

```

#gamma prior for lambda
alpha0 <- 5
beta0 <- 0.5
#data model parameters
lambdad <- 9
nd <- 20

mcsiz=2000
mcnt <-0
mcestg <- c()
mcesti<- c()
CIcovg <- rep(0,times=mcsiz)
CIcovi <- rep(0,times=mcsiz)

#Monte Carlo Simulation of CIs
repeat{
  mcnt <- mcnt+1

  #draw lambda from gamma prior
  lam0 <- rgamma(1, shape=alpha0, rate=beta0)

  #generate from data model
  poisi <- rpois(n=nd, lambda=lambdad)

  #generate estimates of lambda
  mcestg[mcnt] <- (alpha0+sum(poisi))/(nd+beta0)
  mcesti[mcnt] <- (sum(poisi)+1)/nd

  #CI coverage estimates
  CIlg <- qgamma(.05, shape=alpha0+sum(poisi),rate=nd+beta0)
  CIug <- qgamma(.95, shape=alpha0+sum(poisi),rate=nd+beta0)
  if(lambdad<=CIug & lambdad >= CIlg){CIcovg[mcnt]=1}
  CILI <- qgamma(.05, shape=1+sum(poisi),rate=nd)
  CIUI <- qgamma(.95, shape=1+sum(poisi),rate=nd)
  if(lambdad<=CIUI & lambdad >= CILI){CIcovi[mcnt]=1}

  if(mcnt==mcsiz){break}
}
#####
####CI coverages####
mean(CIcovg)
mean(CIcovi)

#####
##### MC CIs #####
#gamma prior
#MC variance and standard error
mcvarg <- (mcsiz*(mcsiz-1))^( -1)*sum((mcestg-mean(mcestg))^2)
mcseg <- sqrt(mcvarg)
#lower and upper bounds for the MC conf. interval
mcgCIL <- mean(mcestg) - qnorm(.975)*mcseg
mcgCIU <- mean(mcestg) + qnorm(.975)*mcseg
#center of the interval
mean(mcestg)
mcgCIL; mcgCIU
#CI width
widthg <- mcgCIU-mcgCIL; widthg

#uniform prior
#MC variance and standard error
mcvari <- (mcsiz*(mcsiz-1))^( -1)*sum((mcesti-mean(mcesti))^2)
mcsei <- sqrt(mcvari)
#lower and upper bounds for the MC conf. interval
mciCIL <- mean(mcesti) - qnorm(.975)*mcsei
mciCIU <- mean(mcesti) + qnorm(.975)*mcsei
#center of the interval
mean(mcesti)
mciCIL; mciCIU
#CI width
widthi <- mciCIU-mciCIL; widthi
#####
#####
#####
#Bayesian Data Analysis
#####
library(R2jags)
library(coda)

clouds <- read.csv("CloudData.csv")

#This is the WinBUGS code for the priors and model specification
model{
  for(i in 1:n){
    y[i]~dgamma(alpha, beta[i])
    x[i]~dgamma(alpha, betac[i])
    beta[i] = alpha/mu
    betac[i] = alpha/muc
  }
  mu ~ dgamma(.001, .001)
  muc ~ dgamma(.001, .001)
  alpha ~ dt(0, pow(2.5,-2), 1)T(0,)
}

#running the model in rJAGS
JMfit <- jags( data = list(y = clouds$SeededRain, x=clouds$UnseededRain, n = length(clouds$SeededRain)), inits = NULL, model.file = "hw5wb.R",
n.chains = 3, n.iter = 110000, n.burnin = 50000, parameters.to.save = c("mu", "muc", "alpha"), n.thin = 10)
print(JMfit)

#diagnostics
traceplot(JMfit)
JMmcmc <- as.mcmc(JMfit)
gelman.plot(JMmcmc)
autocorr.plot(JMmcmc)
densplot(JMmcmc)
HPDinterval(JMmcmc)

```