Final Project 2131

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```
library(nlme)
library(orcutt)
## Loading required package: lmtest
## Loading required package: zoo
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
        as.Date, as.Date.numeric
library(DescTools)
set.seed(1122214)
xt <- rep(NA, 50)
xt[1] \leftarrow 0
for (t in 1:length(xt)) {
    if (t == 1) {
         xt[t] <- 0
         next
    }
    xt[t] \leftarrow xt[t - 1] + 0.2
}
eps <- rep(NA, 50)
rho = 0.5
beta0 = 2
beta1 = 0.5
c \leftarrow matrix(mrow = 500, mcol = 2)
b \leftarrow matrix(nrow = 500, ncol = 2)
a <- matrix(nrow = 500, ncol = 1)</pre>
z \leftarrow matrix(mrow = 500, mcol = 1)
resi <- matrix(nrow = 500, ncol = 50)</pre>
dw \leftarrow matrix(mrow = 500, mcol = 4)
rho.hat.mle <- matrix(nrow = 500, ncol = 1)</pre>
rho.hat.reml <- matrix(nrow = 500, ncol = 1)</pre>
rhoyw <- matrix(nrow = 500)</pre>
segmle <- matrix(nrow = 500)</pre>
betagmle <- matrix(nrow = 500)</pre>
sereml <- matrix(nrow = 500)</pre>
betareml <- matrix(nrow = 500)</pre>
secoch <- matrix(nrow = 500)</pre>
betacoch <- matrix(nrow = 500)</pre>
```

```
gmleci <- matrix(nrow = 500, ncol = 2)</pre>
remlci <- matrix(nrow = 500, ncol = 2)</pre>
cochci <- matrix(nrow = 500, ncol = 2)</pre>
counter = 0
counter2 = 0
countergmle = 0
counterreml = 0
countercoch = 0
for (i in 1:500) {
    for (t in 1:length(eps)) {
         if (t == 1) {
             eps[t] \leftarrow rnorm(1) * sqrt(1/(1 - rho^2))
         eps[t] \leftarrow rnorm(1) + rho * eps[t - 1]
    }
    y0 = beta0 + beta1 * xt + eps
    fit1 <- lm(y0 ~ xt)
    conf <- confint(fit1)</pre>
    c[i, 1:2] \leftarrow conf[2, 1:2]
    b[i, 1:2] <- fit1$coefficients</pre>
    resi[i, ] <- fit1$residuals</pre>
    \# (c2/3)
    gmle <- gls(y0 ~ xt, correlation = corAR1(form = ~1), method = "ML")</pre>
    a[i] <- as.numeric(gmle$modelStruct)</pre>
    rho.hat.mle[i] = (exp(a[i]) - 1)/(exp(a[i]) + 1)
    reml <- gls(y0 ~ xt, correlation = corAR1(form = ~1), method = "REML")
    z[i] <- as.numeric(reml$modelStruct)</pre>
    rho.hat.reml[i] = (exp(z[i]) - 1)/(exp(z[i]) + 1)
    coch <- cochrane.orcutt(fit1)</pre>
    rhoyw[i] <- coch$rho</pre>
    dwtemp <- DurbinWatsonTest(fit1, alternative = c("two.sided"))</pre>
    dw[i, 1] <- dwtemp$statistic</pre>
    dw[i, 2] <- dwtemp$method</pre>
    dw[i, 3] <- dwtemp$alternative</pre>
    dw[i, 4] <- dwtemp$p.value</pre>
    # Part (d)
    segmle[i] <- sqrt(gmle$varBeta[2, 2])</pre>
    betagmle[i] <- gmle$coefficients[2]</pre>
    sereml[i] <- sqrt(reml$varBeta[2, 2])</pre>
    betareml[i] <- reml$coefficients[2]</pre>
    secoch[i] <- coch$std.error[2]</pre>
    betacoch[i] <- coch$coefficients[2]</pre>
    gmleci[i, 1] <- betagmle[i] - (1.96 * segmle[i])</pre>
    gmleci[i, 2] <- betagmle[i] + (1.96 * segmle[i])</pre>
    remlci[i, 1] <- betareml[i] - (1.96 * sereml[i])</pre>
    remlci[i, 2] <- betareml[i] + (1.96 * sereml[i])</pre>
    cochci[i, 1] <- betacoch[i] - (1.96 * secoch[i])</pre>
    cochci[i, 2] <- betacoch[i] + (1.96 * secoch[i])</pre>
for (i in 1:500) {
    if (c[i, 1] < 0.5 \&\& c[i, 2] > 0.5) {
         counter = counter + 1
```

```
if (dw[i, 1] < 1.5) {
    counter2 = counter2 + 1
}

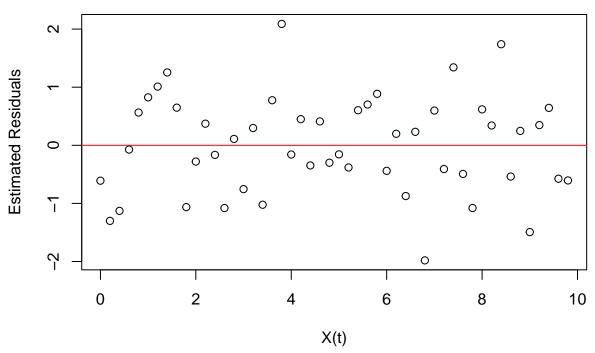
if (gmleci[i, 1] < 0.5 && gmleci[i, 2] > 0.5) {
    countergmle = countergmle + 1
}

if (remlci[i, 1] < 0.5 && remlci[i, 2] > 0.5) {
    counterreml = counterreml + 1
}

if (cochci[i, 1] < 0.5 && cochci[i, 2] > 0.5) {
    countercoch = countercoch + 1
}

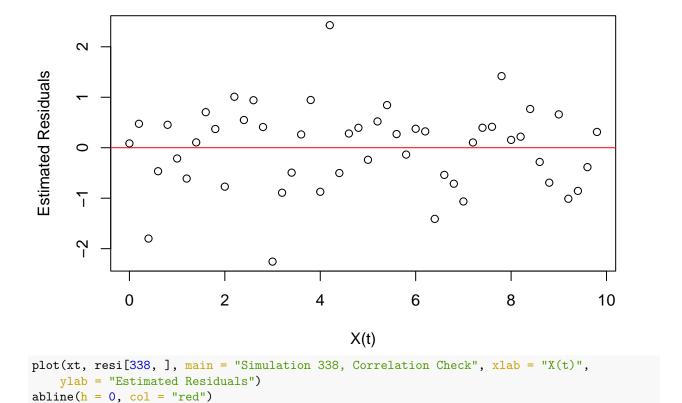
plot(xt, resi[24, ], main = "Simulation 24, Correlation Check", xlab = "X(t)", ylab = "Estimated Residuabline(h = 0, col = "red")
```

Simulation 24, Correlation Check

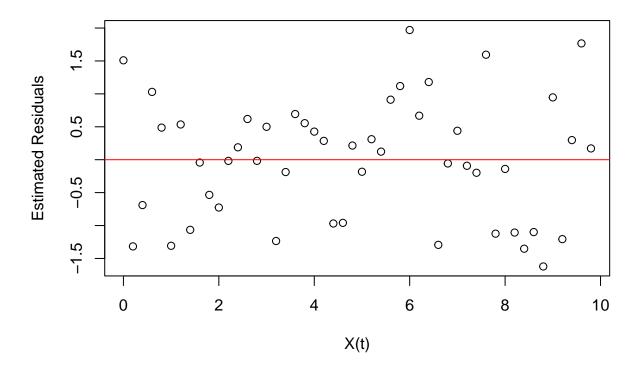


```
plot(xt, resi[154, ], main = "Simulation 154, Correlation Check", xlab = "X(t)",
    ylab = "Estimated Residuals")
abline(h = 0, col = "red")
```

Simulation 154, Correlation Check



Simulation 338, Correlation Check



```
betahat0 <- mean(b[1:500, 1])</pre>
betahat1 <- mean(b[1:500, 2])
rhohatmle <- mean(rho.hat.mle[1:500])</pre>
rhohatreml <- mean(rho.hat.reml[1:500])</pre>
rhohatyw <- mean(rhoyw[1:500])</pre>
print(counter)
## [1] 363
print(counter2)
## [1] 449
print(countergmle)
## [1] 448
print(rhohatmle)
## [1] 0.4197968
print(counterreml)
## [1] 459
print(rhohatreml)
## [1] 0.4825033
print(countercoch)
## [1] 443
print(rhohatyw)
## [1] 0.418544
```