Homework 7

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Problem 1

$$y_t = \rho y_{t-1} + \epsilon_t$$

$$\gamma_k = Cov(y_{t-k}, y_t) = \rho^k \gamma_0$$

$$y_{t-k} \cdot y_t = y_{t-k} \cdot \rho y_{t-1} + y_{t-k} \cdot \epsilon_t$$

$$E(y_{t-k} \cdot y_t) = \rho E(y_{t-k} \cdot y_{t-1}) + E(y_{t-k}) \cdot E(\epsilon_t)$$

by independence.

$$E(y_{t-k} \cdot y_t) = \rho E(y_{t-k} \cdot y_{t-1})$$

and we can keep going , by writing the right hand side as

$$E(y_{t-k} \cdot y_{t-1}) = \rho E(y_{t-k} \cdot y_{t-2}) + 0$$

Pluggin this into the above we get

$$E(y_{t-k} \cdot y_t) = \rho^2 E(y_{t-k} \cdot y_{t-2})$$

So continuing we see that

$$E(y_{t-k} \cdot y_t) = \rho^k E(y_{t-k} \cdot y_{t-k})$$

We know that

$$E(y_{t-k}^2) = Var(y_{t-k}) - E(y_{t-k})^2$$

$$E(y_{t-k}^2) = Var(y_{t-k}) - 0^2$$

because $E(y_{t-k}) = 0$.

We also know that

$$Var(y_{t-k}) = Cov(y_{t-k}, y_{t-k}) = Cov(y_t, y_t) = \gamma_0$$

So plugging back in we see,

$$E(y_{t-k} \cdot y_t) = \rho^k \gamma_0$$

Problem 2

library(R2jags)

Loading required package: rjags

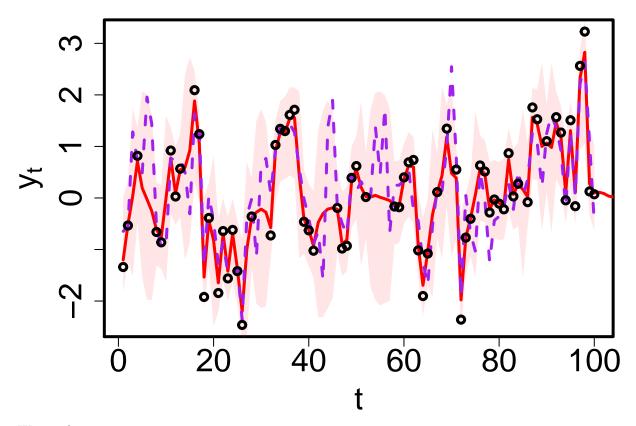
Loading required package: coda

Linked to JAGS 4.3.0

Loaded modules: basemod, bugs

```
##
## Attaching package: 'R2jags'
## The following object is masked from 'package:coda':
##
##
       traceplot
library(rjags)
GetAR <- function(nyears, # length of series</pre>
                   rho, sigma, # AR parameters
                   eps0.t = NULL, # innovations (optional)
                   ystart = NULL # starting value y1 (optional)
                   ){
  if (is.null(eps0.t)){
    set.seed(123)
    eps0.t <- rnorm(nyears, 0, 1)
  }
  y.t <- rep(NA, nyears)
  if (is.null(ystart)){
    y.t[1] <- sigma/sqrt(1-rho^2)*eps0.t[1]
  } else {
    y.t[1] <- ystart
  for (t in 2:nyears){
    y.t[t] \leftarrow rho*y.t[t-1] + sigma*eps0.t[t]
  return(y.t)
}
rho <- 0.5
sigma <- 1
sigma.y <- 0.5
nyears <- 100
mu.t <- GetAR(nyears, rho, sigma)</pre>
set.seed(124)
y.t <- mu.t + rnorm(nyears, 0, sigma.y)</pre>
model <- "
model{
mu.t[1] ~ dnorm(0, tau.stat)
tau.stat <- (1-pow(rho,2))/pow(sigma,2)</pre>
for (t in 2:(nyears+P)){
mu.t[t] ~ dnorm(muhat.t[t], tau)
muhat.t[t] <- rho*mu.t[t-1]</pre>
yhat.t[t] ~ dnorm(muhat.t[t], tau.y)
}
for (t in 1:nyears){
 y.t[t] ~ dnorm(mu.t[t], tau.y)
tau <- pow(sigma,-2)</pre>
```

```
sigma ~ dunif(0,3)
tau.y <- pow(sigma.y,-2)</pre>
sigma.y ~ dunif(0,3)
rho \sim dunif(-1,1)
}
11
P <- 20
set.seed(1234)
t.i \leftarrow sort(sample(seq(1,100), size = 70))
y.t[-t.i] \leftarrow NA
jags.data <- list(y.t = y.t, nyears=nyears,P=P)</pre>
parnames <- c("sigma", "rho", "mu.t", "sigma.y", "muhat.t", "yhat.t")</pre>
mod <- jags(data = jags.data,</pre>
            parameters.to.save= c(parnames), n.chains = 4, n.burnin = 1000, n.iter = 1000+30000, n.thin
           model.file = textConnection(model))
## module glm loaded
## Compiling model graph
      Resolving undeclared variables
      Allocating nodes
##
## Graph information:
##
      Observed stochastic nodes: 70
##
      Unobserved stochastic nodes: 272
##
      Total graph size: 475
##
## Initializing model
mcmc.array <- mod$BUGSoutput$sims.array</pre>
mcmc.list <- mod$BUGSoutput$sims.list</pre>
mean(mcmc.array[,,"mu.t[6]"])
## [1] -0.01087514
quantile(mcmc.array[,,"mu.t[6]"],c(.025,.975))
        2.5%
                  97.5%
## -1.968996 2.008788
```



We can forecast μ_t using

$$\mu_{101}^{(s)}|\mu_{100}^{(s)},\rho^{(s)},(\delta^2)^{(s)} \sim N(\mu_{100}^{(s)},(\delta^2)^{(s)})$$

and forecast y_t using

$$y_{101}^{(s)}|\mu_{101}^{(s)},(\sigma^2)^{(s)} \sim N(\mu_{101}^{(s)},(\sigma^2)^{(s)})$$

and so on as we extend in time.

```
print ("muhat[101]")

## [1] "muhat[101]"

print (mean(mcmc.array[,,"muhat.t[101]"]))

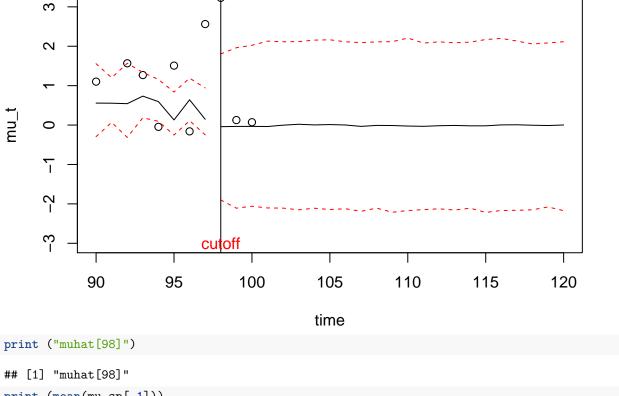
## [1] 0.08308189

print (quantile(mcmc.array[,,"muhat.t[101]"],c(.025,.975)))

## 2.5% 97.5%

## -0.3632181 0.6854923
```

 \mathbf{c}



```
## [1] "muhat[98]"
print (mean(mu.sp[,1]))
## [1] -0.04101947
print (quantile(mu.sp[,1],c(.025,.975)))
## 2.5% 97.5%
```

-1.901172 1.804929 The PI's that result from part b are smaller than those of part c because we are forecasting from later on in the time series.