

## SSC386D / M394C: Monte Carlo Statistical Methods — Midterm 1

Solutions are due **We, March 11** as harcopy, or as pdf per email (please use subject “MCMC Midterm”) by Fr, March 13, 5pm in my mailbox (8th floor RLM, or stop by my office).

Open book, open notes, internet research (wikipedia or whatever you care) is fine. Just no teamwork please, and help from anyone else outside class.

You should plan around 5 hours for working through the problems (w/o implementation). Actual implementation (item (f)) might take longer. However, there is **no formal time limit**.

### Likelihood (sampling model):

Let  $x_i$ ,  $i = 1, \dots, n$ , denote a set of multinomial random variables, with outcomes  $x_i \in \{1, \dots, N\}$ . Let  $y = (y_1, \dots, y_N)$  denote a contingency table summarizing the multinomial experiment, i.e.,  $y_j$  is the frequency of outcome  $j$ . Let  $\pi_j = Pr(x_i = j)$  denote the unknown probability of observing outcome  $j$ ,  $\sum \pi_j = 1.0$ . We write

$$y \sim Mn(n; \pi_1, \dots, \pi_N)$$

### Prior:

Due to the nature of the experiment we believe *a priori* that some of the  $\pi_j$  are much larger than others. We refer to this subset of outcomes with much larger probability as the “prevalent” outcomes  $A_1$ , and to the set of not so likley outcomes as “rare” outcomes  $A_0$ . We will formally define  $A_0$  and  $A_1$  below.

*Parametrization:* To define a prior probability model for  $\theta = (\pi_1, \dots, \pi_N)$  it is convenient to introduce latent indicators  $z_i$ , and a change of variables for the  $\pi_j$ :

$$\pi_j = \begin{cases} \pi^* q_j & \text{if } z_j = 1 \\ (1 - \pi^*) r_j & \text{if } z_j = 0, \end{cases} \quad (1)$$

with  $\sum q_j = 1$  and  $\sum r_j = 1$  (let  $q_j = 0$  if  $z_j = 0$ , and  $r_j = 0$  if  $z_j = 1$ ). In words, the latent variable  $z_j$  is an indicator for outcome  $j$  being a prevalent outcome, i.e.,  $A_1 = \{j : z_j = 1\}$  and  $A_0 = \{j : z_j = 0\}$ . The probability of observing some prevalent outcome is  $\pi^*$ , with  $\pi^*$  close to 1.0;  $q_j$  is the probability of outcome  $j$  given that we observe a prevalent outcome, and  $r_j$  is the probability of  $j$  given a rare outcome. For later reference we define

$$M_1 = \#A_1 \text{ and } M_0 = N - M_1$$

as the number of prevalent and rare outcomes, respectively.

*Prior Probability Model  $p(z, \pi^*, q, r)$ :* We assume

$$Pr(z_j = 1) = \rho, \quad (2)$$

a beta prior on the total probability of prevalent outcomes:

$$\pi^* \sim Be(a^*, b^*), \quad (3)$$

and a Dirichlet prior for the partitioning of  $\pi^*$  into the cell probabilities  $\pi_j$ ,  $j \in A_1$ . Let  $\tilde{q}_h$ ,  $h = 1, \dots, M_1$  denote the non-zero weights  $q_j$ .

$$(\tilde{q}_1, \dots, \tilde{q}_{M_1}) \sim Dir(M_1; a_1, \dots, a_1). \quad (4)$$

The use of equal Dirichlet parameters  $a_1$  reflects the fact that prior believes about  $(q_1, \dots, q_{M_1})$  are *a priori* exchangeable. Similarly for  $r_j$ ,  $j \in A_0$ .

$$(\tilde{r}_1, \dots, \tilde{r}_{M_0}) \sim Dir(M_0; a_0, \dots, a_0). \quad (5)$$

## Hyperparameters:

The hyperparameters  $\rho, a^*, b^*, a_1, a_0$  are fixed. Use, for example,

$$\rho = 0.1, a^* = 9, b^* = 1, a_1 = 0.1 \text{ and } a_0 = 10.$$

## Data

posted on the canvas page as *sage.dta*, together with the pdf file of this exam. The file gives the observed values  $x_i \in \{0, N\}$ ,  $i = 1, \dots, n$ , with  $n = 700$  and  $N = 77$ .

```
22 48 54 96 96 94 96 22 22 102 99 33 87 96 99 96 26 28 26 117
53 116 57 28 40 28 33 96 26 33 96 98 53 102 96 33 96 952 54 96
102 33 102 33 28 85 96 33 102 85 96 25 102 33 26 43 96 2 102 87
96 94 33 96 96 87 28 28 18 54 96 102 102 96 96 96 96 85 33 85
18 33 96 84 83 85 96 57 99 89 26 96 102 96 96 33 96 28 631 748
33 85 25 85 7 96 67 33 85 120 96 22 96 85 33 35 33 96 102 54
26 96 43 22 43 28 98 948 39 33 102 54 33 102 102 85 96 96 96 96
99 48 96 96 85 96 33 96 26 28 102 96 87 33 99 96 28 2 102 22
102 87 102 26 85 57 22 94 102 75 85 28 96 117 85 33 48 28 98 28
35 28 85 33 96 54 28 487 96 503 102 85 33 54 28 96 618 33 98 96
182 96 26 96 94 112 87 96 26 28 397 87 87 99 102 96 33 22 96 57
18 94 28 85 443 28 112 85 102 85 102 87 22 636 85 2 96 96 43 75
22 85 96 96 96 96 33 33 813 854 33 96 87 85 85 96 85 85 33 84
85 102 96 348 420 748 96 22 87 33 96 96 32 96 96 28 102 85 28 54
96 33 98 28 26 35 33 96 96 85 85 102 102 87 40 96 22 33 85 96
85 85 96 85 102 33 96 393 96 28 118 96 120 26 87 85 102 85 96 85
85 87 96 102 96 28 96 85 102 85 96 96 98 26 96 33 33 28 85 118
33 42 39 33 28 26 28 96 102 369 85 102 102 96 96 96 28 117 94 28
96 96 85 85 54 28 96 22 102 102 33 85 112 96 33 87 85 28 102 96
85 30 43 102 33 33 96 960 96 33 118 99 18 28 53 98 33 54 96 96
96 98 113 33 116 22 87 85 96 96 96 26 102 96 22 96 221 71 96 57
96 57 39 96 32 57 96 87 102 85 28 28 102 102 96 94 33 33 96 85
102 54 96 96 85 85 96 43 96 33 102 33 85 102 96 534 33 57 98 28
33 33 33 26 102 96 33 102 26 87 85 33 87 630 26 94 87 958 102 85
96 28 96 96 118 102 116 33 53 102 26 102 882 22 33 102 28 118 83 96
94 96 33 116 102 96 117 96 96 33 26 22 85 85 85 54 32 22 99 723
30 26 22 98 28 85 85 96 26 33 87 28 33 96 28 161 26 85 102 54
28 96 85 102 550 18 28 96 96 96 96 85 85 96 49 28 28 33 96 102
102 85 33 71 96 33 33 116 94 102 85 102 43 689 28 102 102 26 96 366
347 96 96 7 85 96 96 26 96 102 28 40 85 96 94 28 85 96 28 96
102 28 85 87 85 85 87 96 96 28 57 28 85 32 96 96 28 85 587 96
102 85 96 54 28 102 163 87 96 26 96 28 85 96 85 102 22 96 102 87
898 102 54 87 28 33 96 39 57 846 326 96 28 94 22 96 33 33 40 54
48 96 94 96 116 118 28 939 96 96 116 96 28 87 864 102 54 33 96 102
28 33 33 96 33 85 57 26 28 22 121 96 85 102 680 28 87 85 96 102
```

(a)[2pts] **Joint probability model.** Write out the joint probability model

$$p(\pi^*, q_1, r_1, \dots, q_N, r_N, z_1, \dots, z_N, y_1, \dots, y_N).$$

**(b)[2pts] Graphical Model.** Show a graphical model representation of the probability model. Use circles (or whatever shape you like) for each r.v., and connect any two random variables that are *not* conditionally independent given all other variables.<sup>1</sup>

**(c)[4pts] Conditional posterior distributions.** Find the complete conditional posterior distributions

1.  $p(z_i | \dots)$
2.  $p(\pi^* | \dots)$
3.  $p(q | \dots)$
4.  $p(r | \dots)$

Here  $\dots$  denotes “all other parameters and the data  $y$ .”

**(d)[2pts] Marginalizing w.r.t.  $q$  and  $r$ :** Find the posterior distribution  $p(z | \pi^*, y)$ , marginalizing w.r.t.  $q$  and  $r$ .

**(e)[4pts] MCMC I.** Consider a Gibbs sampling scheme based on sampling from the complete conditional posterior distributions found in (c), i.e. an MCMC scheme with steps:

1.  $z_i^{t+1} \sim p(z_i | \dots)$
2.  $\pi^{*(t+1)} \sim p(\pi^* | \dots)$
3.  $q^{t+1} \sim p(q | \dots)$
4.  $r^{t+1} \sim p(r | \dots)$ .

Show that MCMC I violates irreducibility (of course, don't implement MCMC I – it would not work).

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<sup>1</sup>I realize we did not talk much about graphical models in class. Nothing fancy required here - just a representation of conditional independence structure by an undirected graph; or a directed acyclical graph (DAG) for the model (if you know how to do this).

**(f)[4pts] MCMC II.** Implement a Gibbs sampling scheme based on sampling from the conditional posterior distributions found in (d), (c3), (c4) and (c2):

1.  $z_i \sim p(z_i | z_{-i}, \pi^*, y), i = 1, \dots, n$
2.  $q \sim p(q | \dots)$
3.  $r \sim p(r | \dots)$ .
4.  $\pi^* \sim p(\pi^* | \dots)$

Use a convergence diagnostic of your choice to determine when to stop. In your report, please discuss and show:

- (a) Trajectories of simulated values  $\pi^{*(t)}$  against iteration  $t$ .
- (b) Convergence diagnostic used.
- (c) Boxplot of the (marginal) posterior distributions  $p(\pi_j | y)$ . Use one figure with multiple boxplots (using, for example, the R command `boxplot(.)`).
- (d) Plot posterior means  $E(\pi_j | y)$  against the m.l.e.'s  $\hat{\pi}_j = y_j/n$ . Include the 45 degree line and discuss the shrinkage pattern you (should) see.

**(g)[2pts] MCMC III.** Consider the following MCMC scheme. We describe the MCMC by constructive definition of the transition probability, i.e.,  $p(\theta^{t+1} | \theta^t)$ :

1. Metropolis-Hastings step to change  $z_i, q, r, i = 1, \dots, N$ .

- Generate a proposal  $(z'_i, q', r')$ :<sup>2</sup>

$$Pr(z'_i = 1) = 0.5, q' \sim p(q | z', \dots, y), \text{ and } r' \sim p(r | z', \dots, y),$$

where  $z'$  is the currently imputed  $z$ , with  $z_i$  replaced by  $z'_i$ , i.e.,  $z' = (z_1, \dots, z_{i-1}, z'_i, z_{i+1}, \dots, z_N)$ .

- Compute an appropriate acceptance probability  $\alpha$ .
- Set

$$(z_i, q, r) = \begin{cases} (z'_i, q', r') & \text{with prob } \alpha \\ (z_i, q, r) & \text{with prob } 1 - \alpha \end{cases}$$

2.  $\pi^* \sim p(\pi^* | \dots)$

Find the correct expression for the acceptance probability  $\alpha$  (in step 1.).

*No need to implement MCMC III.*

**(z) Mystery:** Can anyone guess what the application behind this probability model could be.

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<sup>2</sup>We use  $q'$  to mark proposals, since we already used the traditional  $\tilde{q}$  for the non-zero  $q_i$ , before.

## R Code

Here is some fragment R code (also on canvas), to get you started..

```
require(gtools) # I found a Dir r.v. generator in this package
## if needed install it with
## install.packages("gtools")

## DATA
x <- scan("sage.dta")          # raw data
y <- table(x)                  # counts
N <- length(y)
names(y) <- 1:N
n <- length(x)

## HYPERPARS:
rho <- 0.1
as <- 0.9
bs <- 0.1
a1 <- .1
a0 <- 10

## initialize
## this function creates a list with
## z=(z1,.. zN); pis=pi*, r=(r~[1],..r~[M0])
## q=(q~[1],..q~[M1])
## Youc an use it to initialize the state of the MC
init <- function()
{ # initialize parameters
  ## z
  z <- ifelse(y<10,0,1)
  ## pi*: empirical frquency
  A0 <- which(z==0);   A1 <- which(z==1)
  M0 <- length(A0);    M1 <- length(A1)
  Y0 <- sum(y[A0]);    Y1 <- sum(y[A1])
  pis <- sum(Y1)/n
  ## r and q: empirical fequencies
  q <- y[A1]/Y1 # this is the q~ of the text
  r <- y[A0]/Y0 # this is the r~ of the text
  return(th=list(z=z,pis=pis,r=r,q=q))
}

## main function for MCMC
gibbs <- function(n.iter=100, verbose=0)
{
  TH <- NULL # initialize - will save pi*,z here
  ## for each iteration
  PI <- NULL # similar - will save (pi1,.., piN) here
  th <- init()
  pis <- th$pis # initialize pis = pi*
  z <- th$z     # initialize z
  for(it in 1:n.iter){ # loop over iterations
    z <- sample.z(pis, z) # 1. z ~ p(z | pis, y)
    q <- sample.q(pis,z)  # 2. q ~ p(q | pis,z,y)
    r <- sample.r(pis,z)  # 3. r ~ p(r | pis,z,y)
    pis <- sample.pis(z)  # 4. pi
    if (verbose > 0){
      if (it %% 10 ==0) # print short summary
        prt.summary(z,q,r,pis)
    }
    ## save iteration
    TH <- rbind(TH, c(pis,z))
  }
}
```

```

        pi <- rep(0,N)
        pi[z==1] <- pis*q
        pi[z==0] <- (1-pis)*r
        PI <- rbind(PI, pi)
    }
    return(list(TH=TH, PI=PI))
}

## run the MCMC :-)
ex <- function()
{ ## RUN these lines to get the plots
  n.iter <- 500
  gbs <- gibbs(n.iter)
  ## assume gbs returns a list with elements
  ## TH = (niter x p) matrix with each row being the
  ##       state (pi, z)
  ## PI = (niter x 1) vector with pi
  TH <- gbs$TH
  PI <- gbs$PI
  its <- 1:n.iter

  ## trajectory plot
  plot(its, TH[,1],xlab="ITER",ylab="PI*",bty="l",type="l")

  ## boxplot
  boxplot(log(PI))

  ## plotting posterior means vs. mle's
  pibar <- apply(PI,2,mean) # posterior means
  pihat <- as.numeric(y)/n
  plot(pihat, pibar, type="p",
       pch=19, bty="l",xlab="MLE pihat", ylab="E(pi | y)")
  abline(0,1)

  ## same thing, zoom in to left lower corner
  plot(pihat, pibar, type="p", xlim=c(0,0.03), ylim=c(0,0.03),
       pch=19, bty="l",xlab="MLE pihat", ylab="E(pi | y)")
  abline(0,1)
}

#####
## aux functions

... ah the rest you have to do :-). Write functions sample.z(), sample.q(), sample.r() and sample.pis() to carry
out one step of the Gibbs sampler.

```