

# Mathematical Modeling of Infectious Diseases, UCSF

## Homework 3

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**Exercise 1.** Prove  $\text{cov}(X, X) = \text{var}(X)$ . Prove  $\text{cov}(X, aY) = a\text{cov}(X, Y)$  for  $a$  any constant. Prove  $\text{cov}(X, Y) = \text{cov}(Y, X)$  for any random variables  $X$  and  $Y$  (as long as  $E[X]$ ,  $E[Y]$  exist!) ■

$$\text{cov}(X, X) = E[X^2] - E[X]E[X] = \text{var}(X)$$

$$\text{cov}(X, aY) = E[XaY] - E[X]E[aY] = aE[XY] - aE[X]E[Y] = a(E[XY] - E[X]E[Y]) = a\text{cov}(X, Y)$$

$$\text{cov}(X, Y) = E[XY] - E[X]E[Y] = E[YX] - E[Y]E[X] = \text{cov}(Y, X)$$

**Exercise 2.** Prove for any  $X$  and  $Y$  independent or not,  $\text{var}(X + Y) = \text{var}(X) + 2\text{cov}(X, Y) + \text{var}(Y)$ . ■

From lecture 1:

$$\text{var}[X + Y] = E[X^2] + 2E[XY] + E[Y^2] - (EX)^2 - 2E[X]E[Y] - (EY)^2$$

$$= E[X^2] + 2E[XY] + E[Y^2] - (EX)^2 - 2E[X]E[Y] - (EY)^2$$

$$\text{var}(X + Y) = \text{var}(X) + 2E[XY] - 2E[X]E[Y] + \text{var}(Y)$$

$$\text{var}(X + Y) = \text{var}(X) + 2\text{cov}[X, Y] + \text{var}(Y)$$

**Exercise 3.** Pick  $a = 1/3$ , and try adding  $(1/3)^0 + (1/3)^1 + (1/3)^2 + \dots = 1 + 1/3 + 1/9 + 1/27 + \dots$ . In other words, add successive terms and note the behavior. What does the formula say the limit ought to be? ■

Use R:

```
sum <- 0
sums <- c()
for(ii in 0:15){
  sum <- sum + (1/3)^ii
  sums <- c(sums, sum)
}
```

| Number of Terms | Sum    |
|-----------------|--------|
| 1               | 1.0000 |
| 2               | 1.3333 |
| 3               | 1.4444 |
| 4               | 1.4815 |
| 5               | 1.4938 |
| 6               | 1.4979 |
| 7               | 1.4993 |
| 8               | 1.4998 |
| 9               | 1.4999 |
| 10              | 1.5000 |
| 11              | 1.5000 |
| 12              | 1.5000 |
| 13              | 1.5000 |
| 14              | 1.5000 |
| 15              | 1.5000 |
| 16              | 1.5000 |

The limit ought to be  $1/(1 - 1/3) = 1.5$

**Exercise 4.** Use the geometric series formula to show that the sum of the geometric probabilities is 1. Pay attention to the fact that the geometric probabilities start at 1, not 0. ■

$$\sum_y P(Y = y) = \sum_{y=1}^{\infty} p(1-p)^{y-1} = p \sum_{u=0}^{\infty} (1-p)^u = p \frac{1}{1 - (1-p)} = 1$$

**Exercise 5.** Suppose that each time a case of disease is introduced into a region,  $a < 1$  secondary cases are produced from each case on average. Show that the size of the  $n$  generation is  $a^n$ . Don't worry about what fractions of a person mean; interpret these in some average sense. Show that the total size of the outbreak—adding all cases together including the index case—is  $1/(1 - a)$ . ■

**Proof by induction:**

**Step 1:** Show the base case is true.

In the first generation the expected number of cases is  $a$  (given in the problem).

**Step 2:** If the expected size of the  $n$ th generation is  $a^n$ , show the expected size of the  $n + 1$ th generation is  $a^{n+1}$ .

Therefore, we have proved that the size of the  $n$ th generation is  $a^n$ .

The total outbreak size is given by:

$$\sum_{n=0}^{\infty} a^n = \frac{1}{1-a}$$

**Exercise 6.** Suppose that on average a case of disease causes  $R > 1$  secondary cases when no one is vaccinated or immune. Suppose now that a fraction  $f$  is vaccinated with complete protection. Discuss the epidemiological credibility of the assumption that now each case would cause (on average)  $R(1-f)$  secondary cases (i.e. when the coverage fraction is  $f$ ). (Do you see the expectation of a binomial possibly concealed in here?) Assume that with coverage  $f$ , each case causes  $R(1-f)$  secondary cases, and that  $R(1-f) < 1$ ; compute the total expected size of an outbreak, counting all generations. What is the critical coverage level needed for  $R(1-f) < 1$ ? ■

We are making the assumption that each person's vaccination status is independent of another's. This may not be a valid assumption in the real world, where non-vaccinated individuals may cluster at certain schools or in certain neighborhoods.

$$R(1-f) < 1$$

$$1-f < \frac{1}{R}$$

$$f > 1 - \frac{1}{R}$$

The critical coverage level is  $1 - \frac{1}{R}$ .

**Exercise 7.** Use Newton's formula

$$(a+b)^N = \sum_{i=0}^N \binom{N}{i} a^i b^{N-i}$$

to prove that the probability generating function for a binomial is  $(up + (1-p))^N$ . ■

Let  $g(u)$  be the PGF for a binomial.

$$\begin{aligned} g(u) &= E[u^X] = \sum_{x=0}^N u^x \binom{N}{x} p^x (1-p)^{N-x} \\ &= \sum_{x=0}^N \binom{N}{x} (up)^x (1-p)^{N-x} \\ &= (up + (1-p))^N \end{aligned}$$

**Exercise 8.** Suppose a needle is discarded with probability 0.2 after every usage; each disposal decision is independent of the past. What is the expected number of usages? ■

$$1/0.2 = 5$$

**Exercise 9.** Let  $Y$  be geometric with parameter  $p$ . Prove that the conditional distribution of  $Y - Y_0$  given  $Y > Y_0$  for any  $Y_0$  is geometric with parameter  $p$  also. ■

We can do this by showing:  $E[u^Y] = E[u^{Y-Y_0} | Y > Y_0]$ .

$$E[u^Y] = \sum_{i=0}^{\infty} (1-p)^{i-1} p u^i = \frac{up}{1 - (1-p)u}$$

$$E[u^{Y-Y_0} | Y > Y_0] = \sum_{i=Y_0+1}^{\infty} (1-p)^{i-Y_0-1} p u^{i-Y_0}$$

$$= \sum_{j=1}^{\infty} (1-p)^{j-1} p u^j$$

$$= \frac{up}{1 - (1-p)u}$$

Since both distributions have the same generating function, they are the same distribution.

**Exercise 10.** Intuitively we know the mean of the Poisson should be  $\lambda$ , from the derivation. But what if something untoward happened when we took the limit? Show that if  $Y$  is Poisson with parameter  $\lambda$ , then  $E[Y] = \lambda$ . ■

$$E[Y] = \sum_{i=0}^{\infty} i \frac{e^{-\lambda} \lambda^i}{i!} = \lambda e^{-\lambda} \sum_{i=0}^{\infty} \frac{i \lambda^{i-1}}{i!}$$

$$= \lambda e^{-\lambda} \sum_{i=0}^{\infty} \frac{d}{d\lambda} \frac{\lambda^i}{i!}$$

$$= \lambda e^{-\lambda} \frac{d}{d\lambda} \sum_{i=0}^{\infty} \frac{\lambda^i}{i!}$$

$$= \lambda e^{-\lambda} \frac{d}{d\lambda} e^{\lambda} = \lambda$$

**Exercise 11.** Suppose that accidents occur according to a Poisson distribution with mean 1. What is the probability of zero accidents? Suppose the mean is 0.1 or 10; what is the probability of zero accidents? ■

Use R:

```
dpois(0,1)
```

```
## [1] 0.3678794
```

```
dpois(0,.1)

## [1] 0.9048374

dpois(0,10)

## [1] 4.539993e-05
```

**Exercise 12.** Suppose that HIV-negative participants in a trial acquire new partners at random according to a Poisson distribution with mean  $\lambda$ . Supposing that the prevalence of infection is  $p$  and the probability of infection given an infected partner is  $\beta$ , what is the distribution of the number of *sufficient exposures*? What is the probability of zero sufficient exposures? What is the probability of the person becoming infected (i.e. of having one or more sufficient exposures)? ■

$$P(X = i) = \frac{e^{-\lambda} \lambda^i}{i!} = \frac{e^{-\beta p} (\beta p)^i}{i!}$$

Probability of 0 sufficient exposures:

$$P(X = 0) = \exp(-\beta p)$$

Probability of at least 1 sufficient exposure:

$$1 - P(X = 0) = 1 - \exp(-\beta p)$$

**Computer Exercise 13.** Please use the above R code to simulate the IAR(1) process with (a)  $\lambda = 10$  and  $p = 0$  (no survival),  $p = 0.3$ , and  $p = 0.8$ . Part (b): Also, use  $\lambda = 200$ , with  $p = 0.1$  in one run and  $p = 0.99$  in another; please comment on the differences. Please try, in an exploratory way, different values of  $\lambda$  and  $p$ ; feel free to set `xz <- 1:500` to go out to time 500 or even longer. Plot your results; comment on the appearance of the graph (smoothness, initial rise). Only turn in 3–4 graphs. ■

Load in given code:

```
iarlsim <- function(len,lambda,pp,starting=0) {
  if (len <= 0 || len > 65536) {
    stop("length bad")
  }
  if (pp<0 || pp>1.0) {
    stop("p bad")
  }
}
```

```

if (lambda < 0) {
  stop("lambda bad")
}
answer <- rep(NA,len)
answer[1] <- starting
cur <- starting
for (ii in 2:len) {
  cur <- pp*cur + rpois(1,lambda=lambda)
  answer[ii] <- cur
}
answer
}

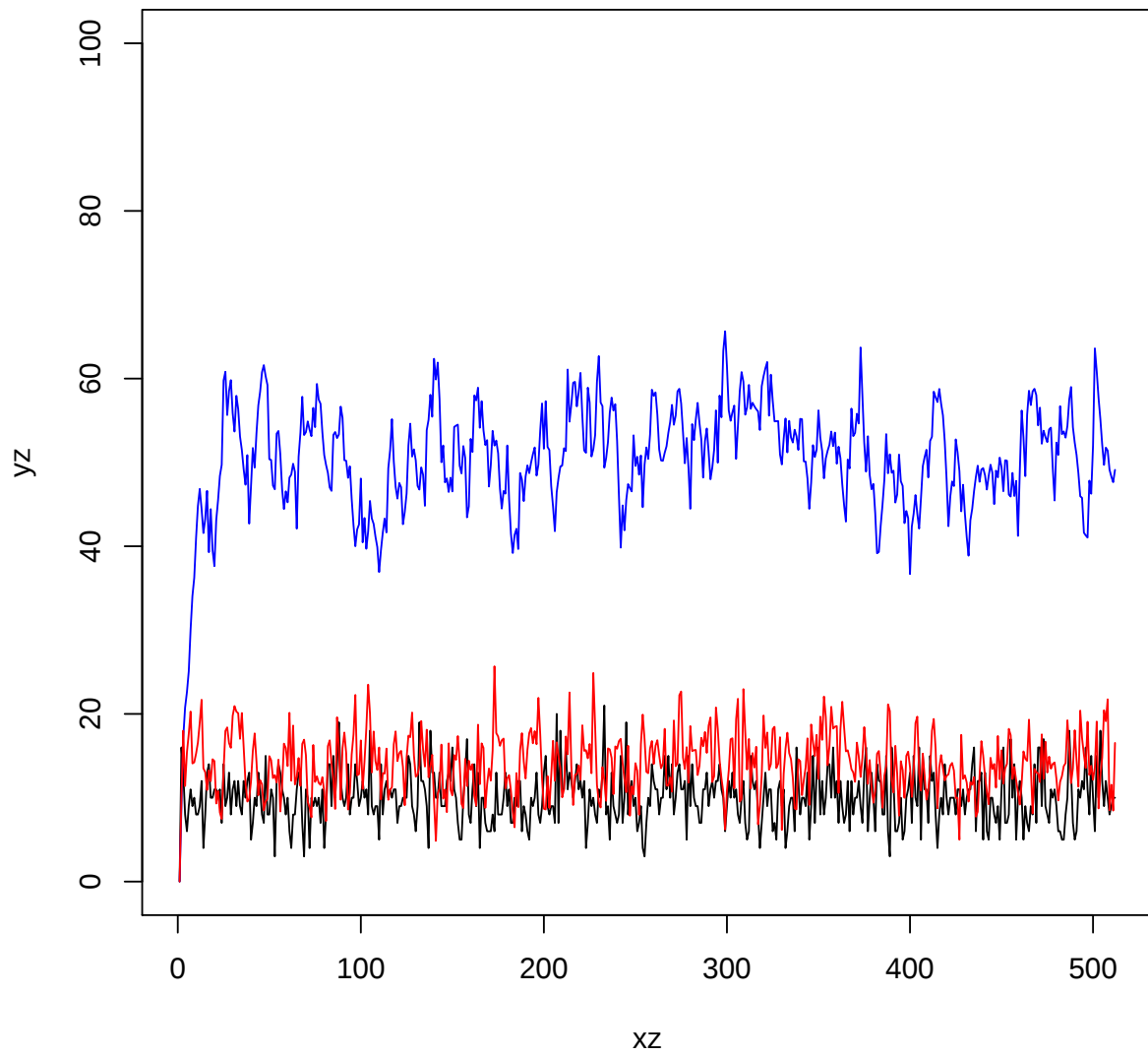
```

a)

```

xz <- 1:512
yz <- iar1sim(length(xz), 10, 0)
yz2 <- iar1sim(length(xz), 10, 0.3)
yz3 <- iar1sim(length(xz), 10, 0.8)
plot(xz,yz,type="l",ylim=c(0,100))
points(xz,yz3,type="l",col="blue")
points(xz,yz2,type="l",col="red")

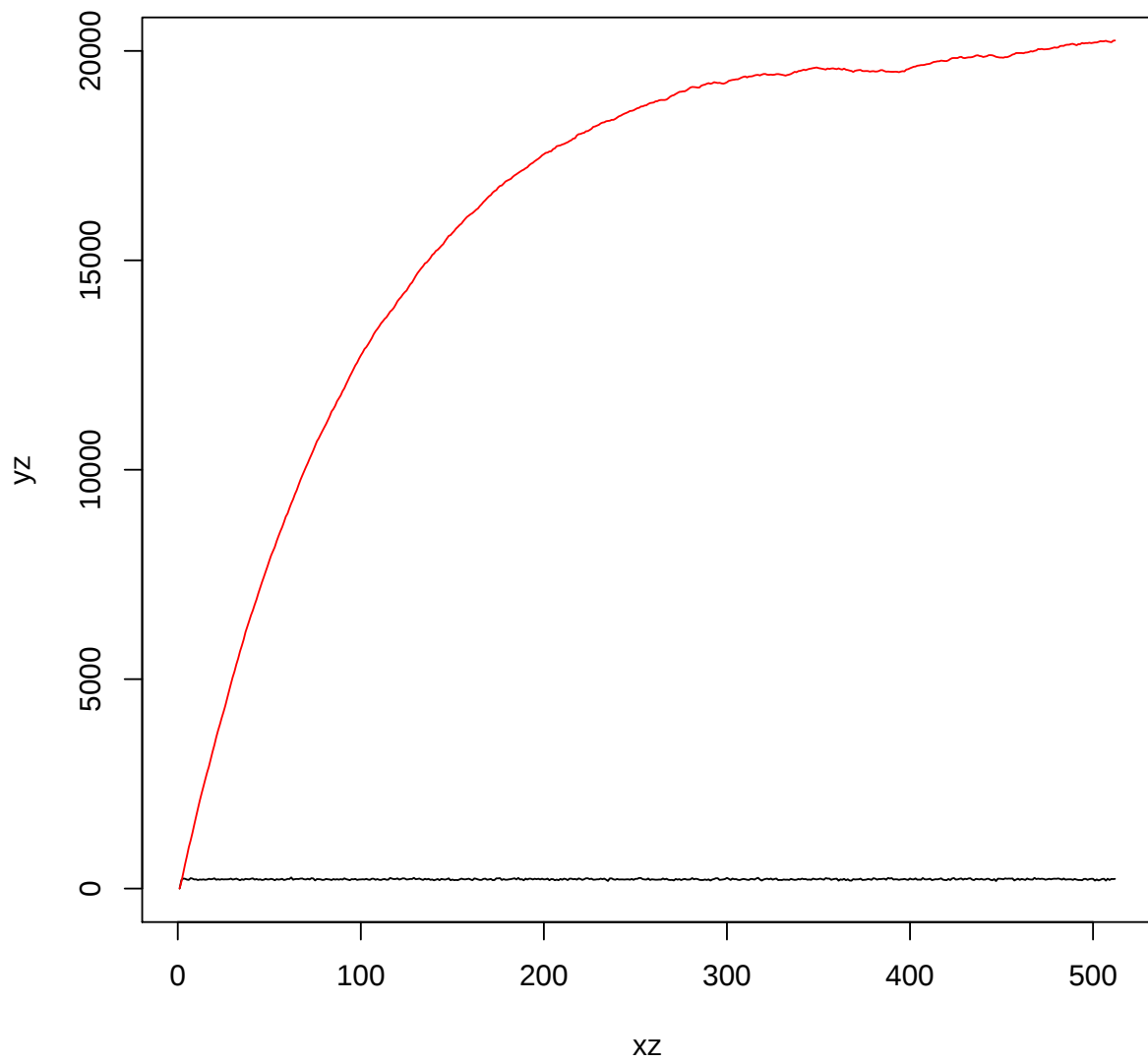
```



b)

```
xz <- 1:512  
yz <- iar1sim(length(xz), 200, 0.1)  
yz2 <- iar1sim(length(xz), 200, 0.99)  
plot(xz,yz,type="l",ylim=c(0,2e4))
```

```
points(xz,yz2,type="l",col="red")
```



**Exercise 14.** How long did this assignment take you? ■