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BI 471/571: Population Ecology

Spring 2016

## Homework 2: Single species dynamics

Due 4/14/2016

### 1. Hastings Problems 4.1 & 4.3

**4.1** model:  $\frac{dN}{dt} = rN \left[ 1 - (N/K)^\theta \right]$

a) Find equilibria + determine stability

$$\frac{dN}{dt} = 0, \text{ so } 0 = rN \left[ 1 - (N/K)^\theta \right]$$

solve for N:

$$0 = rN - rN(N/K)^\theta$$

$$\frac{rN}{rN} = \frac{rN(N/K)^\theta}{rN}$$

$$1 = (N/K)^\theta$$

$$N = K^{1/\theta}$$

when  $N = 0$ , unstable

$N = K$ , stable

graph using  $\theta = 0.5, \theta = 1, \theta = 2$

$$\frac{dN/dt}{N} = r \left[ 1 - (N/K)^\theta \right]$$

??

**4.3**  $\frac{dN}{dt} = rN(N-a) \left[ 1 - (N/K) \right]$

a) equilibrium:

$$0 = rN(N-a) \left[ 1 - (N/K) \right]$$

$$0 = (rN^2 - rNa) \left[ 1 - (N/K) \right]$$

$$0 = rN^2 - \frac{rN^3}{K} - rNa + \frac{rN^2a}{K}$$

$$\frac{rN^3}{K} - \frac{rN^2a}{K} = rN^2 - rNa$$

$$\frac{rN^2}{K} (N-a) = rN(N-a)$$

$$rN^2 = rNK$$

$$N^2 = NK$$

$$N = K$$

b) stability.?

c) graph?

### 2. (a)

## write logistic growth function from exp.growth, note changes to dN.dt

```
log.growth <- function(t, y, p) {  
  N <- y[1]  
  with(as.list(p), {  
    dN.dt <- r * N * (1 - (N/K))  
    return(list(dN.dt))  
  })  
}
```

## make vector of parameters

```
p <- c('r' = 0.25, 'K' = 100)
```

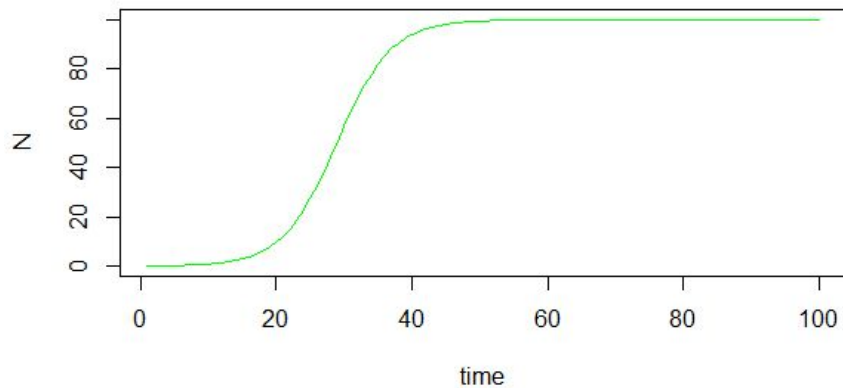
```
## using runif, write for interval
0.01 to 0.1, initial conditions
y0 <- c('N' = runif(1, min = 0.01,
max = 0.1))
```

```
## time
t <- 1:100
```

```
## load library if needed
library(deSolve)
```

```
## simulation
sim <- ode(y = y0, times = t,
func = log.growth, parms = p,
method = 'lsoda')
sim <- as.data.frame(sim)
```

```
## plot my simulation
plot(N ~ time, data = sim, type =
'l', col = 'green')
```



**(b)**

```
## write logistic growth function from exp.growth, note changes to dN.dt
```

```
log.growth <- function(t, y, p) {
  N <- y[1]
  with(as.list(p), {
    dN.dt <- r * N * (1 - (N/K))
    return(list(dN.dt))
  })
}
```

```
## make vector of
parameters
p <- c('r' = 0.25, 'K' =
100)
```

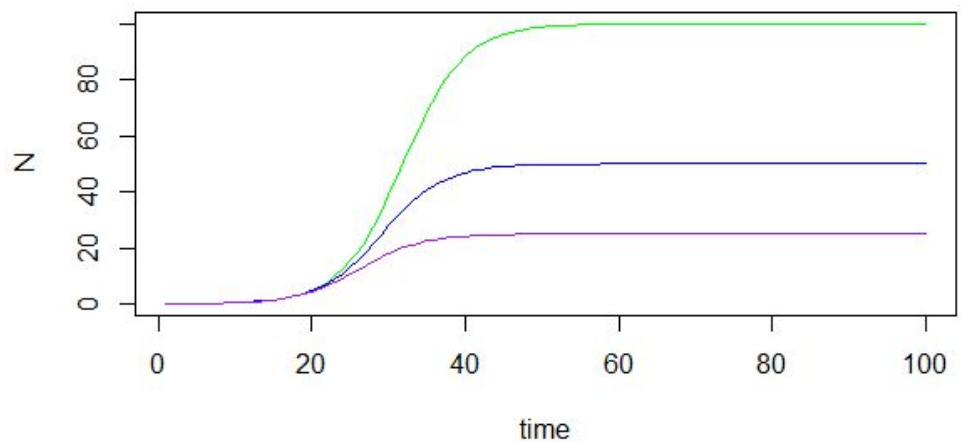
```
## using runif, write for
interval 0.01 to 0.1,
initial conditions
y0 <- c('N' = runif(1,
min = 0.01, max =
0.1))
```

```
## time
t <- 1:100
```

```
## load library if
needed
library(deSolve)
```

```
## simulation
sim <- ode(y = y0,
times = t, func =
log.growth, parms = p,
method = 'lsoda')
sim <- as.data.frame(sim)
```

```
## plot my simulation
plot(N ~ time, data = sim, type = 'l', col = 'green')
```



```
## add plots for k = 50 and k = 25
## first for k=50 use p.2 and sim.2
p.2 <- c('r' = 0.25, 'K' = 50)
sim.2 <- ode(y = y0, times = t, func = log.growth, parms = p.2, method = 'lsoda')
sim.2 <- as.data.frame(sim.2)

## next k=25 use p.3 and sim.3
p.3 <- c('r' = 0.25, 'K' = 25)
sim.3 <- ode(y = y0, times = t, func = log.growth, parms = p.3, method = 'lsoda')
sim.3 <- as.data.frame(sim.3)

## plot on same figure, ylim will help with line
plot(N ~ time, data = sim, type = 'l', col = 'green')

## using points, it allows us to use above plot and add in the other data frames
points(N ~ time, data = sim.2, type = 'l', col = 'blue')
points(N ~ time, data = sim.3, type = 'l', col = 'purple')
```

(c)

Code for 'r' = 0.25, 'K' = 100:

```
## theta log growth function
log.growth <- function(t, y, p) {
  N <- y[1]
  with(as.list(p), {
    dN.dt <- r * N * (1 - (N / K))
    return(list(dN.dt))
  })
}

## parameters
p <- c('r' = 0.25, 'K' = 100)
y0 <- c('N' = runif(1, min = 0.01, max = 0.1))
t <- 1:100

## simulation
sim <- ode(y = y0, times = t, func =
log.growth, parms = p, method =
'lsoda')
sim <- as.data.frame(sim)

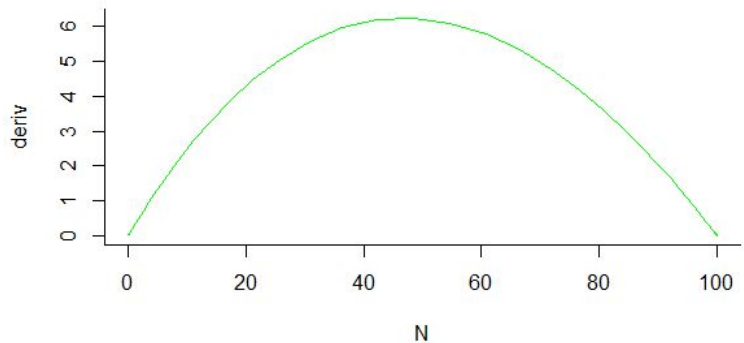
## plot
plot(N ~ time, data = sim, type = 'l', lwd = 2, bty = 'l', col = 'green')

## compute derivatives
sim$deriv <- c(diff(sim$N), NA)

## plot vs pop abundance
plot(deriv ~ N, data = sim, type = 'l', col = 'blue', bty = 'l')

## max value
max(sim$deriv, na.rm = TRUE)
## Output: 6.235844
which(sim$deriv == max(sim$deriv, na.rm = TRUE))
## Output: 30

## value of N in sim with the same index as the biggest deriv value
```



```
sim$N[which(sim$deriv == max(sim$deriv, na.rm = TRUE))]
## Output: 48.43107
```

**Code for 'r' = 0.25, 'K' = 50:**

```
## theta log growth function
log.growth <- function(t, y, p) {
  N <- y[1]
  with(as.list(p), {
    dN.dt <- r * N * (1 - (N / K))
    return(list(dN.dt))
  })
}

## parameters
p.2 <- c('r' = 0.25, 'K' = 50)
y0 <- c('N' = runif(1, min = 0.01, max = 0.1))
t <- 1:100

## simulation
sim.2 <- ode(y = y0, times = t, func = log.growth, parms = p.2, method = 'lsoda')
sim.2 <- as.data.frame(sim.2)

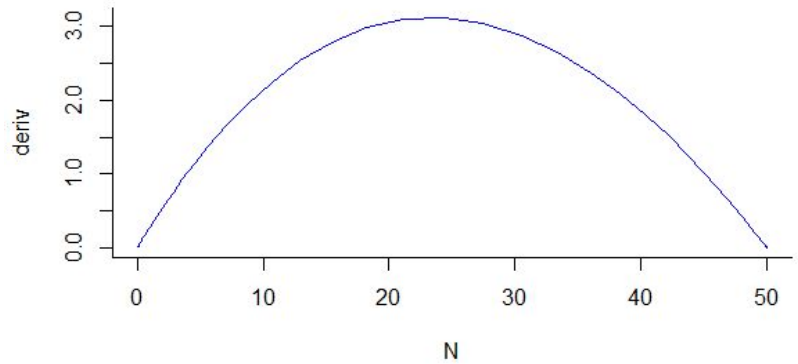
## plot
plot(N ~ time, data = sim.2, type = 'l',
     lwd = 2, bty = 'l', col = 'blue')

## compute derivatives
sim.2$deriv <- c(diff(sim.2$N), NA)

## plot vs pop abundance
plot(deriv ~ N, data = sim.2, type = 'l', col = 'blue', bty = 'l')

## max value
max(sim.2$deriv, na.rm = TRUE)
##Output: 3.117201
which(sim.2$deriv == max(sim.2$deriv, na.rm = TRUE))
## Output: 27

## value of N in sim with the same index as the biggest deriv value
sim.2$N[which(sim.2$deriv == max(sim.2$deriv, na.rm = TRUE))]
## Output: 24.30376
```

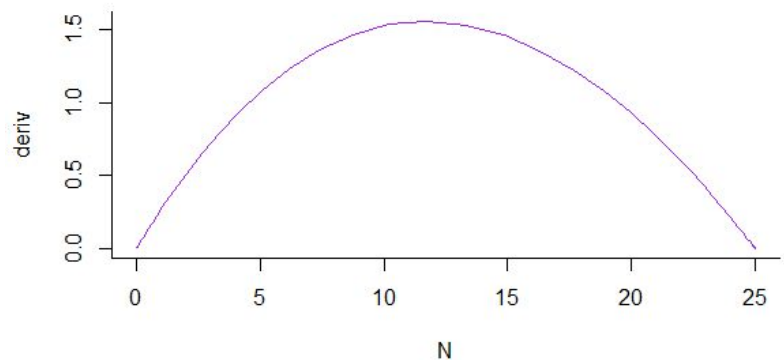


**Code for 'r' = 0.25, 'K' = 25:**

```
## theta log growth function
log.growth <- function(t, y, p) {
  N <- y[1]
  with(as.list(p), {
    dN.dt <- r * N * (1 - (N / K))
    return(list(dN.dt))
  })
}

## parameters
p.3 <- c('r' = 0.25, 'K' = 25)
y0 <- c('N' = runif(1, min = 0.01, max = 0.1))
t <- 1:100

## simulation
```



```
sim.3 <- ode(y = y0, times = t, func = log.growth, parms = p.3, method = 'lsoda')
sim.3 <- as.data.frame(sim.3)
```

```
## plot
plot(N ~ time, data = sim.3, type = 'l', lwd = 2, bty = 'l', col = 'purple')
```

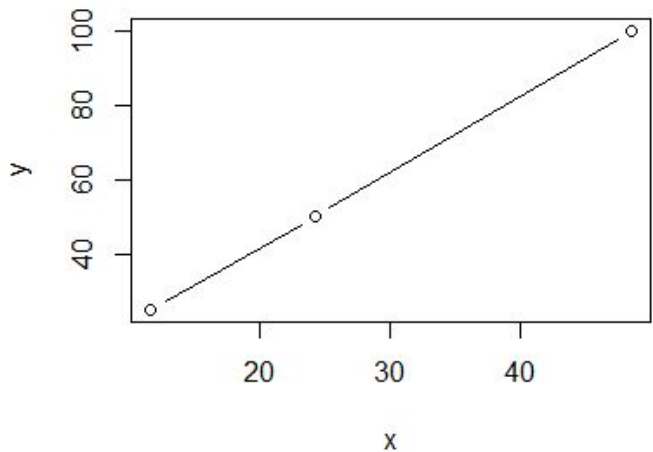
```
## compute derivatives
sim.3$deriv <- diff(sim.3$N)
sim.3$deriv <- c(diff(sim.3$N), NA)
```

```
## plot vs pop abundance
plot(deriv ~ N, data = sim.3, type = 'l', col = 'purple',
bty = 'l')
```

```
## max value
max(sim.3$deriv, na.rm = TRUE)
## Output: 1.560464
which(sim.3$deriv == max(sim.3$deriv, na.rm =
TRUE))
##Output: 29
```

```
## value of N in sim with the same index as the
biggest deriv value
sim.3$N[which(sim.3$deriv == max(sim.3$deriv,
na.rm = TRUE))]
## Output: 11.74914
```

```
## plot these values against their corresponding K
parameter
x <- c(48.43107, 24.30376, 11.74914) ## N values
y <- c(100, 50, 25) ## K values
plot(x, y, type = 'b')
```



3.

A) Theta value: 0.5

```
log.growth <- function(t, y, p) {
  N <- y[1]
  with(as.list(p), {
    dN.dt <- r * N * (1 - (N / K)^theta)
    return(list(dN.dt))
  })
}
```

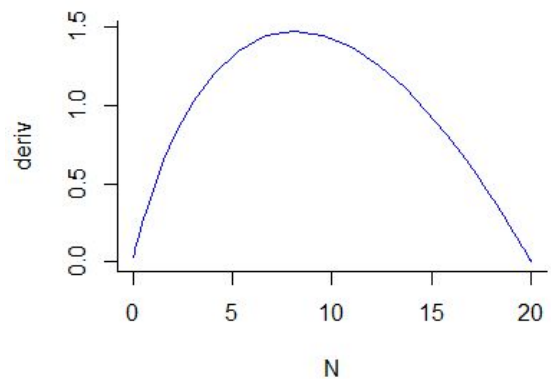
```
p <- c('r' = 0.5, 'K' = 20, 'theta' = 0.5)
y0 <- c('N' = 0.05)
t <- 1:100
```

```
sim <- ode(y = y0, times = t, func = log.growth, parms = p, method =
'lsoda')
sim <- as.data.frame(sim)
plot(N ~ time, data = sim, type = 'l', lwd = 2, bty = 'l', col = 'blue')
```

```
sim$deriv <- c(diff(sim$N), NA)
plot(deriv ~ N, data = sim, type = 'l', col = 'blue', bty = 'l')
```

```
max(sim$deriv, na.rm = TRUE)
##Output: 1.478818
```

```
which(sim$deriv == max(sim$deriv, na.rm = TRUE))
```

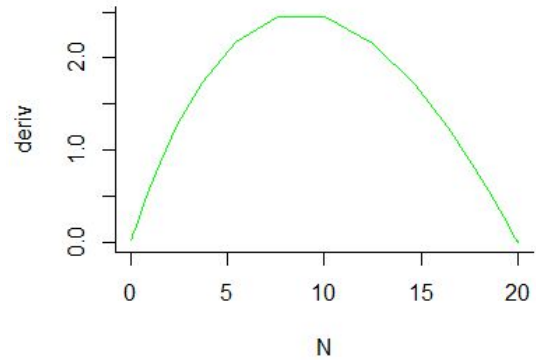


##Output:15

```
sim$N[which(sim$deriv == max(sim$deriv, na.rm = TRUE))]  
##Output: 8.075343
```

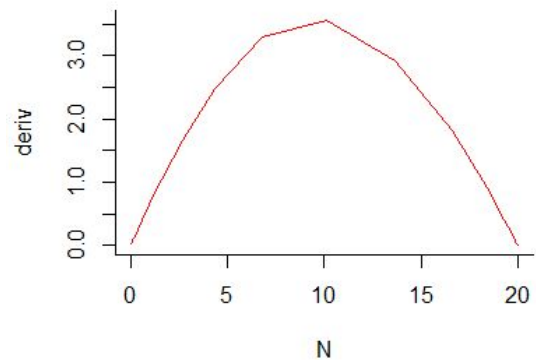
## B) Theta value: 1

```
log.growth <- function(t, y, p) {  
  N <- y[1]  
  with(as.list(p), {  
    dN.dt <- r * N * (1 - (N / K)^theta)  
    return(list(dN.dt))  
  })  
}  
  
p <- c('r' = 0.5, 'K' = 20, 'theta' = 1)  
y0 <- c('N' = 0.05)  
t <- 1:100  
  
sim <- ode(y = y0, times = t, func = log.growth, parms = p, method =  
'lsoda')  
sim <- as.data.frame(sim)  
plot(N ~ time, data = sim, type = 'l', lwd = 2, bty = 'l', col = 'green')  
  
sim$deriv <- c(diff(sim$N), NA)  
plot(deriv ~ N, data = sim, type = 'l', col = 'green', bty = 'l')  
  
max(sim$deriv, na.rm = TRUE)  
##Output: 2.452442  
  
which(sim$deriv == max(sim$deriv, na.rm = TRUE))  
##Output: 12  
  
sim$N[which(sim$deriv == max(sim$deriv, na.rm = TRUE))]  
##Output: 7.602863
```



## C) Theta value: 1.8

```
log.growth <- function(t, y, p) {  
  N <- y[1]  
  with(as.list(p), {  
    dN.dt <- r * N * (1 - (N / K)^theta)  
    return(list(dN.dt))  
  })  
}  
  
p <- c('r' = 0.5, 'K' = 20, 'theta' = 1.8)  
y0 <- c('N' = 0.05)  
t <- 1:100  
  
sim <- ode(y = y0, times = t, func = log.growth, parms = p, method =  
'lsoda')  
sim <- as.data.frame(sim)  
plot(N ~ time, data = sim, type = 'l', lwd = 2, bty = 'l', col = 'red')  
  
sim$deriv <- c(diff(sim$N), NA)  
plot(deriv ~ N, data = sim, type = 'l', col = 'red', bty = 'l')  
  
max(sim$deriv, na.rm = TRUE)  
##Output: 3.568666  
  
which(sim$deriv == max(sim$deriv, na.rm = TRUE))
```



*##Output: 12*

```
sim$N[which(sim$deriv == max(sim$deriv, na.rm = TRUE))]
```

*##Output: 10.09747*

According to the calculations, species C maintained the highest population abundance in the fishery (10.09747).