## Homework 2

1. (book #2.2.) A monkey types on a 26-letter keyboard that has lowercase letters only. Each letter is chosen independently and uniformly at random from the alphabet. If the monkey types n = 1,000,004 letters, what is the expected number of times the sequence "proof" appears?

The expected number of times is equal to (up to 2 decimals is fine):

2. (book #2.4) Prove that  $\underbrace{E[X^n]}_L \ge \underbrace{(EX)^n}_R$  for n = 2k.

Which of the following arguments proves the claim?

- (a)  $f''(x) = n(n-1)x^{2(k-1)} \ge 0 \implies L \ge R$  by Jensen's inequ.
- (b) by the mean value theorem we have  $L \geq R$
- (c)  $L \ge (E[X^n])^{1/n} = E(X) \ge [E(X)]^n = R$
- (d)  $L \ge \operatorname{Var}(X^k) \ge [\operatorname{Var}(X)]^k \ge R$
- (e) none of these
- 3. (book # 2.14) The geometric distribution arises as the distribution of the number of times we flip a coin until it comes up heads. Consider now the distribution of the number of flips X until the kth head appears, where each coin flip comes up heads independently with probability p. This is known as the negative binomial distribution, and we also write  $X \sim NBin(k, p)$ . Find  $p_X(n)$ . Hint: Here X counts the number of flips including the flip with the k-th head.

Mark the correct expression for  $p_X(n)$ , below.

- (a)  $\binom{n-1}{k-1} p^k (1-p)^{n-k}$  (b)  $\binom{n}{k} p^k (1-p)^{n-k}$  (c)  $p^k (1-p)^{n-k}$  (d)  $\binom{n}{k} p^{k-1} (1-p)^{n-k}$  (e) none of these
- 4. (book # 2.15) For a coin that comes up heads independently with probability p on each flip, what is the expected number (X) of flips until the kth heads?

Hint: here x counts the number of flips including the flip with the k-th head.

Mark the correct expression for E(X):

- (a) 1/p (b) kp (c) k/p (d) np/k (e) none of these
- 5. (book #2.24) We roll a standard fair die over and over. What is the expected number of rolls (N) until the first pair of consecutive sixes appears?

Hint: Let  $X_i$  denote the face on the i-th roll. Set up Lemma 2.5 twice, once for EN, conditioning on  $X_1$ ; and a second time for  $E(N \mid X_1 = 6)$ , now (additionally) conditioning on  $X_2$ . For the second application of Lemma 2.5, a version of the lemma as:  $E(X \mid Y = y, Z = z) = \sum_z E(X \mid Y = y, Z = z) \Pr(Z = z \mid Y = y)$  (that is, use the lemma for the probability function  $q(N) = \Pr(N \mid X_1 = 6)$ ). Solve the two equations in a = EN and  $b = E(N \mid X_1 = 6)$ .

EN =

(book # 2.25) A blood test is being performed for  $n = N \cdot k$  individuals by pooling the samples for k people, and analyzing them together. If the test is negative, this pooled test suffices for all k individuals.

If the test is positive, we test each of the k persons separately, i.e., we carry out a total of k+1 tests.

Suppose that we create N disjoint groups of k people and use the pooling method. Let  $A_i = i$ -th person has a positive result, and assume  $Pr(A_i) = p$ , independently,  $i = 1, \ldots, n$ .

**6.** Let  $B_j$  denote the event of a positive pooled sample, for the j-th pooled sample consisting of samples from a batch of k individuals.

For p = 0.01, k = 4 and n = 40, find  $Pr(B_i)$ .

$$\Pr(B_j) =$$

7. Let X = # of tests. What is the expected number of tests necessary, E(X)?

For p = 0.01, k = 4 and n = 40, find E(X).

$$E(X) =$$

**8.** Describe how to find the best value of k.

Nothing to turn in for this problem.

9. Prove the Poi approximation of binomial probabilities.

Claim: If  $X \sim \text{Bin}(n,p)$  with  $np = \lambda$  (or  $p = \lambda/n$ ) and large n, then  $p_X(i) \approx e^{-\lambda} \frac{\lambda^i}{i!}$ , i.e., approximated by a Poisson probability with rate  $\lambda$ .

Hint: Use  $(1-\lambda/n)^i \to 1$ ;  $(1-\lambda/n)^n \to e^{-\lambda}$ ; and  $(n-i+1)/n \to 1$  as  $n \to \infty$ . The last assumption implies we use the approximation for moderate i only.

Which of the following arguments justifies the Poisson approximation? All limits are as  $n \to infty$ .

- (a)  $\lim np = \lambda$ , by assumption
- (b)  $\lim_{n \to \infty} {n \choose i} p^i (1-p)^{(n-i)} = \lim_{n \to \infty} (1+i/n)^n = e^{-\lambda} \frac{\lambda^i}{i!}$
- (c)  $\lim_{i \to \infty} {n \choose i} p^i (1-p)^{(n-i)} = \lim_{i \to \infty} \frac{n(n-1)\cdots(n-i+1)}{n^i} = e^{-\lambda} \frac{\lambda^i}{i!}$ (d)  $\lim_{i \to \infty} {n \choose i} p^i (1-p)^{(n-i)} = \lim_{i \to \infty} \frac{n(n-1)\cdots(n-i+1)}{n^i} \frac{\lambda^i}{i!} \frac{(1-\lambda/n)^n}{(1-\lambda/n)^i} = e^{-\lambda} \frac{\lambda^i}{i!}$
- (e) none of these
- 10. Recall the birthday problem which we discussed in Chapter 1. Assume that each of n people is equally likely to have any of the 365 days of the year as his or her birthday. Let  $A_2$  = "no two persons share the same birthday". In Chapter 1 we used a combinatorial argument to find  $Pr(A_2)$ . Alternatively we could use a Poisson approximation as follows. Consider all  $L = \binom{n}{2}$  possible (unordered) pairs  $\{i, j\}$ . Let  $E_{ij} = 1$  if i and j have the same birthdays and  $E_{ij} = 0$  otherwise. Then  $p = \Pr(E_{ij} = 1) = 1/365$ . Although the  $E_{ij}$  are not mutually independent, they are only weakly dependent. In fact, they are pairwise independent. Let  $X = \sum_{i < j} E_{ij}$  denote the number of pairs with shared birthdays. Let  $Y \sim$ Bin(L,p). Then  $p_X(0) \approx p_Y(0)$ , and  $p_Y(0)$  in turn we can approximate by a Poisson approximation.

Use a similar approximation to find the probability of  $A_3 =$  "no 3 same birthdays among n people". That is, letting X = # of triples  $\{i, j, k\}$  with same birthday, approximate  $p_X(0)$ .

Hint: Let  $K = \binom{n}{3}$  and let  $p = (1/365)^2$  and proceed similar to before.

For n = 50, find the described approximation for  $Pr(A_3)$ :

$$\Pr(A_3) =$$

11. Prove Lemma 5.2 for two Poi r.v.'s. That is, assuming  $X \sim \text{Poi}(\mu_1)$  and  $Y \sim \text{Poi}(\mu_2)$ , independently, show that  $S = X + Y \sim \text{Poi}(\mu_1 + \mu_2)$ .

*Hint*: Use the law of total probability with  $E_k = \{X_1 = k\}$  to find  $p_S(j) = \sum_{k=0}^{j} \dots$ 

Starting with the law of total probability for  $p_S(j)$ , and then moving all factors that do not involve j outside the sum we get

$$p_S(j) = \sum_{k=0}^{j} p_{X_1}(k) \, p_{X_2}(j-k) = \sum_{k=0}^{j} \frac{e^{-\mu_1} \mu_1^k}{k!} \cdot \frac{e^{-\mu_2} \mu_2^{j-k}}{(j-k)!} = \frac{e^{-(\mu_1 + \mu_2)}}{j!} \, \sum_{k=0}^{j} \binom{j}{k} \, \mu_1^k \mu_2^{j-k}$$

Let RHS denote the final right hand side above. Completing the proof from here, which of the following argument proves the claim.

(a) 
$$RHS = \frac{e^{-(\mu_1 + \mu_2)}(\mu_1 + \mu_2)^j}{j!} \sum_{k=0}^j {j \choose k} p^k (1-p)^{j-k}$$
 with  $p = \mu_1/(\mu_1 + \mu_2)$  and  $\sum_k \dots = 1$ .  
(b)  $RHS = \frac{e^{-(\mu_1 + \mu_2)}}{j!}$  since  $\sum_{k=0}^j {j \choose k} \mu_k^k \mu_2^{j-k} = 1$ 

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 since  $\sum_{k=0}^{j} {j \choose k} \mu_1^k \mu_2^{j-k} = 1$ 

(c) Use 
$$np = \lambda = \mu_1 + \mu_2$$
 for  $p = \mu_1/(\mu_1 + \mu_2)$   
(d)  $\lim_{j \to \infty} RHS = \frac{e^{-(\mu_1 + \mu_2)}(\mu_1 + \mu_2)^j}{j!}$ 

(d) 
$$\lim_{j\to\infty} RHS = \frac{e^{-(\mu_1+\mu_2)}(\mu_1+\mu_2)^j}{j!}$$

(e) none of these

## Simulation

The following problem requires some programming using R, or any comparable programming language Kullback-Leibler divergence (KL) of two distributions  $p_X$  and  $p_Y$  is defined as

$$D(p_X, p_Y) = \sum_{x} p_X(x) \log \left( \frac{p_X(x)}{p_Y(y)} \right).$$

KL divergence is a measure of discrepancy between two distributions with D=0 for  $p_X=p_Y$  and D>0 for  $p_X\neq p_Y$ . Using KL divergence, we then define mutual information (MI) for two jointly distributed r.v.'s as

$$I(X,Y) = D(p_{X,Y}, p_X p_Y),$$

that is KL divergence between the joint distribution  $p_{X,Y}$  and the product of the marginals  $p_X$  and  $p_Y$ . The latter is the hypothetical joint distribution under independence.

In this example we use MI to judge whether two variables X, Y are independent or not. We record npairs  $(X_i, Y_i) \sim p_{X,Y}$ , i = 1, ..., n, independently (data). Here  $X_i \in \{1, 2, 3, 4\}$  and  $Y_i \in \{1, 2, 3\}$ . The following  $(4 \times 3)$  table summarizes the data by reporting  $n_{X,Y}(x,y) = \#\{(X_i,Y_i) = (x,y)\}$ , the count of observations with  $X_i = x$  and  $Y_i = y$ .

Table 1. counts  $n_{X,Y}(x,y)$  (center block),  $n_X(x)$  (right column) and  $n_Y(y)$  (bottom row).

The row totals are  $n_X(x) = \#\{X_i = x\}$  and similarly the column totals report  $n_Y(y) = \#\{Y_i = y\}$ . Let  $f_{X,Y}(x,y) = n_{X,Y}(x,y)/n$  denote the (relative) frequencies, and similarly for  $f_X(x)$  and  $f_Y(y)$ . We use  $f_X \approx p_X$  as an estimate for  $p_X$ , and  $f_Y \approx p_Y$  as an estimate for  $p_Y$ . Then

$$\hat{I} = \sum_{x,y} f_{X,Y}(x,y) \log \left( \frac{f_{X,Y}(x,y)}{f_X(x)f_Y(y)} \right),$$

serves as estimate for I(X,Y). <sup>1</sup> In the following questions we implement a possible approach to decide whether to report that  $X \perp Y$ , or  $X \not\perp Y$ . The logic is

- (a) If  $X \perp Y$  were true, then I(X,Y) = 0.
- (b) Instead of I(X,Y) we can only evaluate  $\hat{I} \approx I(X,Y)$ . It is okay for  $\hat{I} > 0$ , but it should not be "too large" if  $X \perp Y$  were true;
- (c) To judge how much is "too large", we carry out a small simulation: We generate a hypothetical repeat of the experiment, generating  $X_i' \sim f_X$ ,  $Y_i' \sim f_Y$ , independently,  $i=1,\ldots,n$ , and evaluate  $\hat{I}'$ . Repeat this simulation M=100 times and record the M evaluations of  $\hat{I}'$  (see the footnote about zero counts). Let  $\hat{I}'_m$ ,  $m=1,\ldots,M$ , denote the ordered list of those M evaluations. We use  $\hat{I}^{\star} \equiv \hat{I}'_{05}$  to draw the line and decide what is "too large."
- (d) If  $\hat{I}$  is "too large", i.e.,  $\hat{I} > \hat{I}^*$  report  $X \not\perp Y$ . Otherwise we report  $X \perp Y$ .

The logic of our algorithm is an indirect argument: if in fact  $X \perp Y$  were true, then  $\hat{I} > \hat{I}^*$  is unlikely. Therefore we accept  $\hat{I} > \hat{I}^*$  as evidence "beyond reasonable doubt" against  $X \perp Y$ .

In this setup, answer the following questions:

12. Evaluate  $\hat{I}$  (step 2, above). You can use the Rmacro Ih() below this problem (you need not use them).

 $\hat{I} =$ 

13. Carry out the simulation described in step 3. Find  $\hat{I}^*$ . See the Rcode fragments sim() and Ihmstar() shown below this problem (you need not use them).

 $\hat{I}^{\star} =$ 

<sup>&</sup>lt;sup>1</sup> We lucked out here with all counts being non-zero. If any count were 0, we could just replace it by 0.5.

**14.** If  $X \perp Y$  is true, then what is  $\Pr(\hat{I} > \hat{I}^*)$ ? An approximate argument is okay.

$$\Pr(\hat{I} > \hat{I}^\star) =$$

15. In the light of (b) and (c), and following steps 1-4 above, what do you report?

```
(a) X \perp Y (b) X \not\perp Y (c) can not decide (d) need more data (e) none of these
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Note: In this problem we carried out a test of the hypothesis  $H_0: X \perp Y$ . We used  $\hat{I}(X,Y)$  as a test statistic. The decision rule to reject  $H_0$  when  $\hat{I} > \hat{I}^*$  defined the rejection region. We will talk much more about the concept of hypothesis tests later in the course.

Below are three R scripts to evaluate  $\hat{I}$  and to carry out the simulation in **10b**. The argument of Ih(nxy) is the  $(4 \times 3)$  table of counts  $n_{XY}$ , and function returns  $\hat{I}(X,Y)$ . The arguments of sim(M,fxh,fyh) are the simulation size M,  $f_X$  and  $f_Y$ , and the function returns an  $(M \times 1)$  vector  $\hat{I}'$  of  $\hat{I}(X,Y)$  for the M simulations.

```
Ih <- function(nxy)</pre>
                                                        sim <- function(M,fxh,fyh)</pre>
{ # input: f = nx x ny table of counts
                                                        { ## input: M=#simulations,
  # output: MI I(X,Y)
                                                                      fxh = marginal on X, fyh = ... on Y
    n \leftarrow sum(nxy)
                                                           ## output: Ihm = (M \times 1) vector of Ih(X,Y)
    f <- nxy/n
                                                             n \leftarrow 120 # to match the table
    fx <- apply(f,1,sum)</pre>
                                                             nx <- length(fxh)
    fy <- apply(f,2,sum)</pre>
                                                             ny <- length(fyh)</pre>
    fx[fx==0] <- 0.01
                                                             Ihm <- rep(0,M) # initialize</pre>
    fy[fy==0] <- 0.01
                                                             for(m in 1:M){
    f[f==0] \leftarrow 0.01
                                                                  xm <- sample(1:nx,n,prob=fxh,replace=T)</pre>
    fxfy <- fx %*% t(fy)
                                                                  ym <- sample(1:ny,n,prob=fyh,replace=T)</pre>
    I <- sum(f*log(f/fxfy))</pre>
                                                                  nxym <- table(xm,ym)</pre>
                                                                  Ihm[m] <- Ih(nxym)</pre>
    return(I)
}
                                                             }
                                                             return(Ihm)
                                                        }
Ihmstar <- function(Ihm)</pre>
{ ## input: list Ihm[1..M] of Ih(X,Y)
  ## output: threshold Ih* for test
    M <- length(Ihm)
    istar <- round(0.95*M)</pre>
     Is <- sort(Ihm)[istar]</pre>
    return(Is)
}
```