STAT 651 Mini Project 3

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1 Bootstrap Importance Sampling

1.1 Likelihood

I used a four parameter Beta function in order to align the support of the likelihood with the support of the data. I believe this distribution can appropriately capture probable data values, and can do so over the correct range of possible values (1-7). The pdf of this distribution is given as follows:

$$f(y;\alpha,\beta,a,c) = \frac{f(x;\alpha,\beta)}{c-a} = \frac{\left(\frac{y-a}{c-a}\right)^{\alpha-1} \left(\frac{c-y}{c-a}\right)^{\beta-1}}{(c-a)B(\alpha,\beta)} = \frac{(y-a)^{\alpha-1} (c-y)^{\beta-1}}{(c-a)^{\alpha+\beta-1}B(\alpha,\beta)}$$

Where c is fixed at 7 and a is fixed at 1. This is the pdf, and the likelihood would of course be the product of the pdf over the data values. A possible visualization of the likelihood can be seen in Figure 1.

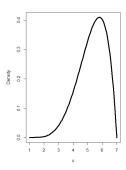


Figure 1: This four-parameter Beta Distribution, for example, represents my belief of how the rating data could be distributed.

1.2 Importance Function

I used a multivariate normal distribution as the importance function. This function works well as $I(\theta)$ because it mimics the values of the likelihood, as seen in Figure 2, and it covers the tails well. It is also a pdf that is easy to sample from and preserves the parameter space, as long as it remains greater than zero. Specifically, this distribution is:

$$MVN\left(\begin{bmatrix}17\\5\end{bmatrix},\begin{bmatrix}25&5\\5&2\end{bmatrix}\right)$$

1.3 Prior Distributions

Since the support of α and β is greater than zero, I used a gamma distribution for both parameters. The shape and scale values were selected with consideration to the equations for the mean and variance of the beta likelihood. I believed the mean of teacher ratings at BYU would be above average, with a somewhat substantial amount of variance, and my choices for prior parameters values reflected that belief.

$$\alpha \sim Gamma(shape=0.5, scale=16)$$

$$\beta \sim Gamma(shape = 0.5, scale = 4)$$

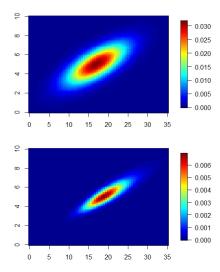


Figure 2: The Importance function above covers all values of $g(\theta)$

1.4 Posterior Distribution

The algorithm produces the bivariate posterior distribution in Figure 3.

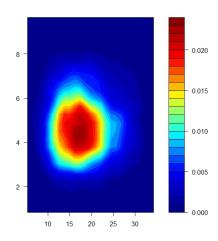


Figure 3: Bivariate posterior

1.5 Summary Statistics

• Mean,
$$E(\theta|Y) = \begin{bmatrix} 18.0421 \\ 4.7701 \end{bmatrix}$$

• Variance,
$$V(\theta|Y) = \begin{bmatrix} 26.3348 & 0.0493 \\ 0.0493 & 1.6479 \end{bmatrix}$$

• Standard Deviation,
$$\sqrt{V(\theta|Y)} = \begin{bmatrix} 5.1317 \\ 1.2837 \end{bmatrix}$$

1.6 Posterior Predictive Distribution

Figure 4 shows the distribution of the "next" score.

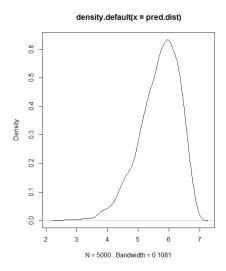


Figure 4: Posterior Predictive Distribution

 Probability next score is greater than 5, $P(\tilde{Y}>5)=.8544$

2 Metropolis-Hastings Algorithm

Figure 5 shows 10000 draws generated through the M-H algorithm from the distribution of X where:

$$f^*(x) \propto (1 + (x - 10)^2/3)^{-2}$$

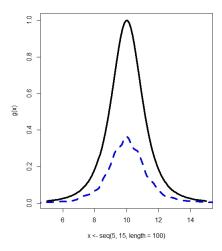


Figure 5: The dashed line density of the draws model the unnormalized distribution of f^* , displayed as a solid line

3 Gibbs Sampler

3.1 Likelihood

I used a Weibull likelihood, defined as follows:

$$f(x) = (\gamma/\beta)x^{\gamma-1}exp(-x^{\gamma})\beta$$

$$x \in (0, \infty), \gamma \in (0, \infty), \beta \in (0, \infty)$$

Where γ is the shape parameter, and β is the scale parameter. Figure 6 shows a possible visualization from this likelihood.

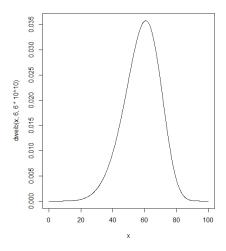


Figure 6: Here, $\gamma = 6$ and $\beta = 6e + 10$

3.2 Prior Distributions

$$\beta \sim InverseGamma(a, b)$$

Which is conjugate and allows for the posterior to be expressed in a closed form as:

$$\beta|x \sim InverseGamma(n+a, \sum x_i^{\gamma} + b)$$

$$\gamma \sim InverseGamma(c, d)$$

Figure 7 shows a plot of the prior distributions, Figure 8 shows a prior predictive plot.

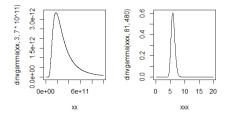


Figure 7: γ is on the right and β the left

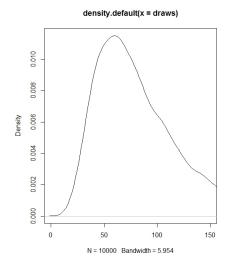


Figure 8: The prior distributions accurately represent the data.

3.3 Posterior Distribution

Figure 9 shows the marginal prior and posterior distributions. Figure 10 shows the bivariate contour plots.

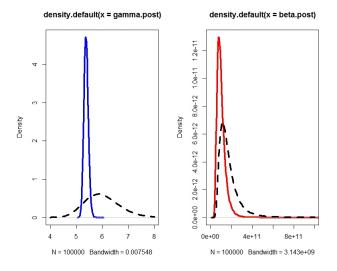


Figure 9: The dashed lines are the prior, and the solid lines are the posterior.

The 97% HPD Interval values are:

 $\gamma = (5.203, 5.574)$

 $\beta = (39637287078, 207119767272)$

Figure 11 shows the posterior distributions with the interval lines.

3.4 Summary Statistics

- Mean, $E(\gamma|x) = 5.3854$
- Variance, $V(\gamma|x) = 0.0073$
- Mean, $E(\beta|x) = 103629763271$

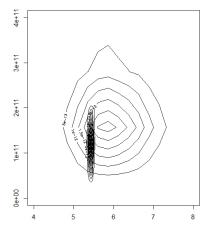


Figure 10: The prior distribution is the widely spaced contour and the posterior is the tightly clustered contour.

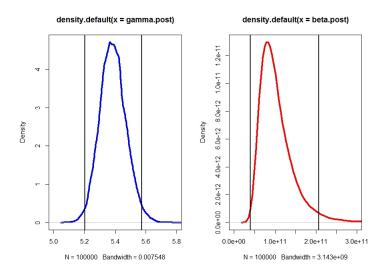


Figure 11: Posterior distributions with 97% HPD intervals

• Variance, $V(\beta|x) = 1.95482e + 21$

3.5 Posterior Predictive Distribution

Figure 12 shows the distribution of the "next" ball bearing.

• Probability next ball bearing will last 120 (10⁶ revolutions) $P(\tilde{Y} > 120) = .19881$

3.6 MCMC Settings

- Burn = 100
- $\sigma_{cand} = 0.25$
- Length = 100000

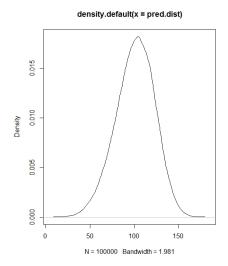


Figure 12: Posterior Predictive Distribution

- Candidate density for γ : Normal (centered at previous draw, σ_{cand}
- Figure 13 shows trace plots
- Acceptance rate for γ draws = 0.21963

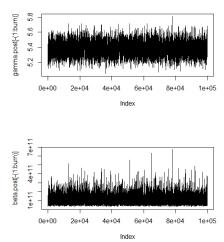


Figure 13: Trace plots suggest satisfactory convergence of the parameters.

4 Appendix: Code

```
score <-read.table("http://madison.byu.edu/bayes/faculty.dat")
library(ExtDist)
library(mvtnorm)</pre>
```

```
g \leftarrow function(a,b) {
      J <- numeric (M)
      for (i in 1:M) {
            J[i] \leftarrow prod(dBeta_ab(score[,1],a[i],b[i],1,7))*dgamma(a[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=16)*dgamma(b[i],.5,scale=1
      J
}
 I \leftarrow function(a,b) {
      J <- numeric (M)
      for (i in 1:M) {
            J[i] \leftarrow dmvnorm(x=c(a[i],b[i]), mean=c(17,5), sigma=matrix(c(25,5,5,2), nrow=2))
      J
}
M < -5000
 vals < -rmvnorm(M, mean=c(17,5), sigma=matrix(c(25,5,5,2), nrow=2))
w \leftarrow g(vals[,1], vals[,2]) / I(vals[,1], vals[,2])
q \ll w/sum(w)
out.a <- sample(vals[,1], size=M, replace=T, prob=q)
out.b <- sample(vals[,2], size=M, replace=T, prob=q)
mean(out.a)
var (cbind (out.a,out.b))
 sqrt(var(out.a))
mean(out.b)
 var (out.b)
 sqrt(var(out.b))
 hist (out.a, nclass=50, prob=T, xlab="")
 lines (density (out.a), lwd=4, col="blue")
 hist (out.b, nclass=50, prob=T, xlab="")
 lines (density (out.b), lwd=4, col="blue")
 library (MASS)
 filled.contour(kde2d(out.a,out.b),color.palette=colorRampPalette(c('white','blue','yellow'
 filled.contour(kde2d(out.a,out.b),color.palette=colorRampPalette(c('darkblue','blue','cyan
#predictive distribution
pred.dist <- numeric (M)
 for (i in 1:M) {
      index \leftarrow sample(1:M,1)
      a.samp <- out.a[index]
      b.samp <- out.b[index]
      pred.dist[i] \leftarrow rBeta_ab(1,a.samp,b.samp,1,7)
plot (density (pred.dist))
\#\text{prob} > 5
```

```
length (which (pred. dist \geq = 5))/M
#2
g \leftarrow function(x) {
  (1+((x-10)^2)/3)^--2
M < -10000
draws<-numeric (M)
draws[1] < -10
csig < -5
acc < -0
for (i in 2:M) {
  u \leftarrow runif(1)
  \operatorname{candy} < \operatorname{-rnorm}(1, \operatorname{draws}[i-1], \operatorname{csig})
  alpha < -g(candy)/g(draws[i-1])
  draws[i] < -draws[i-1]
  if (u<alpha) {
     draws [i] < - candy
     acc < -acc + 1
  }
}
acc < -acc/M
plot (draws, type="l")
plot(x < -seq(5, 15, length = 100), g(x), type = "l", lwd = 4)
lines (density (draws), lwd=4, lty=2, col="blue")
#3
ball <- read.table("http://madison.byu.edu/bayes/ballbearing2.dat")
dweib <- function (x,gamma, beta) {
  (gamma/beta)*x^(gamma-1)*exp((-x^gamma)/beta)
x < - seq(0,100,by=.1)
plot (x, dweib(x, 6, 6*10^10), type='l')
#prior for beta inv-gamma(a,b)
#conjugate is IV(n+a,sum(x^gamma)+b)
library (invgamma)
par(mfrow=c(1,2))
xx < - seq(10^8, 10^12, by=10^8)
plot(xx, dinvgamma(xx, 3, 7*10^11), type='l')
#prior for gamma inv-gamma(a,b)
xxx < - seq(0,20,by=.1)
plot (xxx, dinygamma (xxx, 81,480), type='1')
par(mfrow=c(1,1))
#prior-predictive
draws.beta \leftarrow rinvgamma(M, 3, 7*10^11)
draws.gamma <- rinvgamma (M, 81, 480)
draws <- rweibull (M, scale = (draws.beta^(1/draws.gamma)), shape=draws.gamma)
plot (density (draws), xlim=c (0,150))
```

```
#gibbs sampler
#prior values
a.prior.beta <- 5
b.prior.beta <- 7*10^11
a.prior.gamma <- 81
b.prior.gamma <- 480
# starting values
M < -100000
beta.post <--gamma.post <--numeric (M)
gamma. post [1] \leftarrow 6
beta.post[1] <-6*10^10
burn <\!\!- 100
csig \leftarrow 0.25
acc < -0
n \leftarrow length(ball[,1])
for (i in 2:M) {
  cand \leftarrow rnorm(1, gamma. post[i-1], csig)
  gamma.post[i] <- gamma.post[i-1]
  if (cand > 0)
  {
    lpo \leftarrow sum(dweibull(ball[,1],gamma.post[i-1],beta.post[i-1]^(1/gamma.post[i-1]),log=T)
    lpn \leftarrow sum(dweibull(ball[,1],cand,beta.post[i-1]^(1/cand),log=T))+dinygamma(cand,shape=T)
    if((lpn-lpo)>log(runif(1))){
      gamma.post[i] <- cand
       acc \leftarrow acc+1
  beta.post[i] <- rinvgamma(1,(n+a.prior.beta),(sum(ball^gamma.post[i]))+b.prior.beta)
accept <- acc/M
par(mfrow=c(2,1))
plot (gamma. post [-(1:burn)], type="l")
plot (beta. post [-(1:burn)], type="l")
par(mfrow=c(1,2))
plot (density (gamma. post), lwd=4, col="blue", xlim = c(4,8))
curve (dinygamma (x, shape = a. prior.gamma, rate = b. prior.gamma), xlim = c (4,8), add = TRUE, lwd=
plot (density (beta.post), lwd=4, col="red", xlim = c(0,10^12))
curve(dinvgamma(x, shape = a.prior.beta, rate = b.prior.beta), xlim = c(0,10^12), add = TRUE, lv
mean (gamma. post)
mean (beta.post)
var (gamma. post)
var (beta.post)
```

```
library (MASS)
conts.post <- kde2d(gamma.post, beta.post)
filled.contour(conts.post,color.palette=colorRampPalette(c('white','blue','yellow','red','
xx \leftarrow seq(4,8, length.out=M)
gamma.pre <- rinvgamma(xx,shape = a.prior.gamma,rate = b.prior.gamma)
xxx \leftarrow seq(0,10^11, length.out=M)
beta.pre <- rinvgamma(xxx, shape = a.prior.beta, rate = b.prior.beta)
conts.pre <- kde2d(gamma.pre, beta.pre)
filled.contour(conts.pre,color.palette=colorRampPalette(c('white','blue','yellow','red','d
contour (conts.pre$x,conts.pre$y,conts.pre$z,xlim=c(4,8),ylim=c(0,4e+11))
contour (conts.post$x, conts.post$y, conts.post$z, add=TRUE)
#predictive distribution
pred.dist <- numeric(M)
for (i in 1:M) {
  index <- sample (1:M,1)
  beta.samp <- beta.post[index]
  gamma.samp <- gamma.post[index]
  pred.dist[i] <- rweibull(1, scale=(beta.samp^(1/gamma.samp)), shape=gamma.samp)
par (mfrow=c(1,1))
plot (density (pred.dist))
length (which (pred.dist >= 120))/M
#hpd
library (coda)
HPDinterval (as.mcmc(beta.post), prob=0.97)
HPDinterval (as.mcmc(gamma.post), prob=0.97)
par(mfrow=c(1,2))
plot (density (gamma. post), lwd=4, col="blue", xlim = c(5,5.8))
abline (v=5.203147, lwd=2)
abline (v=5.573943, lwd=2)
plot (density (beta.post), lwd=4, col="red", xlim = c(0, 3e+11))
abline (v=39637287078, lwd=2)
abline(v=207119767272, lwd=2)
```