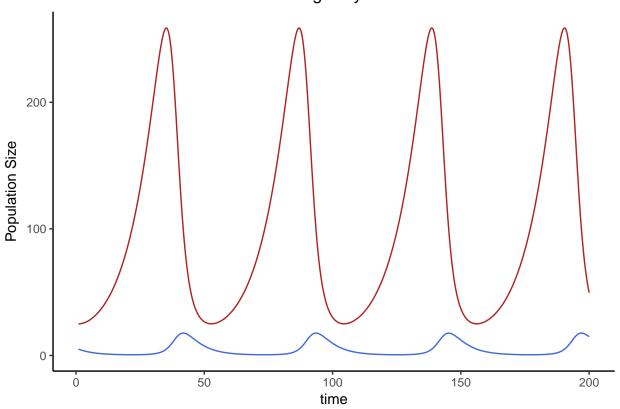
```
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,5)
p = c(0.1,0.02,0.1,0.2)
times=seq(from=1,to=200,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)

# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=predator))</pre>
```

# **Decreasing Prey Birthrate**



# #show result summary of the data frame summary(modelSim)

```
##
            herbivore
                          predator
## Min.
             24.99999
                         0.5784130
             34.98699
## 1st Qu.
                         0.8090509
## Median
             72.61223
                         2.0053024
## Mean
            102.69778
                         4.8304693
## 3rd Qu.
            161.48185
                         7.2861356
            258.66634
## Max.
                        17.6770091
```

```
## N
           1991.00000 1991.0000000
## sd
             77.14174
                          5.4445853
Change prey birthrate from 0.5 to 2.5
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,5)
p = c(2.5, 0.02, 0.1, 0.2)
times=seq(from=1,to=100,by=0.1)
# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"</pre>
colnames(modelSim)[3] <- "predator"</pre>
# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)</pre>
# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=pr
                                     Increasing Prey Birthrate
    3000
Population Size
    2000
    1000
       0
                                                                      75
                               25
                                                  50
                                                                                         100
                                                  time
#show result summary of the data frame
summary(modelSim)
##
               herbivore
                           predator
           9.095405e-12
                           4.693959
## 1st Qu. 9.918855e-11 15.682254
```

## Median 8.119030e-08 60.056245

```
## Mean 1.090432e+02 135.143648
## 3rd Qu. 1.259708e-02 199.813250
## Max. 3.338002e+03 613.872554
## N 9.910000e+02 991.000000
## sd 4.676766e+02 161.949298
```

Changing prey birthrate has significant effects on both max/min of predator prey and of the frequency of both also. Decreasing prey birthrate significantly decreased both frequency and max/min of both predator/prey, much more than changing the inital numbers of prey/predator. Additionally, the prey birthrate seems to have a strong effect on the relative numbers between prey/predator, as shown by the differences in mean between the two. Additionally, increasing prey birthrate causes an interesting phenomenon where most of the times the prey are mostly dead but shoot up exponentially every now and then, whilst decreasing birthrate causes a more conservative change in the number of prey, more like a sinusoidal curve.

#### Changing predator attack rate

Change predator attack rate from 0.02 to 0.1

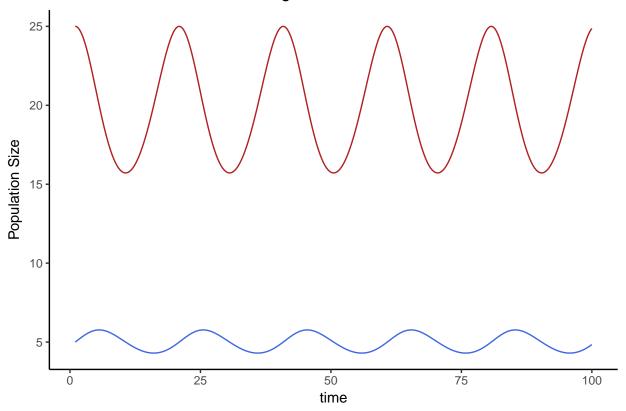
```
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,5)
p = c(0.5,0.1,0.1,0.2)
times=seq(from=1,to=100,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)

# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=pr</pre>
```





# #show result summary of the data frame summary(modelSim)

```
##
            herbivore
                          predator
## Min.
            15.715773
                         4.3024738
## 1st Qu.
            16.744698
                         4.4830608
## Median
            19.618941
                         4.9686235
## Mean
            19.972108
                         5.0004559
## 3rd Qu.
            23.192277
                         5.5175409
## Max.
            25.000000
                         5.7691227
## N
           991.000000 991.0000000
             3.264623
                         0.5194991
## sd
```

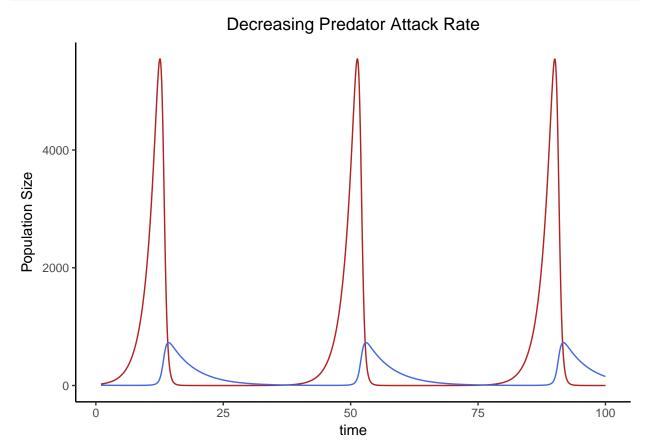
Change predator attack rate from 0.02 to 0.004

```
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,5)
p = c(0.5,0.004,0.1,0.2)
times=seq(from=1,to=100,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)</pre>
```

```
# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=pr
```



#show result summary of the data frame
summary(modelSim)

```
##
              herbivore
                          predator
## Min.
              0.0839023
                           2.156335
## 1st Qu.
              0.3204918
                           4.319488
              9.4939063
## Median
                         27.365309
## Mean
            585.7889308 139.188351
## 3rd Qu.
            313.3331253 203.998745
## Max.
           5549.1553765 730.282173
## N
            991.0000000 991.000000
## sd
           1287.3686222 202.618168
```

Changing predator attack rate has significant effects on the overall model. Both the max/min and frequecy of predator/prey are significantly effected. Increasing the attack rate significantly decreases both the number of predators and prey and also increases the frequency of both predator/prey cycle, whilst decreasing the attack rate does the opposite. Additionally, decreasing the attack rate brings the prey to near extinction levels, as evidenced by the 0.08 minimum of prey, perhaps making it an unsustainable model. However, the relationship between the means does not change much from the original model (~1:4 ratio for predator:prey).

#### Changing conversion efficiency

Change conversion efficiency from 0.1 to 0.5

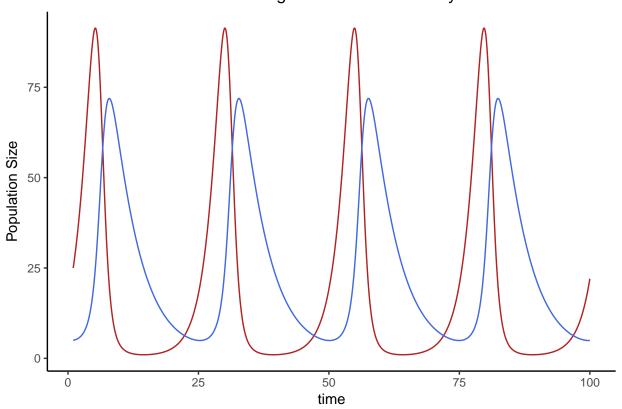
```
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,5)
p = c(0.5,0.02,0.5,0.2)
times=seq(from=1,to=100,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)

# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=prediction))</pre>
```

# **Increasing Conversion Efficiency**



# #show result summary of the data frame summary(modelSim)

```
predator
##
             herbivore
## Min.
             0.9950496
                         4.933426
                         7.059209
## 1st Qu.
             1.5542510
## Median
             5.2672934 15.703786
## Mean
            19.9923981
                        25.043921
## 3rd Qu.
           28.4812166
                        39.363239
            91.3990109 71.922212
## Max.
```

```
## N 991.000000 991.000000
## sd 27.1966963 21.480734
```

Change conversion efficiency from 0.1 to 0.02

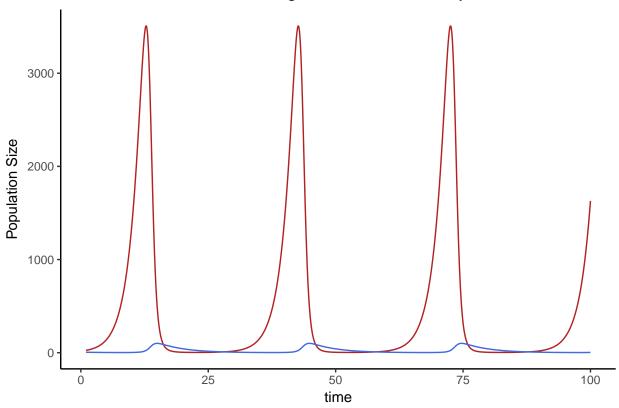
```
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,5)
p = c(0.5,0.02,0.02,0.2)
times=seq(from=1,to=100,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)

# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=prediction))</pre>
```

# **Decreasing Conversion Efficiency**



# #show result summary of the data frame summary(modelSim)

##	herbivore	predator
## Min.	3.163315	1.952785
## 1st Qu.	7.168682	2.905456
## Median	50.566255	8.216081

```
## Mean 488.707225 22.871278
## 3rd Qu. 473.439118 32.804981
## Max. 3508.812000 100.467086
## N 991.000000 991.000000
## sd 885.037682 28.422646
```

Changing conversion efficiency also changes min/max, frequency, and predator/prey relationships. However, the most interesting aspect of changing the conversion efficiency is the change in relative means between the prey/predator relationship. Increasing conversion efficiency brings the relative number of prey much closer to the relative mean of the predator, as would be expected because the predator can more efficiently convert the prey into momre predators.

#### Changing predator death rate

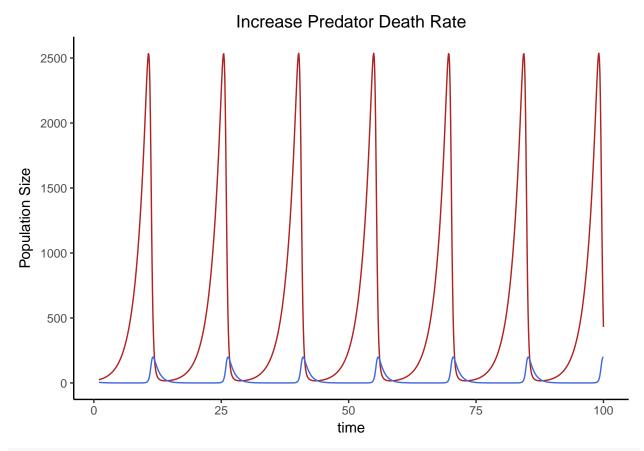
Change predator death rate from 0.2 to 1

```
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,5)
p = c(0.5,0.02,0.1,1)
times=seq(from=1,to=100,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)

# plot simulation
ggplot(data=modelSim2) + geom_line(aes(x=time,y=herbivore),color="firebrick")+geom_line(aes(x=time,y=pr</pre>
```



# #show result summary of the data frame summary(modelSim)

```
predator
##
            herbivore
             16.39540
                        0.06861493
## Min.
## 1st Qu.
             36.67678
                        0.14542813
## Median
            154.09082
                        1.00662302
## Mean
            518.29438 23.64104318
## 3rd Qu. 722.83426 15.08256137
## Max.
           2536.99386 199.42709869
## N
            991.00000 991.00000000
            707.82100 48.74699762
## sd
```

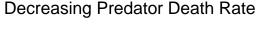
Change predator death rate from 0.2 to 0.04

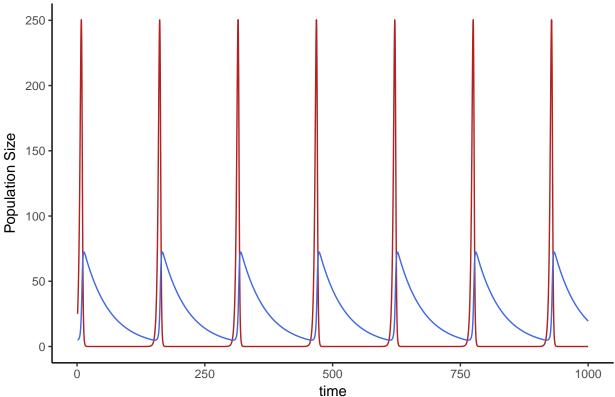
```
# simulate dynamics by resetting parameters and intitial conditions
y = c(25,5)
p = c(0.5,0.02,0.1,0.02)
times=seq(from=1,to=1000,by=0.1)

# Simulate the model using ode
modelSim=ode(y=y,times=times,func=predPreySim,parms=p)
colnames(modelSim)[2] <- "herbivore"
colnames(modelSim)[3] <- "predator"

# save model as a data frame for plotting purposes
modelSim2 <- as.data.frame(modelSim)</pre>
```

# # plot simulation ggplot(data=modelSim2) + geom\_line(aes(x=time,y=herbivore),color="firebrick")+geom\_line(aes(x=time,y=pro





#show result summary of the data frame
summary(modelSim)

```
##
                            predator
              herbivore
## Min.
           3.340911e-09
                            4.856713
## 1st Qu. 1.776776e-08
                            9.078601
## Median 3.437706e-06
                           19.836287
## Mean
           1.068199e+01
                           26.101291
## 3rd Qu. 1.702700e-02
                           39.313460
## Max.
           2.503980e+02
                           72.404377
           9.991000e+03 9991.000000
## N
## sd
           4.043063e+01
                           19.789750
```

Predator death rate also has significant effects on the max/min and frequency in predator/prey. Additionally, the relative means between the predator/prey relationships is also effected. Increasing predator death rate increases prey counts (both max and mean) but also increases the max predator count. Decreasing predator death rate has the opposite effect while both increase the frequency.

#### Conclusion

Overall, all of the state variable and parameters have some effect on the model, whether they change the max/min of the prey/predator, change the overall frequency, change the rate of increase/decrease, or change the relative means of the two populations. What is interesting to note is the models in which the predator/prey

minimun numbers reach <1. Realistically, it seems impossible to have half a predator or half a prey, suggesting these models are unsustainable. This suggests, that especially when changing the parameters, one must be careful that the parameters are not changed significantly since the model may become unrealistic, unless that is the purpose.

#### Questions

What can you say about the "role" of each parameter?

#### What is the relationship between parameter values and predator-prey cycle length?

Each parameter affects the model in different ways, with some parameters having larger effects than others depending on which parameters you are interested in. Dramatically changing some of the parameters makes the model almost unsustainable, as evidenced by the minimum numbers of prey/predator in each summary statement that are less than 1. Otherwise, however, the role of each parameter is fairly straight forward. For starters, what needs to be understood is that predator/prey relationships are intertwined, as evidenced by their rate equations being dependent on the population size of the other.

The prey birth rate has a significant effect on the number of prey in the model, which inherently makes sense. Since the prey birthrate is the only factor that increases the number of prey (assuming the second factor will always be negative or zero), increasing/decreasing this number understandably increase/decreases the number of prey. However, increasing the prey birth rate by a factor of five does not increase the number of prey by a factor of five, because the change in the number of prey (dp/dt), is limited by the number of prey also. Since the number of predators is reliant on the number of prey avialable, understandably increaseing the prey birth rate also increases the predator count. However, increasing the prey birthrate causes an exponential increase and sudden crash in population numbers caused by the exponential increase in predator numbers, suggesting that increasing the prey birthrate may be unsustainable. Decreasing the prey birthrate has the opposite effect, however, the model seems more sustainable. Additionally, we do not see the exponential increase and crash of the prey but instead see a more steady increase decrease in prey numbers and a similar trend with the predator numbers. Prey birth rate seems to significantly affect the frequency of the prey/predator model, where decreasing prey birth rate decreases the frequency whilst increasing prey birth rate increases the frequency.

Changing the predator attack rate has significant effect on both prey and predator numbers in the model, as expected. Increasing attack rate significantly signicificantly decreases the number of prey, and as a result the maximum number of predators decreases. However, the mean predator count is very close to the original model, most likely due to the dependance of the predator numbers on the prey numbers. Increased predator attact rate increases the frequency of prey/predator turnaround and also creates a steady, sinusodial model. Decreasing attack rate has the opposite effect with exponential increase/decrease of predator/prey population size whilst also lowering the frequency.

The conversion efficiency is interesting, as it is one of the few parameters that significantly affects the ratios between predator and prey population sizes. Increasing conversion efficiency significantly increases the number of predators relative to the number of prey whilst limiting the population size of the prey, and vice versa. The conversion efficiency does not seem to have a significant effect on the frequency of the predator/prey cycle.

Finally, the predator death rate also has effects as expected. Increasing predator death rate causes a surge in prey numbers until such a point that we see a significant enough population of predators to bring the prey population down. Increasing predator death rate also increases the predator/prey cycle frequency. Decreasing the predator death rate has the opposite effect, where we observe a sudden spike in prey numbers when the predator numbers decrease. Due to the predator lingering (because of their lower death rate), prey essentially go extinct until there are low enough numbers of predators that the prey can recover. The lingering of the predators causes a decrease in frequency of the predator/prey cycle, as expected.

#### What can you say about the role of predators in the simulations?

The role of predators in the simulations is to keep in check the population of prey when the number of prey get too high. Without a limiting constraint on the prey, such as the predator, the prey would hypothetically expand exponentially unless there were to be a limiting resource or outside competition. This fact can be observed from the fact that in any of the given models above, the predator numbers only rise when the prey number rises, which inherently makes sense if we observe the equations. Because in the Lotka-Volterra model the only factor that allows predator numbers to increase is e \* a \* P \* H (assuming s \* P is always negative), the number of prey in the model (H) is the limiting factor in the model. And only when the number of predators rises the prey numbers start to decrease, and thus are kept in check. Once the number of prey die down, e \* a \* P \* H goes down and s \* P takes over, causing the overall number of predators to decrease.

# Group Project Question 2 - Rosenzweig-MacArthur Model

## **Model Description**

The Rosenzweig-MacArthur model describes predator-prey dynamics. It is based off of the simpler Lotka-Volterra Model, but it adds two elements:

Prey become self-limiting in absence of predators.

Predator kill density cannot grow indefinitely; there reaches a maximum no matter how large the prey density is.

## **Model Equations**

Rosenzweig and MacArthur described the prey population (H) and the predator population (P) with the following equations:

```
dh/dt = b * H * (1 - alpha * H) - w * H * P/(d + H)

dP/dt = e * w * H * P/(d + H) - s * P
```

#### Qualitative Results

There are three qualitative results for this model:

1. Prey go to carrying capacity, predators go extinct.

```
H^* = 1/aplha = k and P^* = 0
```

2. Prey and predators coexist in stable equilibrium.

```
H^* = s * d/(e * w - s) and P^* = (b * H^* - b * d * alpha - b * alpha * H^* * H^* + d * b)/w
```

3. Prey and predators coexist in an oscillatory manner.

Let's begin by writing a function that describes the model:

```
### Function for Rosenzweig-MacArthur Predator/Prey Equations ###
ddRM <- function (t,y,p){

#Unpack State Variables
H = y[1] #Prey
P = y[2] #Predator</pre>
```

```
#Unpack Model Parameters
b = p[1] #Prey birth rate
e = p[2] #Conversion efficency of prey to predator
s = p[3] #Predator death rate
w = p[4] #Predator attack rate
d = p[5] #Density of prey when predator's kill rate is 1/2*max
alpha = p[6] #Prey self-limiting term

#Defining Rates of Change
dHdt = b*H*(1 - alpha*H) - w*H/(d + H)*P
dPdt = e*w*H/(d + H)*P- s*P

return(list(c(dHdt, dPdt)))
}
```

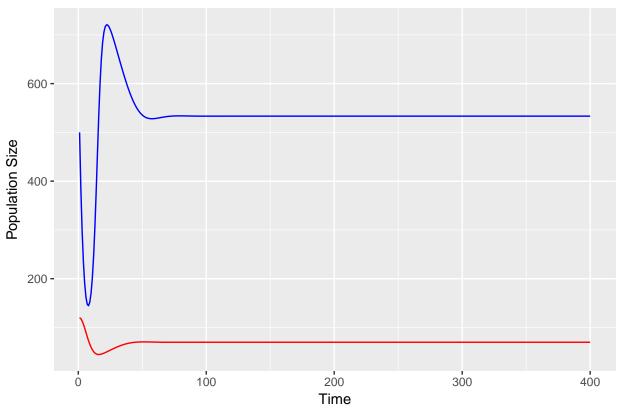
#### **Model Solution**

```
Now, we will solve the following initial parameters:
```

```
b = 0.8
e = 0.07
s = 0.2
w = 5
d = 400
alpha = 0.001
H0 = 500
P0 = 120
### Defining Initial Parameters ###
b = 0.8
e = 0.07
s = 0.2
w = 5
d = 400
alpha = 0.001
params = c(b, e, s, w, d, alpha)
HO = 500
P0 = 120
YO = c(HO, PO)
times = 1:400
modelSim = ode(y=Y0, times = times, func = ddRM, parms = params)
#Subsetting the Results
modelOutput = data.frame(time=modelSim[,1],H=modelSim[,2],P = modelSim[,3])
#Graphing the Results
ggplot(modelOutput) +
```

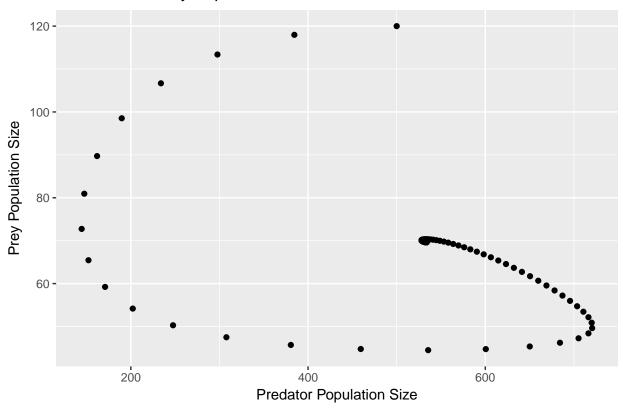
```
geom_line(aes(x = time, y = H), colour = "blue") +
geom_line(aes(x = time, y = P), colour = "red") +
ggtitle("Initial Parameters") +
labs(y="Population Size", x = "Time")
```

# **Initial Parameters**



```
ggplot(modelOutput, aes(H,P)) + geom_point() +
ggtitle("Predator vs. Prey Population") +
labs(y="Prey Population Size", x = "Predator Population Size")
```

# Predator vs. Prey Population



# **Describing Role of Parameters**

Let's now change individual parameters in order to show the role that each one has

## **Predator Attack Rate**

#### Low Attack Rate:

Low attack rate means that predators cannot sustain themselves, and their population goes to zero Prey population reach self-limiting equilibrium

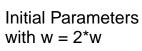
#### High Attack Rate

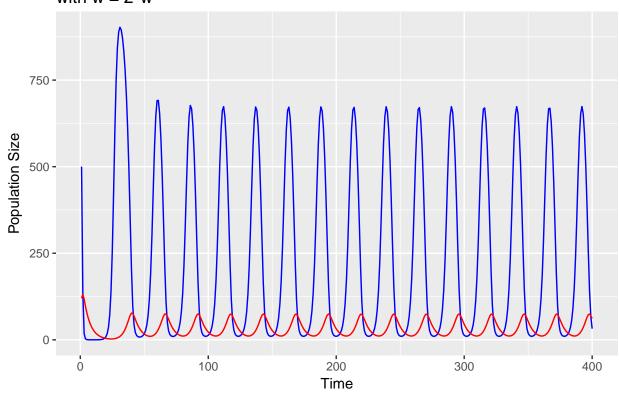
Predators are able to survive

High attack rate implies that prey will be driven to a small population. Predators can't survive if prey population is small, so they decreases

As predator population decreases, prey population is able to increase. The cycle repeats

Long-term behaivor is stable equilibrium





# Initial Parameters with w = 1/2\*w 1000 750 250 0 1000 200 300 400

Time

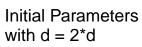
# Prey Denisty to Maximize Predator Attack

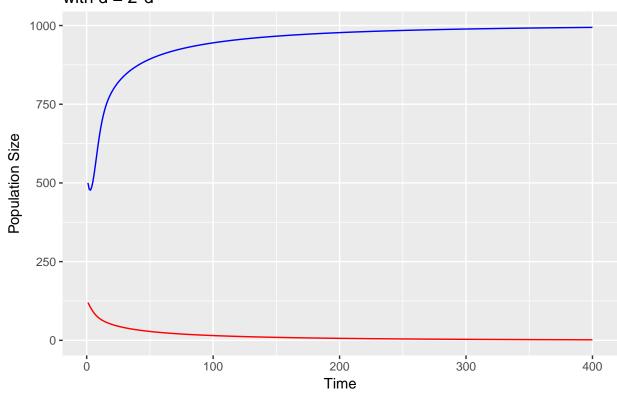
## Low d values

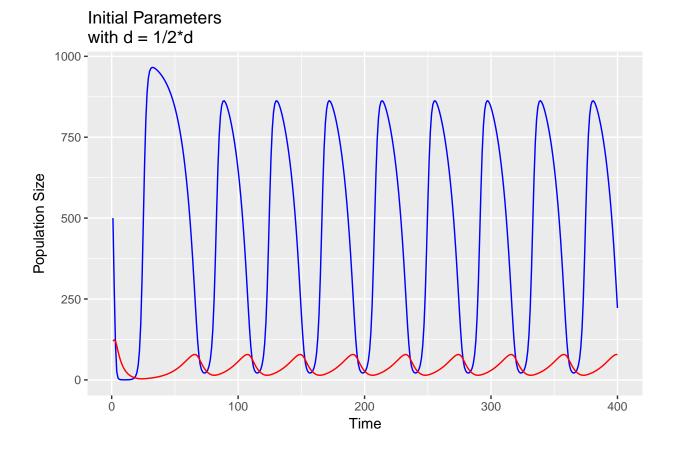
Predator population can maximize attack rate more easily
Maximizing attack rate drives prey population down
Small prey population hurts the predators, so predators decrease
The system oscillates

# High d values

Predators cannot reach an attack rate that sustains their population Predator population goes to zero; prey go to carrying capacity







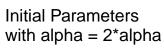
# Carrying Capacity/Alpha (alpha = 1/k)

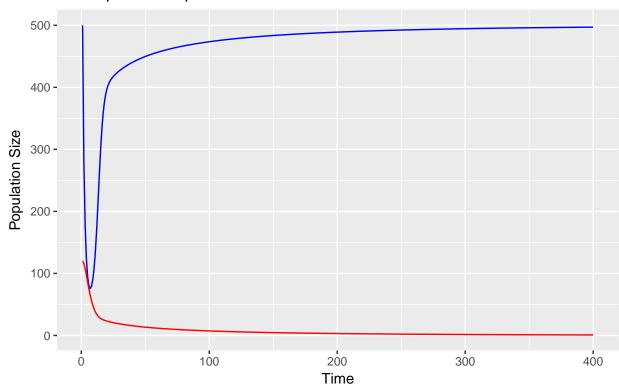
## Low alpha

Low alpha implies higher carrying capacity of prey Higher carrying capacity implies higher population; this sustains predators System oscillates

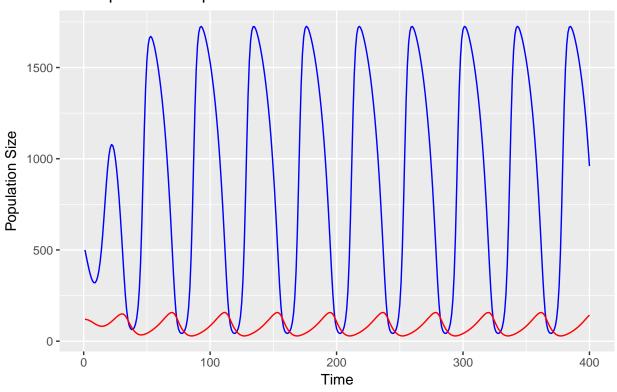
## High alpha

Carrying Capacity is lower, and predators cannot sustain themseleves when there are less prey Prey go to their carrying capacity





# Initial Parameters with alpha = 1/2\*alpha



# **Prey Birthrate**

## At low birth rate (0.2-1.6)

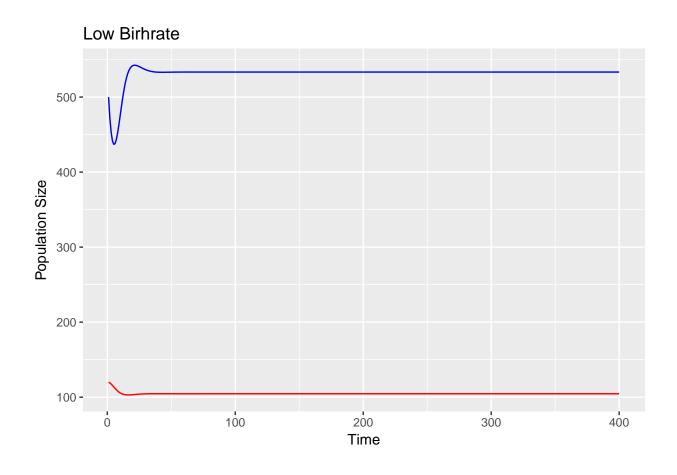
Prey and Predator population will decrease initially before increase in prey then increase in predator to equilibrium.

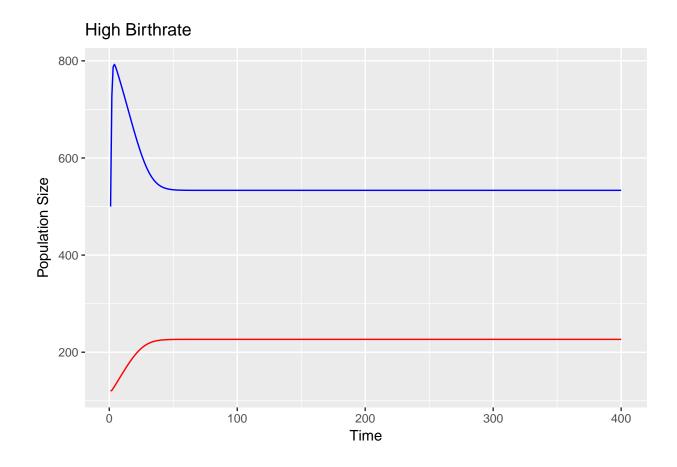
Predator death due to starvation allowing Prey population to increase to self-limiting density. Increase in Prey population allows Predator population to increase to capacity while Prey population decreases to equilibrium

## At high birth rate (2.0-3.2)

Prey birthrate increases to self-limiting density before Predator decreases Prey rate to equilibrium

Initial Predator population does not require as much Prey to sustain itself, therefore Prey population reaches self-limiting density. Predator population will continue to increase until capacity is reached. Capacity for Prey and Predator will increase as birth rate of prey increases





## Conversion efficiency of Prey to Predator

#### Low efficiency conversion (0.0175-0.0525)

Rate of Prey will initially decrease because Predator is present in system.

Rate of Predator decreases because of efficiency to convert energy to more Predator, the population will decrease to 0 as Prey population increases to self-limiting density. Rate to capacity determined by other parameters in the function

## Medium efficiency conversion and mutual equilibrium (0.07-0.0875)

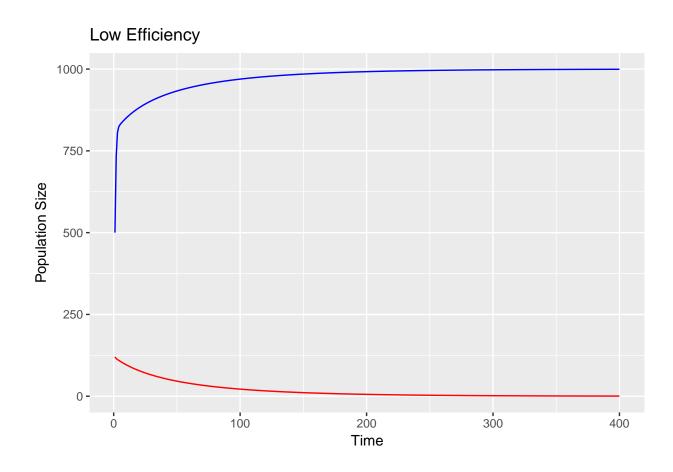
Rate of Prey and Predator will oscillate.

Prey population drops as Predator starts consuming. Predator reaches a population size that is not sustained by the amount of Prey and population will decrease. Prey will bounce back and cycle will begin again until reaching a stable equilibrium.

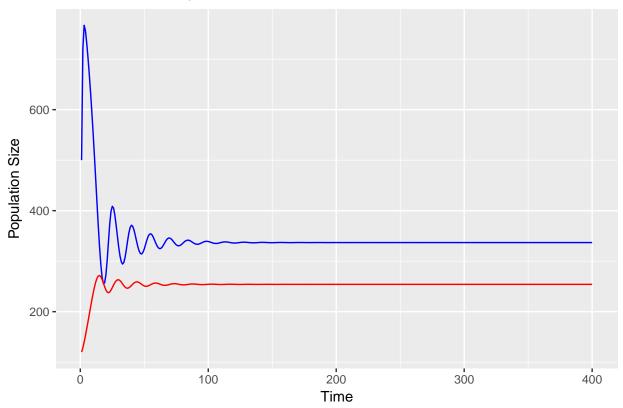
#### High efficiency conversion (0.1050-0.28)

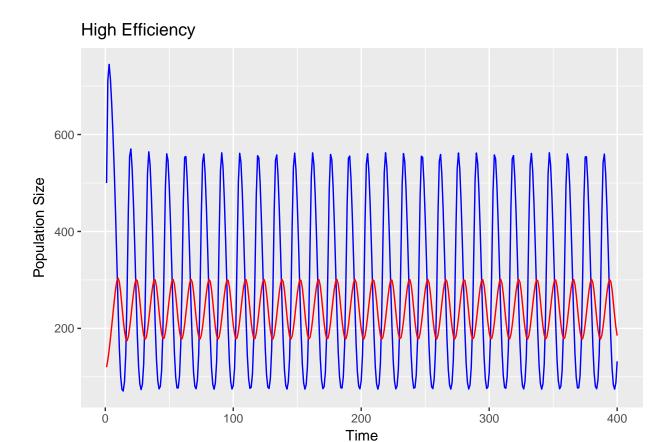
Rate of exchange from Prey to Predator occurs too quickly.

System will change from decreasing Prey and increasing Predator until Predator density is reached. Predators will decrease from starvation or lack of Prey to support system until Prey numbers bounch back. Increase in efficiency will increase maximum capacity of predator









## Predator Death Rate

## Low Death Rate (0.05-0.2)

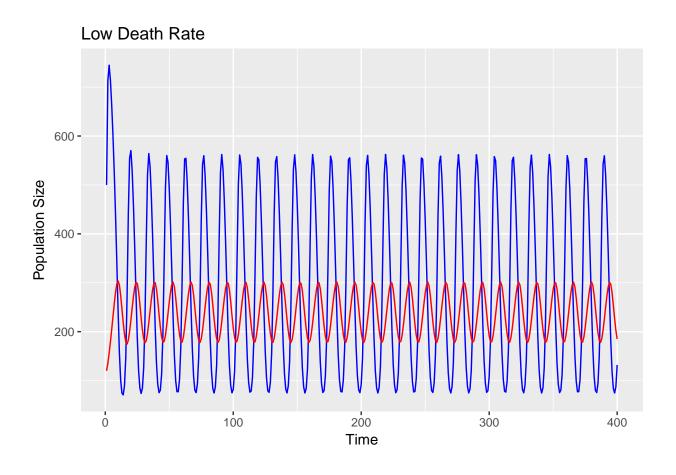
Low death rate leads to depletion of Prey population

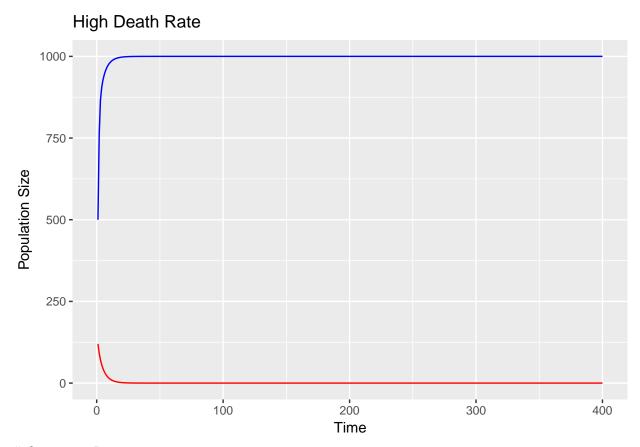
Decrease in Prey population decreases Predator density limit. Oscillation of both population until equilibrium is reached, or will continue to oscillate between capacity and 0

## High Death Rate (0.25-0.8)

Initial dip in Prey population results from initial Predator population

High death rate results in eradication of Predator and increase of Prey Population to self-limiting density





# Question 2 Responses

#### How do the dynamics differ from Lotka-Volterra?

Dynamics in R-M models allow for an equilibrium population. The L-V population conistantly oscillates between Prey depletion and Predator starvation to Prey

# What can you say about the role of each parameter, especially what causes the dyamics to differ between the L-V and R-M models?

The extra parameters seen in the R-M model adds complexity that allows for both populations to reach a population size in which co-habitation can be possible within some range of the parameters.

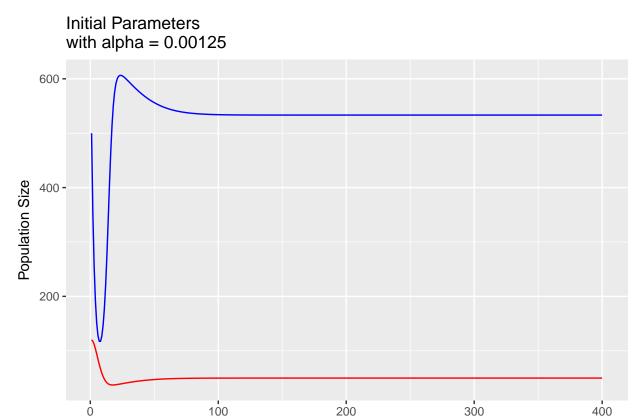
# What is the relationship between parameter values and predator abundance?

Predator abundance has a higher capacity when Prey has a higher birthrate, higher PRey self-limiting capacity, and lower conversion of Predator to Prey.

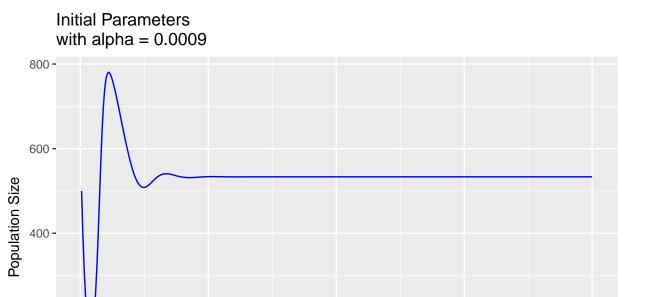
## Paradox of Enrichment

Conceptually, higher carrying capacity (lower alpha) should lead to predator stability because more prey will buffer the population from extinction.

However, the RM model shows that there is a negative relationship between prey carrying capacity and predator stability: as alpha decreases (carrying capacity increases), the population of predators becomes less stable. This is termed the *Paradox of Enrichment*. Let's see some examples of this:

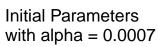


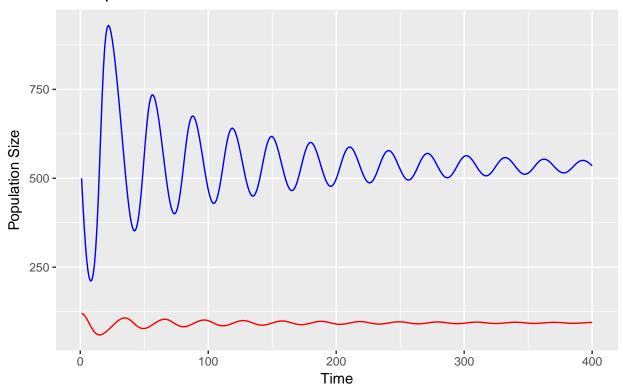
Time



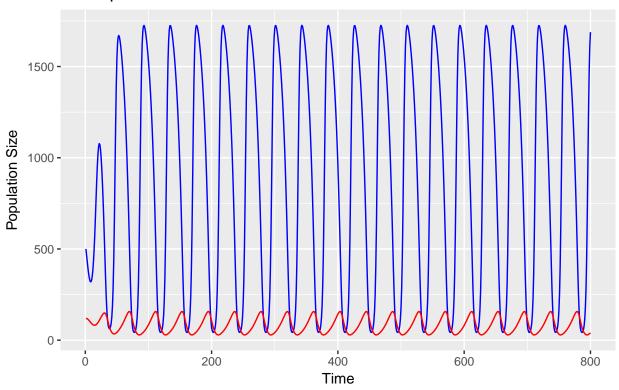
Time

200 -





# Initial Parameters with alpha = 0.0005



At low carrying capacities, the self-limiting aspect of the prey population leads to a stable equilibrium between predators and prey. However, as carrying capacity increases, a certain stability threshold is passed. Prey population is able to grow to such a point that predators can eat enough prey, driving the prey population down, which in turn drives the predator population down, leading to oscillation.