

# STAT 8700 Homework 4

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1. Consider data from a Normal population with unknown mean  $\mu$  and variance  $\sigma^2$ . A random sample of 100 observations is taken from this population, and the sample mean and variance were calculated to be 50 and 25 respectively.

(a) If we choose to use a  $N - Inv - \chi^2(40, 0.64, 1, 16)$  prior distribution, write down the corresponding posterior distribution.

We are given the following:

$$n = 100$$

$$\bar{y} = 50$$

$$s^2 = 25$$

$$\mu_0 = 40$$

$$\kappa_0 = 25$$

$$\nu_0 = 1$$

$$\sigma_0^2 = 16$$

Now we can use these values to calculate the joint posterior distribution,  $N - Inv - \chi^2(\mu_n, \sigma_n^2 / \kappa_n; \nu_n, \sigma_n^2)$ :

$$\begin{aligned}\mu_n &= \frac{\kappa_0}{\kappa_0 + n} \mu_0 + \frac{n}{\kappa_0 + n} \bar{y} \\ &= \frac{25}{25 + 100} 40 + \frac{100}{25 + 100} 50 = 48\end{aligned}$$

$$\begin{aligned}\kappa_n &= \kappa_0 + n \\ &= 25 + 100 = 125\end{aligned}$$

$$\begin{aligned}\nu_n &= \nu_0 + n \\ &= 1 + 100 = 101\end{aligned}$$

$$\begin{aligned}\nu_n \sigma_n^2 &= \nu_0 \sigma_0^2 + (n - 1) s^2 + \frac{\kappa_0 n}{\kappa_0 + n} (\bar{y} - \mu_0)^2 \\ &= 1(16) + (100 - 1)25 + \frac{25(100)}{25 + 100} (50 - 40)^2 = 4491 \\ \sigma_n^2 &\approx 44.4653465347\end{aligned}$$

And thus our joint posterior distributin is  $N - Inv - \chi^2(48, 0.355722772278; 101, 44.4653465347)$ .

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(b) Either analytically or via simulation, construct 95% credible intervals for  $\sigma^2$  and  $\mu$ .

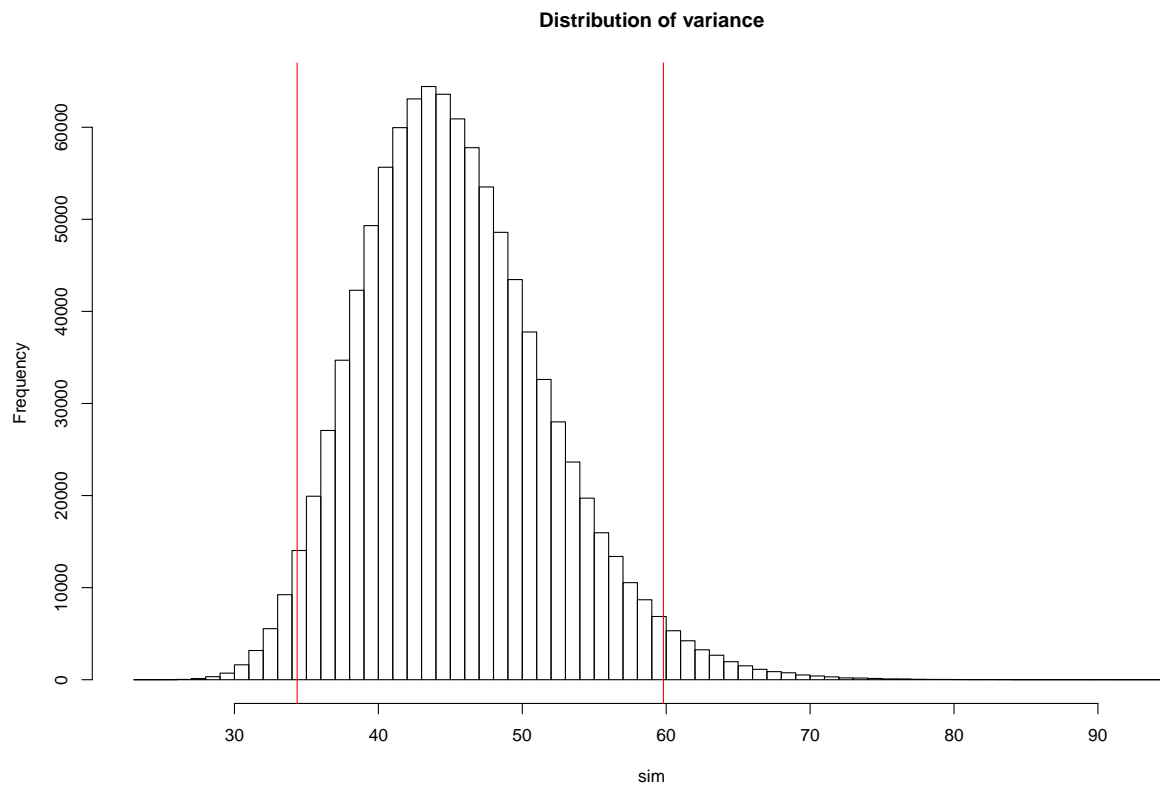
To simulate this, we first draw  $\sigma^2$  from its marginal posterior distribution,  $\sigma^2|y \sim \text{Inv} - \chi^2(\nu_n, \sigma_n^2)$

```
library(geoR)

set.seed(124)

x <- seq(0, 100, by = 0.001)
nu_n <- 101
sigma_n_2 <- 44.4653465347

sim <- rinvchisq(n = 1000000, df = nu_n, scale = sigma_n_2)
hist(sim, breaks = 90, main = 'Distribution of variance')
lower <- sort(sim)[25000]
upper <- sort(sim)[975000]
abline(v=lower, col='red')
abline(v=upper, col='red')
```



A 95% credible interval for  $\sigma^2$  is (34.3564188, 59.8085225).

Then we sample from  $N\left(\frac{\frac{\kappa_0}{\sigma^2} \mu_0 + \frac{n}{\sigma^2} \bar{y}}{\frac{\kappa_0}{\sigma^2} + \frac{n}{\sigma^2}}, \frac{1}{\frac{\kappa_0}{\sigma^2} + \frac{n}{\sigma^2}}\right)$  using the previous values for  $\sigma^2$ .

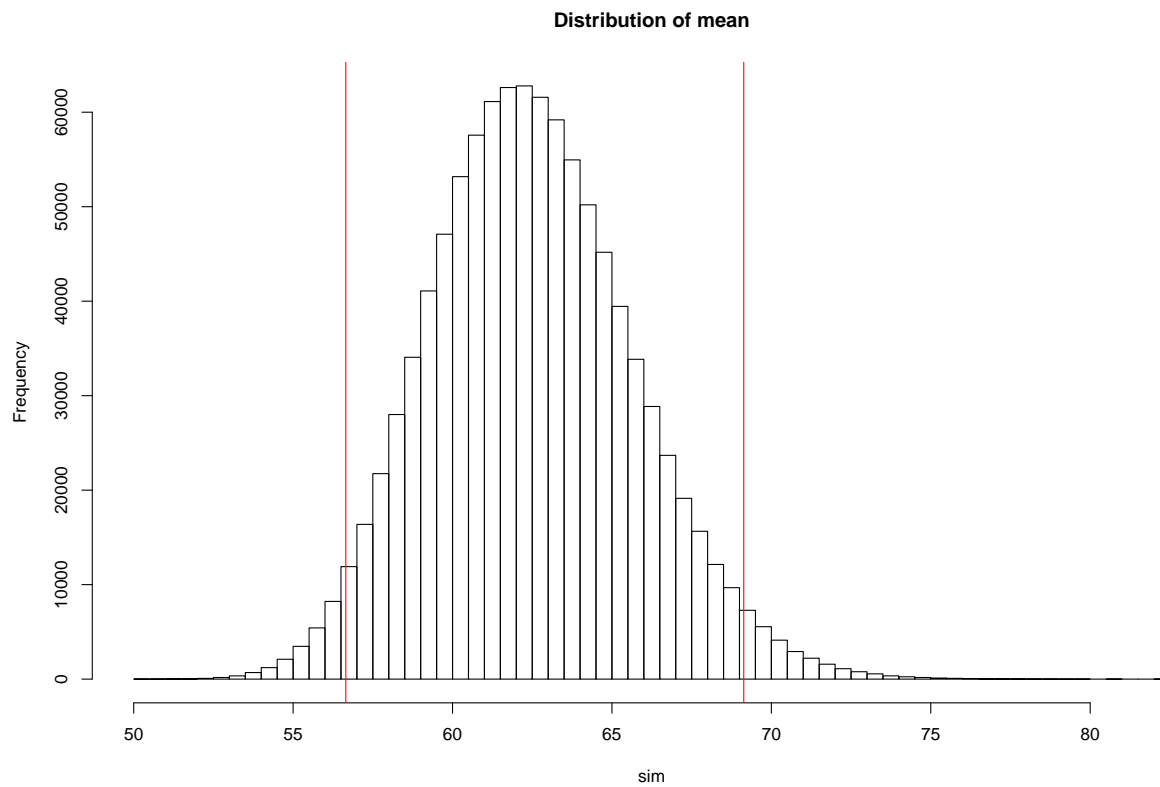
```
sigma_2 <- sim
kappa_0 <- 25
```

```

mu_0 <- 40
n <- 100
y_bar <- 50
mu_n <- ((kappa_0 / sigma_2) * mu_0) + ((n / sigma_2) * y_bar) / ((kappa_0 / sigma_2) + (n / sigma_2))
sigma_2_kappa_n <- 1 / ((kappa_0 / sigma_2) + (n / sigma_2))
sim <- rnorm(n = 1000000, mu_n, sigma_2_kappa_n)
hist(sim, breaks = 90, main = 'Distribution of mean')

lower <- sort(sim)[25000]
upper <- sort(sim)[975000]
abline(v=lower, col='red')
abline(v=upper, col='red')

```



A 95% credible interval for  $\mu$  is (56.6518857, 69.1340067).



2. Two random variables are said to have a bivariate normal distribution with parameters,  $\mu_U, \mu_V, \sigma_U^2, \sigma_V^2$ , and  $\rho$  if they have the following density function:

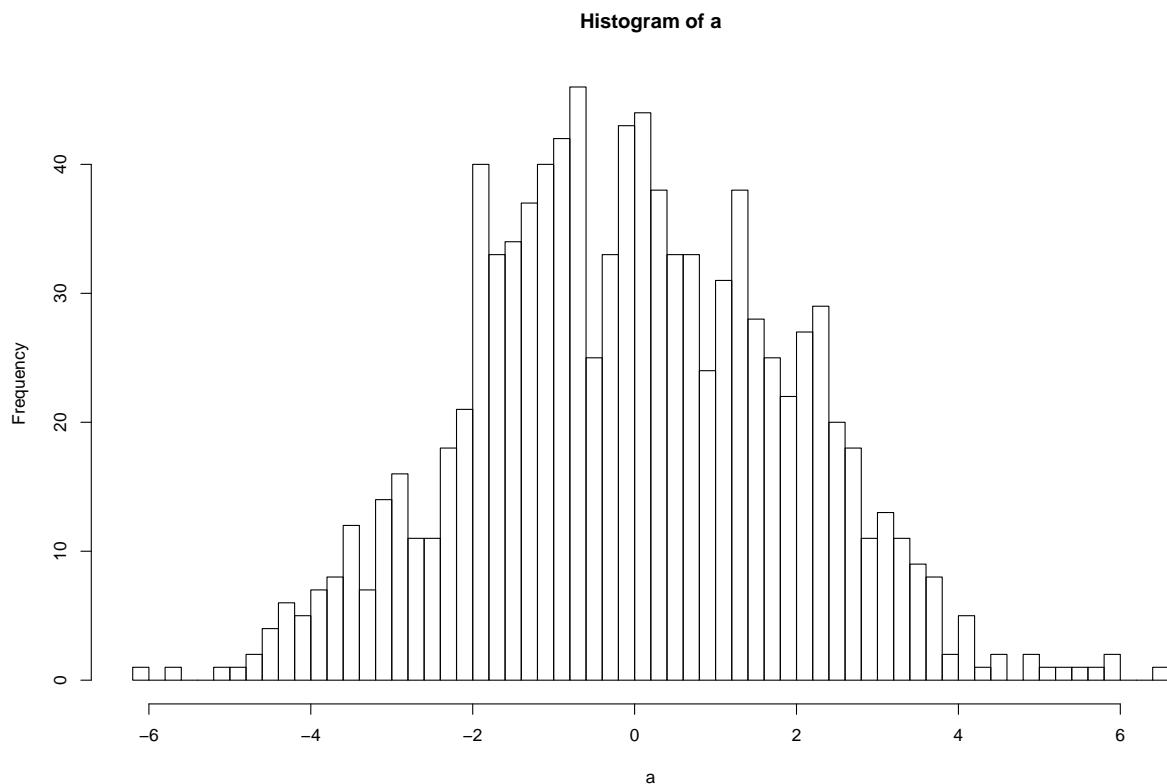
$$f(u, v) = \frac{1}{2\pi\sigma_U\sigma_V\sqrt{1-\rho^2}} e^{-\frac{1}{2(1-\rho^2)} \left[ \frac{(u-\mu_U)^2}{\sigma_U^2} + \frac{(v-\mu_V)^2}{\sigma_V^2} - \frac{2\rho(u-\mu_U)(v-\mu_V)}{\sigma_U\sigma_V} \right]}$$

where  $\mu_U$  and  $\sigma_U^2$  are the mean and variance of  $U$ ,  $\mu_V$  and  $\sigma_V^2$  are the mean and variance of  $V$ , and  $\rho$  is the correlation between  $U$  and  $V$ .

Replace the uniform prior on  $\alpha$  and  $\beta$  in the analysis of the bioassay by a bivariate normal prior with  $\alpha \sim Normal(0, 4), \beta \sim Normal(10, 100)$ , and  $corr(\alpha, \beta) = 0.5$ . Repeat all the computations and plots discussed in section 3.7 and in class.

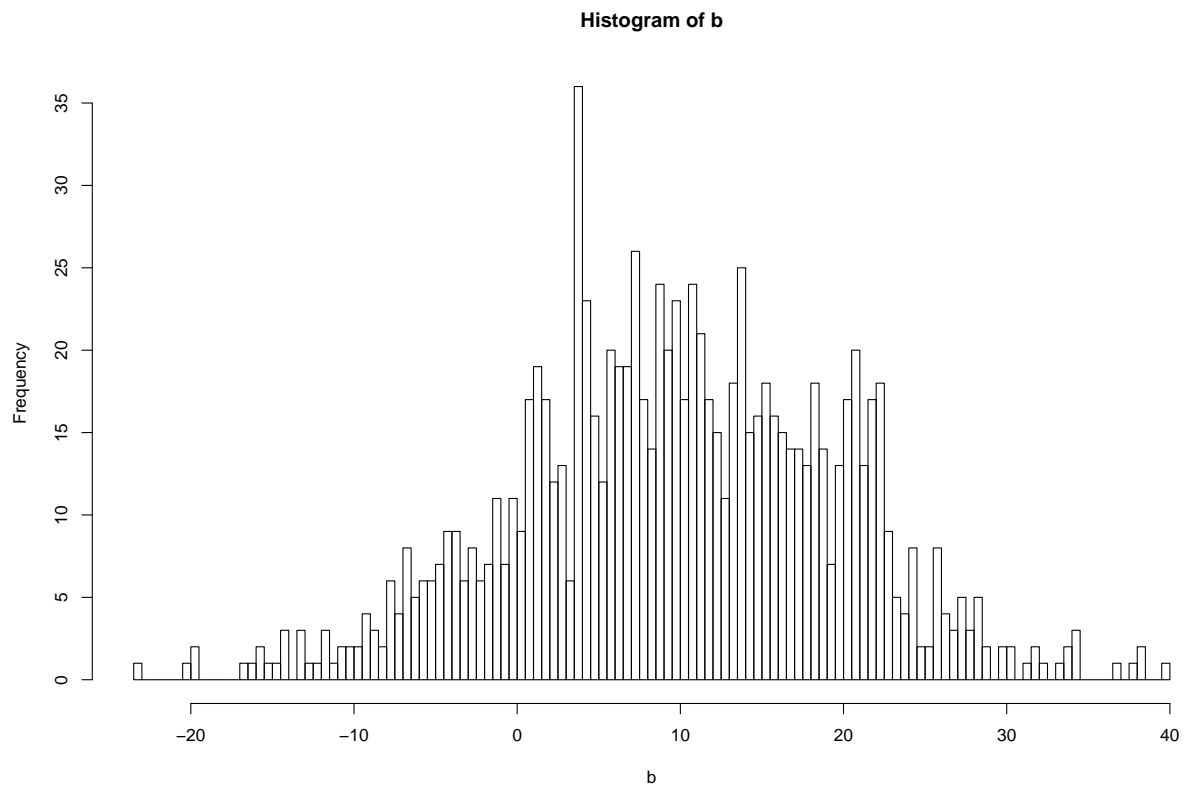
```
mu <- 0
sigma_2 <- 4

a <- rnorm(n = 1000, mu, sqrt(sigma_2))
hist(a, breaks = 90)
```

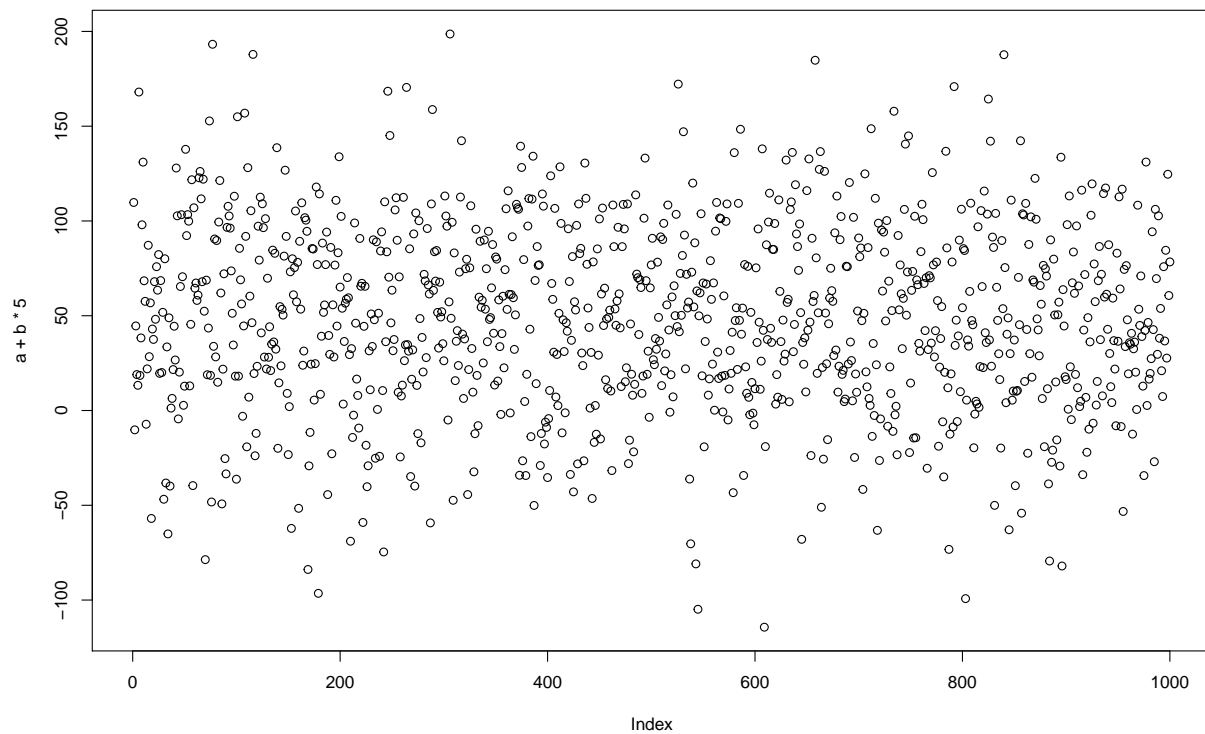


```
mu <- 10
sigma_2 <- 100

b <- rnorm(n = 1000, mu, sqrt(sigma_2))
hist(b, breaks = 90)
```



```
plot(a + b*5)
```



```
f <- function(u, v) {
  mu_U <- 0
  mu_V <- 10

  sigma_2_U <- 4
  sigma_2_V <- 100
  sigma_U <- sqrt(sigma_2_U)
  sigma_V <- sqrt(sigma_2_V)

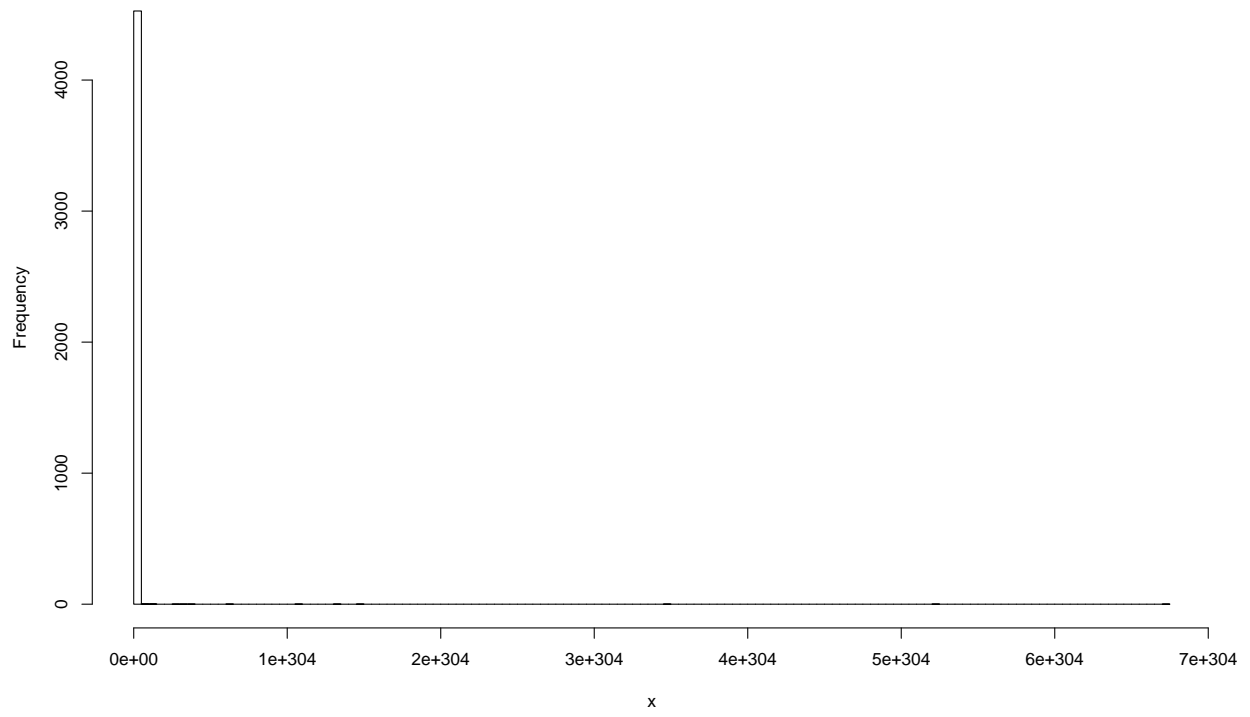
  rho <- 0.5

  first <- 1 / (2 * pi * sigma_2_U * sigma_2_V * sqrt(1 - rho^2))
  second <- exp(-(1/2*(1 - rho^2)) * (((u - mu_U)^2 / sigma_2_U
    + ((v - mu_V)^2 / sigma_2_V)
    - (2 * rho * (u - mu_U) * (v - mu_V)) / sqrt(sigma_U) * sqrt(sigma_V)))
  rval <- first * second
  return(rval)
}

x <- c()

for (i in 0:100) {
  for (j in 0:100) {
    x <- c(x, f(i, j))
  }
}
hist(x, breaks=100)
```

Histogram of x



3. Consider the airline fatalities data discussed in the previous exercise. Let us suppose that we now assume that the number of fatal accidents in year  $t$  follows a Poisson distribution with mean  $\alpha + \beta t$ .

(a) If we let  $y_t$  represent the number of fatal accidents in year  $t$ , write down  $p(y_t|\alpha, \beta)$  the likelihood for year  $t$  in terms of the parameters  $\alpha$ , and  $\beta$ .

(b) If we assume uniform priors on  $\alpha$  and  $\beta$ , write the posterior density for  $(\alpha, \beta)$ .

(c) Following the same idea as the boassay example (and the previous question) create a grid of possible  $\alpha$  and  $\beta$  values on which to evaluate the joint posterior and plot the contours. Start with large ranges for  $\alpha$  and  $\beta$  and refine based on the countour plot. Include all your iterations in your answer, not just you final grid and contour plot.

(d) Simulate 100,000 values of  $\alpha$  and  $\beta$  from the joint posterior and plot the histogram of the posterior density of the expected number of fatal accidents in 1986,  $\alpha + 1986\beta$ .

(e) Use your simulated values of  $\alpha$  and  $\beta$  to simulate the number of fatal accidents in 1986. Use your simulations to construct a 95% predictive (credible) interval.

(f) Return to your simulated values of  $\beta$ , calculate (well, estimate)  $P(\beta < 0)$ , that is, the probability that the number of fatal accidents per year is decreasing.