hw7.R.

mwilde

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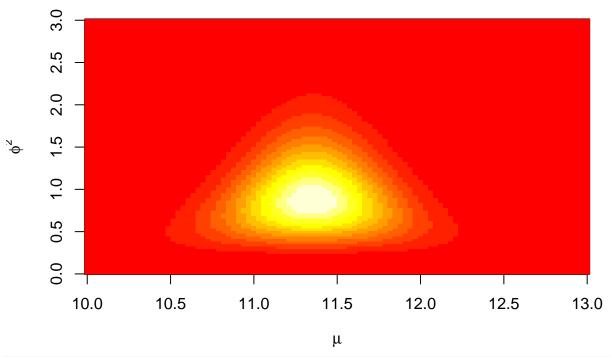
```
### 1
### (e)
### posterior mean
print(mean(rgamma(10000, shape=36, rate=11)))
## [1] 3.278267
### (f)
quantile(rgamma(10000, shape=36, rate=11),c(0.025,0.5,0.975))
                 50%
                         97.5%
## 2.293547 3.245206 4.410697
print(c("lambda left = ", qgamma(0.025, shape = 36, rate=11)))
## [1] "lambda left = " "2.29217794702866"
print(c("lambda r = ", qgamma(0.975, shape = 36, rate=11)))
## [1] "lambda r = "
                           "4.42513885173482"
### 2
### (a)
### Use discrete approximation method to get posterior
y = c(9.95, 10.47, 10.68, 10.90, 11.11, 11.22, 11.53, 11.86, 12.31, 13.51)
mu0 = 10 #mean of prior distribution
tau20 <- 0.01 # precision (or 1/variance) of prior distribution
nu0 <- 1
phi20 <- 1
# compute sample statistics
mean.y <- mean(y)</pre>
var.y <- var(y)</pre>
n <- length(y)
print(c("mean"=mean.y, "var"=var.y, "n"=n))
       mean
                 var
## 11.35400 1.03776 10.00000
G <- 100
H <- 100
mu.grid <- seq(10,13,length=G)</pre>
phi2.grid \leftarrow seq(0.01,3,length=H)
# Note: trial and error may be needed to find the
# right values for these sequences, so that the
# posterior is visible.
post.grid <- matrix(nrow=G,ncol=H) #to store posterior</pre>
for(g in 1:G){
```

```
for(h in 1:H){
   post.grid[g,h] =
        dnorm(mu.grid[g],mu0,1/sqrt(tau20))*
        dgamma(phi2.grid[h],nu0/2,nu0/(2*phi20))*
        prod(dnorm(y,mu.grid[g],1/sqrt(phi2.grid[h])))
}

post.grid <- post.grid / sum(post.grid) # normalizing

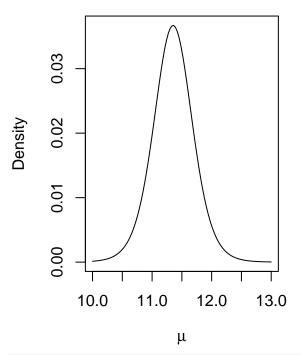
### (i)
### heat map of joint posterior
image(mu.grid,phi2.grid,post.grid,
        main=expression(paste("Heat map for Posterior on (", mu, ",", phi^2, ")")),
        xlab=expression(mu),ylab=expression(phi^2))</pre>
```

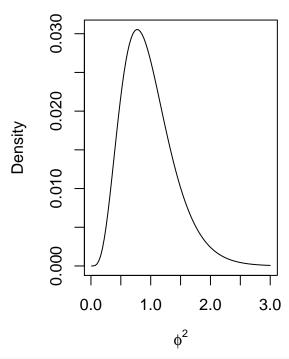
Heat map for Posterior on (μ, ϕ^2)



Marginal posterior for μ

Marginal posterior for ϕ^2

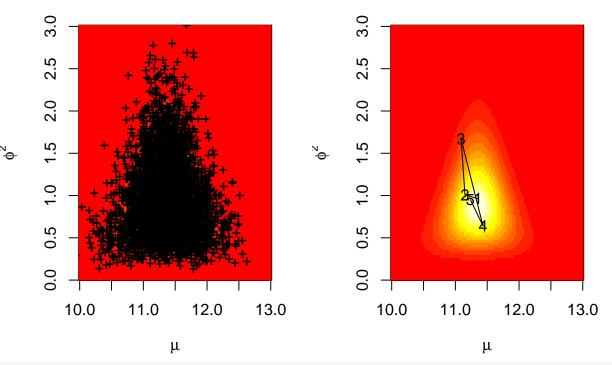




```
### b) Gibbs sampler:
# Specify chain length
S <- 4000 # generate 4000 dependent samples
samples <- matrix(nrow=S,ncol=2) # to store samples</pre>
# col 1 is mu samples
# col 2 is phi^2 samples
### Starting values
samples[1,] <- c(mean.y, 1/var.y)</pre>
set.seed(1) #to ensure random numbers are always the same
for(s in 2:S){
  ### Generate a new value of mu from f(mu | prev value of phi2, data)
  prev.phi2 <- samples[s-1,2]</pre>
  mustar <- (mu0*tau20+n*mean.y*prev.phi2)/(tau20 +n*prev.phi2)</pre>
  phi2star <- tau20 +n*prev.phi2
  mu.new <- rnorm(1,mustar,1/sqrt(phi2star))</pre>
  ### Generate a new value of phi2 from f(phi2 | new value of mu, data)
  astar <- (nu0 + n)/2
  bstar \leftarrow (nu0/phi20 + (n-1)*var.y + n*(mean.y-mu.new)^2)/2
  phi2.new <- rgamma(1,astar,bstar)</pre>
  samples[s,] <- c(mu.new,phi2.new)</pre>
}
### scatter plot showing these 4000 draws
image(mu.grid,phi2.grid,post.grid,
```

Heat map for Posterior on (μ, ϕ^2)

Heat map for Posterior on (μ, ϕ^2)



```
### (ii)
### Compute quantiles from marginal posterior distributions:
quantile(samples[,1],c(0.025,0.5,0.975)) # For mu

## 2.5% 50% 97.5%
## 10.63261 11.35859 12.06882

### (iii)
### Compute quantiles from marginal posterior distributions:
quantile(samples[,2],c(0.025,0.5,0.975)) # For phi ?2

## 2.5% 50% 97.5%
## 0.3078316 0.8912881 1.9898418
```

```
### (iv)
### Quantiles for the population sd
quantile(1/sqrt(samples[,2]),c(0.025,0.5,0.975)) # For sd
```

2.5% 50% 97.5% ## 0.7089094 1.0592317 1.8023682