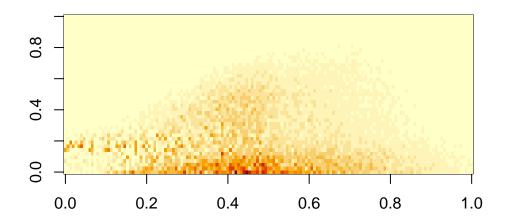
Homework Help

```
# load libraries
  library(tidyverse)
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr 1.1.3
                   v readr 2.1.4
v forcats 1.0.0
                     v stringr 1.5.0
v ggplot2 3.4.4
                   v tibble 3.2.1
v lubridate 1.9.3
                     v tidyr 1.3.0
           1.0.2
v purrr
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag()
               masks stats::lag()
i Use the conflicted package (<a href="http://conflicted.r-lib.org/">http://conflicted.r-lib.org/</a>) to force all conflicts to become
  library(mvtnorm)
  library(coda)
  library(rstanarm)
Loading required package: Rcpp
This is rstanarm version 2.26.1
- See https://mc-stan.org/rstanarm/articles/priors for changes to default priors!
- Default priors may change, so it's safest to specify priors, even if equivalent to the def
- For execution on a local, multicore CPU with excess RAM we recommend calling
  options(mc.cores = parallel::detectCores())
  yX = readRDS(
    url("http://www2.stat.duke.edu/~pdh10/Teaching/360/Materials/yXSS.rds"))
  y < -yX[,1]
```

X < -yX[,-1]

```
# image(matrix(y,151,43))
image(matrix(y,151,43))
```



$$y \sim N(X\beta, \sigma^2 I)$$

- $y: 6493 \times 1$
- X: 6493 × 9
- β : 9×1
- σ^2 : 1×1
- *I*: 6493 × 6493

$$y_i = \beta_1 x_1 + \beta_2 x_2 + \dots \beta_p x_p + \epsilon_i$$

```
set.seed(360)
# prior hyperparameters
p = 9 # number of covariates
Sigma0 = 1 * diag(rep(1, p)) # p dimension since no intercept.
b0 = rep(1/9, p)
nu0 = 2
sigma02 = 1
n = length(y)

# starting values
## note: gamma = 1 / sigma^2
beta = t(rep(1/9, 9))
```

```
# values we should compute just once
  SigmaInv = solve(Sigma0)
  X2 = t(X) %*% X
  Xy = t(X) \%*\% y
  SIBO = SigmaInv %*% b0
  a = (nu0 + n) / 2
  nu0s02 = nu0 * sigma02
  ## empty objects to fill
  BETA = NULL
  GAMMA = NULL
  S = 2000
  for (s in 1:S) {
    ### UPDATE SIGMA
    SSR1 = (y - (X %*% t(beta)))
    SSRB = t(SSR1) %*% SSR1
    gamma = rgamma(1, a, ((nu0s02 + SSRB) / 2))
    ### UPDATE BETA
    V = solve(SigmaInv + (gamma * X2))
    m = V \%*\% (Xy * gamma) # simplified since b0 = 0
    beta = rmvnorm(1, mean = m, sigma = V)
    ### SAVE STATES
    GAMMA = c(GAMMA, gamma)
    BETA = rbind(BETA, beta)
  effectiveSize(BETA)
    var1
             var2
                      var3
                               var4
                                         var5
                                                  var6
                                                           var7
                                                                     var8
2000.000 2000.000 2000.000 2000.000 2000.000 2040.626 2000.000 2000.000
2000.000
  effectiveSize(GAMMA)
```

```
var1
1810.087

# hist(1 / GAMMA)
s2 = 1 / GAMMA
sum(s2 > 100)

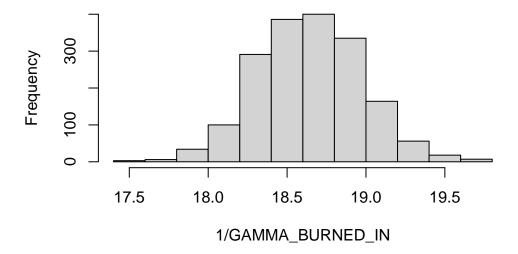
[1] 1

s2[1:5]

[1] 3086.86079   19.68145   18.75163   18.19946   18.33743

GAMMA_BURNED_IN = GAMMA[-c(1:S/10)]
hist(1 / GAMMA_BURNED_IN)
```

Histogram of 1/GAMMA_BURNED_IN



```
BETA_BURNED = BETA[-c(1:S/10),]

posteriorMeanBeta = apply(BETA_BURNED, 2, mean)
posteriorCIBeta = apply(BETA_BURNED, 2, quantile, probs = c(0.025, 0.975))
rbind(posteriorMeanBeta, posteriorCIBeta)
```

```
[,1]
                                    [,2]
                                                   [,3]
                                                               [,4]
                                                                             [,5]
posteriorMeanBeta 0.5466149 -0.01868557 0.0002884716 0.049511233 -0.0122083365
2.5%
                  0.4300168 - 0.05692517 - 0.0163582932 \ 0.004122091 - 0.0255515678
97.5%
                  0.6608110 0.02140099 0.0168541549 0.096441935 0.0009236346
                        [,6]
                                  [,7]
                                              [,8]
                                                           [,9]
posteriorMeanBeta 0.2741874 0.1606993 0.004890124 -0.08407046
                  0.1969602 0.1133077 0.002942036 -0.22878357
97.5%
                  0.3453285 0.2089253 0.006761198 0.06937184
```

Predictors 1, 6, 7.

```
colnames(X)[c(1, 6, 7)]
[1] "Effluent" "Soil"
                           "Street"
```

Alternate method

- rstanarm uses software stan
- stan lets us sample from posteriors without writing a sampler ourselves each time.
- stan uses "HMC": Hamiltonian Monte Carlo to sample from the posterior.
- one of the drawbacks of stan is that the specific prior specification we want may be difficult or impossible to obtain in the package.

```
yX_df = data.frame(yX)
post1 = stan_glm(V1 ~ 0 + ., data = yX_df, # remove the intercept
                 family = gaussian(link = "identity"),
                 seed = 360,
                 prior = normal(1/9, 1), # sets the beta prior
                 prior_aux = NULL) # set a flat prior on sigma
```

```
SAMPLING FOR MODEL 'continuous' NOW (CHAIN 1).
Chain 1:
Chain 1: Gradient evaluation took 6.6e-05 seconds
Chain 1: 1000 transitions using 10 leapfrog steps per transition would take 0.66 seconds.
Chain 1: Adjust your expectations accordingly!
Chain 1:
Chain 1:
Chain 1: Iteration:
                       1 / 2000 [ 0%]
                                         (Warmup)
Chain 1: Iteration: 200 / 2000 [ 10%]
                                         (Warmup)
Chain 1: Iteration: 400 / 2000 [ 20%]
                                         (Warmup)
```

```
Chain 1: Iteration: 600 / 2000 [ 30%]
                                         (Warmup)
Chain 1: Iteration: 800 / 2000 [ 40%]
                                         (Warmup)
Chain 1: Iteration: 1000 / 2000 [ 50%]
                                         (Warmup)
Chain 1: Iteration: 1001 / 2000 [ 50%]
                                         (Sampling)
Chain 1: Iteration: 1200 / 2000 [ 60%]
                                         (Sampling)
Chain 1: Iteration: 1400 / 2000 [ 70%]
                                         (Sampling)
Chain 1: Iteration: 1600 / 2000 [ 80%]
                                         (Sampling)
Chain 1: Iteration: 1800 / 2000 [ 90%]
                                         (Sampling)
Chain 1: Iteration: 2000 / 2000 [100%]
                                         (Sampling)
Chain 1:
Chain 1: Elapsed Time: 0.75 seconds (Warm-up)
                        0.993 seconds (Sampling)
Chain 1:
Chain 1:
                        1.743 seconds (Total)
Chain 1:
SAMPLING FOR MODEL 'continuous' NOW (CHAIN 2).
Chain 2:
Chain 2: Gradient evaluation took 1.3e-05 seconds
Chain 2: 1000 transitions using 10 leapfrog steps per transition would take 0.13 seconds.
Chain 2: Adjust your expectations accordingly!
Chain 2:
Chain 2:
Chain 2: Iteration:
                       1 / 2000 [ 0%]
                                         (Warmup)
Chain 2: Iteration: 200 / 2000 [ 10%]
                                         (Warmup)
Chain 2: Iteration: 400 / 2000 [ 20%]
                                         (Warmup)
Chain 2: Iteration: 600 / 2000 [ 30%]
                                         (Warmup)
Chain 2: Iteration: 800 / 2000 [ 40%]
                                         (Warmup)
Chain 2: Iteration: 1000 / 2000 [ 50%]
                                         (Warmup)
Chain 2: Iteration: 1001 / 2000 [ 50%]
                                         (Sampling)
Chain 2: Iteration: 1200 / 2000 [ 60%]
                                         (Sampling)
Chain 2: Iteration: 1400 / 2000 [ 70%]
                                         (Sampling)
Chain 2: Iteration: 1600 / 2000 [ 80%]
                                         (Sampling)
Chain 2: Iteration: 1800 / 2000 [ 90%]
                                         (Sampling)
Chain 2: Iteration: 2000 / 2000 [100%]
                                         (Sampling)
Chain 2:
Chain 2: Elapsed Time: 1.109 seconds (Warm-up)
Chain 2:
                        1.072 seconds (Sampling)
Chain 2:
                        2.181 seconds (Total)
Chain 2:
SAMPLING FOR MODEL 'continuous' NOW (CHAIN 3).
Chain 3:
Chain 3: Gradient evaluation took 1.4e-05 seconds
```

```
Chain 3: 1000 transitions using 10 leapfrog steps per transition would take 0.14 seconds.
Chain 3: Adjust your expectations accordingly!
Chain 3:
Chain 3:
Chain 3: Iteration:
                       1 / 2000 [ 0%]
                                         (Warmup)
Chain 3: Iteration: 200 / 2000 [ 10%]
                                         (Warmup)
Chain 3: Iteration: 400 / 2000 [ 20%]
                                         (Warmup)
Chain 3: Iteration: 600 / 2000 [ 30%]
                                         (Warmup)
Chain 3: Iteration: 800 / 2000 [ 40%]
                                         (Warmup)
Chain 3: Iteration: 1000 / 2000 [ 50%]
                                         (Warmup)
Chain 3: Iteration: 1001 / 2000 [ 50%]
                                         (Sampling)
Chain 3: Iteration: 1200 / 2000 [ 60%]
                                         (Sampling)
Chain 3: Iteration: 1400 / 2000 [ 70%]
                                         (Sampling)
Chain 3: Iteration: 1600 / 2000 [ 80%]
                                         (Sampling)
Chain 3: Iteration: 1800 / 2000 [ 90%]
                                         (Sampling)
Chain 3: Iteration: 2000 / 2000 [100%]
                                         (Sampling)
Chain 3:
Chain 3: Elapsed Time: 0.862 seconds (Warm-up)
Chain 3:
                        1.101 seconds (Sampling)
Chain 3:
                        1.963 seconds (Total)
Chain 3:
SAMPLING FOR MODEL 'continuous' NOW (CHAIN 4).
Chain 4:
Chain 4: Gradient evaluation took 3.8e-05 seconds
Chain 4: 1000 transitions using 10 leapfrog steps per transition would take 0.38 seconds.
Chain 4: Adjust your expectations accordingly!
Chain 4:
Chain 4:
Chain 4: Iteration:
                       1 / 2000 [ 0%]
                                         (Warmup)
Chain 4: Iteration: 200 / 2000 [ 10%]
                                         (Warmup)
Chain 4: Iteration: 400 / 2000 [ 20%]
                                         (Warmup)
Chain 4: Iteration: 600 / 2000 [ 30%]
                                         (Warmup)
Chain 4: Iteration: 800 / 2000 [ 40%]
                                         (Warmup)
Chain 4: Iteration: 1000 / 2000 [ 50%]
                                         (Warmup)
Chain 4: Iteration: 1001 / 2000 [ 50%]
                                         (Sampling)
Chain 4: Iteration: 1200 / 2000 [ 60%]
                                         (Sampling)
Chain 4: Iteration: 1400 / 2000 [ 70%]
                                         (Sampling)
Chain 4: Iteration: 1600 / 2000 [ 80%]
                                         (Sampling)
Chain 4: Iteration: 1800 / 2000 [ 90%]
                                         (Sampling)
Chain 4: Iteration: 2000 / 2000 [100%]
                                         (Sampling)
Chain 4:
Chain 4: Elapsed Time: 1.309 seconds (Warm-up)
```

```
Chain 4: 1.013 seconds (Sampling)
Chain 4: 2.322 seconds (Total)
Chain 4:
```

cbind(post1\$coefficients, posteriorMeanBeta)

posterior Mean BetaEffluent 0.5489818902 0.5466148619 Influent -0.0193440302 -0.0186855712 Poultry 0.0004494069 0.0002884716 Reference 0.0511734579 0.0495112329 Septic -0.0117853543 -0.0122083365 Soil 0.2742320104 0.2741873716 Street 0.1588910894 0.1606993061 Swine 0.0048560389 0.0048901240 Wetland -0.0855007899 -0.0840704585

```
mean(sqrt(1 / GAMMA_BURNED_IN))
```

[1] 4.317466