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882 Homework 4

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#### Problem 1:

1) Consider a generalized linear model framework, where the response variables  $Y_i$  have distribution within the exponential family, i.e.

$$f(y_i \mid \theta_i) = \exp\left\{\frac{y_i\theta_i - b(\theta_i)}{\phi} + c(y_i, \phi)\right\}$$

where the scale parameter  $\phi$  is assumed to be known and  $\theta_i$  is some function of the regressors and their coefficients which is specified by the data model and the link function. Denote the true mean of the responses as  $\mu_i = E[y_i \mid \theta_i]$ . The link function g relates the covariates to the mean by

$$g(\mu_i) = \mathbf{X}'_i \boldsymbol{\beta}$$

where  $\mathbf{X}'_i$  denotes a *p*-dimensional vector of covariates for the *i*th observation. In order perform model fitting, we complete the model specification by placing a normal prior distribution on the regression coefficients  $\boldsymbol{\beta}$ , i.e. *apriori* 

$$\boldsymbol{\beta} \sim N(\mathbf{a}, \mathbf{R}).$$

By specifying the prior and likelihood in this fashion, the posterior distribution is

$$f(\boldsymbol{\beta} \mid \mathbf{Y}, \mathbf{X}) \propto \exp\left\{-\frac{1}{2}(\boldsymbol{\beta} - \mathbf{a})'\mathbf{R}^{-1}(\boldsymbol{\beta} - \mathbf{a}) + \sum_{i=1}^{n} \frac{y_i\theta_i - b(\theta_i)}{\phi}\right\}.$$

The idea is to approximate this posterior distribution with a normal distribution that will be used for the proposal distribution. To achieve this, a second order Taylor expansion of the likelihood term

$$\ell(\boldsymbol{\beta}) = \sum_{i=1}^{n} \frac{y_i \theta_i - b(\theta_i)}{\phi}$$

is carried out around some value of  $\beta$ , say  $\beta^{(t-1)}$ , in order to combine with the prior term. The result is a normal distribution with mean vector

$$\mathbf{m}^{(t)} = \left(\mathbf{R}^{-1} + \frac{1}{\phi}\mathbf{X}'\mathbf{W}(\boldsymbol{\beta}^{(t-1)})\mathbf{X}\right)^{-1} \times \left(\mathbf{R}^{-1}\mathbf{a} + \frac{1}{\phi}\mathbf{X}'\mathbf{W}(\boldsymbol{\beta}^{(t-1)})\mathbf{\tilde{y}}(\boldsymbol{\beta}^{(t-1)})\right)$$

and covariance matrix

$$\mathbf{C}^{(t)} = \left(\mathbf{R}^{-1} + \frac{1}{\phi}\mathbf{X}'\mathbf{W}(\boldsymbol{\beta}^{(t-1)})\mathbf{X}\right)^{-1},$$

where  $\mathbf{W}(\boldsymbol{\beta}^{(t-1)})$  is a diagonal weight matrix with entries

$$W_{ii}(\boldsymbol{\beta}^{(t-1)}) = \frac{1}{b''(\theta_i)g'(\mu_i)^2}$$

and  $\widetilde{\mathbf{y}}(\boldsymbol{\beta}^{(t-1)})$  is a vector transformed observations with entries

$$\widetilde{y}_i(\boldsymbol{\beta}^{(t-1)}) = \mathbf{X}'_i \boldsymbol{\beta} + (y_i - \mu_i)g'(\mu_i)$$

Then,  $J_{\beta^{(t)}} = N(\mathbf{m}^{(t)}, \mathbf{C}^{(t)})$  approximates the true posterior distribution and thus leads to high acceptance ratios in a metropolis-hastings algorithm. This method is summarized as follows:

- 1. Initialize  $\beta^{(0)}$  and set t = 1;
- 2. Propose  $\beta^{\star}$  from  $J_{\beta^{(t)}}$  and accept it with probability  $p = \min\{r, 1\}$ , where

$$r = \frac{f(\boldsymbol{\beta}^{\star} \mid \mathbf{Y}, \mathbf{X})}{f(\boldsymbol{\beta}^{(t)} \mid \mathbf{Y}, \mathbf{X})} \frac{J_{\boldsymbol{\beta}^{\star}}(\boldsymbol{\beta}^{(t)})}{J_{\boldsymbol{\beta}^{(t)}}(\boldsymbol{\beta}^{\star})};$$

3. Increment t and return to step 2.

The construction of the proposal parameters  $\mathbf{m}^{(t)}$  and  $\mathbf{C}^{(t)}$  approximates the posterior mode and covariance matrix for  $\boldsymbol{\beta}$ . Now, we describe this approach under a logistic regression model.

2) Consider the following logistic data model

$$y_i \sim Bin(\pi_i, 1)$$
  
 $\log\left(\frac{\pi_i}{1 - \pi_i}\right) = \beta_0 + \beta_1 x_i.$ 

First, the distribution of the observed data is a member of the exponential family since

$$f(y_i \mid \pi_i) = \pi_i^{y_i} (1 - \pi_i)^{1 - y_i} = \left(\frac{\pi}{1 - \pi_i}\right)^{y_i} (1 - \pi_i)$$
$$= \exp\left\{y_i \log \frac{\pi_i}{1 - \pi_i} + \log(1 - \pi_i)\right\}.$$

Therefore, we see that  $\theta_i = \log \frac{\pi_i}{1 - \pi_i}$ ,  $b(\theta_i) = \log(1 + e^{\theta_i})$ , and  $\phi = 1$ . For notational ease,

$$\boldsymbol{\beta} = (\beta_0, \beta_1)', \quad \mathbf{Y} = (y_1, ..., y_n)', \text{ and } \mathbf{X} = (\mathbf{1}_n, (x_1, ..., x_n)').$$

Placing a normal prior distribution on  $\beta$ , the posterior distribution is

$$f(\boldsymbol{\beta} \mid \mathbf{Y}, \mathbf{X}) \propto \prod_{i=1}^{n} f(y_i \mid \pi_i) f(\boldsymbol{\beta}) \propto \exp\left\{-\frac{1}{2}(\boldsymbol{\beta} - \mathbf{a})' \mathbf{R}^{-1}(\boldsymbol{\beta} - \mathbf{a}) + \sum_{i=1}^{n} y_i \log \frac{\pi_i}{1 - \pi_i} + \sum_{i=1}^{n} \log(1 - \pi_i)\right\}$$

where  $\pi_i$  is obtained by inverting the link function, i.e.

$$\pi_i = \frac{\exp\{\beta_0 + \beta_1 x_i\}}{1 + \exp\{\beta_0 + \beta_1 x_i\}}.$$

Now we compute the necessary components for the proposal distribution, namely

$$b''(\theta_i) = \frac{e^{\theta_i}}{\left(1 + e^{\theta_i}\right)^2}$$
$$g'(\mu_i) = \frac{1}{\pi_i(1 - \pi_i)} \qquad \text{since } \mu_i = \pi_i.$$

It now follows that the diagonal weight matrix  $\mathbf{W}(\boldsymbol{\beta}^{(t-1)})$  has entries

$$W_{ii}(\boldsymbol{\beta}^{(t-1)}) = \pi_i(1-\pi_i)$$

and the transformed observations take the form

$$\widetilde{y}_i(\boldsymbol{\beta}^{(t-1)}) = \beta_0 + \beta_1 x_i + \frac{y_i - \pi_i}{\pi_i(1 - \pi_i)}.$$

Finally, the proposal distribution in the metropolis-hastings algorithm is  $N(\mathbf{m}^{(t)}, \mathbf{C}^{(t)})$ , where

$$\mathbf{m}^{(t)} = \mathbf{C}^{(t)} \times \left( \mathbf{R}^{-1} \mathbf{a} + \mathbf{X}' \mathbf{W} (\boldsymbol{\beta}^{(t-1)}) \widetilde{\mathbf{y}} (\boldsymbol{\beta}^{(t-1)}) \right)$$
$$\mathbf{C}^{(t)} = \left( \mathbf{R}^{-1} + \mathbf{X}' \mathbf{W} (\boldsymbol{\beta}^{(t-1)}) \mathbf{X} \right)^{-1}.$$

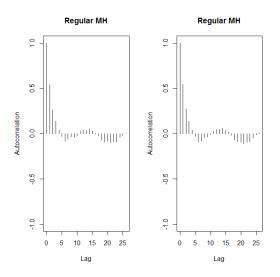
Now the full algorithm summarized in part 1 of this problem can be implemented.

- 3) The code can be found in the appendix.
- 4) The estimates from both algorithms can be found below:

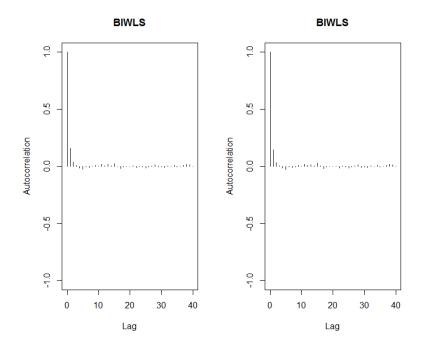
Parameter	BIWLS	Regular MH
$\beta_0$	-4.055	-4.031
$\beta_1$	0.100	0.099

Therefore, both algorithms seem to approximate the parameters in a similar fashion. We will analyze these two approaches further. Firstly, the autocorrelation using a regular MH

algorithm is:



which is the autocorrelation plot for  $\beta_0$  and  $\beta_1$ , respectively. The autocorrelation plot using the BIWLS algorithm is



for  $\beta_0$  and  $\beta_1$ , respectively. The autocorrelation for the BIWLS is far less, and this was even without any thinning. Lastly, the effective sample size for  $\beta_0$  was 7270 out of 10000 samples and for  $\beta_1$  was 7898 out of 10000 samples with the BIWLS. Comparatively, the effective sample size was 311 for  $\beta_0$  and 305 for  $\beta_1$  out of 1000 samples, which also is not as good as the BIWLS method. The last note worthy thing to mention is the acceptance rate for ( $\beta_0, \beta_1$ ) was about 95% while the acceptance rate for  $\beta_0$  and  $\beta_1$  under the regular MH algorithm was about 24% each.

## Problem 2:

1) The idea of a conditional means prior (CMP) is with p predictors, i.e. p parameters, we include our prior information to construct an informative prior distribution to be used in a generalized linear model framework. This is typical tough to do in GLMs. In a GLM, we have

$$m_i := E[Y_i \mid \mathbf{X}_i] = g^{-1}(\mathbf{X}'_i \boldsymbol{\beta})$$

where g is a specified link function that relates the covariates to the mean response. The informative prior for  $\beta$  is induced by a CMP on the new mean vector  $\tilde{\mathbf{m}} = (\tilde{m}_1, ..., \tilde{m}_n)$ , where

$$\widetilde{m}_i = E[\widetilde{Y}_i \mid \widetilde{\mathbf{X}}_i]$$

where  $\widetilde{Y}_i$  is some observable response at covariate  $\widetilde{\mathbf{X}}_i$ , where the selected p covariate vectors  $\widetilde{\mathbf{X}}_i$  are linearly independent. Now, for each new mean  $\widetilde{m}_i$ , elicit the independent prior  $\pi(\widetilde{m}_i)$ . Then, the CMP is given by

$$\pi(\widetilde{\mathbf{m}}) = \prod_{i=1}^{p} \pi(\widetilde{m}_i).$$

Define  $\mathbf{G}(\widetilde{\mathbf{m}})$  that applies g to each element, i.e. entries  $g(\widetilde{m}_i)$ . The link function gives the relationship  $\boldsymbol{\beta} = \widetilde{\mathbf{X}}^{-1} \mathbf{G}(\widetilde{\mathbf{m}})$ , and so through a transformation the induced prior is

$$\pi(\boldsymbol{\beta}) \propto \prod_{i=1}^{p} \pi(\widetilde{m}_{i})/g'(\widetilde{m}_{i})$$
$$\propto \prod_{i=1}^{p} \pi\left(g^{-1}(\widetilde{\mathbf{X}}_{i}\boldsymbol{\beta})\right)/g'\left(g^{-1}(\widetilde{\mathbf{X}}_{i}\boldsymbol{\beta})\right)$$

2) Now we outline the details to develop the CMP for a logistic regression model. Consider the model of the form

$$Y_i \sim Bin(m_i, 1)$$
$$g(m_i) = \log\left(\frac{m_i}{1 - m_i}\right) = \mathbf{X}_i \boldsymbol{\beta}$$

where

$$m_i = E[Y_i \mid \mathbf{X}_i] = g^{-1}(\mathbf{X}_i \boldsymbol{\beta}) = \frac{\exp\{\mathbf{X}_i \boldsymbol{\beta}\}}{1 + \exp\{\mathbf{X}_i \boldsymbol{\beta}\}}$$

which is a logistic regression model. Notice that

$$g'(m_i) = m_i^{-1}(1-m_i)^{-1} \quad \Longrightarrow \quad g'\left(g^{-1}\left(\widetilde{\mathbf{X}}_i\boldsymbol{\beta}\right)\right) = \frac{\left(1+\exp\{\mathbf{X}_i\boldsymbol{\beta}\}\right)^2}{\exp\{\mathbf{X}_i\boldsymbol{\beta}\}}.$$

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Then, once specifying the independent priors  $\pi(\tilde{m}_i)$ , the induced prior is

$$\pi(\boldsymbol{\beta}) \propto \prod_{i=1}^{p} \pi\left(g^{-1}\left(\widetilde{\mathbf{X}}_{i}\boldsymbol{\beta}\right)\right) \frac{\exp\{\mathbf{X}_{i}\boldsymbol{\beta}\}}{\left(1 + \exp\{\mathbf{X}_{i}\boldsymbol{\beta}\}\right)^{2}}.$$

In logistic regression, a reasonable choice for the independent priors are

$$\widetilde{m}_i \sim \text{Beta}(a_i, b_i)$$

and therefore the induced prior for the regression coefficients is

$$\pi(\boldsymbol{\beta}) \propto \prod_{i=1}^{p} \left( g^{-1}(\widetilde{\mathbf{X}}_{i}\boldsymbol{\beta}) \right)^{a_{i}-1} \left( 1 - g^{-1}(\widetilde{\mathbf{X}}_{i}\boldsymbol{\beta}) \right)^{b_{i}-1} \frac{\exp\{\mathbf{X}_{i}\boldsymbol{\beta}\}}{\left( 1 + \exp\{\mathbf{X}_{i}\boldsymbol{\beta}\} \right)^{2}}$$
$$\propto \prod_{i=1}^{p} \left( g^{-1}(\widetilde{\mathbf{X}}_{i}\boldsymbol{\beta}) \right)^{a_{i}} \left( 1 - g^{-1}(\widetilde{\mathbf{X}}_{i}\boldsymbol{\beta}) \right)^{b_{i}}.$$

3) Now we develop a sampling strategy to draw posterior samples for  $\beta$  under the CMP prior using the technique discussed in problem 1. Consider a binomial data model of the form

$$\widetilde{Y}_i \sim \operatorname{Bin}(n_i, p_i)$$
  
 $\log\left(\frac{p_i}{1-p_i}\right) = \widetilde{\mathbf{X}}_i \boldsymbol{\beta}.$ 

The likelihood function for a single observation in this situation is given by

$$f(\widetilde{Y}_i \mid \widetilde{\mathbf{X}}_i) \propto p_i^{Y_i} (1-p_i)^{n_i-Y_i}.$$

Comparing this function to the induced prior in part 2), we define the parameters in the following way:

$$a_i = \widetilde{Y}_i$$
 and  $b_i = n_i - \widetilde{Y}_i$ .

Now, for sampling, we implement the method discussed in problem 1. Combine variables as

$$\begin{split} \mathbf{Y}^{\star} &= (\mathbf{Y}, \widetilde{Y}_1, ..., \widetilde{Y}_p)' \\ \mathbf{X}^{\star} &= (\mathbf{X}, \widetilde{\mathbf{X}}_1, ..., \widetilde{\mathbf{X}}_p)' \end{split}$$

where  $\mathbf{Y} = (Y_1, ..., Y_n)$  defined in part 2). Then, using the derivations from problem 1,

$$\mathbf{C} = \left( \mathbf{X}^{\star T} \mathbf{W}(\boldsymbol{\beta}) \mathbf{X}^{\star} \right)^{-1}$$
$$\mathbf{m} = \mathbf{C} \times \left( \mathbf{X}^{\star T} \mathbf{W}(\boldsymbol{\beta}) \overline{\mathbf{Y}}^{\star}(\boldsymbol{\beta}) \right)$$

where  $\overline{\mathbf{Y}}^{\star}(\boldsymbol{\beta})$  is a vector of transformed observations, i.e. transformation of  $\mathbf{Y}^{\star}$ , with entries

$$\begin{split} \overline{Y}_{j}^{\star}(\boldsymbol{\beta}) &= \mathbf{X}_{j}\boldsymbol{\beta} + \frac{Y_{j} - m_{j}}{m_{j}(1 - m_{j})} \quad \text{for } j = 1, ..., n\\ \overline{Y}_{i}^{\star}(\boldsymbol{\beta}) &= \widetilde{\mathbf{X}}_{i}\boldsymbol{\beta} + \frac{\widetilde{Y}_{i} - n_{i}p_{i}}{n_{i}p_{i}(1 - p_{i})} \quad \text{for } i = 1, ..., p \end{split}$$

and  $\mathbf{W}(\boldsymbol{\beta})$  is the diagonal weight matrix with entries

$$W_{ii}(\boldsymbol{\beta}) = n_i p_i (1 - p_i).$$

Then, we use the algorithm discussed in problem 1.

### Problem 3:

Here we develop a metropolis algorithm to sample the  $\rho$  term in the CAR model from problem 2 of homework 3. Namely, consider the spatial regression model given by

$$Y_i = \beta_0 + b_i + \epsilon_i$$

where  $\epsilon_i \stackrel{iid}{\sim} N(0, \sigma^2)$  for i = 1, ..., n. We assume that the spatial random effects follow the  $CAR(\sigma^2 \tau^2, \rho)$  model, i.e.

$$\mathbf{b} = (b_1, ..., b_n)' \sim N(\mathbf{0}, \sigma^2 \tau^2 (\mathbf{D} - \rho \mathbf{W}))$$

It can be shown that  $\rho$  must be between -1 and 1, and as a result we will assume  $\rho \sim \text{Unif}(-1, 1)$ . Then, the posterior distribution for  $\rho$  is simply

$$\pi(\rho \mid \mathbf{b}, \sigma^2, \tau^2) \propto \exp\left\{-\frac{1}{2\sigma^2\tau^2}\mathbf{b}'(\mathbf{D}-\rho\mathbf{W})\mathbf{b}\right\}.$$

Now, for the proposal distribution, we will consider a symmetric random walk, namely

$$J(\rho \mid \rho^{(t)}) = \text{Unif}(\rho^{(t)} - c, \rho^{(t)} + c)$$

where c is a tuning parameter to achieve an acceptance probability of roughly 35%. Denote  $\theta_{-\rho}$  as the set of parameters to update, excluding  $\rho$ . Then, the algorithm is as follows:

- 1. Initialize  $\boldsymbol{\theta}^{(0)}$  and set t = 1.
- 2. Update  $\boldsymbol{\theta}_{-\rho}^{(t)}$  via the Gibbs sampling scheme of problem 2 of homework 3.
- 3. Propose  $\rho^*$  from  $J(\rho \mid \rho^{(t)})$  and accept it with probability  $p = \min\{r, 1\}$ , where

$$r = \frac{\pi(\rho^{\star} \mid \mathbf{b}, \sigma^2, \tau^2)}{\pi(\rho^{(t)} \mid \mathbf{b}, \sigma^2, \tau^2)}$$

#### APPENDIX

###### Chase Joyner <del>#####</del> <del>||-||-||-||-||-</del> 882 Homework 4 ## Required libraries ## library (MASS) library (mvtnorm) library (coda) ###### Problem 1 BIWLS = function(Y, X, iter = 1e4)## Preliminaries ## n = length(Y) $p = \dim(X)[2]$ R = 100 \* diag(p)a = rep(1, p)## Initial values ## beta = rep(0, p)acc = 0## Save records ## Beta = matrix (-99, nrow = iter, ncol = p)## Compute only once ## IR = solve(R)IRa = IR %%a ## Link function for pi ## link = function(u) $\exp XB = \exp (X \% \% u)$ val = expXB / (1 + expXB)return(val) } llik = function(s)

```
pi = link(s)
                return (val)
        }
        for(i in 1:iter){
                ## Update beta ##
                pi.t = as.vector(link(beta))
                newY.t = X \%*% beta + (Y - pi.t) / (pi.t * (1 - pi.t))
                Wt = diag(pi.t * (1 - pi.t))
                Ct = solve(IR + t(X) \% Wt \% X)
                mt = Ct \% \% (IRa + t(X) \% \% Wt \% \% newY.t)
                beta.s = as.vector(rmvnorm(1, mt, Ct))
                pi.s = as.vector(link(beta.s))
                newY.s = X \%*% beta.s + (Y - pi.s) / (pi.s * (1 - pi.s))
                Ws = diag(pi.s * (1 - pi.s))
                Cs = solve(IR + t(X) \% \% Ws \% \% X)
                ms = Cs \%*\% (IRa + t(X) \%*\% Ws \%*\% newY.s)
                r = \exp(11ik(beta.s) - 11ik(beta) + dmvnorm(beta, ms, Cs, log = TRI)
                z = rbinom(1, 1, min(r, 1))
                if(z = 1){
                        beta = beta.s
                        acc = acc + 1
                }
                Beta[i,] = beta
                print(c(i))
        }
        return(list(Beta = Beta, accept = acc / iter))
}
\# Generate some data and run \#
\# par(m frow = c(2, 1))
\#n = 1000
\#beta.true = c(-1, 0.5)
\#X = cbind(rep(1, n), rnorm(n, 2, 1))
\# \text{probs} = \exp(X \% \% \text{ beta.true}) / (1 + \exp(X \% \% \text{ beta.true}))
\#Y = rbinom(n, 1, probs)
\#res = BIWLS(Y, X)
#apply(res$Beta, 1, mean)
## Analyze diabetes data ##
library (MASS)
data(Pima.tr)
data (Pima.te)
pima = rbind(Pima.tr, Pima.te)
```

```
Y = as.vector(pima[, 8])
Y[Y == "Yes"] = 1
Y[Y == 'No'] = 0
Y = as.numeric(Y)
X = cbind(1, as.matrix(pima[,5]))
res = BIWLS(Y, X)
apply(res$Beta, 2, mean)
Beta.mcmc1 = as.mcmc(res$Beta[1,])
Beta.mcmc2 = as.mcmc(res$Beta[2,])
autocorr.plot(Beta.mcmc1)
autocorr.plot(Beta.mcmc2)
effectiveSize(Beta.mcmc1)
effectiveSize(Beta.mcmc2)
```