

Math 142 Homework1

Chen Li – 705426669

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Problem 1

- (a) $N_{k+1} = N_k + \#KakapoBorn - \#KakapoCaptured + \#KakapoReintroduced - \#KakapoKilled$
- (b) $N_{Blood_t} = N_{Blood_{t-1}} + \text{AmongAbsorbedFromGut}_{t-1} - \text{AmongtAbsorbedIntoTissue}_{t-1} - \text{AmountFilteredByKidneys}_{t-1} + \text{AmountReabsorbed}_{t-1}$

Problem 2

(a) Derive model one

- (i) The number of Female is half of the population, $Female = 0.5 * N_k$. Since A female lays one egg every four year. The expected Amount of female give birth in year k is a quarter of number of females, so $Birth_k = \frac{1}{4} \times Female_k$. And only 29% of new born can survive to year k+1. So the new born in year k+1 is, $Newborn_k = 0.29 \times Birth_k$. Put together,

$$Newborn_k = 0.29 * 0.25 * 0.5 * N_k = 0.03625 * N_k$$

- (ii) Based on the information, one-fiftyth of the population would die every year.

$$Death_k = 0.02 * N_k$$

- (iii) Assume that the starting population size on this island is 50 birds

```
#Define the function to stimulate recurrence.
N_nextyear <- function(N_thisyear, birthrate = 0.25, surrate = 0.29) {
  N_thisyear + surrate * birthrate * 0.5 * N_thisyear - 0.02 * N_thisyear
}

#Run stimulation
N <- 50
PopulationArray <- rbind(0:5, rep(50, 6))
for (i in 2:6) {
  PopulationArray[2, i] <- N_nextyear(PopulationArray[2, i-1])
}
apply(PopulationArray, 2, function(x) {
  (paste("The population in year", x[1], "is", x[2]))
})

## [1] "The population in year 0 is 50"
## [2] "The population in year 1 is 50.8125"
```

```
## [3] "The population in year 2 is 51.638203125"
## [4] "The population in year 3 is 52.4773239257813"
## [5] "The population in year 4 is 53.3300804395752"
## [6] "The population in year 5 is 54.1966942467183"
```

- (iv) Mathmatically, we realize the recurrence can be reduce to $N_k = 1.01625^k \times 50$, which is exponential explode to inf. So we can simple solve $1.01625^k \times 50 = 100$, solution is 43.0008536 years round up. And $1.01625^k \times 50 = 200$, solution is 86.0017072 years round up

(b) Comparing two strategeis

- (i) **Strategy 1** changes birth rate and affect Newborn. $\text{Newborn}_k = 0.29 \times \frac{1}{3} \times 0.5 \times N_k = 0.0483 \times N_k$, and

$$N_{k+1} = N_k + 0.0483 \times N_k - 0.02 \times N_k = 1.0283 \times N_k$$

Simply change the parameter birthrate in N_nextyear. Run the same stimulation. echo = FALSE

```
## [1] "The population in year 0 is 50"
## [2] "The population in year 1 is 51.4166666666667"
## [3] "The population in year 2 is 52.8734722222222"
## [4] "The population in year 3 is 54.3715539351852"
## [5] "The population in year 4 is 55.9120812966821"
## [6] "The population in year 5 is 57.4962569334214"
```

- (ii) **Strategy 2** changes survive rate and affect Newborn. $\text{Newborn}_k = 0.5 \times \frac{1}{4} \times 0.5 \times N_k = 0.0625 \times N_k$, and

$$N_{k+1} = N_k + 0.0625 \times N_k - 0.02 \times N_k = 1.0425 \times N_k$$

Simply change the parameter surrate in N_nextyear. Run the same stimulation. echo = FALSE

```
## [1] "The population in year 0 is 50"
## [2] "The population in year 1 is 52.125"
## [3] "The population in year 2 is 54.3403125"
## [4] "The population in year 3 is 56.64977578125"
## [5] "The population in year 4 is 59.0573912519531"
## [6] "The population in year 5 is 61.5673303801611"
```

- (iii) Obsviously, they are both exponential growth and $1.0425 > 1.0283$, so **Strategy 2** gives the biggest increase.

Problem 3

(a)

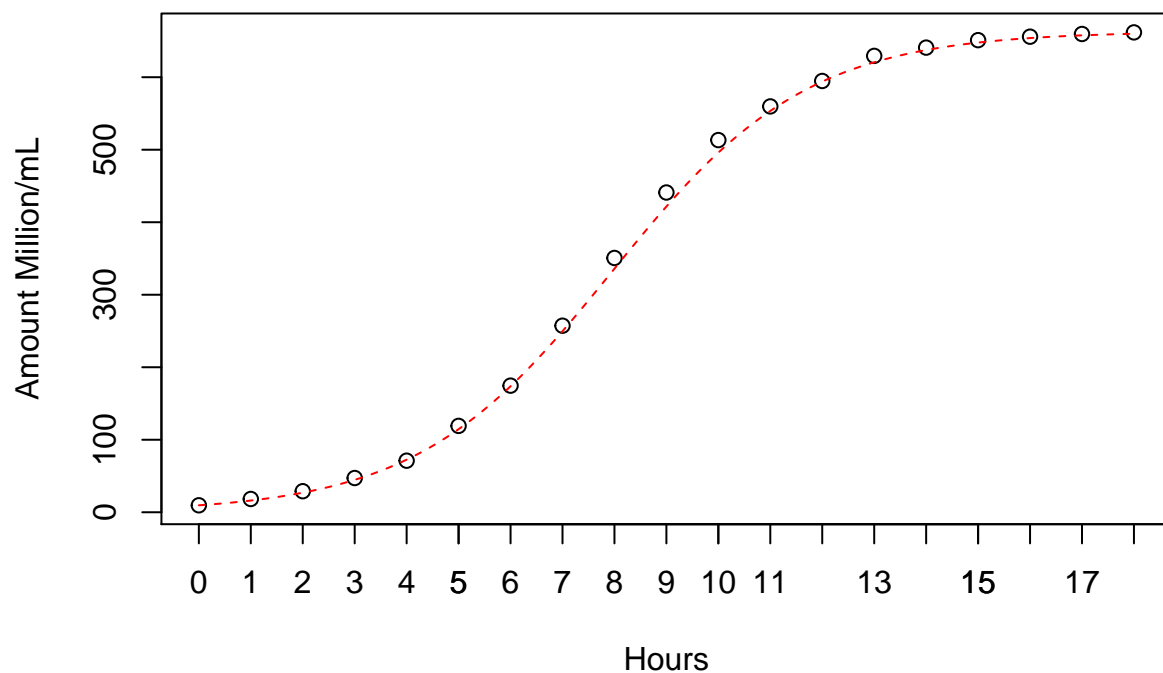
Start from the equation: $N(t + \Delta t) = N(t) + \Delta t(b - m)N(t)$, if we can find the value of birth rate, I can estimate mortality rate. Or if we can find out the distribution of life span, such as average, or pmf, we can infer motality rate.

Moreover, it's likely that in the early hours, the change rate is dominated by birth rate, and death rate play in a role in later hours. So we can estimate the birth rate with data from first few hours and calculate coresponding death rate with data in later hours.

(b)

```
data <- read.table("31395.txt", header = TRUE)
plot(data, main="Yeast Cell Number against Time in Carlson's Experiment", ylab="Amount Million/mL")
axis(1, at = seq_along(data[[1]]))
timevalues <- seq(0, 18, by=0.1)
y <- seq_along(timevalues)
for (i in seq_along(timevalues)) {
  y[i] <- (9.741*exp(0.530877*timevalues[i])) / (1+0.01469*exp(0.530877*timevalues[i]))
}
lines(timevalues, y,lty=2,col=2)
```

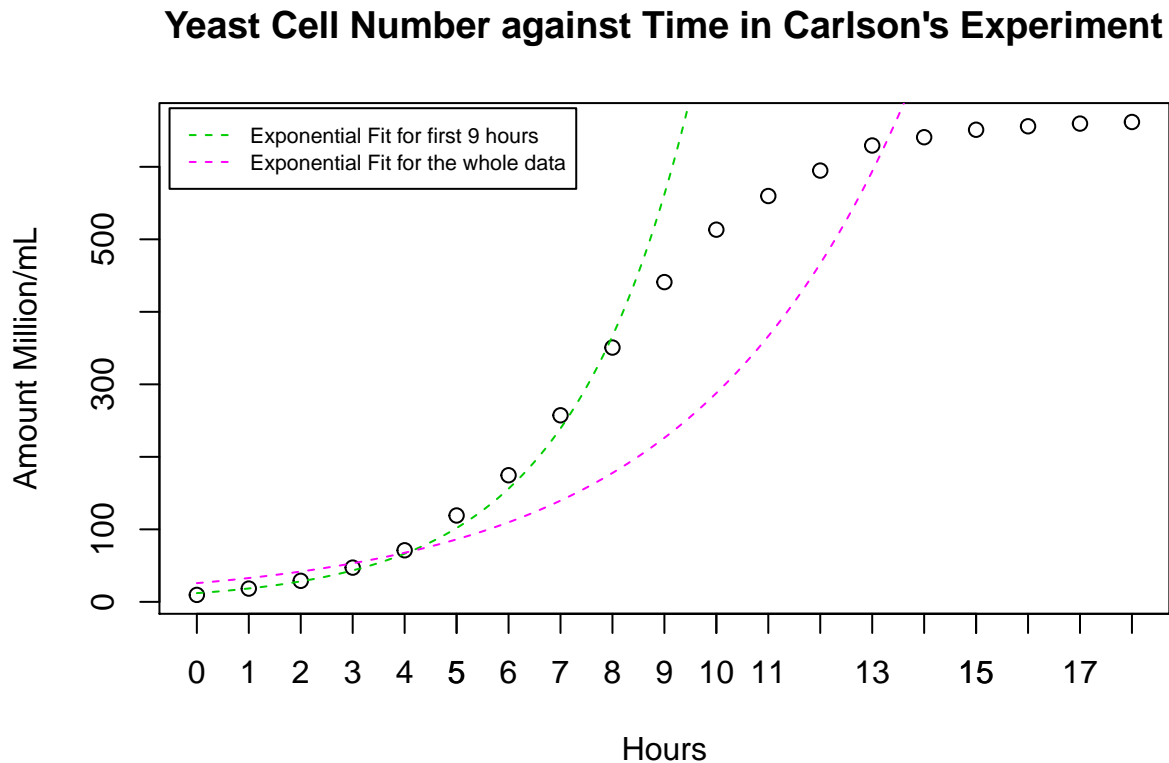
Yeast Cell Number against Time in Carlson's Experiment



Logistic growth is similar to exponential growth model when N is small. Exponential growth model is valid under ideal condition where there is no environment adverse force. However, there is environment limit apply upon every system. So exponential growth would only work when population is small. On the other hand, logistic growth model consider environment force, and fits better than exponential.

```
data <- read.table("31395.txt", header = TRUE)
plot(data, main="Yeast Cell Number against Time in Carlson's Experiment", ylab="Amount Million/mL")
axis(1, at = seq_along(data[[1]]))
for (i in c(10, 18)) {
  exponential.model <- lm(log(Amount) ~ Hours, data = data[1:i,])
  timevalues <- seq(0, 18, by=0.1)
  Counts.exponential2 <- exp(predict(exponential.model,list(Hours=timevalues)))
  lines(timevalues, Counts.exponential2,lty=2,col=(i%/%3))
}
```

```
}
legend("topleft", inset = 0.01, cex = 0.7 ,c("Exponential Fit for first 9 hours", "Exponential Fit for the whole data"))
```



From the plot, it's obvious that the growth dose not fit exponential, but it fits well in first 9 hour when the population is relatively small. From this perspective, it is very likely that the growth was limited by enrivoment resources. The growth has limited by enviroment capacity.

(c)

This is shown in plot above.

Also, we can take log to Amount and run a least square method.

```
exponential.model <- lm(log(Amount) ~ Hours, data = data[1:10,])
```

The R Square of the model is 0.9890276, fits very well

(d)

print $\frac{N_{K+1}}{N_k}$ of first couple estimates:

```
esp <- exp(predict(exponential.model,list(Hours=seq(5))))
print(esp[-1] / esp[-5])
```

```
##          2          3          4          5
## 1.532808 1.532808 1.532808 1.532808
```

Therefore, the least square model produces $\frac{N_{K+1}}{N_K} = 1.532808$, so an estimate using least square method for growth rate is $R_0 = 0.532808$

Problem 4

(a)

Consider time interval of 1 year. Population is N_k , where half are women, so number of Women $W_k = \frac{1}{2} \times N_k$. Each have 6 children through 35 year, so on average, every women is expected to have $\frac{6}{35}$ children each year. Putting together there are $\frac{1}{2} \times N_k \times \frac{6}{35} = \frac{3}{35} \times N_k$ new born each year. Thus $b = \frac{3}{35}$

(b)

Based on the information, the population is in exponential decay with half-life period of 35 years. Put this information into equation, we get:

$$N_{k+35} = \frac{1}{2} \times N_k$$

$$(1 - m)^{35} N_k = \frac{1}{2} \times N_k$$

$$m = 1 - \frac{1^{\frac{1}{35}}}{2} = 0.01960939$$

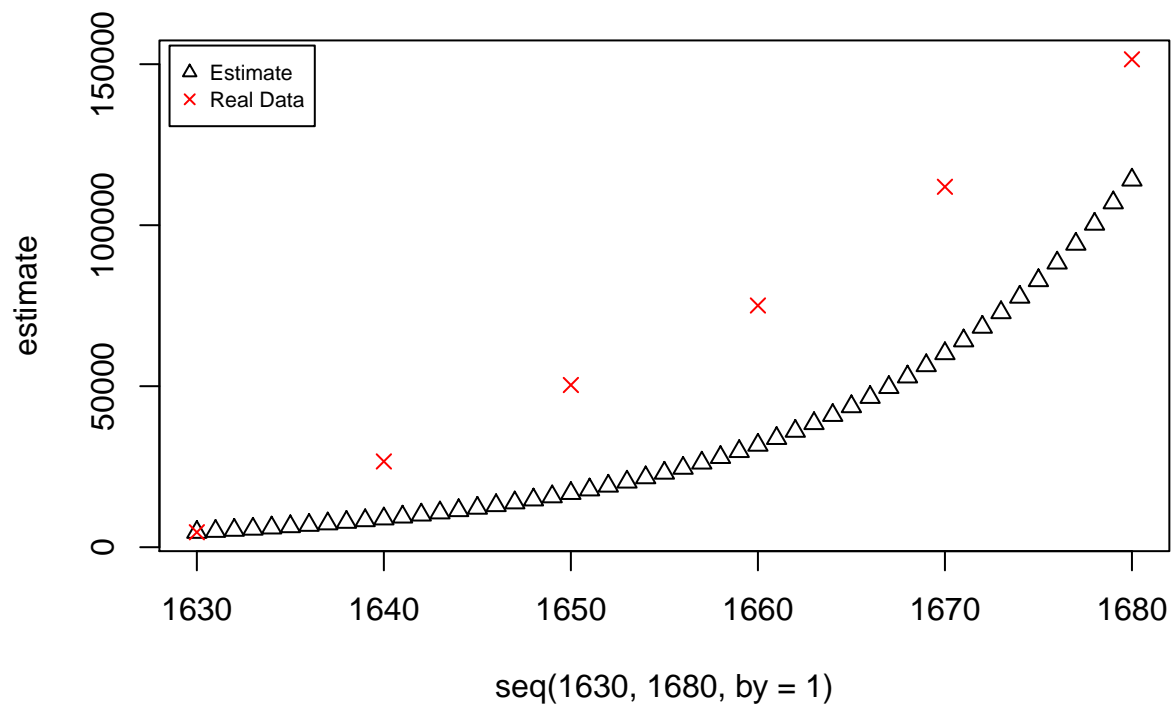
(c)

- (1) This model omits the fact there are people move in and move out. We can add another term to count immigrant.
- (2) There are some infant can't survive through first year, based on the medical condition at that time. We can add a survive rate.
- (3) Since the United States is a new settler, the environment is harsh and initial population is more likely to contain more male. We need to separate population for man and women, then find leslie matrix.

(d)

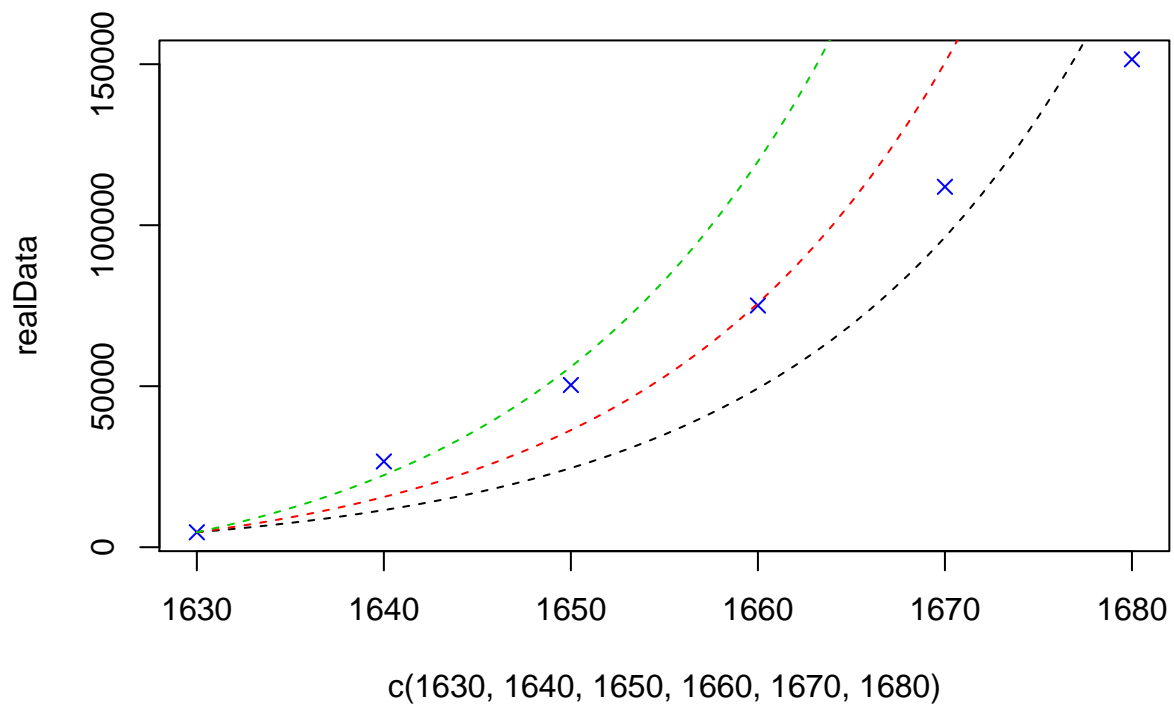
```
population <- function (year, Ni = 4646, b = 3/35, m = 0.01960939) {
  Ratio <- 1 + b - m
  Ratio ^ (year - 1630) * Ni
}
estimate <- population(seq(1630,1680,by=1))
realData <- c(4646, 26634, 50368, 75058, 111935, 151507)

plot(seq(1630,1680,by=1), estimate, col=1, pch=2,ylim=range(realData))
points(c(1630, 1640, 1650, 1660, 1670, 1680), realData, col=2, pch=4)
legend("topleft", inset = 0.01, cex = 0.7 ,c("Estimate", "Real Data"), pch=c(2,4),col=c(1, 2))
```



(e)

```
Ratio <- 1+3/35-0.01960939
population_next <- function(N, r, I) {
  N * r + I
}
year <- seq(1630,1680,by=1)
estimate <- matrix(0,nrow=3,ncol=length(year))
estimate[,1] <- 4646
plot(c(1630, 1640, 1650, 1660, 1670, 1680), realData, col=4, pch=4)
for (i in 1:3) {
  I <- c(200, 500, 1000)[i]
  for (j in 2:length(year)) {
    estimate[i, j] <- population_next(estimate[i, j-1], Ratio, I)
  }
  lines(seq(1630,1680,by=1), lty=2,estimate[i,], col=i, pch=2)
}
```



Seems like none fits very well.
 1000 fits early years.
 500 fits middle years. 200 fits one later years.

Problem 5

(a)

```
Ct <- function(c) {
  (1-0.2425) * c
}
concentration <- rep(40,8)
concentration[1] <- 0
names(concentration) <- paste("C", 0:7, sep="")
for (i in 3:8) {
  concentration[i] <- Ct(concentration[i-1])
}
concentration[8] = concentration[8] + 40
print(concentration)
```

```
##          C0          C1          C2          C3          C4          C5          C6          C7
## 0.000000 40.000000 30.300000 22.952250 17.386329 13.170145  9.976384 47.557111
```

(b)

When take n-th pill, time is $6 \times (n - 1)$, time after take in Pill_{n-1} is 6×1 , time after Pill_{n-2} is $6 \times 2 \dots$ time after take in Pill_{n-k} is $6k$, where k ranges from 1 to n.

The concentration caused by 1 pill k hours after take in is

$$\text{Concentration}(k) = \begin{cases} 0 & k = 0 \\ (1 - 0.2425)^{k-1} \times 40 & k \neq 0 \end{cases}$$

Based on this information, we get the recurrence

$$C_n = (1 - 0.2425)^6 \cdot C_{n-1} + 40$$

(c)

```
Ratio <- (1-0.2425)^6
CP <- function (c) {
  Ratio * c + 40
}
concentration <- rep(40, 5)
names(concentration) <- paste("C", 1:5, sep="")
for (i in 2:5) {
  concentration[i] <- CP(concentration[i-1])
}
print(concentration)
```

```
##          C1          C2          C3          C4          C5
## 40.00000 47.55711 48.98486 49.25460 49.30556
```

(d)

Solve $C = (1 - 0.2425)^6 \cdot C + 40$, solution is 49.3174

(e) Solve Recurrence

- Simplify recurrence

$$C_n - (1 - 0.2425)^6 \cdot C_{n-1} - 40 = 0$$

- Solve Characteristic polynomial

$$r - (1 - 0.2425)^6 = 0 \Rightarrow r = (1 - 0.2425)^6$$

- Therefore,

$$C_n = A \cdot (1 - 0.2425)^{6n} + C_{par}$$

where C_{par} is particular solution, in this case should be an constant B. And we will solve B in next step.

- Plug C_{par} into recurrence, we get

$$B = (1 - 0.2425)^6 \cdot B + 40 \Rightarrow B = 49.3174$$

- Then

$$C_n = A \cdot (1 - 0.2425)^{6n} + 49.3174$$

, and solve for initial condition

$$C_1 = 40 = A \cdot (1 - 0.2425)^6 + 49.3174 \Rightarrow A = -49.3173$$

- Which gives us

$$C_n = -49.3173 \cdot (1 - 0.2425)^{6n} + 49.3174$$

- First part of this formula exponential decay to 0 and second part is constant. So

$$\lim_{n \rightarrow \infty} C_n = 49.3174$$