Exercise 2.2 - Survey of ecological models, Part 2

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2023-11-09 16:34:44

Instructions

Create either an R script (.R file) or R Markdown document (.Rmd) to save all of your work for today.

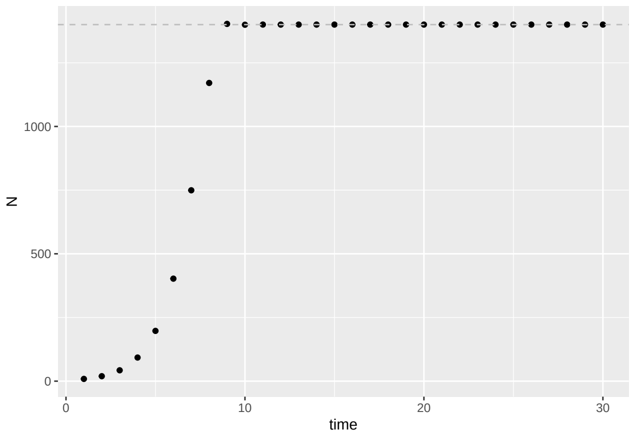
Exercise 1. Work with simulation of discrete-time logistic growth

In class, I discussed how making simulations of the models we learn in class can help us understand how our system works. Let's take a closer look at the *discrete-time model of logistic population growth*. The model formula is:

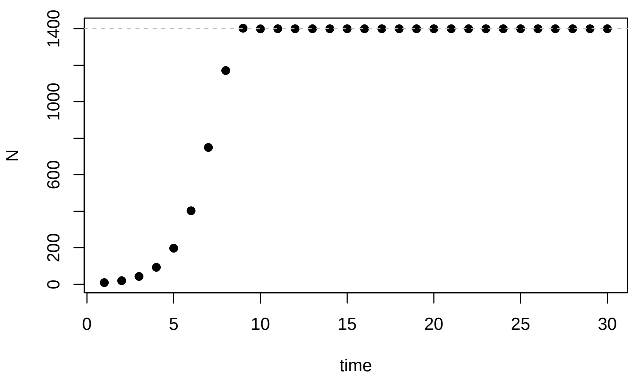
$$n_{t+1} = n_t + rn_t(1 - \frac{n_t}{K})$$

And the simulation works as follow:

```r
# Parameter values to use for simulation
r <- 1.21
K <- 1400
N0 <- 4
t <- 30
# model function
disc_log <- function(r, N0, K) {
    Nt1 <- N0 + r * N0 * (1 - N0/K)
    return(Nt1)
}
# Simulation of model for t time steps
N <- rep(NA, t)
for (i in 1:t) {
    N[i] <- disc_log(r, N0, K)
    N0 <- N[i]
}
# Plot simulation: ggplot
dat <- as.data.frame(N)
dat$time <- as.numeric(rownames(dat))
ggplot2::ggplot(dat, aes(time, N)) + geom_point() + geom_hline(yintercept = K,
    linetype = "dashed", color = "gray")
```

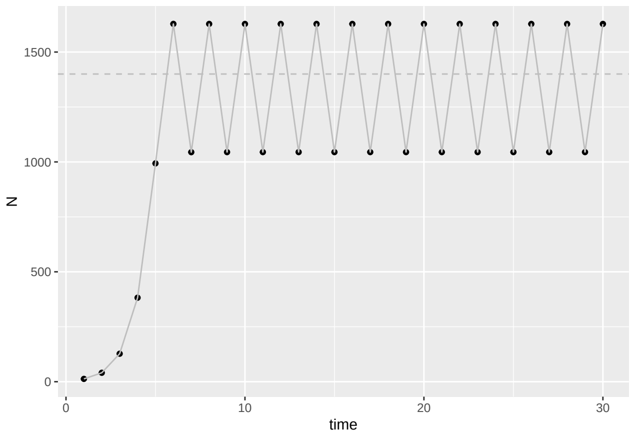



```
# Plot simulation: base R
plot(N, xlab = "time", ylab = "N", pch = 19, col = "black")
abline(h = K, col = "grey", lty = "dashed")
```

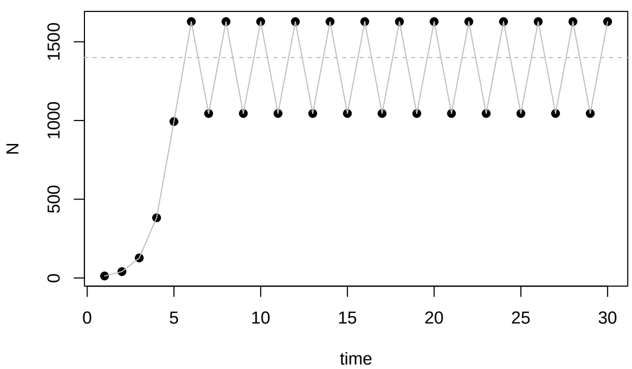


Question 1. Please change the simulation to run with a growth rate of 2.2, and show me a plot of the population dynamics over time.

```
# Parameter values to use for simulation
r <- 2.2
K <- 1400
N0 <- 4
t <- 30
# Simulation of model for t time steps
N <- rep(NA, t)
for (i in 1:t) {
    N[i] <- disc_log(r, N0, K)
    N0 <- N[i]
}
# Plot simulation: ggplot
dat <- as.data.frame(N)
dat$time <- as.numeric(rownames(dat))
ggplot2::ggplot(dat, aes(time, N)) + geom_point() + geom_line(color = "grey") +
    geom_hline(yintercept = K, linetype = "dashed", color = "grey")
```




```r
# Plot simulation: base R
plot(N, xlab = "time", ylab = "N", pch = 19, col = "black")
lines(N, col = "grey")
abline(h = K, col = "grey", lty = "dashed")
```



Question 2. A. Use R's index syntax `var[i]` - where *var* is the name of your variable and `[i]` is the *i*th position of that variable - to show me the value of population size `N` at time 5, 6, and 7.

```r
N[5:7]
```

```
## [1]  993.4435 1628.1292 1044.4632
```

B. Use the 'disc_log' function to calculate the value of N at time 6 using the value of N at time 5 for N0

```r
disc_log(r, N[5], K)
```

```
## [1] 1628.129
```

C. Use the 'disc_log' function to calculate the value of N at time 7 using the value of N at time 6 for N0

```r
disc_log(r, N[6], K)
```

```
## [1] 1044.463
```

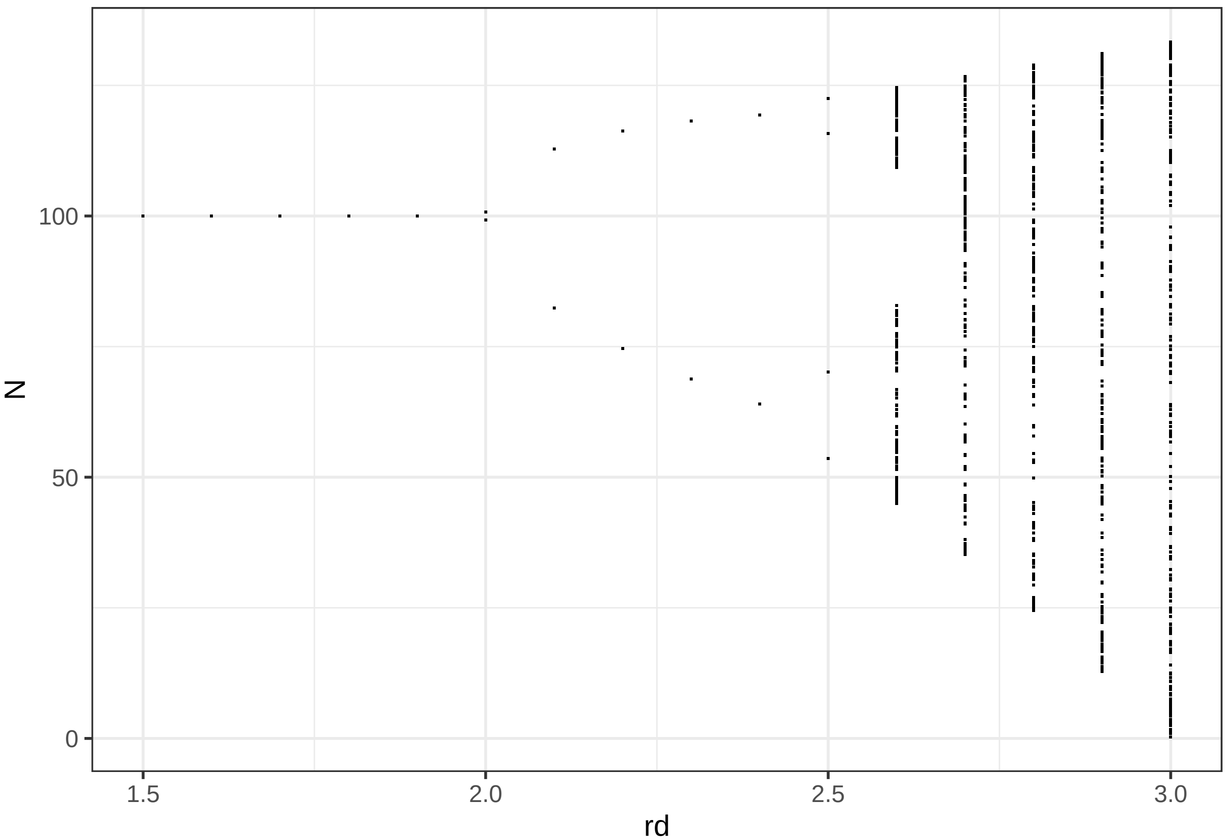
D. Use the formula above for discrete-time logistic growth to calculate population size at time 7 using the value of N at time 6 for N0

```
N[6] + r * N[6] * (1 - (N[6]/K))
```

```
## [1] 1044.463
```

E. Why is the population not going to its carrying capacity K? How many *periodic attractors* (values of N for which the system is stably drawn towards) are there in this system? In other words, how many values does N cycle between when $r = 2.2$, and what are those values of N?

Question 3. Look at the *bifurcation plot* for this discrete-time logistic growth system:



We see that as r increases the number of attractors continues to double, growing geometrically. Eventually, we reach a point when there becomes an infinite number of unique points, that are determined by r. This completely deterministic, non-repeating pattern in N is a property of *chaos*. *Chaos* is not a random phenomenon; rather it is the result of deterministic mechanisms generating non-repeating patterns.

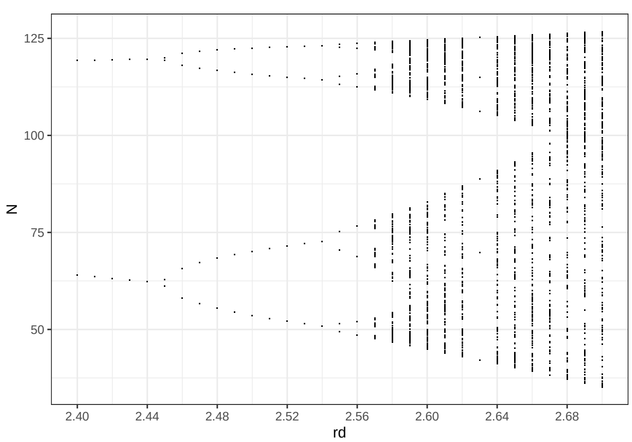
A. How many periodic attractors are there at $r = 2$, 2.1, 2.2, 2.3, 2.4, and 2.5?

B. How many periodic attractors are there at $r = 2.55$? I share the code below to produce a finer scale of r values:

```
rd.s <- seq(2.4, 2.7, by = 0.01)  # the actual rd values
t <- 0:1000  # how long to run each population
N0 <- c(N = 99)  # N-zero
## do all the dynamical simulations putting each run into a
## column of a matrix.
Ns <- sapply(rd.s, function(r) {
    p <- c(rd = r, alpha = 0.01)
    outd <- deSolve::ode(y = N0, times = t, func = dlogistic,
```

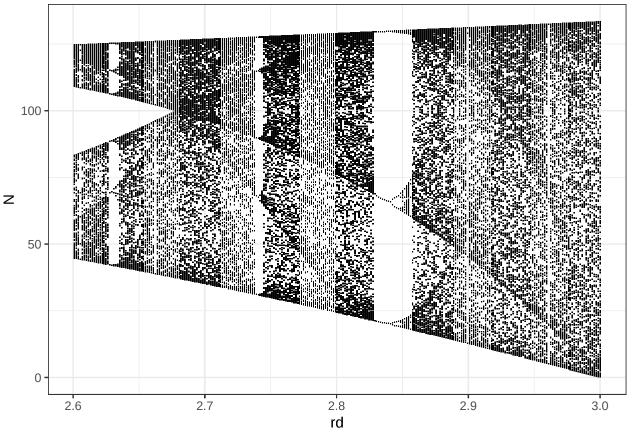
```r
        parms = p, method = "euler")[, 2]
})
out <- data.frame(t = t, Ns)  # put times to the front of the matrix
names(out) <- c("t", paste("rd=", rd.s, sep = ""))  # relabel columns
out.last <- subset(out, t > 0.8 * max(t))  # keep only the last 20%
## put the data in the 'long format'
out.l <- pivot_longer(out.last, cols = -1, names_to = "r.d",
    values_to = "N")
out.l <- dplyr::arrange(out.l, r.d, t)  # re-order the data
## extract text out of the 'r.d' label that is the numeric
## value
text.values <- substr(out.l$r.d, regexpr("=", out.l$r.d) + 1,
    100)
## convert the characters of '1.5' to the number 1.5
out.l$rd <- as.numeric(text.values)

## plot the stable limits to show the bifurcations
ggplot(out.l, aes(rd, N)) + geom_point(pch = ".") + theme_bw() +
    scale_x_continuous(breaks = seq(2.4, 2.7, by = 0.04))
```



C. Let's look closer at the chaotic portion of the graph, with a much finer range of r values. Convince yourself about chaos - that a TINY change in the parameter value r can lead to very different observed dynamics of N. You can read a bit more about chaos at this link: (https://geoffboeing.com/2015/03/chaos-theory-logistic-map/)

```r
out.l.chaotic <- subset(out.l, rd > 2.6)
ggplot(out.l.chaotic, aes(rd, N)) + geom_point(pch = ".") + theme_bw()
```

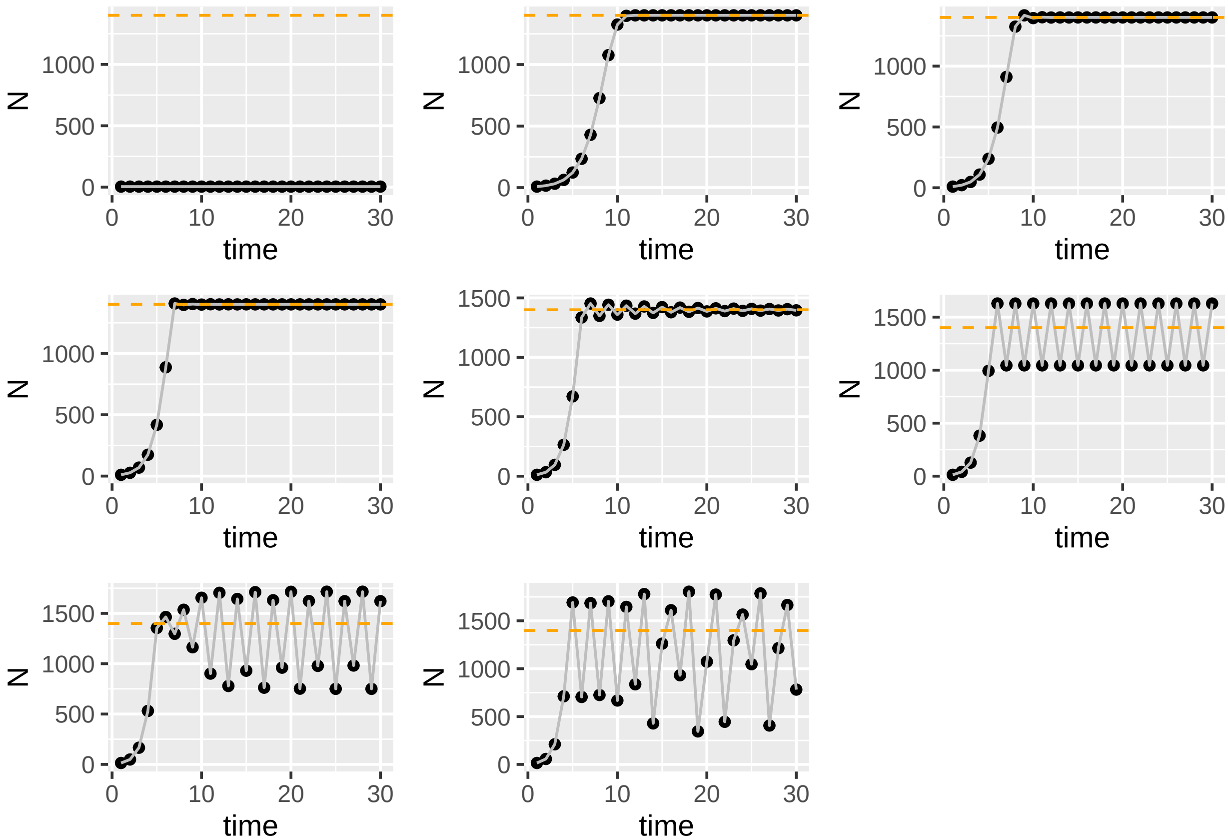


Question 4. Plot dynamics in the discrete-time logistic growth system.

In class, I showed a plot of population dynamics then ran the simulation for 8 different values of the population growth rate r :

```
# Parameter values to use for simulation
r_range <- c(0, 1, 1.3, 1.6, 1.9, 2.2, 2.5, 2.8)
K <- 1400
N0 <- 4
t <- 30
# Simulation of model for t time steps
N.g <- numeric()
plist <- list()
for (i in 1:length(r_range)) {
    N <- rep(NA, t)
    for (k in 1:t) {
        N[k] <- disc_log(r_range[i], N0, K)
        N0 <- N[k]
    }
    dat <- as.data.frame(N)
    dat$time <- as.numeric(rownames(dat))
    dat$r <- rep(r_range[i], t)
    N.g <- rbind(N.g, dat)
    N0 <- 4
    p <- ggplot(dat, aes(time, N)) + geom_point() + geom_line(color = "gray") +
        geom_hline(yintercept = K, linetype = "dashed", color = "orange")
    plist[[i]] <- p
}

gridExtra::grid.arrange(grobs = plist, nrow = round(length(r_range)/3))
```



Please use this plotting code to show dynamics for $r = 2$, 2.2, 2.4, 2.5, 2.55, and 2.8, for 50 time steps. Please use RMarkdown to write an informative caption for this figure, explaining what the graphs show. You can use this link: (https://uoepsy.github.io/scs/rmd-bootcamp/06-figs.html#captions)

```
# Parameter values to use for simulation
r_range <- c(2, 2.2, 2.4, 2.5, 2.55, 2.8)
K <- 1400
N0 <- 4
t <- 50
# Simulation of model for t time steps
N.g <- numeric()
plist <- list()
for (i in 1:length(r_range)) {
    N <- rep(NA, t)
    for (k in 1:t) {
        N[k] <- disc_log(r_range[i], N0, K)
        N0 <- N[k]
    }
    dat <- as.data.frame(N)
    dat$time <- as.numeric(rownames(dat))
    dat$r <- rep(r_range[i], t)
    N.g <- rbind(N.g, dat)
    N0 <- 4
    p <- ggplot(dat, aes(time, N)) + geom_point() + geom_line(color = "gray") +
        geom_hline(yintercept = K, linetype = "dashed", color = "orange")
    plist[[i]] <- p
}
```

```r
gridExtra::grid.arrange(grobs = plist, nrow = 3)
```

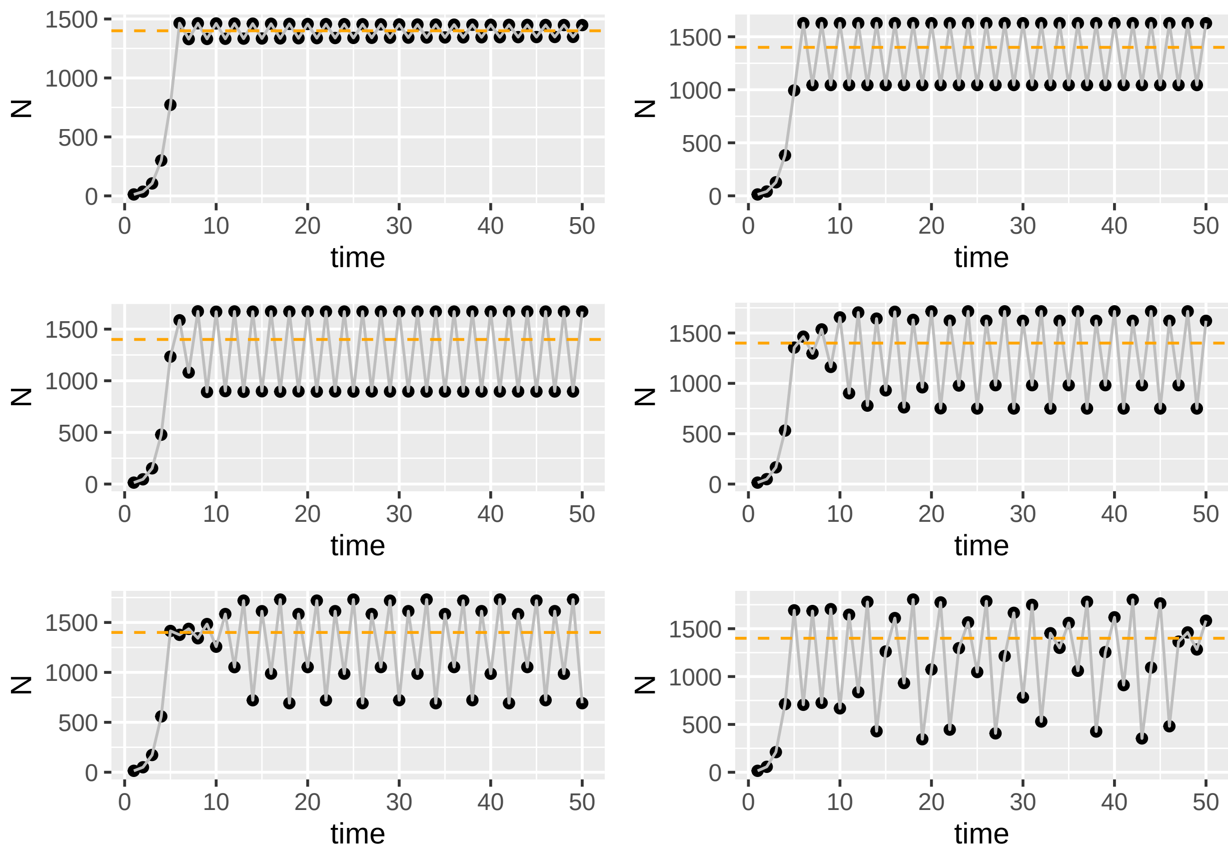


Figure 1: A figure of population size over time using the discrete-time logistic growth model

Exercise 2. Create simulation of continuous-time Lotka-Volterra competition model

Here is code for a continuous-time logistic growth model:

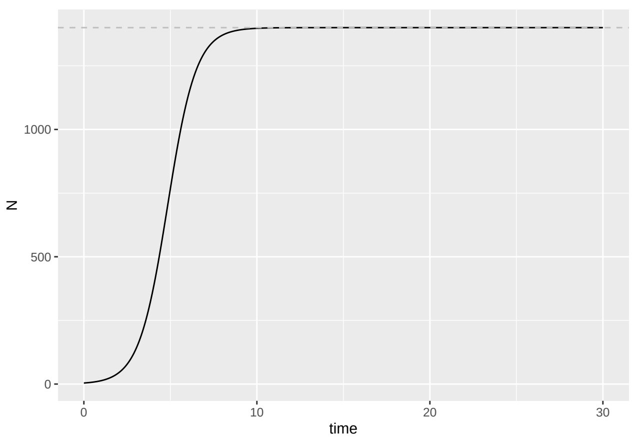
The formula:

$$\frac{dn}{dt} = rn(1 - \frac{n}{K})$$

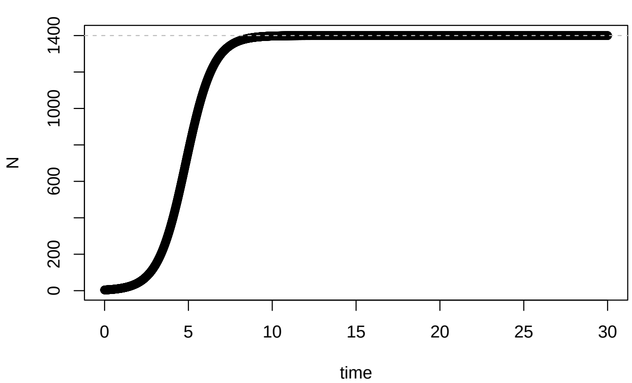
The code:

```r
# Parameter values to use for simulation
parameters <- c(r = 1.21, K = 1400)
state <- c(N = 4)
times <- seq(0, 30, by = 0.01)
# model function
cont_log <- function(t, state, parameters) {
    with(as.list(c(state, parameters)), {
        dN <- r * N * (1 - N/K)
```

```r
        return(list(dN))
    })
}
# Simulation of model for t time steps
out <- deSolve::ode(y = state, times = times, func = cont_log,
    parms = parameters)
# Plot simulation: ggplot
out.g <- as.data.frame(out)
ggplot2::ggplot(out.g, aes(time, N)) + geom_line() + geom_hline(yintercept = K,
    linetype = "dashed", color = "gray")
```




```r
# Plot simulation: base R
plot(out.g$time, out.g$N, xlab = "time", ylab = "N", pch = 19,
    col = "black")
abline(h = K, col = "grey", lty = "dashed")
```



A. Please modify this to create a simulation of the continuous time Lotka-Volterra competition equations, seen as equations 3.15 in Otto & Day Chapter 3.

The formula:

$$\frac{dn_1}{dt} = r_1 n_1 (1 - \frac{n_1 t + \alpha_{12} n_2}{K_1})$$
$$\frac{dn_2}{dt} = r_2 n_2 (1 - \frac{n_2 t + \alpha_{21} n_1}{K_2})$$

The parameters you will need are:

`r1, r2, K1, K1, alpha_12, alpha_21, n1_0, n2_0`

Use these values:

`r1 = 1.3, r2 = 1.5, K1 = 1,400, K2 = 1,000, alpha_12 = 0.4, alpha_21 = 0.6, N1_0 = 5, N2_0 = 5`

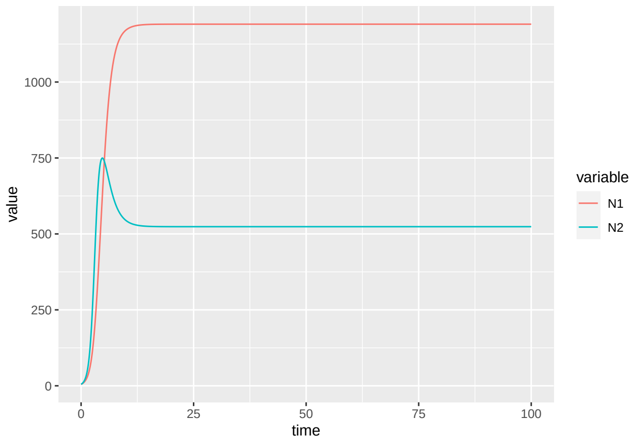
Note: be very careful with your parentheses

Here I show with prompts what code you need to fill in. Then I show my code for running the continuous simulation and plotting.

```
# Parameter values to use for simulation
parameters <- 1  ## fill in param values where the 1 is
state <- 1  ## fill in initial values for state variables where the 1 is
times <- 1  ## fill in values for time intervals where the  1 is
# model function
cont_LV <- function(t, state, parameters) {
    with(as.list(c(state, parameters)), {
        ## fill in equation for N1 fill in equation for N2
        return(list(c(dN1, dN2)))
    })
}
```

```
# Parameter values to use for simulation
parameters <- c(r1 = 1.3, r2 = 1.7, K1 = 1400, K2 = 1000, alpha_12 = 0.4,
    alpha_21 = 0.6)
state <- c(N1 = 5, N2 = 5)
times <- seq(0, 100, by = 0.1)
# model function
cont_LV <- function(t, state, parameters) {
    with(as.list(c(state, parameters)), {
        dN1 <- r1 * N1 * (1 - ((N1 + alpha_12 * N2)/K1))
        dN2 <- r2 * N2 * (1 - ((N2 + alpha_12 * N1)/K2))
        return(list(c(dN1, dN2)))
    })
}
```

```
# Simulation of model for t time steps
out <- deSolve::ode(y = state, times = times, func = cont_LV,
    parms = parameters)
# Plot simulation: ggplot
out.g <- as.data.frame(out)
out.melt <- reshape2::melt(out.g, id.var = "time")
ggplot2::ggplot(out.melt, aes(x = time, y = value, col = variable)) +
    geom_line()
```




```r
# Plot simulation: base R
plot(out.g$time, out.g$N1, xlab = "time", ylab = "N", col = "black",
    type = "l")
lines(out.g$time, out.g$N2, col = "skyblue")
```