

# Population & Community Ecology Homework 1

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## Problem 1

In this problem we look to find the density of duckweed plants per  $m^2$  after 47 days. We will estimate the future population using the exponential growth formula,  $N_t = N_0 e^{rt}$  where the initial population,  $N_0$ , is 11.1 duckweed plants per  $m^2$ . We can calculate the rate of unitless population change,  $r$ , by dividing the estimate after 20 day from the initial population.

```
r = 23.5/11.1  
r
```

```
## [1] 2.117117
```

Thus with  $r = 2.12$ , we can calculate  $N_{27}$  as  $N_{27} = 11.1e^{2.12 \times 47}$ .

```
n27 <- 11.1*exp(2.12*47)  
n27
```

```
## [1] 2.081734e+44
```

Thus after a total of 47 days, the population density will be  $2.08 \times 10^{44}$  duckweed plants per  $m^2$ .

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## Problem 2

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## Problem 3

### Part a

Given the information we would like to solve for  $N_7$ , the population in seven years.

```
n_7 <- (1.037^7)*112  
n_7
```

```
## [1] 144.434
```

Thus the population density will be 144 (rounded, no units given) at year 7.

### Part b

Using the exponential growth formula,  $N_t = N_0 e^{rt}$ , we would like to solve for an  $r$  value where  $N_0 = 112$  and  $N_t = 144$ . Thus we solve for  $r$  as  $\ln(\frac{N_t}{N_0})/t = r$ .

```
r <- log(144/112)/7  
r
```

```
## [1] 0.03590206
```

Thus  $r$  is equal to .036 in the exponential model to achieve the same results as the discrete population growth model.

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## Problem 4

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## Problem 5

Survivorship, as defined by  $\frac{S_x}{S_0}$ , is calculated in the vector below.

```
age <- 0:4
sx <- c(740,280,105,32,0)
bx <- c(0,0.4,1.3,3,8,NA)
lx <- sx/740
print(lx)
```

```
## [1] 1.00000000 0.37837838 0.14189189 0.04324324 0.00000000
```

I assume that  $g(x)$  was meant to be  $p(x)$

Solving for  $p(x)$

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## Problem 6

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## Problem 7

The following is the setup for the population project matrix:

```
stages <- c('seedling','juvenile','small adult', 'medium adult', 'large adult')
A <- matrix(c(
  0, 0,.015,.063,.189,
  .294,.25, 0 , 0 , 0 ,
  .176,.35,.385,.059,.012,
  0 ,.35,.461,.568,.171,
  0 , 0 ,.108,.356,.805
), nrow = 5, byrow = TRUE,
  dimnames = list(stages,stages)
)
print(A)
```

```
##           seedling juvenile small adult medium adult large adult
## seedling      0.000      0.00      0.015      0.063      0.189
## juvenile      0.294      0.25      0.000      0.000      0.000
## small adult    0.176      0.35      0.385      0.059      0.012
```

## medium adult	0.000	0.35	0.461	0.568	0.171
## large adult	0.000	0.00	0.108	0.356	0.805

## Part a

To calculate  $\lambda$  from the popbio package we do the following:

```
library(popbio)
lambda(A)
```

```
## [1] 1.037396
```

Thus  $\lambda = 1.04$ .

## Part b

To calculate the stable stage distribution we do the following:

```
stable.stage(A)
```

##	seedling	juvenile	small adult	medium adult	large adult
##	0.10633963	0.03970536	0.08489524	0.28816968	0.48089009