

This report is automatically generated with the R package [knitr](#) (version [1.24](#)) .

```

---
title: "STAT 457 - FINAL"
author: "Martha Eichlersmith"
date: "2019-12-12"
output:
  pdf_document:
    fig_caption: yes
#   number_sections: true
header-includes:
- \usepackage{color}
- \usepackage{mathtools}
- \usepackage{bbm} #for \mathbb{for} numbers
- \usepackage{amssys}
- \usepackage{caption} #to remove automatic table name and number - \captionset[table]{labelformat=empty}, put code under YAML
- \usepackage{booktabs}
- \usepackage{geometry}
- \usepackage{float} #to hold things in place
- \floatplacement{figure}{H}
- \usepackage{lastpage}
- \usepackage{fancyhdr}
- \pagestyle{fancy}
- \fancyhf{}
- \fancyhead[L]{STAT 457 Fall 2019 \ Final}
- \fancyhead[R]{Martha Eichlersmith \ Page \thepage\ of\ \pageref*{LastPage}}
- \setlength{headheight}{22.5pt} #to remove \fancyhead error for head height
geometry: "left=0.75in,right=0.75in,top=1.1in,bottom=1in"
---

\captionset[table]{labelformat=empty}
```{r setup, echo=FALSE, results="hide", warning=FALSE, message=FALSE}
library(ggplot2) #ggplot
library(gridExtra) #organize plots
library(grid) #organize plots
library(latex2exp) #latex in ggplot titles
library(dplyr) #for piping
library(MASS)
library(invgamma)
knitr::opts_chunk$set(fig.width = 10, fig.height = 4)
knitr::opts_chunk$set(echo=FALSE)
decimal <- function(x, k) trimws(format(round(x, k), nsmall=k))
dec <- 5
```

\newpage
# Problem 1
## Problem 1a
For the data  $Y = (125, 18, 20, 34)$ , implement the Gibbs sampler algorithm. Use a flat prior on  $\theta$ .
Plot  $\theta^i$  versus iteration  $i$ .
 $Y = (y_1, y_2, y_3, y_4)$  \propto  $(2 + \theta, 1 - \theta, 1 - \theta, \theta)$ 
1. Draw a starting value,  $t \sim \text{Uniform}(0, 1)$ 
2. Draw a latent value,  $Z \sim \text{Binomial}(\frac{y_1}{2 + \theta}, \theta)$ 
3. Draw a parameter,  $\theta \sim \text{Beta}(Z + y_4 + 1, y_2 + y_3 + 1)$ 

```{r p1.func_chain}
func_chain <- function(startseed, Y){
 set.seed(startseed)
 t <- runif(1)
 theta <- t
 chain <- c()#rep(NA, 10000)
 chain[1] <- theta
 Z.i <- rbinom(1, Y[1], (theta/(theta+2)))
 theta.i <- rbeta(1, Z.i+Y[4]+1, Y[2]+Y[3]+1)
 chain[2] <- theta.i
 Z.i <- rbinom(1, Y[1], (theta.i/(theta.i+2)))
 theta.i <- rbeta(1, Z.i+Y[4]+1, Y[2]+Y[3]+1)
 chain[3] <- theta.i
 k <- 3
 while(abs(chain[k]-chain[k-1]) >= 0.000001){
 Z.i <- rbinom(1, Y[1], (theta.i/(theta.i+2)))
 theta.i <- rbeta(1, Z.i+Y[4]+1, Y[2]+Y[3]+1)
 k <- k+1
 chain[k] <- theta.i
 }
 chain <- chain[!is.na(chain)]
 chain
}
```

```{r p1.scale_funcs}
func_scalelike <- function(x, y1, y2, y3, y4){
 like <- (2+x)^y1*(1-x)^(y2+y3)*(x)^y4
 like.max <- max(like)
 like/like.max #normalized likelihood (on scale from 0 to 1)
}

func_scalenormal <- function(x, mean, sd){
 scales::rescale(dnorm(x, mean, sd), to=c(0, 1)) #normal (on scale from 0 to 1)
}
```

```{r p1.func_p1AB}
func_problem1AB <- function(startseed, Y, number){

```

```

chain <- func_chain(startseed, Y)

df.chain <- data.frame(
 "theta.i" = chain,
 "i"=c(1:length(chain))
)

chain.mu <- mean(chain)
chain.sd <- sd(chain)
table <- data.frame(
 "Name" = c("Mean", "SD", "it", "Start"),
 "Value" = c(
 decimal(chain.mu, dec),
 decimal(chain.sd, dec),
 decimal(length(chain), 0),
 decimal(chain[1], dec)
)
)
tg <- tableGrob(table)

print.Y <- paste(Y, collapse = ",")

#plot theta_i versus i (iterations)
plot.Gibbs <- ggplot(df.chain, aes(x=i, y=theta.i))+
 geom_line(alpha=0.4)+
 ggtitle("Gibbs Sampler")

colors <- c("navy", "maroon")
norm.like <-stat_function(
 fun = func_scalelike
 , args = list(y1=Y[1], y2=Y[2], y3=Y[3], y4=Y[4])
 , lwd = 1
 , linetype="solid"
 , aes(col="Normalized Like.")
)
normal.approx <-stat_function(
 fun = func_scalenormal
 , args = list(mean=chain.mu, sd=chain.sd)
 , lwd = 1.5
 , linetype="dotted"
 , aes(col="Normal Approx.")
)

name <- paste("Normal Likelihood and Normal Approx")

#print normalized likelihoods
x <- seq(0, 1, 0.001)
df.x <- data.frame("X"=x)
plot.Like <- ggplot(data=df.x, aes(x=X))+
 norm.like+normal.approx+
 ggtitle(paste(name))+
 theme(axis.title.x = element_blank()
 ,axis.title.y = element_blank()
 ,axis.text.x = element_blank()
 ,axis.text.y = element_blank()
)+
 scale_colour_manual("", values = c(colors[1], colors[2])) +
 theme(legend.position = c(.2,.9))

main <- paste("Chain", number, "for data for Y=", print.Y, ")")

gs <- list(plot.Gibbs, tg, plot.Like)
grid.arrange(grobs=gs, nrow=1, widths=c(2, 1, 2),
 top = textGrob(main, vjust = .5, gp = gpar(fontface = "bold", cex = 1.2))
)
}
...

```{r p1aRESULT, fig.height=2}
Y.A <- c(125, 18, 20, 34)

func_problem1AB(111, Y.A, 1)
func_problem1AB(112, Y.A, 2)
func_problem1AB(113, Y.A, 3)
func_problem1AB(114, Y.A, 4)
func_problem1AB(115, Y.A, 5)
```

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Problem 1b
Repeat 1a for $Y = (14, 0, 1, 5)$.
```{r p1bRESULT, fig.height=2}
Y.B <- c(14,0,1,5)

func_problem1AB(121, Y.B, 1)
func_problem1AB(122, Y.B, 2)
func_problem1AB(123, Y.B, 3)
func_problem1AB(124, Y.B, 4)
func_problem1AB(125, Y.B, 5)
```

There is a lack of fit for the data in 1b, where the fit appears to be better for data in 1a. Convergence was assessed when values were had a difference

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Problem 1c
```{r p1.func_chaininfo}
func_chaininfo <- function(startseed, Y){
  chain <- func_chain(startseed, Y)

```

```

it <- length(chain)
chain.mu <- mean(chain)
chain.sd <- sd(chain)
chain.se <- chain.sd/sqrt(it)
vec <- c(chain.mu, chain.sd, chain.se, it)
return(vec)
}...

```{r p1cRESULT20chains}
Y.C <- c(125, 18, 20, 34)
print.Y.C <- paste(Y.C, collapse=",")
cnames <- c("Mean", "Standard Deviation", "Standard Error", "Iterations")
info.20chain <- mapply(func_chaininfo, startseed=c(1:20), Y=rep(list(Y.C), 20))
info.20chain <- t(info.20chain)
colnames(info.20chain) <- cnames

knitr::kable(info.20chain, align='rrrr', digits=dec, caption=paste("20 Chains for Y=", print.Y.C, "))
```

```{r p1cRESULTavgsd.vs.se}
sd.of.avgs <- sd(info.20chain[,1])
avg.of.se <- mean(info.20chain[,3])

paste(round(sd.of.avgs, dec+1), "is the standard deviation of the 20 averages of theta")
paste(decimal(avg.of.se, dec+1), "is the average of the standard errors of the 20 chains of theta")
```

```

You would expected these values to be similar but it appears that our standard error average is under-estimating the variation in the chain means of θ

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Problem 2

Problem 2a

For the genetic linkage model applied to $Y = (125, 18, 20, 34)$, implement the Metropolis algorithm. (use a flat prior on θ). Use one long chain and plot θ^i versus i . Try several driver functions:

```

```{r p2.func_pi,func_Metropolis,func_drivers}
func_pi <- function(theta, Y){
 (2+theta)^(Y[1]) * (1 - theta)^(Y[2]+Y[3]) * (theta)^(Y[4])
}

m <- 10

#theta.i = X (old value)
#theta.j = Y (new value)
theta.i <- 0.1
Y <- c(125, 18, 20, 34)

func_MetroFix <- function(startseed, m, Y, driver){
 #fixed driver
 set.seed(startseed)
 theta.i <- runif(1)
 chain <- c()
 alpha <- c()
 chain[1] <- theta.i
 for (k in 2:m){
 theta.j <- driver(1, theta.i)
 alpha.ij <- min(c(1, func_pi(theta.j, Y)/func_pi(theta.i, Y)))
 if (theta.j > 1 | theta.j < 0){alpha.ij <- 0}
 u <- runif(1)
 if (u < alpha.ij){theta.i <- theta.j}
 chain[k] <- theta.i
 }
 chain
}

func_MetroDyn <- function(startseed, m, Y, driver){
 #dynamic driver
 set.seed(startseed)
 theta.i <- runif(1)
 chain <- c()
 alpha <- c()
 chain[1] <- theta.i
 for (k in 2:m){
 theta.j <- driver(1, theta.i)
 alpha.ij <- min(c(1, func_pi(theta.j, Y)/func_pi(theta.i, Y)))
 if (theta.j > 1 | theta.j < 0){alpha.ij <- 0}
 u <- runif(1)
 if (u < alpha.ij){theta.i <- theta.j}
 chain[k] <- theta.i
 }
 chain
}

func_driver1.Uniform <- function(n){runif(n, min=0, max=1) }
func_driver2.Norm.sd.01 <- function(n, mu){rnorm(n, mean=mu, sd=0.01)}
func_driver3.Norm.sd.1 <- function(n, mu){rnorm(n, mean=mu, sd=0.1)}
func_driver4.Norm.sd.5 <- function(n, mu){rnorm(n, mean=mu, sd=0.5)}
func_driver5.Norm.mu.4sd.5 <- function(n){rnorm(n, mean=0.4, sd=0.5)}
```

```{r p2.func_p2ABC}
func_problem2ABC <- function(chain, driver.name){
 #main title
 print.Y <- paste(Y, collapse=",")
}

```

```

main <- paste("Metropolis for data Y=(", print.Y, ") using the driver", driver.name)

df <- data.frame(
 "theta.i"=chain
 , "i"=c(1:length(chain))
)

#table grob
chain.mu <- mean(chain)
chain.sd <- sd(chain)
table <- data.frame(
 "Name" = c("Mean", "SD", "it", "Start"),
 "Value" = c(
 decimal(chain.mu, dec),
 decimal(chain.sd, dec),
 decimal(length(chain), 0),
 decimal(chain[1], dec)
)
)
tg <- tableGrob(table)

#plot theta_i versus i (iterations)
plot <- ggplot(df, aes(x=i, y=theta.i))+
 geom_line(alpha=0.4)

grid.arrange(plot, tg, widths=c(4,1)
 , top = textGrob(main, vjust = .5, gp = gpar(fontface = "bold", cex = 1.1))
)

...

```{r p2aRESULT, fig.height=2}
Y.A <- c(125, 18, 20, 34)

func_problem2ABC(func_MetroFix(211, 10000, Y.A, func_driver1.Uniform), "Uniform(0,1)")
func_problem2ABC(func_MetroDyn(212, 10000, Y.A, func_driver2.Norm.sd.01), "Normal(theta.i, 0.01)")
func_problem2ABC(func_MetroDyn(213, 10000, Y.A, func_driver3.Norm.sd.1 ), "Normal(theta.i, 0.10)")
func_problem2ABC(func_MetroDyn(214, 10000, Y.A, func_driver4.Norm.sd.5 ), "Normal(theta.i, 0.50)")
func_problem2ABC(func_MetroFix(215, 10000, Y.A, func_driver5.Norm.mu.4sd.5), "Normal(0.40, 0.10)")
...

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## Problem 2b
Repeat 2a for $Y = (14, 0, 1, 5)$
```{r p2bRESULT, fig.height=2}
Y.B <- c(14, 0, 1, 5)

func_problem2ABC(func_MetroFix(221, 10000, Y.B, func_driver1.Uniform), "Uniform(0,1)")
func_problem2ABC(func_MetroDyn(222, 10000, Y.B, func_driver2.Norm.sd.01), "Normal(theta.i, 0.01)")
func_problem2ABC(func_MetroDyn(223, 10000, Y.B, func_driver3.Norm.sd.1), "Normal(theta.i, 0.10)")
func_problem2ABC(func_MetroDyn(224, 10000, Y.B, func_driver4.Norm.sd.5), "Normal(theta.i, 0.50)")
func_problem2ABC(func_MetroFix(225, 10000, Y.B, func_driver5.Norm.mu.4sd.5), "Normal(0.40, 0.10)")
...

Problem 2c
Compute both the posterior mean and standard deviation for both data sets.
Compare to results from the previous problem.

In problem 1 the means were similar, but in problem 2 the means vary depending on the driver. Some of the means in problem 2 are close to the means in

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Problem 2d

```{r p2.func_Metroinfo}
func_Metroinfo <- function(startseed, m, Y, driver, metro.func){
  chain <- metro.func(startseed, m, Y, driver)
  it <- length(chain)
  chain.mu <- mean(chain)
  chain.sd <- sd(chain)
  chain.se <- chain.sd/sqrt(it)
  vec <- c(chain.mu, chain.sd, chain.se)
  return(vec)
}
...

```{r p2.func_20chains}
func_20metrochains <- function(Y, driver, metro.func){
 print.Y <- paste(Y, collapse=",")
 cnames <- c("Mean", "SD", "SE")
 info.20chain <- mapply(func_Metroinfo
 , startseed=c(1:20)
 , m =rep(10000, 20)
 , Y=rep(list(Y), 20)
 , driver=rep(list(driver), 20)
 , metro.func=rep(list(metro.func), 20)
)
 info.20chain <- t(info.20chain)
 colnames(info.20chain) <- cnames
 info.20chain
}
...

```{r p2dRESULT20chains}
Y.D <- c(125, 18, 20, 34)

d1 <- func_20metrochains(Y.D, func_driver1.Uniform, func_MetroFix)
d2 <- func_20metrochains(Y.D, func_driver2.Norm.sd.01, func_MetroDyn)

```

```

d3 <- func_20metrochains(Y.D, func_driver3.Norm.sd.1, func_MetroDyn)
d4 <- func_20metrochains(Y.D, func_driver4.Norm.sd.5, func_MetroDyn)
d5 <- func_20metrochains(Y.D, func_driver5.Norm.mu.4sd.5, func_MetroFix)

table2d <- cbind(d1, d2, d3, d4, d5)

print.Y.D <- paste(Y.D, collapse=",")

knitr::kable(table2d, digits=4, booktabs=TRUE, 'latex'
  , caption=paste("20 Chains for Y=(",print.Y.D, ") with Different Drivers")
  ) %>%
  kableExtra::kable_styling(latex_options=c("hold_position", "scale_down") ) %>%
  kableExtra::add_header_above(c( "Uniform(0, 1)" =3
    , "Normal(theta.i, 0.01)"=3
    , "Normal(theta.i, 0.10)"=3
    , "Normal(theta.i, 0.50)"=3
    , "Normal(0.40, 0.10)"=3
  ))
...

```{r p2dRESULTavgsd.vs.se}
sd.of.avgs <- c(
 sd(table2d[,1])
 ,sd(table2d[,4])
 ,sd(table2d[,7])
 ,sd(table2d[,10])
 ,sd(table2d[,13])
)

avg.of.se <- c(
 mean(table2d[,3])
 ,mean(table2d[,6])
 ,mean(table2d[,9])
 ,mean(table2d[,12])
 ,mean(table2d[,15])
)

driver.name <- c(
 "Uniform(0,1)"
 , "Normal(theta.i, 0.01)"
 , "Normal(theta.i, 0.10)"
 , "Normal(theta.i, 0.50)"
 , "Normal(0.40, 0.10)"
)

table2d.sdse <- cbind(sd.of.avgs, avg.of.se)
rownames(table2d.sdse) <- driver.name

knitr::kable(table2d.sdse, 'latex', booktabs=TRUE
 , caption=paste("SD of Average theta's versus Average SE")
) %>%
 kableExtra::kable_styling(latex_options="hold_position")
...

```

Similar to problem 1, it appears that our estimation of the variation in the mean of theta is lower than the actual variation between the means.

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# Problem 3

## Problem 3a

Consider the 1-way variance components model

$Y_{ij} = \theta_i + \epsilon_{ij}$

where  $Y_{ij}$  is the  $j$ th observation from the  $i$ th group,  $\theta_i$  is the effect,  $\epsilon_{ij}$  is error,  $i=1, \dots, K$  and  $j=1, \dots, J$ .

It is assumed that  $\epsilon_{ij} \stackrel{\text{iid}}{\sim} \mathcal{N}(0, \sigma^2_{\epsilon})$

and  $\theta_i \stackrel{\text{iid}}{\sim} \mathcal{N}(\mu, \sigma^2_{\theta})$ .

Under the prior specification  $p(\sigma^2_{\epsilon}, \sigma^2_{\theta}, \mu) =$

$p(\sigma^2_{\epsilon})p(\sigma^2_{\theta})p(\mu)$ , with

$p(\sigma^2_{\theta}) = \text{InverseGamma}(a_1, b_1)$ ,

$p(\sigma^2_{\epsilon}) = \text{InverseGamma}(a_2, b_2)$ , and

$p(\mu) = \mathcal{N}(\mu_0, \sigma^2_0)$ .

Let  $\bar{Y}_i = \frac{1}{J} \sum_{j=1}^J Y_{ij}$  and  $\theta = (\theta_1, \dots, \theta_K)$ .

Show the following:

```

$$
\begin{aligned}
p(\mu, \sigma^2_{\epsilon}, \sigma^2_{\theta}) &= p(\mu) p(\sigma^2_{\theta}) p(\sigma^2_{\epsilon}) \\
&= \mathcal{N}(\mu_0, \sigma^2_0) \text{IG}(a_1, b_1) \text{IG}(a_2, b_2) \\
&= \left[\frac{1}{\sqrt{2\pi}} \frac{1}{\sigma_0} \exp \left\{ -\frac{1}{2} \frac{(\mu - \mu_0)^2}{\sigma_0^2} \right\} \right] \\
&\quad \cdot \left[\frac{1}{\sigma^2_{\theta}} \exp \left\{ -\frac{b_1}{\sigma^2_{\theta}} \right\} \right] \\
&\quad \cdot \left[\frac{1}{\sigma^2_{\epsilon}} \exp \left\{ -\frac{b_2}{\sigma^2_{\epsilon}} \right\} \right] \\
&= \left[\frac{1}{\sqrt{2\pi}} \frac{1}{\sigma_0} \exp \left\{ -\frac{1}{2} \frac{(\mu - \mu_0)^2}{\sigma_0^2} \right\} \right] \\
&\quad \cdot \left[\frac{1}{\sigma^2_{\theta}} \exp \left\{ -\frac{b_1}{\sigma^2_{\theta}} \right\} \right] \\
&\quad \cdot \left[\frac{1}{\sigma^2_{\epsilon}} \exp \left\{ -\frac{b_2}{\sigma^2_{\epsilon}} \right\} \right]
\end{aligned}

```

```

-\frac{1}{2} \frac{(\mu - \mu_0)^2}{\sigma_0^2} + b_1 \sigma^2_{\theta} + b_2 \sigma^2_{\epsilon}
\right)
\\[2ex]
p(\mu, \theta, \sigma^2_{\theta}, \sigma^2_{\epsilon} \mid Y) \propto p(Y \mid \mu, \theta, \sigma^2_{\theta}, \sigma^2_{\epsilon}) \cdot p(\theta \mid \mu,
\\
& = \left[\prod_{i=1}^K \prod_{j=1}^J \text{left}(p(y_{ij}) \mid \mathcal{N}(\theta_i, \sigma^2_{\epsilon}) \right) \cdot
\left[\prod_{i=1}^K \text{left}(p(\theta_i) \mid \mathcal{N}(\mu, \sigma^2_{\theta}) \right)
\cdot \prod_{i=1}^K p(\mu, \sigma^2_{\theta}, \sigma^2_{\epsilon})
\\[0.5ex]
& = \left[
\sigma_{\epsilon}^{-(KJ)} \cdot \exp \left\{ - \frac{1}{2} \frac{\sum_{i=1}^K \sum_{j=1}^J (y_{ij} - \theta_i)^2}{\sigma^2_{\epsilon}} \right\}
\right]
\\[0.5ex]
& \cdot \left[\sigma_{\theta}^{-2K} \exp \left\{ - \frac{1}{2} \frac{\sum_{i=1}^K (\theta_i - \mu)^2}{\sigma^2_{\theta}} \right\} \right]
\\[0.5ex]
& \cdot \left[
\sigma_{\theta}^{-2K(a_1 - 1)} \cdot
\sigma_{\epsilon}^{-2K(a_2 - 1)} \cdot
\exp \left\{ K \text{left}(
- \frac{1}{2} \frac{(\mu - \mu_0)^2}{\sigma_0^2} + b_1 \sigma^2_{\theta} + b_2 \sigma^2_{\epsilon}
\right) \right\}
\right]
\end{aligned}
$$

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Problem 3a(1)
$$
\begin{aligned}
& p(\mu \mid \theta, \sigma^2_{\epsilon}, \sigma^2_{\theta}, Y) \propto
& \exp \left\{ - \frac{1}{2} \frac{\sum_{i=1}^K (\theta_i - \mu)^2}{\sigma^2_{\theta}} \right\}
& \cdot
& \exp \left\{ K \text{left}(
& - \frac{1}{2} \frac{(\mu - \mu_0)^2}{\sigma_0^2}
& \right) \right\}
\\[0.5ex]
& & = \mathcal{N} \left(\text{left}(
& \frac{\sigma^2_{\theta} \sum_{i=1}^K \theta_i}{\sigma^2_{\theta} + K \sigma^2_{\epsilon}}
& , \frac{\sigma^2_{\theta} \sigma^2_{\epsilon}}{\sigma^2_{\theta} + K \sigma^2_{\epsilon}}
& \right)
& \end{aligned}
$$

Problem 3a(2)
$$
\begin{aligned}
& p(\theta_i \mid \mu, \sigma^2_{\epsilon}, \sigma^2_{\theta}, Y) \propto
& \exp \left\{ - \frac{1}{2} \frac{\sum_{j=1}^J (y_{ij} - \theta_i)^2}{\sigma^2_{\epsilon}} \right\}
& \cdot
& \exp \left\{ - \frac{1}{2} \frac{(\theta_i - \mu)^2}{\sigma^2_{\theta}} \right\}
& \cdot
& \frac{J \sigma^2_{\theta}}{J \sigma^2_{\theta} + \sigma^2_{\epsilon}} \cdot \overline{Y}_i
& +
& \frac{\sigma^2_{\epsilon}}{J \sigma^2_{\theta} + \sigma^2_{\epsilon}} \cdot \mu
& , \frac{\sigma^2_{\theta} \sigma^2_{\epsilon}}{J \sigma^2_{\theta} + \sigma^2_{\epsilon}}
& \right)
& \end{aligned}
$$

Problem 3a(3)
$$
\begin{aligned}
& p(\sigma^2_{\epsilon} \mid \mu, \theta, \sigma^2_{\theta}, Y) \propto
& =
& \sigma_{\epsilon}^{-(KJ)} \cdot \exp \left\{ - \frac{1}{2} \frac{\sum_{i=1}^K \sum_{j=1}^J (y_{ij} - \theta_i)^2}{\sigma^2_{\epsilon}} \right\}
& \cdot
& \sigma_{\epsilon}^{-2K(a_2 - 1)} \cdot
& \exp \left\{ K \text{left}(
& b_2 \sigma^2_{\epsilon}
& \right) \right\}
\\[0.5ex]
& & = \text{IG}
& \left(
& a_2 + \frac{KJ}{2}, \frac{
b_2 + \frac{1}{2} \sum_{i=1}^K \sum_{j=1}^J (Y_{ij} - \theta_i)^2
}{
}
& \right)
& \end{aligned}
$$

Problem 3a(4)
$$
\begin{aligned}
& p(\sigma^2_{\theta} \mid \mu, \theta, \sigma^2_{\epsilon}, Y) \propto
& \sigma_{\theta}^{-2K} \exp \left\{ - \frac{1}{2} \frac{\sum_{i=1}^K (\theta_i - \mu)^2}{\sigma^2_{\theta}} \right\}
& \cdot
& \sigma_{\theta}^{-2K(a_1 - 1)} \cdot
& \exp \left\{ K \text{left}(
& b_1 \sigma^2_{\theta}
& \right) \right\}
& = \text{IG}
& \left(
& \right)
& \end{aligned}

```

```

a_1 + \frac{K}{2}
, \
b_1 + \frac{1}{2} \sum_{i=1}^K (\theta_i - \mu)^2
\right)
\end{aligned}
$$

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Problem 3b
Run the Gibbs sampler for the data below.
```{r p3b.data, eval=TRUE}
j1 <- c( 7.298,  3.846,  2.434,  9.566,  7.990)
j2 <- c( 5.220,  6.556,  0.608, 11.788, -0.892)
j3 <- c( 0.110, 10.386, 13.434,  5.510,  8.166)
j4 <- c( 2.212,  4.852,  7.092,  9.288,  4.980)
j5 <- c( 0.282,  9.014,  4.458,  9.446,  7.198)
j6 <- c( 1.722,  4.782,  8.106,  0.758,  3.758)

Y <- cbind(j1, j2, j3, j4, j5, j6)
y_j. <- apply(Y, 2, mean)

rbind(Y, c(rep(" ", 6)), y_j.)
paste("y_." =, mean(Y))
...

Use one chain of length 75,000.
Take  $p(\mu) = \text{mathcal{N}}(0, 10^{12})$ ,  $p(\sigma^2, \epsilon) = \text{IG}(0, 0)$ , and  $p(\sigma^2, \theta) = \text{IG}(1, 1)$ .
For each  $\theta_i$ , for  $\sigma, \epsilon$ , and for  $\theta$ , plot the simulated value at iteration  $j$  versus  $j$ .
Summarize each posterior marginal.

```{r p3b.func_gibbsdf}
func_gibbsdf <- function(startseed, it, mu.o, sig2.o, a1, b1, a2, b2, Y){
 set.seed(startseed)
 J <- dim(Y)[2] #ROW, observations
 K <- dim(Y)[1] #COLUMN, group
 Y.col <- apply(Y, 2, mean) #col means
 sig.o <- sqrt(sig2.o)
 chain.mu <- c()
 chain.sig2.0 <- c()
 chain.sig2.e <- c()
 chain.theta <- matrix(0, nrow=it, ncol=K)

 #Iteration ONE - STARTING VALUES
 mu <- rnorm(1, mu.o, sig.o) #generate \mu
 sig2.0 <- rinvgamma(1, a1, b1) #generate \sigma^2, \theta
 sig2.e <- rinvgamma(1, a2, b2) #generate \sigma^2, \epsilon
 theta <- rnorm(K, mu, sqrt(sig2.0)) #generate \theta = (\theta_1, ..., \theta_K)
 chain.mu[1] <- mu
 chain.sig2.0[1] <- sig2.0
 chain.sig2.e[1] <- sig2.e
 chain.theta[1,] <- theta

 #ITERATION TWO+
 for (l in 2:it){
 #values for respective distributions
 mean.mu <- (sig2.0 * mu.o + sig2.o * sum(theta)) / (sig2.0 + K * sig2.o)
 sig2.mu <- (sig2.0 * sig2.o) / (sig2.0 + K * sig2.o)
 A.sig2.0 <- a1 + K/2
 B.sig2.0 <- b1 + .5 * sum((theta - mu)^2)
 A.sig2.e <- a2 + (K*J)/2
 B.sig2.e <- b2 + .5 * sum((Y - matrix(theta, K, J, byrow=TRUE))^2)
 mean.theta <- Y.col * (J * sig2.0) / (J * sig2.0 + sig2.e) + mu * (sig2.e) / (J * sig2.0 + sig2.e)
 sig2.theta <- (sig2.0 * sig2.e) / (J * sig2.0 + sig2.e)

 #new values
 mu <- rnorm(1, mean.mu, sqrt(sig2.mu))
 sig2.0 <- rinvgamma(1, A.sig2.0, B.sig2.0)
 sig2.e <- rinvgamma(1, A.sig2.e, B.sig2.e)
 theta <- rnorm(K, mean.theta, sqrt(sig2.theta))

 #put new values into chains
 chain.mu[l] <- mu
 chain.sig2.0[l] <- sig2.0
 chain.sig2.e[l] <- sig2.e
 chain.theta[l,] <- theta
 }

 df.chain <- data.frame(
 "iteration" = c(1:it)
 , "mu" = chain.mu
 , "sig2.theta" = chain.sig2.0
 , "sig2.epsilon" = chain.sig2.e
 , "theta.1" = chain.theta[,1]
 , "theta.2" = chain.theta[,2]
 , "theta.3" = chain.theta[,3]
 , "theta.4" = chain.theta[,4]
 , "theta.5" = chain.theta[,5]
 , "theta.6" = chain.theta[,6]
)

 return(df.chain)
}

...

```{r 3b.func_plotchain}
func_plotchain <- function(df.chain, throw.out, c, yval.tex){

```

```

df.chain <- df.chain[throw.out:nrow(df.chain), ] #throw out first _ values
yval <- df.chain[, c]
post.mean <- mean(yval)
xname <- colnames(df.chain)[1]
ggplot(df.chain, aes(x=iteration, y=df.chain[,c])) + geom_line(alpha=0.4)+
  ggtitle(TeX(paste(yval.tex, "versus iteration")))+ ylab(TeX(yval.tex)) + xlab("Iteration")+
  annotate('text', x=mean(df.chain$iteration), y=Inf,
    label=paste("Mean=", decimal(post.mean, dec)), vjust=2, fontface="bold")
}
...

```{r p3b.func_problem3b}

func_problem3b <- function(startseed, it, mu.o, sig2.o, a1, b1, a2, b2, Y, throw.out){
df.chain <- func_gibbsdf(startseed, it, mu.o, sig2.o, a1, b1, a2, b2, Y)
plot.mu <- func_plotchain(df.chain, throw.out, 2, "\\mu$")
plot.sig2theta <- func_plotchain(df.chain, throw.out, 3, "\\sigma^2_{\\theta}$")
plot.sig2epsilon <- func_plotchain(df.chain, throw.out, 4, "\\sigma^2_{\\epsilon}$")
plot.theta1 <- func_plotchain(df.chain, throw.out, 5, "\\theta_1$")
plot.theta2 <- func_plotchain(df.chain, throw.out, 6, "\\theta_2$")
plot.theta3 <- func_plotchain(df.chain, throw.out, 7, "\\theta_3$")
plot.theta4 <- func_plotchain(df.chain, throw.out, 8, "\\theta_4$")
plot.theta5 <- func_plotchain(df.chain, throw.out, 9, "\\theta_5$")
plot.theta6 <- func_plotchain(df.chain, throw.out, 10, "\\theta_6$")

gs <- list(plot.mu
 , plot.sig2theta
 , plot.sig2epsilon
 , plot.theta1
 , plot.theta2
 , plot.theta3
 , plot.theta4
 , plot.theta5
 , plot.theta6)

#gs <- list(plot.theta1, plot.theta2, plot.theta3)
grid.arrange(grobs=gs, nrow=3, ncol=3
 , top = textGrob(paste("Chains versus Iterations, throwing out first", throw.out,"values on", it, "chain"), vjust = .5, gp = gpar(fontface
)
)
}
...

Cannot use $a_1, b_1 = 0$, otherwise $\\sigma^2_{\\theta} = \\infty$, resulting in N/A so will use 0.01 instead.
```{r p3bRESULT, fig.height=8}
it <- 75000
#prior for mu
mu.o <- 0
sig2.o <- 10^12
#prior for sig2.theta
a1 <- .01
b1 <- .01
#prior for sig2.epsilon
a2 <- 1
b2 <- 1

Y <- cbind(j1, j2, j3, j4, j5, j6)
startseed <- 32

func_problem3b(startseed, it, mu.o, sig2.o, a1, b1, a2, b2, Y, 1000)
...

\\newpage
## Problem 3c
Repeat 3b using the prior specification  $p(\\mu) = \\mathcal{N}(0, 10^{12})$ ,  $p(\\sigma_{\\epsilon}^2) = \\text{IG}(0, 0)$ , and
 $p(\\sigma_{\\theta}^2) = \\text{IG}(0, 0)$ . Does this specification violate the Hobart-Casella conditions?
Describe what happens to the Gibbs sampler chain in this case.

This violates the Hobart-Casella conditions. If left at zero, the result will "blow up" (i.e., the chains are full of NA).
Cannot use $a_1, b_1, a_2, b_2 = 0$, otherwise $\\sigma^2_{\\theta} = \\sigma^2_{\\epsilon} = \\infty$, resulting in N/A so will use 0.01 instead.
```{r p3cRESULT, fig.height=8}
it <- 75000
#prior for mu
mu.o <- 0
sig2.o <- 10^12
#prior for sig2.theta
a1 <- .01
b1 <- .01
#prior for sig2.epsilon
a2 <- .01
b2 <- .01

Y <- cbind(j1, j2, j3, j4, j5, j6)
startseed <- 32

func_problem3b(startseed, it, mu.o, sig2.o, a1, b1, a2, b2, Y, 1000)
...

\\newpage
Problem 5
Suppose that X and Y have exponential conditional distributions restricted over the interval $(0, B)$, i.e.
 $p(x | y) \\propto y \\exp \\left\\{ -yx \\right\\}$ for $0 < x < B < \\infty$ and
 $p(y | x) \\propto x \\exp \\left\\{ -xy \\right\\}$ for $0 < y < B < \\infty$, where B is known constant.

Problem 5a
Take $m=1$ and $B=3$. Run the data augmentation algorithm using these conditionals.
(Hint: Reject the exponential deviates that lie outside $(0, B)$.)
Obtain the marginal for X using the mixture of conditionals $p(x | y)$, mixed over the simulated Y deviates in your chain.

```



```

```{r p5.func_gibbs.prob5A}
func_gibbs.p5 <- function(B, it){
  x <- c(rep(B+1, it))
  y <- c(rep(B+1, it))
  x[1] <- runif(1, 0, B)
  y[1] <- runif(1, 0, B)
  for(k in 2:it) {
    while(x[k] > B){ x[k]<-rexp(1,y[k-1]) }
    while(y[k] > B){ y[k]<-rexp(1,x[k-1]) }
  }
  df <- as.data.frame(cbind(x, y))
  return(df)
}

marginal <- function(k, rate){
  (1-exp(-rate*k))/(k)
}

func_marginal <- function(k, rate){
  marginal(k, rate)/rate #to normalize to 0 to 1
}

func_problem5A <- function(startseed, B, it){
  set.seed(startseed)
  df <- func_gibbs.p5(B, it)

  true.cdf <- stat_function(fun = func_marginal, args = B, lwd = 1, linetype="solid", col="maroon")

  plot.x <- ggplot(df, aes(x)) + xlim(0, B)+
    geom_histogram(aes(y=..density..), alpha=0.4) +
    true.cdf +xlab("X Chain Values")+
    ggtitle("Marginal of X: Histogram and True Curve")

  plot.y <- ggplot(df, aes(y)) + xlim(0, B)+
    geom_histogram(aes(y=..density..), alpha=0.4) +
    true.cdf +xlab("Y Chain Values")+
    ggtitle("Marginal of Y: Histogram and True Curve")

  grid.arrange(plot.x, plot.y, nrow=1)
}
...

```{r p5aRESULTS, message=FALSE, warning=FALSE}
func_problem5A(510, 3, 10000)
```

```

Problem 5b

Show that the marginal for x is proportional to $(1 - \exp(-Bx))/x$. Compare your results in 5a to this curve.

```

$$
\begin{aligned}
p_{X|Y}(x|y) &= \frac{p_{XY}(x,y)}{p_Y(y)} \\
&= \frac{\int_0^B p_{XY}(x,y) dy}{\int_0^B p_Y(y) dy} \\
&= \frac{\int_0^B \frac{1}{x} e^{-xy} dy}{\int_0^B \frac{1}{y} e^{-y} dy} \\
&= \frac{1}{x} \left[ -e^{-xy} \right]_{y=0}^{y=B} \\
&= \frac{1}{x} (1 - e^{-Bx})
\end{aligned}

```

See above for the histogram of the marginals with the curves. Note the curves are normalized by dividing by B so that both the histogram density and

\newpage

Problem 5c

Repeat 5a and 5b using $B = \infty$. Describe what happens. Is the marginal for x a proper density in this case?

```

$$
\lim_{B \rightarrow \infty} \frac{1 - e^{-Bx}}{B} = \frac{1}{x}

```

This is not a proper marginal.

```

$$
\int_0^\infty \frac{1}{x} dx = \infty

```

```

```{r p5.func_prob5B}

```

```

func_margInfy <- function(k, B){

```

```

 scales::rescale(marginal(k, B), to=c(0, 1/B))
}

func_problem5C <- function(startseed, B, it){
 set.seed(startseed)
 df <- func_gibbs.p5(B, it)

true.cdf <- stat_function(fun = func_margInfty, args = B, lwd = 1, linetype="solid", col="maroon")

plot.x <- ggplot(df, aes(x)) +
 xlim(0, B)+
 ylim(0, 1/B)+
 geom_histogram(aes(y=..density..), alpha=0.4) +
 true.cdf +
 xlab("X Chain Values")+
 ggtitle("Marginal of X: Histogram and True Curve")

plot.y <- ggplot(df, aes(y)) +
 xlim(0, B)+
 ylim(0, 1/B)+
 geom_histogram(aes(y=..density..), alpha=0.4) +
 true.cdf +
 xlab("Y Chain Values")+
 ggtitle("Marginal of Y: Histogram and True Curve")

grid.arrange(plot.x, plot.y, nrow=1)

}

...

```{r p5RESULTS, message=FALSE, warning=FALSE}
func_problem5C(530, 1e08, 100000)
```

```{r PRINTCODE}
#PRINTING THE CODE
#knitr::stitch("HW06.Rmd") to go to latex
#knitr::stitch(  script="STAT457-FINAL.Rmd" , system.file("misc", "knitr-template.Rhtml", package="knitr")) #code to HTML
```

Error: <text>:10:3: unexpected input
9: header-includes:
10: - \
^

```

The R session information (including the OS info, R version and all packages used):

```

sessionInfo()

R version 3.6.1 (2019-07-05)
Platform: x86_64-w64-mingw32/x64 (64-bit)
Running under: Windows 10 x64 (build 17763)
##
Matrix products: default
##
locale:
[1] LC_COLLATE=English_United States.1252 LC_CTYPE=English_United States.1252
[3] LC_MONETARY=English_United States.1252 LC_NUMERIC=C
[5] LC_TIME=English_United States.1252
##
attached base packages:
[1] grid stats graphics grDevices utils datasets methods base
##
other attached packages:
[1] invgamma_1.1 MCMCpack_1.4-4 MASS_7.3-51.4 coda_0.19-3
[5] gtable_0.3.0 numDeriv_2016.8-1.1 matlab_0.9.2 readr_1.3.1
[9] dplyr_0.8.3 latex2exp_0.4.0 gridExtra_2.3 ggplot2_3.2.1
[13] knitr_1.24
##
loaded via a namespace (and not attached):
[1] httr_1.4.1 jsonlite_1.6 viridisLite_0.3.0
[4] carData_3.0-2 shiny_1.3.2 assertthat_0.2.1
[7] highr_0.8 cellranger_1.1.0 yaml_2.2.0
[10] pillar_1.4.2 backports_1.1.4 lattice_0.20-38
[13] quantreg_5.51 glue_1.3.1 digest_0.6.20
[16] manipulateWidget_0.10.0 promises_1.0.1 rvest_0.3.4
[19] colorspace_1.4-1 htmltools_0.3.6 httpuv_1.5.1
[22] Matrix_1.2-17 pkgconfig_2.0.2 SparseM_1.77
[25] haven_2.1.1 purrr_0.3.2 xtable_1.8-4
[28] scales_1.0.0 webshot_0.5.1 openxlsx_4.1.0.1
[31] later_0.8.0 rio_0.5.16 MatrixModels_0.4-1
[34] tibble_2.1.3 car_3.0-3 withr_2.1.2
[37] lazyeval_0.2.2 magrittr_1.5 crayon_1.3.4
[40] readxl_1.3.1 mime_0.7 mcmc_0.9-6
[43] evaluate_0.14 forcats_0.4.0 xml2_1.2.2
[46] foreign_0.8-71 tools_3.6.1 data.table_1.12.4
[49] hms_0.5.0 stringr_1.4.0 munsell_0.5.0
[52] zip_2.0.4 kableExtra_1.1.0 compiler_3.6.1
[55] rlang_0.4.0 rstudioapi_0.10 htmlwidgets_1.5.1
[58] crosstalk_1.0.0 miniUI_0.1.1.1 labeling_0.3
[61] rmarkdown_1.14 abind_1.4-5 curl_4.2
[64] R6_2.4.0 zeallot_0.1.0 stringi_1.4.3

```

|                                  |            |              |
|----------------------------------|------------|--------------|
| ## [67] Rcpp_1.0.2               | vctr_0.2.0 | rgl_0.100.30 |
| ## [70] tidyselect_0.2.5         | xfun_0.8   |              |
| Sys.time()                       |            |              |
| ## [1] "2019-12-11 12:08:17 CST" |            |              |