

Homework 7

STA 360: Assignment 7, Fall 2020

Due Monday November 2nd, 5 PM Standard Eastern Time

Question 1

Hoff 7. 3

Part A.

```
#Read in the data sets from github
blue_crab_data = as.matrix(read.table(url('https://raw.githubusercontent.com/resteorts/modern-bayes/mas

orange_crab_data = as.matrix(read.table(url('https://raw.githubusercontent.com/resteorts/modern-bayes/m

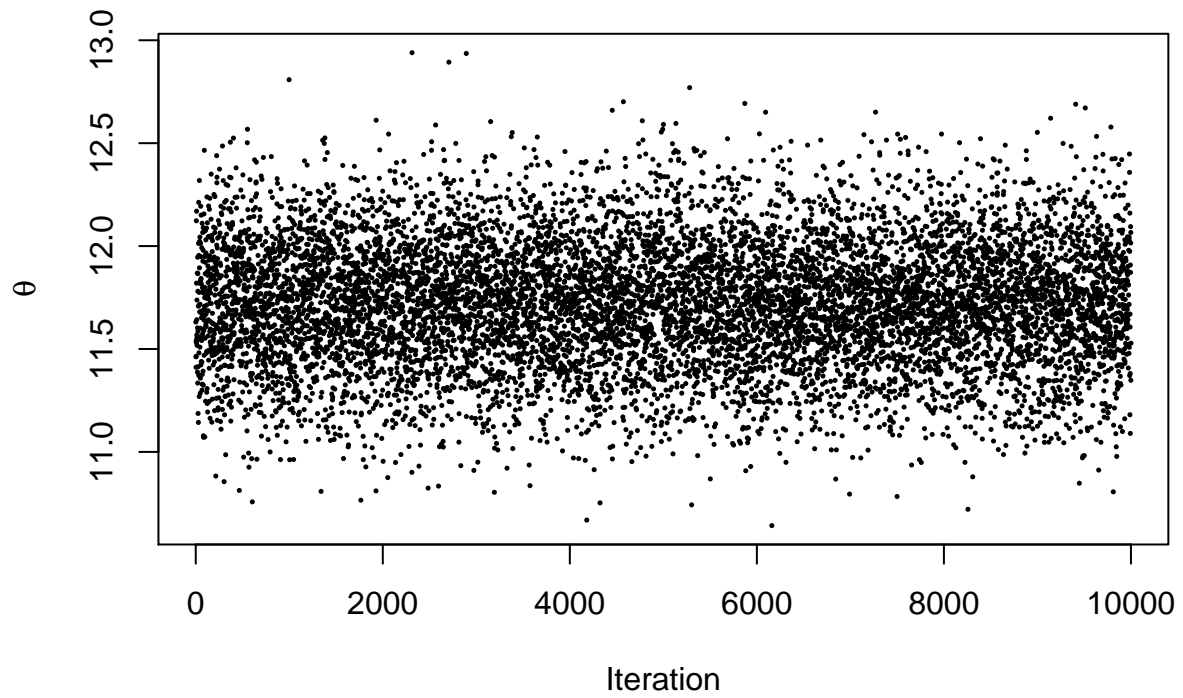
#We need to sample from the posterior for both data sets, so let's use lapply with a function
#The function is heavily based on Rebecca Steort's code in Module 8
crab.mcmc = lapply(list('blue_crab' = blue_crab_data, 'orange_crab' = orange_crab_data), function(crab)
  n = nrow(crab)
  # Set prior parameters according to problem
  mu0 = colMeans(crab)
  lambda0 = cov(crab)
  s0 = cov(crab)
  nu0 = 4
  THETA <- SIGMA <- NULL
  # Start with sigma sample
  sigma = s0
  set.seed(1);
  for (s in 1:10000) {
    # Update theta
    Ln <- solve(solve(lambda0) + n * solve(sigma))
    mun <- Ln %*% (solve(lambda0) %*% mu0 + n * solve(sigma) %*% mu0)
    theta <- rmvnorm(1, mun, Ln)

    ## update sigma
    Sn <- s0 + (t(crab) - c(theta)) %*%          t(t(crab)-c(theta))
    sigma <- solve(rwish(nu0 + n, solve(Sn)))
    ## save results
    THETA <- rbind(THETA, theta)
    SIGMA <- rbind(SIGMA, c(sigma))
  }

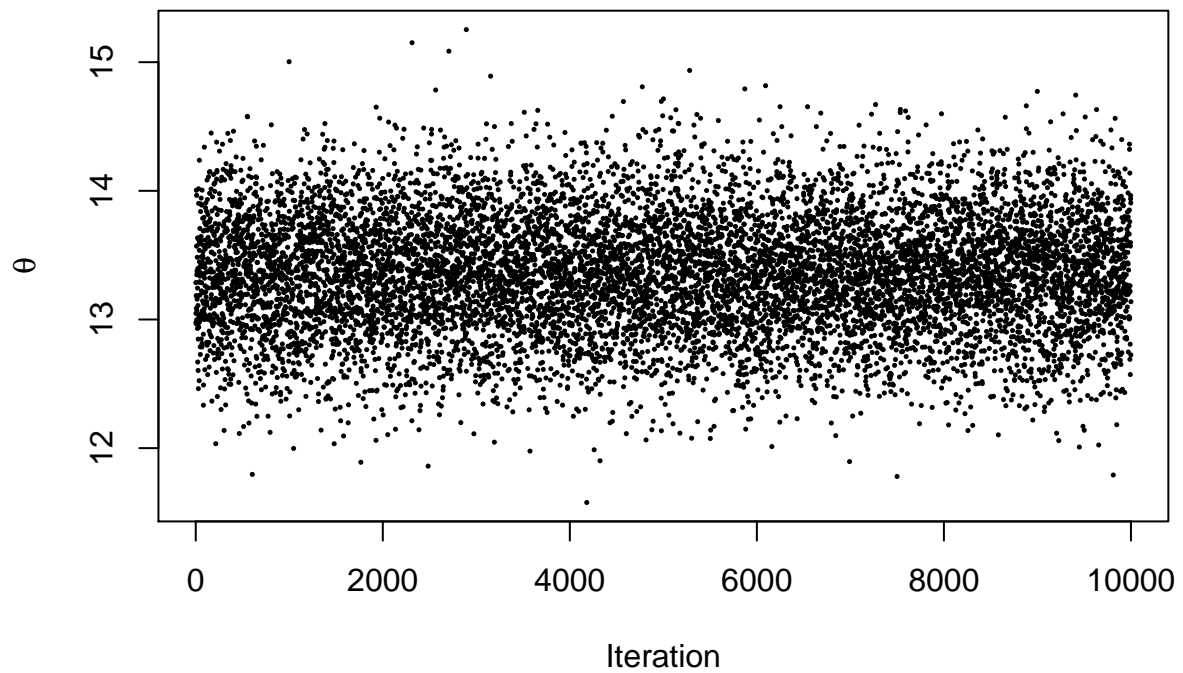
  list(theta = THETA, sigma = SIGMA)
})
```

Part B.

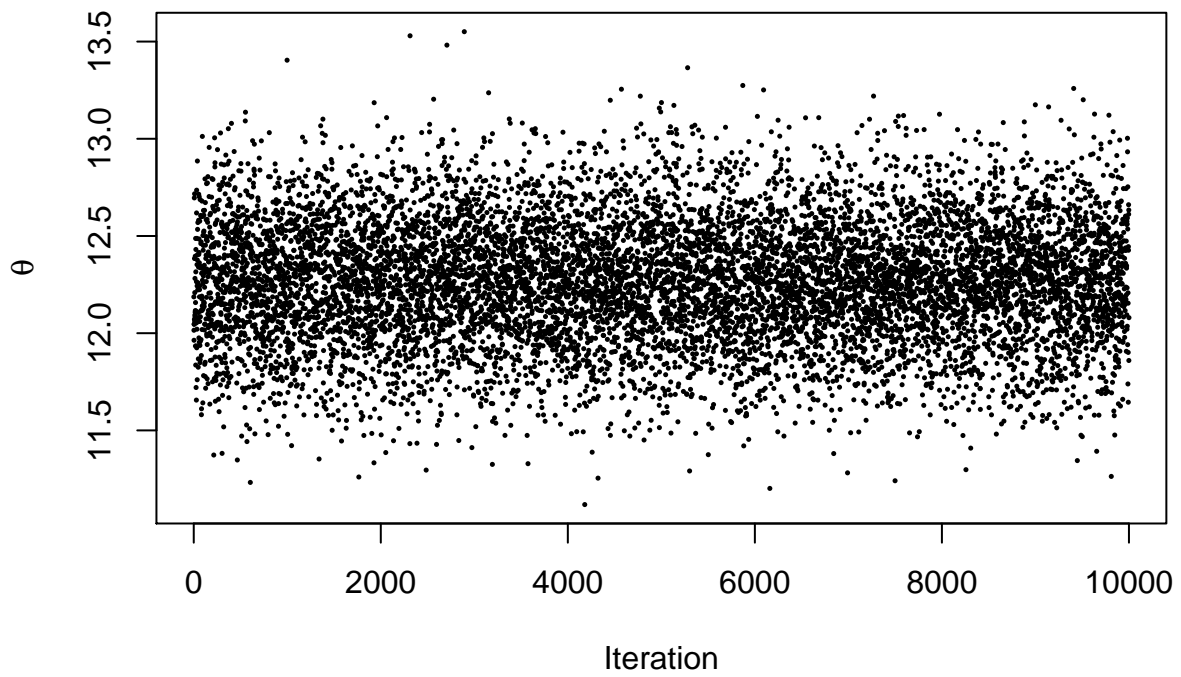
Traceplot of θ_1 for Blue Crab



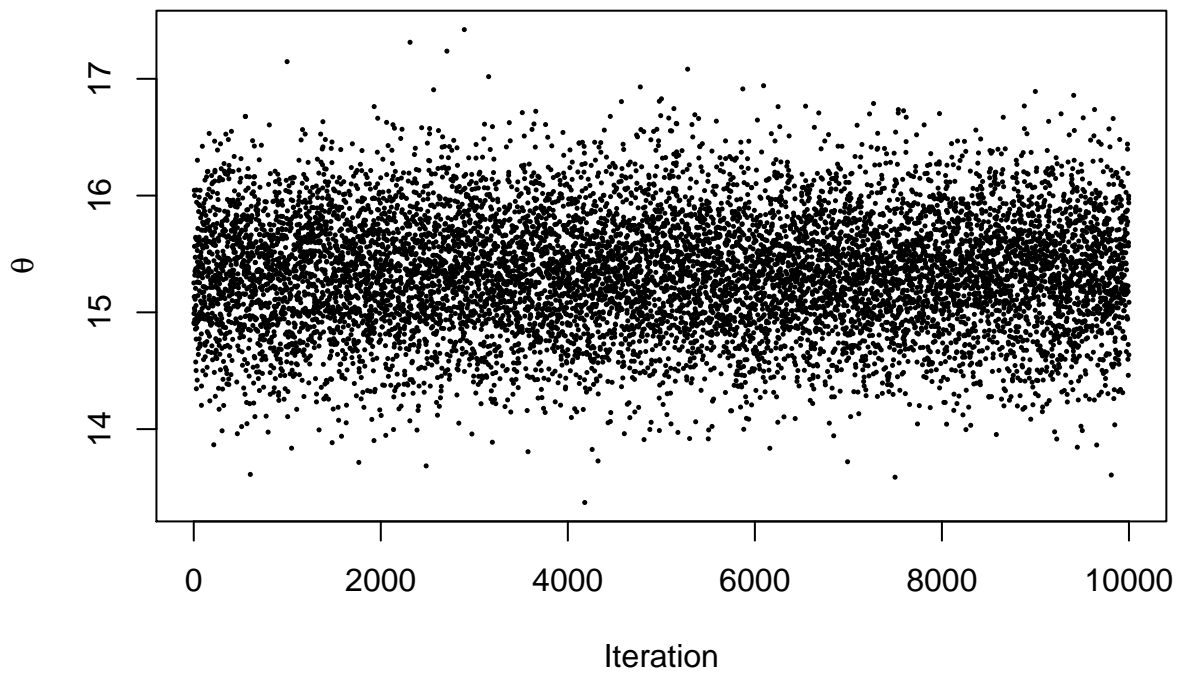
Traceplot of θ_2 for Blue Crab



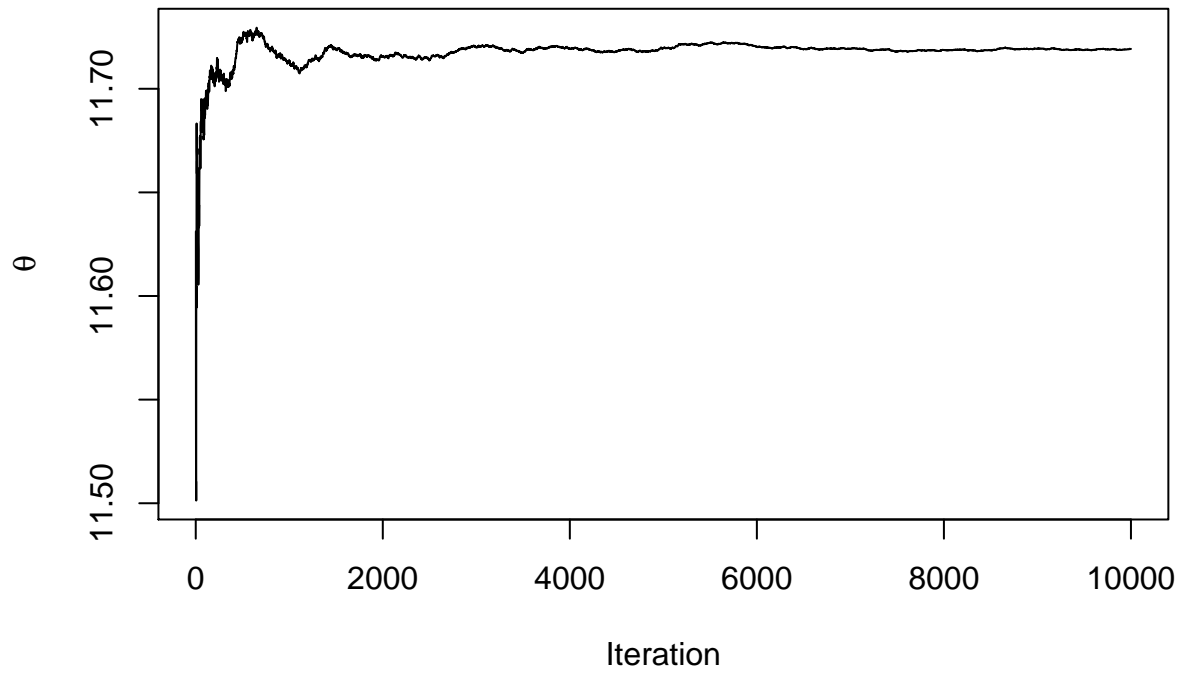
Traceplot of θ_1 for Orange Crab



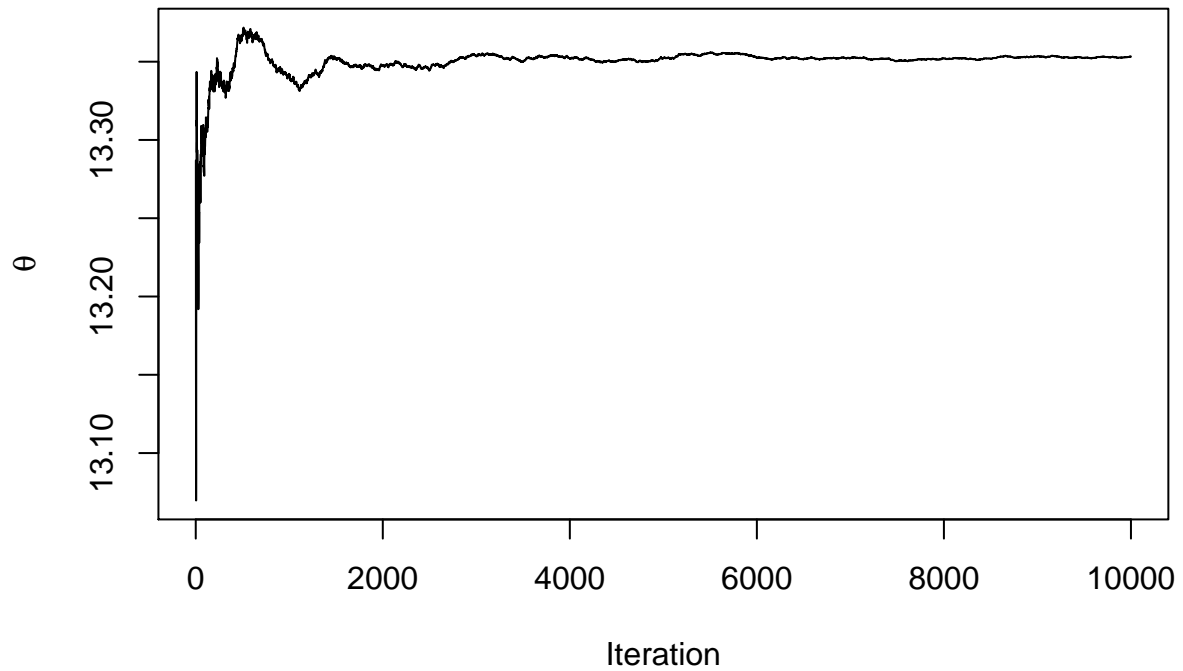
Traceplot of θ_2 for Orange Crab



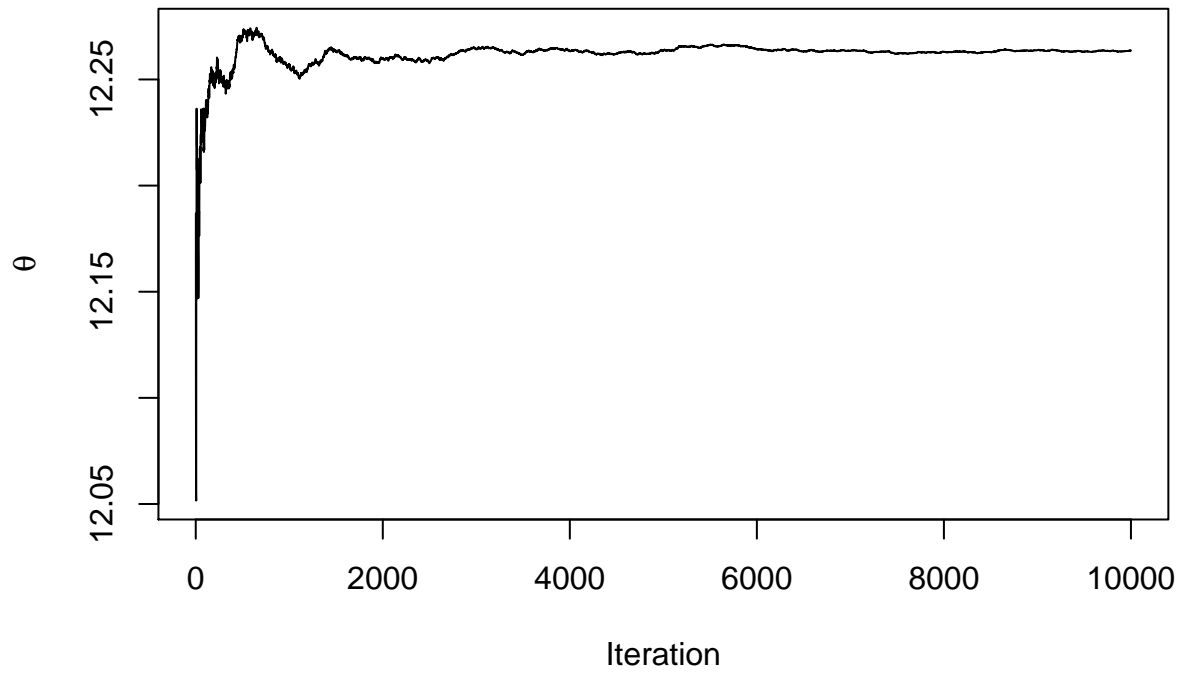
Running Average Plot of θ_1 for Blue Crab



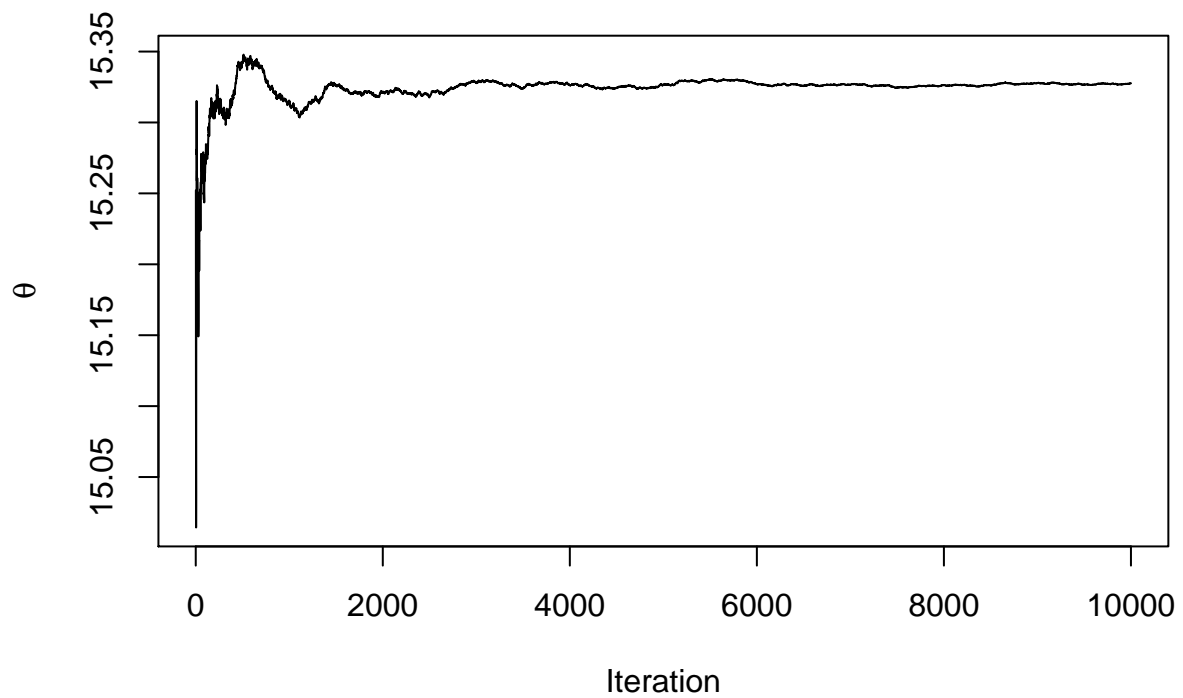
Running Average Plot of θ_2 for Blue Crab



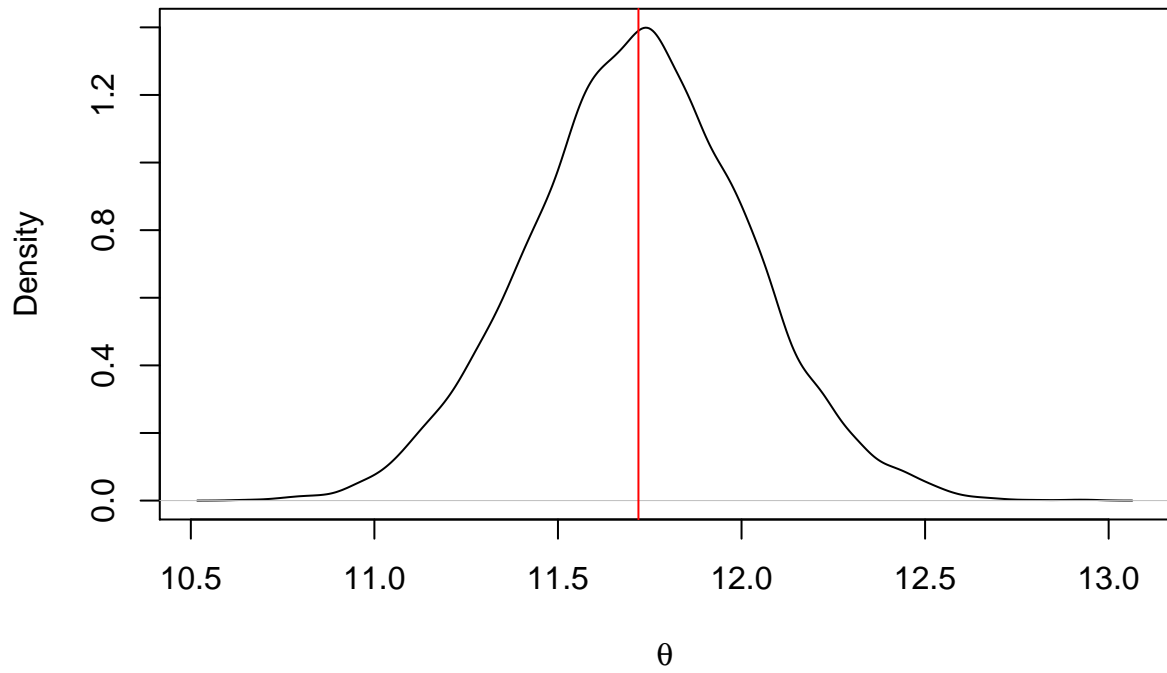
Running Average Plot of θ_1 for Orange Crab



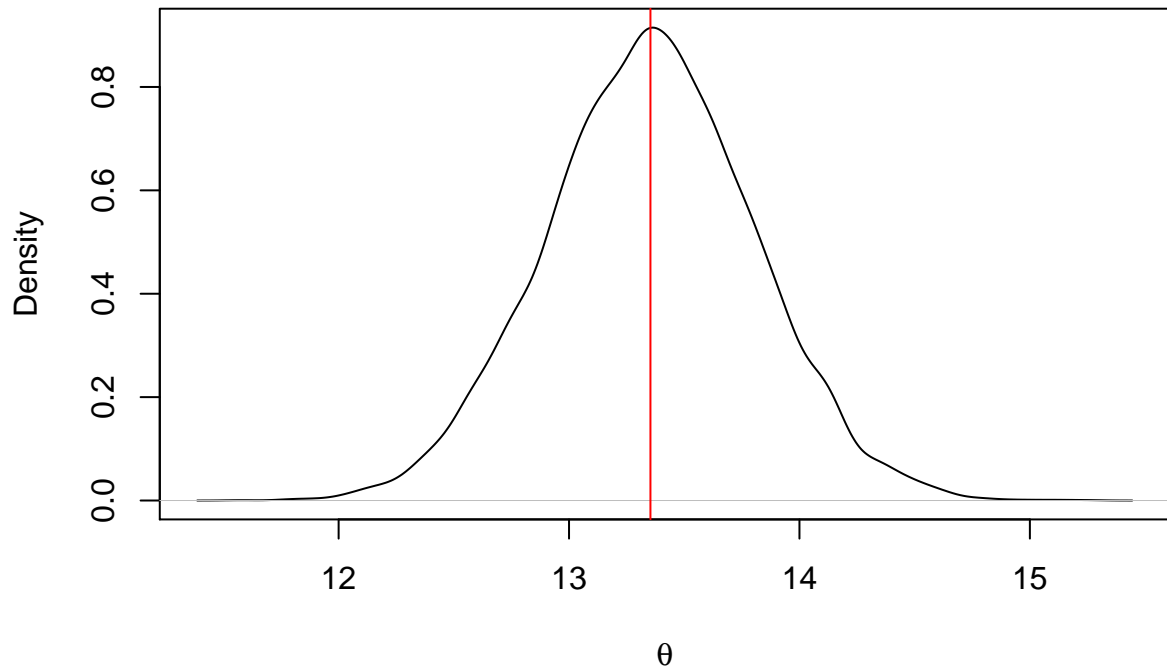
Running Average Plot of θ_2 for Orange Crab



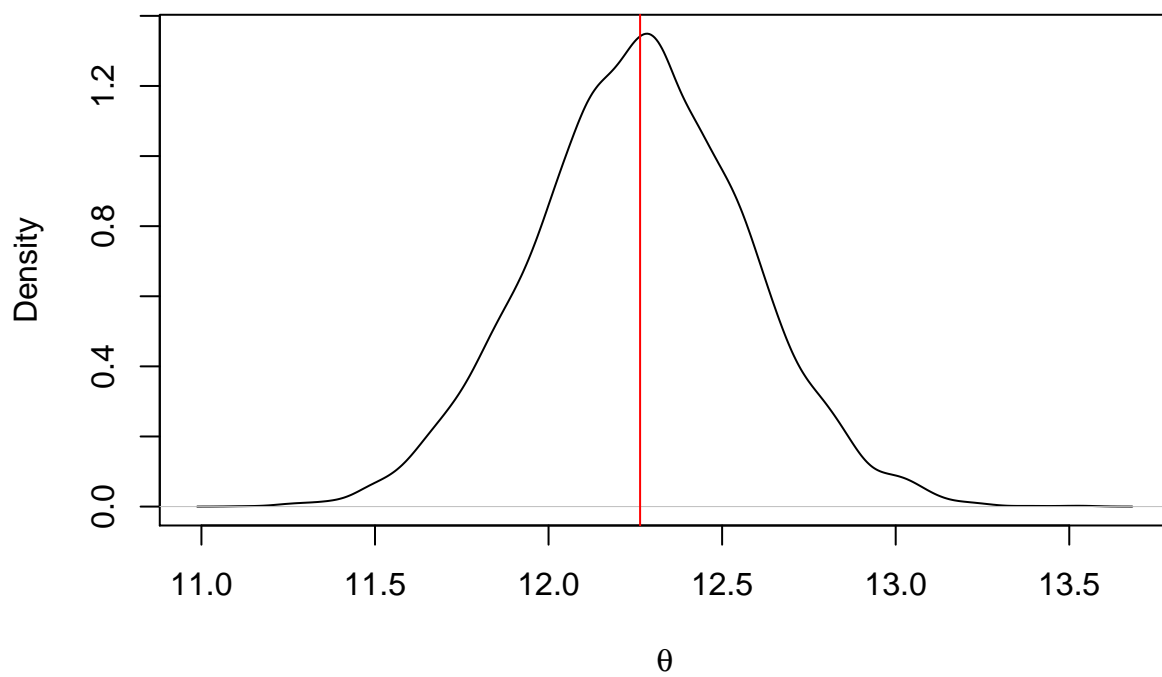
Density of θ_1 for Blue Crab



Density of θ_2 for Blue Crab



Density of θ_1 for Orange Crab



Density of θ_2 for Orange Crab

