

Introduction to Bayesian Data Analysis

Problem Set 6 - Solutions

- (1) (a) $\theta_B = \theta \times \gamma$ and $p(\theta_B|\theta, \gamma) = p(\theta) \times p(\gamma)$ because θ and γ are independent, and $p(\theta_B|\theta, \gamma) \neq p(\theta_B|\gamma)$. Now $\theta_A = \theta$, and so $p(\theta_B|\theta, \gamma) = p(\theta_B|\theta_A, \gamma) = p(\theta_A) \times p(\gamma) \neq p(\theta_B|\gamma)$, hence θ_B and θ_A are not independent under this prior distribution.

This prior is justified if we believe there exists an underlying rate θ for the population and it is modified by γ for those men that do not have a Bachelor's degree.

$$(b) \quad p(\mathbf{y}_A|\theta) = \prod_{i=1}^{n_A} \frac{e^{-\theta} \theta^{y_{A,i}}}{y_{A,i}!}; \quad p(\mathbf{y}_B|\theta, \gamma) = \prod_{i=1}^{n_B} \frac{e^{-\theta\gamma} (\theta\gamma)^{y_{B,i}}}{y_{B,i}!}$$

$$p(\theta) = \frac{b_\theta^{a_\theta}}{\Gamma(a_\theta)} \theta^{a_\theta-1} e^{-b_\theta\theta}; \quad p(\gamma) = \frac{b_\gamma^{a_\gamma}}{\Gamma(a_\gamma)} \gamma^{a_\gamma-1} e^{-b_\gamma\gamma}$$

$$\text{Note that } p(\theta, \gamma|\mathbf{y}_A, \mathbf{y}_B) = \frac{p(\theta, \gamma, \mathbf{y}_A, \mathbf{y}_B)}{p(\mathbf{y}_A, \mathbf{y}_B)} \propto p(\mathbf{y}_A|\theta) p(\mathbf{y}_B|\theta, \gamma) p(\theta, \gamma);$$

so

$$\begin{aligned} p(\theta|\mathbf{y}_A, \mathbf{y}_B, \gamma) &\propto p(\mathbf{y}_A|\theta) p(\mathbf{y}_B|\theta, \gamma) p(\theta) \\ &\propto \left[\prod_{i=1}^{n_A} e^{-\theta} \theta^{y_{A,i}} \right] \left[\prod_{i=1}^{n_B} e^{-\theta\gamma} (\theta\gamma)^{y_{B,i}} \right] \theta^{a_\theta-1} e^{-b_\theta\theta} \\ &\propto e^{-n_A\theta} \theta^{\sum_{i=1}^{n_A} y_{A,i}} e^{-n_B\theta\gamma} (\theta\gamma)^{\sum_{i=1}^{n_B} y_{B,i}} \theta^{a_\theta-1} e^{-b_\theta\theta} \\ &\propto e^{-(n_A+n_B\gamma+b_\theta)\theta} \theta^{n_A\bar{y}_A+n_B\bar{y}_B+a_\theta-1} \end{aligned}$$

Therefore, $\theta|\mathbf{y}_A, \mathbf{y}_B, \gamma \sim \text{gamma}(n_A\bar{y}_A + n_B\bar{y}_B + a_\theta, n_A + n_B\gamma + b_\theta)$

(c)

$$\begin{aligned}
 p(\gamma|\mathbf{y}_A, \mathbf{y}_B, \theta) &\propto p(\mathbf{y}_B|\theta, \gamma)p(\gamma) \\
 &\propto \left[\prod_{i=1}^{n_B} e^{-\theta\gamma} (\theta\gamma)^{y_{B,i}} \right] \gamma^{a_\gamma-1} e^{-b_\gamma\gamma} \\
 &\propto e^{-(n_B\theta+b_\gamma)\gamma} \gamma^{n_B\bar{y}_B+a_\gamma-1}
 \end{aligned}$$

Therefore, $\gamma|\mathbf{y}_A, \mathbf{y}_B, \theta \sim \text{gamma}(n_B\bar{y}_B + a_\gamma, n_B\theta + b_\gamma)$

(d) `a_theta<-2`

`b_theta<-1`

`a_gamma<-c(8,16,32,64,128)`

`yA <- read.table("menchild30bach.txt", header = F)`

`yA<-yA[,1]`

`nA<-length(yA)`

`ybarA<-mean(yA)`

`yB <- read.table("menchild30nobach.txt", header = F)`

`yB<-yB[,1]`

`nB<-length(yB)`

`ybarB<-mean(yB)`

`#Gibbs sampling`

`S<-5000`

`PHI<-matrix(nrow=S,ncol=2)`

`PHI[1,]<-phi<-c(ybarA ,ybarB/ybarA)`

`mean.B.M.A<-NULL`

`for (i in 1:5){`

`b_gamma<-a_gamma[i]`

`for(s in 2:S) {`

`# generate a new theta value from its full conditional`

`phi[1]<-rgamma(1,nA*ybarA+nB * ybarB+a_theta,nA+nB*phi[2]+b_theta)`

`# generate a new gamma value from its full conditional`

`phi[2]<- rgamma(1, nB*ybarB+a_gamma[i], nB*phi[1]+b_gamma)`

`PHI[s,]<-phi }`

```
mean.B.M.A<-c(mean.B.M.A,mean(PHI[,2]*PHI[,1]-PHI[,1]))
}
```

```
> mean.B.M.A
```

```
[1] 0.3780609 0.3355366 0.2714495 0.2016046 0.1324830
```

As the hyper-parameters in the prior for gamma increase, the mean difference of $(\theta_B - \theta_A | \mathbf{y}_A, \mathbf{y}_B)$ gets smaller.

```
> effectiveSize(PHI[,1])
```

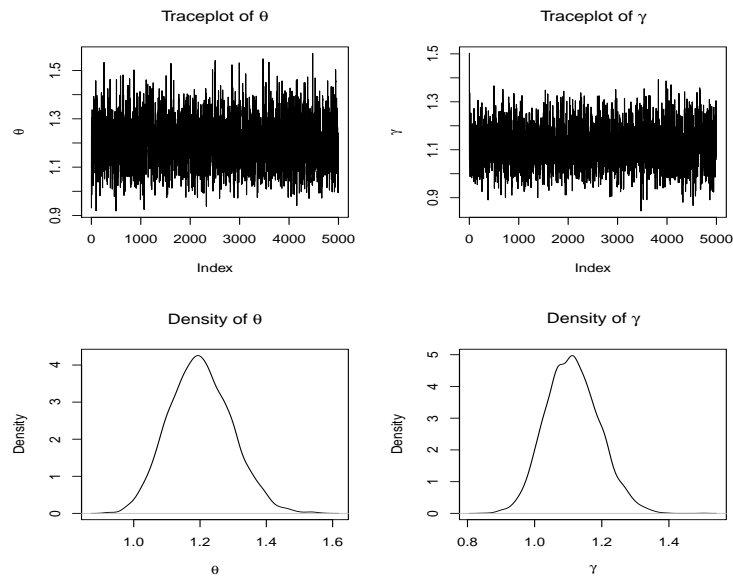
```
var1
```

```
1760.03
```

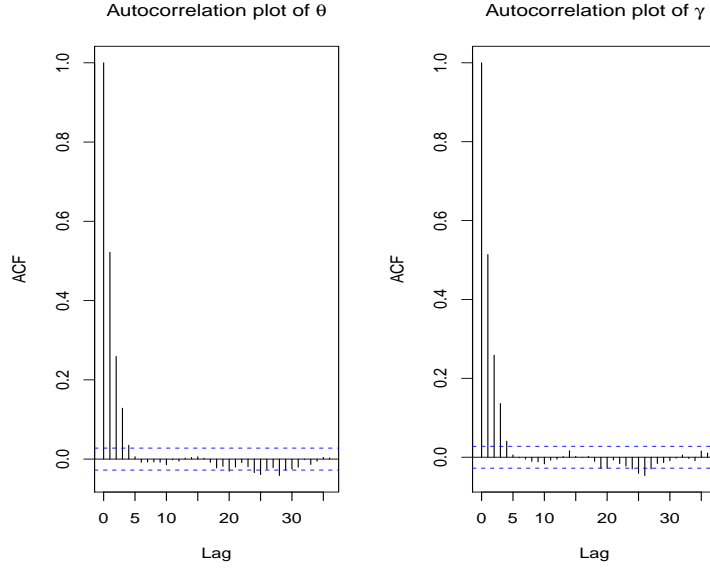
```
> effectiveSize(PHI[,2])
```

```
var1
```

```
1741.798
```



The effective sample sizes are large, the traceplots show the sequence of parameter draws randomly moves around the parameter space, and the autocorrelation plots do not show high autocorrelations to be concerned about.



(2) (a) The empirical distribution appears slightly skewed to the right.

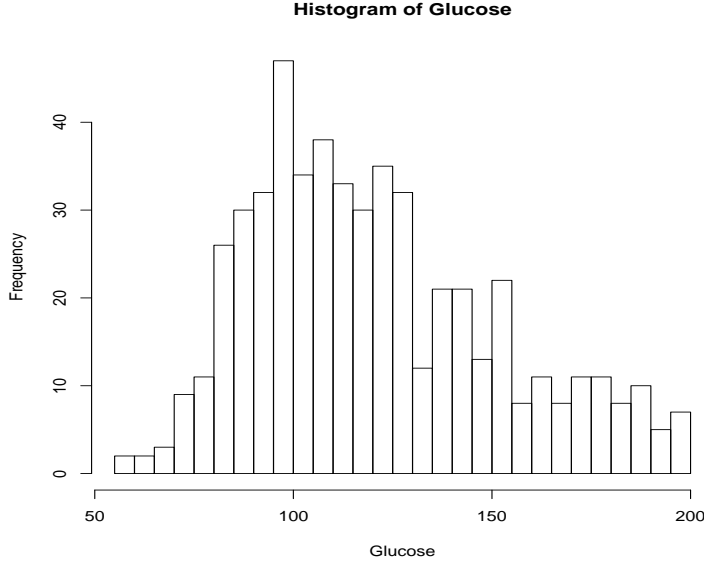
(b) $f(X_1, \dots, X_n, p, \theta_1, \theta_2, \sigma_1^2, \sigma_2^2 | \mathbf{y}) \propto$
 $f(\mathbf{y} | X_1, \dots, X_n, p, \theta_1, \theta_2, \sigma_1^2, \sigma_2^2) f(X_1, \dots, X_n | p) f(p) f(\theta_1) f(\theta_2) f(\sigma_1^2) f(\sigma_2^2).$

Let $\phi = (\theta_1, \theta_2, \sigma_1^2, \sigma_2^2)$. Let $X = (X_1, \dots, X_n)$

The Gibbs sampler cycle is:

1. For $i = 1, \dots, n$ update $X_i \sim f(y_i | \phi, y_i) f(X_i | p)$ (conditional independence)
2. Update $p \sim f(p | X)$
3. For $k = 1, 2$, update $\theta_k \sim f(\theta_k | X, p, \theta_{-k}, \sigma_1^2, \sigma_2^2)$
4. For $k = 1, 2$, update $\sigma_k^2 \sim f(\sigma_k^2 | X, p, \theta_1, \theta_2, \sigma_{-k}^2)$

The joint posterior distribution is



$$\begin{aligned}
f(X, \phi | \mathbf{y}) &\propto \left(\prod_{i=1}^n f(y_i | \phi, x_i) f(x_i | p) \right) f(p) f(\phi) \\
&\propto \left(\prod_{i=1}^n \frac{1}{\sqrt{\sigma_{x_i}^2}} \exp \left(-\frac{1}{2\sigma_{x_i}^2} (y_i - \theta_{x_i})^2 \right) p^{2-x_i} (1-p)^{x_i-1} \right) \\
&\quad p^{\alpha-1} (1-p)^{\beta-1} \exp \left(-\frac{1}{2\tau_0^2} (\theta_1 - \mu_0)^2 \right) \exp \left(-\frac{1}{2\tau_0^2} (\theta_2 - \mu_0)^2 \right) \\
&\quad (\sigma_1^2)^{-(\nu_0/2+1)} \exp \left(-\frac{\nu_0 \sigma_0^2 / 2}{\sigma_1^2} \right) (\sigma_2^2)^{-(\nu_0/2+1)} \exp \left(-\frac{\nu_0 \sigma_0^2 / 2}{\sigma_2^2} \right)
\end{aligned}$$

The conditional distribution on the class labels:

$$\begin{aligned}
f(x_i | y_i, p, \phi) &= \frac{f(y_i | x_i, \theta, p)}{\sum_{k=1}^2 f(y_i | x_i = k, \theta, p)} \\
&\propto \frac{\frac{1}{\sqrt{\sigma_{x_i}^2}} \exp \left(-\frac{1}{2\sigma_{x_i}^2} (y_i - \theta_{x_i})^2 \right) p^{2-x_i} (1-p)^{x_i-1}}{\sum_{k=1}^2 \frac{1}{\sqrt{\sigma_k^2}} \exp \left(-\frac{1}{2\sigma_k^2} (y_i - \theta_k)^2 \right) p^{2-k} (1-p)^{k-1}}
\end{aligned}$$

The conditional distribution on p :

$p| \sim \text{Beta}(\alpha + 2n - \sum_i x_i, \beta + \sum_i x_i - n)$ The conditional distribution on the group means $(\theta_k, (k = 1, 2))$:

Let n_k be the current size of class k ; $n_k = \# \{i : x_i = k\}$

and so $\theta_k|X, \mathbf{y}, p, \theta_{-k}, \sigma_1^2, \sigma_2^2) \sim N(\mu_k, \sigma_k^2)$ where $\mu_k = \frac{\mu_0/\tau_0^2 + n_k \bar{y}_k/\sigma_k^2}{1/\tau_0^2 + n_k/\sigma_k^2}$

where $\bar{y}_k = \frac{1}{n_k} \sum_{i:x_i=k} y_i$

and $\sigma_k^2|X, \mathbf{y}, p, \sigma_{-k}^2, \theta_1, \theta_2) \sim \text{Inv} - \text{gamma}(\nu_k/2, \nu_k \sigma_k^2(\theta_k)/2)$

where $\nu_k = \nu_0 + n_k$, $\sigma^2(\theta_k) = \frac{1}{\nu_k} [\nu_0 \sigma_0^2 + n_k s_k^2(\theta_k)]$ and

$s_k^2(\theta_k) = \sum_{i:x_i=k} (y_i - \theta_k)^2 / n_k$

(c) `alpha<-1`
`beta<-1`

`mu0<-120`

`tau20<-200`

`sigma20<-1000`

`nu0<-10`

```
mcmc.mix<-function(y, niter=10000, theta1=mean(y), theta2=mean(y),
                    sigma12=var(y), sigma22=var(y)){
  n=length(y); K=2; theta=c(theta1, theta2)
  sigma2=c(sigma12, sigma22); p=0.5; x=NULL; nk=NULL
  p.mcmc=NULL; theta.mcmc=NULL; sigma2.mcmc=NULL; x.mcmc=NULL
  for (nit in 1:niter) {
    for (i in 1:n) {
      p.z<-exp(-(y[i]-theta)^2/2/sigma2)/sqrt(sigma2)*c(p, 1-p)
      x[i]<-sample(1:2, 1, prob=p.z/sum(p.z))
      #print(p.z/sum(p.z))
      x.mcmc=rbind(x.mcmc, x)
      for (k in 1:K){
        nk[k]=sum(x==k)
        muk<-(mu0/tau20+nk[k]*mean(y[x==k])/sigma2[k])/(1/tau20+nk[k]/sigma2[k])
        t2n<-1/(1/tau20+nk[k]/sigma2[k])
        theta[k]<-rnorm(1, muk, sqrt(t2n))
        nuk<-nu0+sum(x==k)
        sk2<-(nu0*sigma20+(nk[k]-1)*var(y[x==k])+nk[k]*
              (mean(y[x==k])-theta[k])^2)/nuk
        sigma2[k]<-rinvgamma(1, nuk/2, nuk*sk2/2)
```

```

}

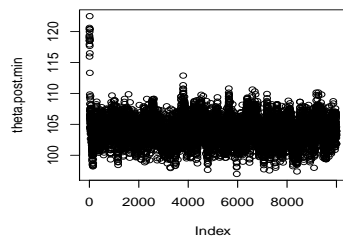
theta.mcmc<-rbind(theta.mcmc,theta)
sigma2.mcmc<-rbind(sigma2.mcmc,sigma2)
p<-rbeta(1,alpha+2*n-sum(x),beta+sum(x)-n)
p.mcmc<-rbind(p.mcmc,p)

}
return(list(x=x.mcmc, theta=theta.mcmc,sigma2=sigma2.mcmc,p=p.mcmc))
}

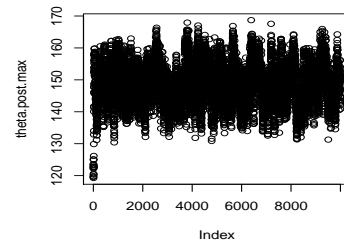
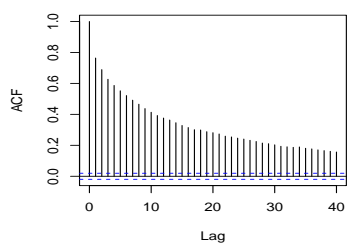
obj<-mcmc.mix(y, niter=10000, theta1=mean(y),theta2=mean(y),
              sigma12=var(y),sigma22=var(y))
> theta.post<-obj$theta
> thin.theta.post.min<-theta.post.min[seq(1,length(theta.post.min),10)]
> thin.theta.post.max<-theta.post.max[seq(1,length(theta.post.max),10)]
> effectiveSize(theta.post.min[9001:10000])
    var1
144.9595
> effectiveSize(theta.post.max[9001:10000])
    var1
45.40116
> effectiveSize(thin.theta.post.min)
    var1
421.3194
> effectiveSize(thin.theta.post.max)
    var1
291.4248

```

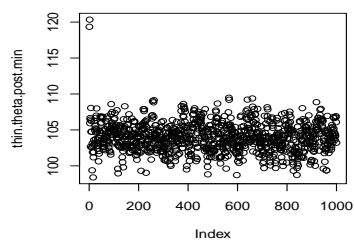
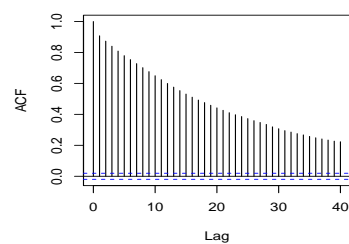
Traceplots of $\theta_{(1)}^{(s)}$ and $\theta_{(2)}^{(s)}$ indicated immediate convergence. However, the autocorrelation plots show a high degree of autocorrelation. If we thin each sequence by taking every tenth draw, the second set of plots show a lower degree of autocorrelation and increased effective sample sizes.



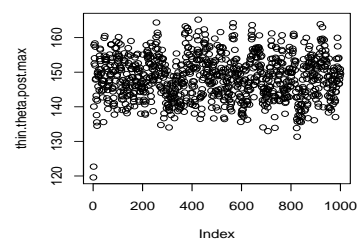
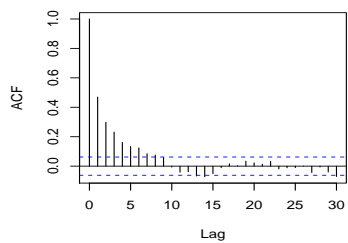
Series theta.post.min



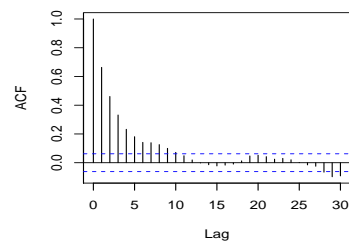
Series theta.post.max



Series thin.theta.post.min



Series thin.theta.post.max



- (d) The histogram of posterior draws of \tilde{Y}_s is quite similar to the empirical data density except that the frequency of higher blood glucose levels is underestimated. Posterior predictive checks on tail-area percentiles show that the empirical tail area probabilities are adequately captured by the two-component mixture model. Hence, we conclude that the two-component mixture model is reasonable for inferential purposes.

```
p.post<-obj$p
sigma.post<-obj$sigma2
x.post<-NULL
for (i in 1:10000){
x.post<-c(x.post,sample(c(1,2),1,prob=c(p.post[i],1-p.post[i])))
}

y.post<-NULL
for (i in 1:10000){
y.post<-c(y.post,rnorm(1,theta.post[i,x.post[i]],
                        sqrt(sigma.post[i,x.post[i]])))
}

#Posterior predictive checks
> result<-NULL
>
> for (i in 1:10000){
+ x<-sample(c(1,2),length(y),prob=c(p.post[i],1-p.post[i]),replace=TRUE)
+ y.s<-rnorm(length(y),theta.post[i,x],sqrt(sigma.post[i,x]))
+ result<-cbind(result,c(min(y.s),quantile(y.s,probs=c(0.75,0.9,0.95,0.975)
+               max(y.s)))
+ }
>

> mean(result[1,]>=min(y))
```

```

[1] 0.2049
> mean(result[2,]>=quantile(y,0.75))
[1] 0.3813
> mean(result[3,]>=quantile(y,0.90))
[1] 0.1458
> mean(result[4,]>=quantile(y,0.95))
[1] 0.318
> mean(result[5,]>=quantile(y,0.975))
[1] 0.5239
> 1-mean(result[6,]>=max(y))
[1] 0.0043
> mean(result[7,]>=mean(y))
[1] 0.4651

```

