Exponential population growth

Matthew Malishev¹*

¹ Department of Biology, Emory University, 1510 Clifton Road NE, Atlanta, GA, USA, 30322

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*Corresponding author: matthew.malishev@gmail.com

This document can be found at https://github.com/darwinanddavis

R session info

params\$session

R version 3.5.0 (2018-04-23)

Platform: x86_64-apple-darwin15.6.0 (64-bit) Running under: OS X El Capitan 10.11.6

Matrix products: default

BLAS: /Library/Frameworks/R.framework/Versions/3.5/Resources/lib/libRblas.0.dylib LAPACK: /Library/Frameworks/R.framework/Versions/3.5/Resources/lib/libRlapack.dylib

locale:

[1] en_US.UTF-8/en_US.UTF-8/en_US.UTF-8/C/en_US.UTF-8/en_US.UTF-8

attached base packages:

[1] stats graphics grDevices utils datasets methods base

loaded via a namespace (and not attached):

[1] compiler_3.5.0 backports_1.1.2 magrittr_1.5 rprojroot_1.3-2 tools_3.5.0 htmltools_0.3.6 [7] pillar_1.2.3 tibble_1.4.2 yaml_2.2.0 Rcpp_0.12.18 stringi_1.2.3 rmarkdown_1.10

[13] knitr_1.20 stringr_1.3.1 digest_0.6.15 rlang_0.2.1 evaluate_0.10.1

Overview

Examples of exponential population growth in R.

Install dependencies

```
packages <- c("dplyr","deSolve","pdftools")
if (require(packages)) {
    install.packages(packages,dependencies = T)
    require(packages)
}
lapply(packages,library,character.only=T)</pre>
```

Section 1

Exponential growth equation

```
N_t = N_0 \cdot e^{rt}
```

 $\mathrm{Nt}=\mathrm{the}\ \mathrm{number}\ \mathrm{of}\ \mathrm{individuals}\ \mathrm{in}\ \mathrm{the}\ \mathrm{population}\ \mathrm{after}\ \mathrm{t}\ \mathrm{units}\ \mathrm{of}\ \mathrm{time}$

No = the initial population size (t = 0)

r = the exponential growth rate

t = time unit (usually in years)

e = the base of the natural logarithms (2.72)

Exponential rate of growth is commonly named the parameter lambda λ

$$\lambda = e^r$$

 $e^r = \text{lambda}$. Exponential growth rate parameter.

The natural log (ln) of e = 1

$$ln(e) = 1$$

because $e^1 = e$.

The natural log of 1 = 0

$$ln(1) = 0$$

because $e^0 = 1$.

Parameters

```
# parameters
N_t <- 0 # expected pop size
N_0 <- 500 # initial pop size
e <- exp
r <- 0.012 # exponetial rate of growth
lambda <- e(1^r)
t <- 10 # time (in years)

# putting the above all together in R
N_t <- N_0 * e(r*t)
N_t</pre>
```

Example

A moose population has a growth rate of 0.02. In 2000, the population was 500. What will the population be in 2020?

```
# input your R code here
```

Instantaneous rate of growth

Equation showing the rate of population increase

$$\frac{dN}{dt} = rN$$

```
\begin{split} \mathrm{d} N &= \mathrm{change\ in\ number} \\ \mathrm{d} t &= \mathrm{change\ in\ time} \\ \mathrm{r} &= \mathrm{the\ per\ head\ maximum\ potential\ growth\ rate} \\ N &= \mathrm{number\ of\ individuals\ in\ a\ population} \end{split}
```

```
# in R
N <- 1000
dNdt <- r*N
dNdt
```

Example

A population of 100 individuals. Each individual can on average contribute 1/4 of an individual (new individual) to the population in a given unit of time. Find the rate of population increase.

```
# your r code
```

Simulating population growth

Set your parameters for the population

```
N_0 <- 20; # initial population size
```

Over time

```
N_1 \leftarrow N_0 * r; # population size at t = 1
```

What does this look like at each time point?

```
N_2 <- # ??
N_3 <- # ??
# etc

Population size
popsize = c(N_0, N_1, N_2, N_3, N_4, N_5)
popsize
popsize[2]</pre>
```

Section 2

Simulation loop

```
r <- 2
N_0 <- 20
t <- 10
popsize <- list() # create empty list to populate in loop
tt <- 2:t # now make t a vector to make a time period

for(i in tt){ # start loop
    N_0[i] <- N_0[i-1]*r # middle of loop
    popsize <- N_0 # put the pop size into the list we created
} # end loop
popsize</pre>
```

Plot

```
# plot it
plot(popsize)
# extend plot params
tta <- c(1,tt)# add the first year onto the time vector
ttm <- max(tt) # get the maximum value of the time vector
xlim <- c(1,ttm) # put this in a vector</pre>
ylim \leftarrow c(0, max(N_0)) \# do the same for growth
par(las=1,bty="n") # set plotting params
plot(tta,popsize, # vectors to plot
     xlim=xlim, # set the x limits
     ylim=ylim, # y limits
     xlab="Time",
     ylab="Population growth",
     pch=20, # set point type
     col="pink"
?plot # help page for plot function
```

Event-based conditions

Changing conditions halfway through a simulation. Using if and else statements.

```
popsize <- list() # reset the list

for (i in tt){
   if((i;\%5) == 0){ #modify growth rate depending on the year
        r = 4
   }else{ # or just an ok year
        r = 2
    }
   N_0[i] <- N_0[i-1]*r # the population growth equation
   popsize <- N_0
}

popsize

par(las=1,bty="n") # set plotting params
plot(popsize)</pre>
```