STA 360/602L: Module 3.9

MCMC AND GIBBS SAMPLING III

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RECAP OF NORMAL MODEL

Sampling model:

$$y_i | \mu, au \stackrel{iid}{\sim} \mathcal{N}\left(\mu, au^{-1}
ight)$$
 .

- Suppose we want to specify our uncertainties about μ and τ independently of each other. That is, we want $\pi(\mu,\tau)=\pi(\mu)\pi(\tau)$.
- For example,

$$egin{aligned} \mu &\sim \mathcal{N}\left(\mu_0, \sigma_0^2
ight). \ & au &\sim \mathrm{Gamma}\left(rac{
u_0}{2}, rac{
u_0}{2 au_0}
ight). \end{aligned}$$

- Then in this form, where σ_0^2 is not proportional to $\frac{1}{\tau}$, the marginal density of τ is not a gamma density (or a density we can easily sample from).
- We need to do Gibbs sampling.

Full conditionals

■ That is, we need

$$\mu|Y, au \sim \mathcal{N}(\mu_n, au_n^{-1}),$$

where

$$\mu_n = rac{rac{\mu_0}{\sigma_0^2} + n auar{y}}{rac{1}{\sigma_0^2} + n au}$$

$$au_n = rac{1}{\sigma_0^2} + n au.$$

Full conditionals

and

$$au|\mu, Y \sim \mathrm{Gamma}\left(rac{
u_n}{2}, rac{
u_n \sigma_n^2(\mu)}{2}
ight),$$

where

$$\nu_n = \nu_0 + n$$

$$\sigma_n^2(\mu) = rac{1}{
u_n} \Bigg[rac{
u_0}{ au_0} + \sum_{i=1}^n (y_i - \mu)^2 \Bigg] = rac{1}{
u_n} \Big[rac{
u_0}{ au_0} + n s_n^2(\mu) \Big]$$

$$ext{with} \;\; s_n^2(\mu) = rac{1}{n} \sum_{i=1}^n (y_i - \mu)^2 \;\; \Rightarrow \;\; n s_n^2(\mu) = (n-1) s^2 + n (ar{y} - \mu)^2.$$

RECALL THE PYGMALION DATA

- Let's implement this Gibbs sampler for the Pygmalion data.
- For now, let's focus only on the accelerated group.
- Data for accelerated group (A): 20, 10, 19, 15, 9, 18.
- Summary statistics: $\bar{y}_A=15.2;\,s_A=4.71.$



RECALL THE PYGMALION DATA

- Suppose we assume, as we did before, that these improvement scores are normal with mean μ and variance $\frac{1}{\tau}$.
- As a reminder, in the conjugate case, we had

$$au \sim \mathrm{Gamma}\left(rac{
u_0}{2},rac{
u_0}{2 au_0}
ight) \ \mu| au \sim \mathcal{N}\left(\mu_0,rac{1}{\kappa_0 au}
ight).$$

- We set
 - ullet $\mu_0=0$, to reflect "no idea whether students would improve IQ on average";
 - $\kappa_0 = 1$, to reflect "little faith in this belief, equivalent to having only 1 prior observation in each group";
 - $\nu_0=1$ and $1/ au_0=100$, based on literature, that is, SD of change scores should be around 10.

RECALL THE PYGMALION DATA

Now, in the non-conjugate case, we have

$$egin{split} \mu &\sim \mathcal{N}\left(\mu_0, \sigma_0^2
ight). \ & au &\sim \mathrm{Gamma}\left(rac{
u_0}{2}, rac{
u_0}{2 au_0}
ight). \end{split}$$

- Suppose for μ , we use a $\mathcal{N}(0,100)$ prior, and for τ we use a $\mathrm{Ga}(\frac{1}{2},50)$ prior.
- Matching with

$$egin{aligned} \mu &\sim \mathcal{N}\left(\mu_0, \sigma_0^2
ight). \ & au &\sim \mathrm{Gamma}\left(rac{
u_0}{2}, rac{
u_0}{2 au_0}
ight), \end{aligned}$$

we have: $\mu_0=0$, $\sigma_0^2=100$, $\nu_0=1$ and $au_0=1/100$.

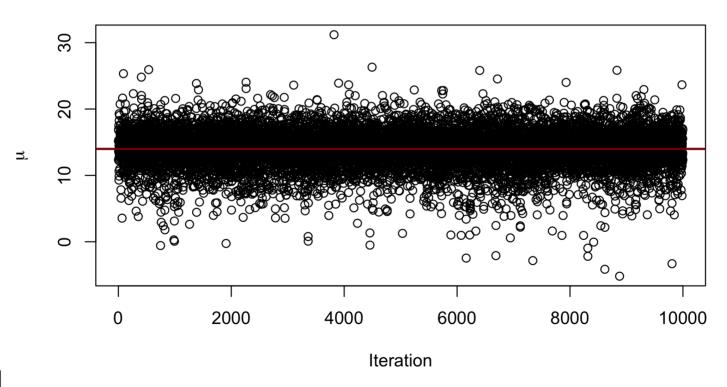
GIBBS SAMPLING FOR PYGMALION DATA

```
y \leftarrow c(20,10,19,15,9,18) #data
v bar <- mean(y); s2 <- var(y); n <- length(y) #sample statistics you'll need</pre>
S <- 10000 # number of samples to draw
PHI <- matrix(nrow=S,ncol=3); #matrix to save results
colnames(PHI) <- c("mu","tau","sigma2")</pre>
PHI[1,] <- phi <- c(v bar,1/s2,s2) #starting values are MLEs
mu0 <- 0; sigma02 <- 100; nu0 <- 1; tau0 <- 1/100 #hyperparameters
###### Gibbs sampler #####
set.seed(1234) #to replicate results exactly
for(s in 2:S) {
#first, draw new mu
taun <- 1/sigma02 + n*phi[2]
mun <- (mu0/sigma02 + n*y_bar*phi[2])/taun</pre>
phi[1] <- rnorm(1,mun,sgrt(1/taun))</pre>
#now, draw new tau/sigma2
nun <- nu0+n
#trick to speed up calculation of sum(v i-\mu)^2
s2nmu < (nu0/tau0 + (n-1)*s2 + n*(y bar-phi[1])^2)/nun
phi[2] <- rgamma(1,nun/2,nun*s2nmu/2)</pre>
phi[3] <- 1/phi[2] #save sigma2</pre>
#save the current joint draws
PHI[s,] <- phi
###### End of Gibbs sampler #####
```

PYGMALION DATA: MEAN

```
plot(PHI[,1],ylab=expression(mu),xlab="Iteration",
    main=expression(paste("Sampled values of ",mu)))
abline(a=mean(PHI[,1]),b=0,col="red4",lwd=2)
```

Sampled values of μ

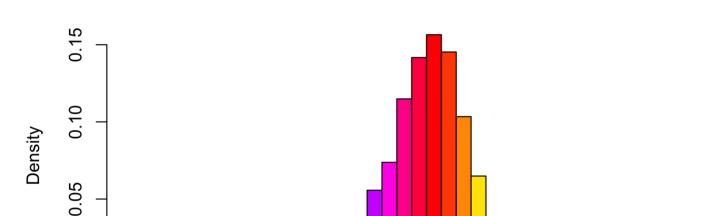




PYGMALION DATA: MEAN

0

Posterior density of µ



10

μ

20



0.00

30

PYGMALION DATA: MEAN

```
round(mean(PHI[,1]),3)

## [1] 13.99

round(quantile(PHI[,1],c(0.025,0.5,0.975)),3)

## 2.5% 50% 97.5%
## 7.520 14.217 19.277
```

Posterior summaries for μ :

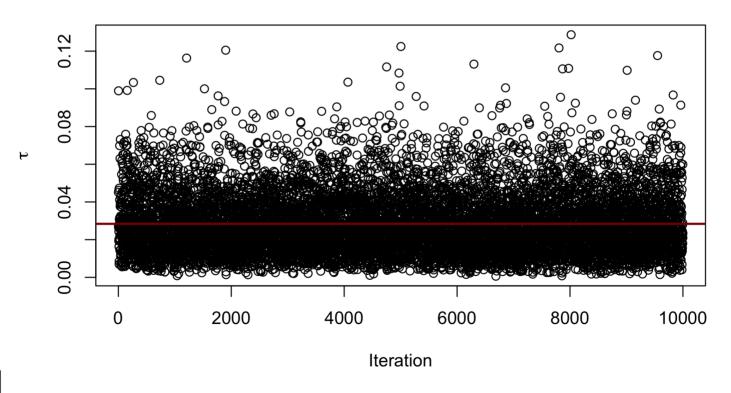
- Posterior mean ≈ 14 .
- Posterior median ≈ 14.22 .
- 95% credible interval $\approx (7.52, 19.28)$.

For context, ${ar y}_A=15.2$, and we used a $\mathcal{N}(0,100)$ prior for $\mu.$

PYGMALION DATA: PRECISION

```
plot(PHI[,2],ylab=expression(tau),xlab="Iteration",
    main=expression(paste("Sampled values of ",tau)))
abline(a=mean(PHI[,2]),b=0,col="red4",lwd=2)
```

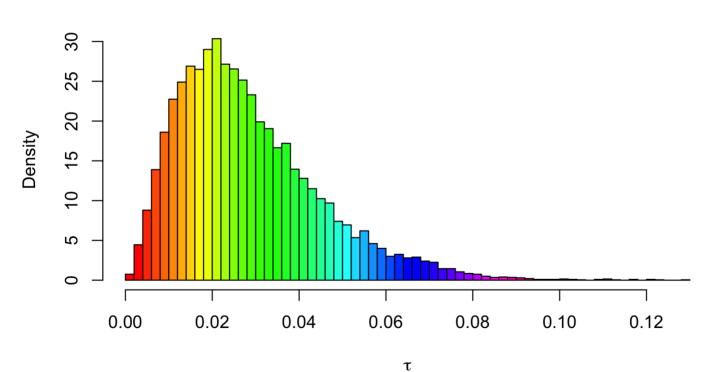
Sampled values of τ





PYGMALION DATA: PRECISION

Posterior density of τ





PYGMALION DATA: PRECISION

```
round(mean(PHI[,2]),3)

## [1] 0.028

round(quantile(PHI[,2],c(0.025,0.5,0.975)),3)

## 2.5% 50% 97.5%
## 0.006 0.025 0.069
```

Posterior summaries for τ :

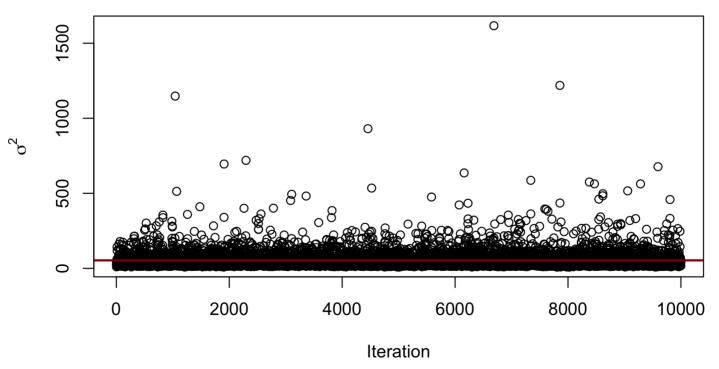
- Posterior mean ≈ 0.028 .
- Posterior median ≈ 0.025 .
- 95% credible interval pprox (0.006, 0.069).

For context, $s_A=4.71$, which means sample precision $=1/4.71^2=0.045$. Also, we used a ${\rm Ga}(\frac{1}{2},50)$ prior for $\tau.$

PYGMALION DATA: VARIANCE

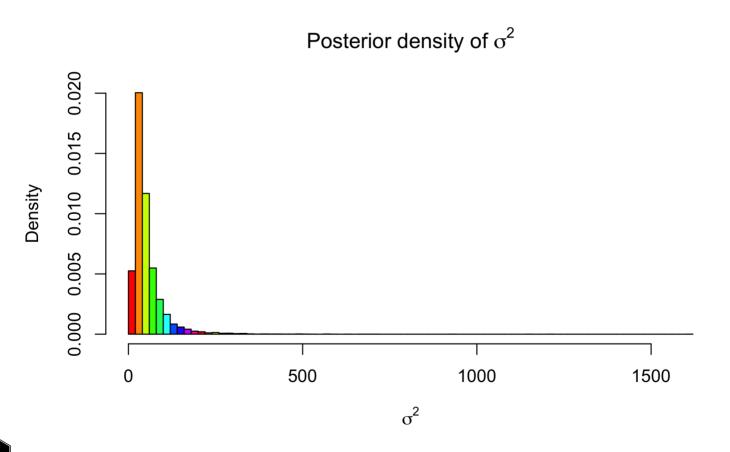
```
plot(PHI[,3],ylab=expression(sigma^2),xlab="Iteration",
    main=expression(paste("Sampled values of ",sigma^2)))
abline(a=mean(PHI[,3]),b=0,col="red4",lwd=2)
```

Sampled values of σ^2





PYGMALION DATA: VARIANCE



PYGMALION DATA: VARIANCE

```
round(mean(PHI[,3]),2)

## [1] 53.34

round(quantile(PHI[,3],c(0.025,0.5,0.975)),2)

## 2.5% 50% 97.5%
## 14.52 39.60 174.11
```

Posterior summaries for σ^2 :

- Posterior mean = 53.34.
- Posterior median = 39.60.
- 95% credible interval = (14.52, 174.11).

For context, $s_A=4.71$, which means sample variance $4.71^2=22.18$. Again, we used a ${\rm Ga}(\frac{1}{2},50)$ prior for $\tau.$

WHAT'S NEXT?

MOVE ON TO THE READINGS FOR THE NEXT MODULE!

