732A62 Lab 2

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Assignment 1

a)

```
library(astsa)
library(kernlab)
library(TSA)
library(forecast)
set.seed(12345)
AR3 \leftarrow arima.sim(1000, model = list(order = c(3,0,0),
                                      ar = c(0.8, -0.2, 0.1))
## The theoretical
AR3.pacf <- pacf(AR3, plot=F)
AR3.data <- ts.intersect(xt = AR3, x1 = lag(AR3, 1), x2 = lag(AR3, 2), x3 = lag(AR3, 3))
AR.lm \leftarrow resid(lm(xt \sim x1 + x2, data = AR3.data))
AR.lm.lag3 \leftarrow resid(lm(x3 \sim x1 + x2 , data = AR3.data))
AR3.pacf[3]
##
## Partial autocorrelations of series 'AR3', by lag
       3
##
## 0.117
cor(AR.lm, AR.lm.lag3)
## [1] 0.1146076
```

b)

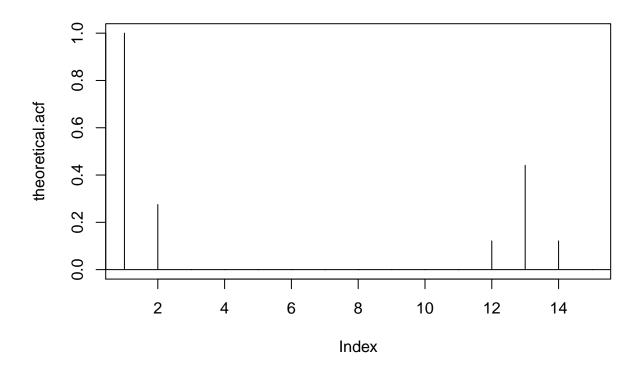
```
##
## Call:
## ar(x = AR2, aic = FALSE, order.max = 2, method = "yw")
## Coefficients:
##
       1
## 0.8029 0.1037
##
## Order selected 2 sigma^2 estimated as 1.267
ar2.ols
##
## Call:
## ar(x = AR2, aic = FALSE, order.max = 2, method = "ols")
## Coefficients:
##
        1
## 0.8067 0.1205
##
## Intercept: -0.04401 (0.1074)
## Order selected 2 sigma^2 estimated as 1.129
ar2.mle
##
## arima(x = AR2, order = c(2, 0, 0), method = "ML")
## Coefficients:
##
                  ar2 intercept
            ar1
         0.7967 0.1189
##
                            0.8290
## s.e. 0.0992 0.1000
                            1.1385
## sigma^2 estimated as 1.126: log likelihood = -148.71, aic = 303.41
Yes, the theoretical value for
is inside the confidence-intervall for the ML estimate.
```

c)

```
set.seed(12345)
ma.coef <- c(0.3, rep(0, 10), 0.6)
ts4 <- arima.sim(n=200, model=list(order=c(0, 0, 12), ma = ma.coef))
theoretical.acf <- ARMAacf(ma=c(ma.coef, 0.3 * 0.6))
theoretical.pacf <- ARMAacf(ma=c(ma.coef, 0.3 * 0.6), pacf=TRUE)

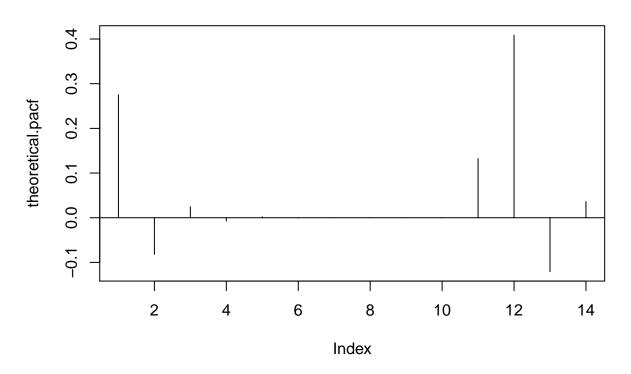
plot(theoretical.acf, type="h", main="Theoretical ACF")
abline(h=0)</pre>
```

Theoretical ACF



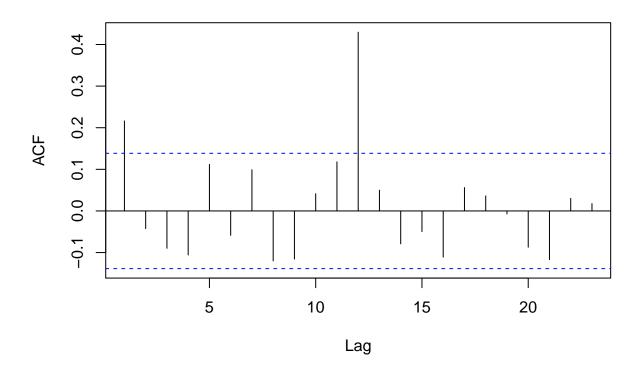
plot(theoretical.pacf, type="h", main="Theoretical PACF")
abline(h=0)

Theoretical PACF



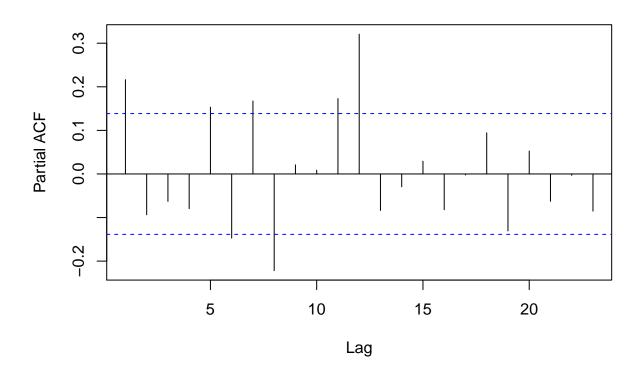
empirical.acf <- acf(ts4)</pre>

Series ts4



empirical.pacf <- pacf(ts4)</pre>

Series ts4

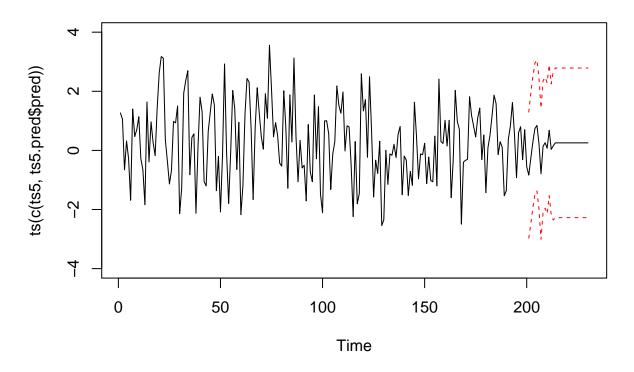


d)

```
set.seed(12345)
ma.coef <- c(0.3, rep(0, 10), 0.6)
ts5 <- arima.sim(n=200, model=list(order=c(0, 0, 12), ma = ma.coef))

ts5.fit <- arima(ts5, order=c(0, 0, 1), seasonal=list(order=c(0, 0, 1), period=12))
ts5.pred <- predict(ts5.fit, n.ahead=30, se.fit=TRUE)

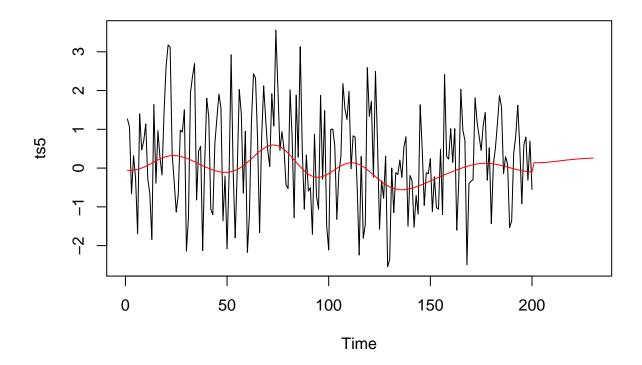
plot(ts(c(ts5, ts5.pred$pred)), ylim=c(-4, 4))
lines(200 + 1:length(ts5.pred$pred), ts5.pred$pred + 1.96 * ts5.pred$se, lty=2, col="red")
lines(200 + 1:length(ts5.pred$pred), ts5.pred$pred - 1.96 * ts5.pred$se, lty=2, col="red")</pre>
```



```
gausspr.data <- data.frame(y=ts5, x=1:200)
gausspr.fit <- gausspr(y ~ x, gausspr.data)

## Using automatic sigma estimation (sigest) for RBF or laplace kernel
gausspr.pred <- predict(gfit, data.frame(x=201:230))

plot(ts5, xlim=c(0, 230))
lines(c(fitted(gausspr.fit), gausspr.pred), , col="red")</pre>
```



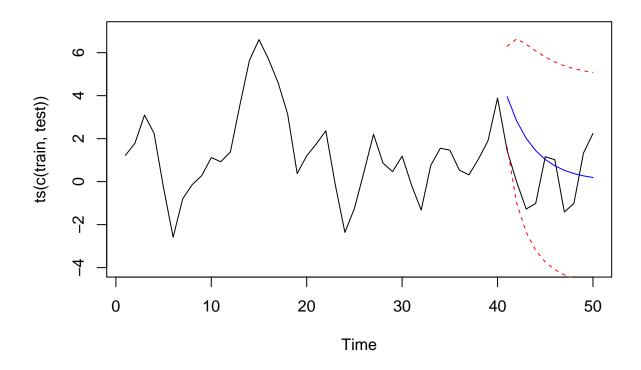
e)

```
set.seed(12345)
ts6 <- arima.sim(model=list(ma=c(0.5), ar=c(0.7)), n=50)

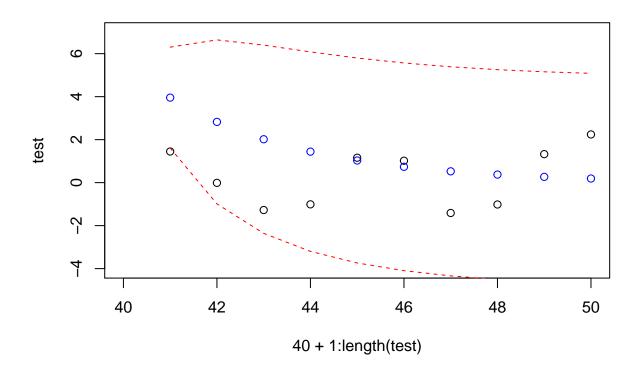
train <- ts(ts6[1:40])
test <- ts(ts6[41:50])

ts6.fit <- arima(train, order=c(1, 0, 1), include.mean = F)
ts6.pred <- predict(ts6.fit, n.ahead=10)

plot(ts(c(train, test)), ylim=c(-4, 7), type="1")
lines(40 + 1:length(test), ts6.pred$pred, col="blue")
lines(40 + 1:length(test), ts6.pred$pred + 1.96 * ts6.pred$se, lty=2, col="red")
lines(40 + 1:length(test), ts6.pred$pred - 1.96 * ts6.pred$se, lty=2, col="red")</pre>
```



```
plot(40 + 1:length(test), test, ylim=c(-4, 7), xlim=c(40, 50), type="p")
points(40 + 1:length(test), ts6.pred$pred, col="blue")
lines(40 + 1:length(test), ts6.pred$pred + 1.96 * ts6.pred$se, lty=2, col="red")
lines(40 + 1:length(test), ts6.pred$pred - 1.96 * ts6.pred$se, lty=2, col="red")
```



Assignment 2

```
assignment2 <- function(data){</pre>
    old \leftarrow par(mfrow = c(2, 2))
    acf(data, lag.max = 40)
    pacf(data, lag.max = 40)
    acf(diff(data, lag = 1), lag.max = 40)
    pacf(diff(data, lag = 1), lag.max = 40)
    par(old)
}
```

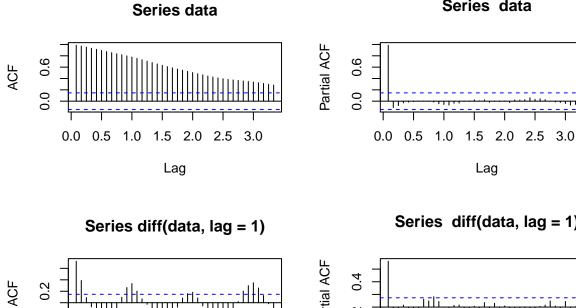
Series data

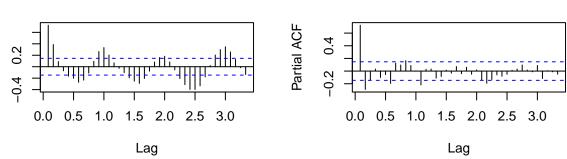
Lag

Series diff(data, lag = 1)

Chicken

assignment2(chicken)



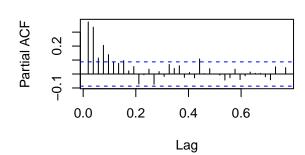


so2

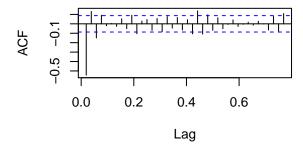
assignment2(so2)

Series data

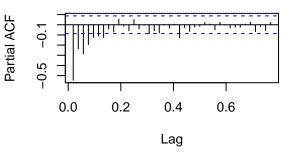
Series data



Series diff(data, lag = 1)

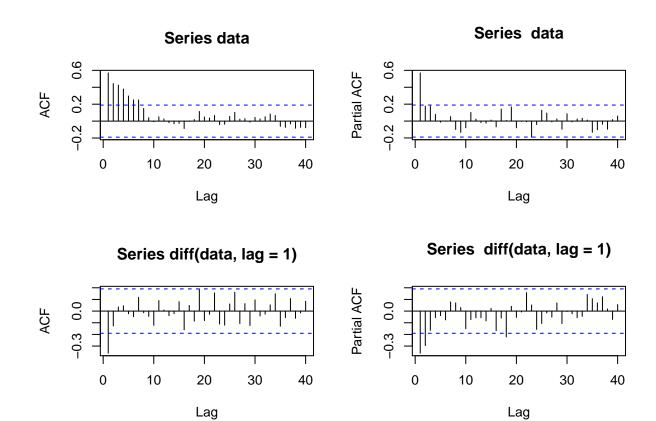


Series diff(data, lag = 1)



EQcount

assignment2(EQcount)

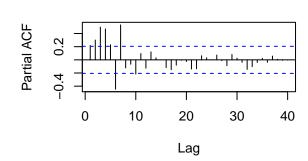


HCT

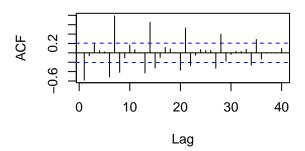
assignment2(HCT)

Series data

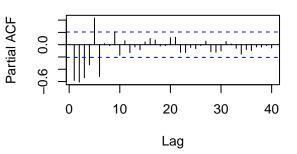
Series data



Series diff(data, lag = 1)



Series diff(data, lag = 1)

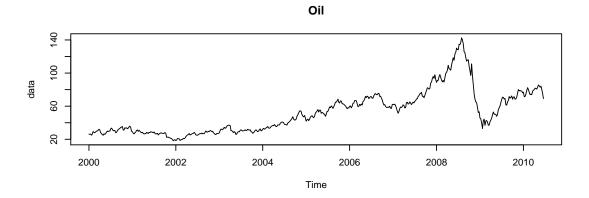


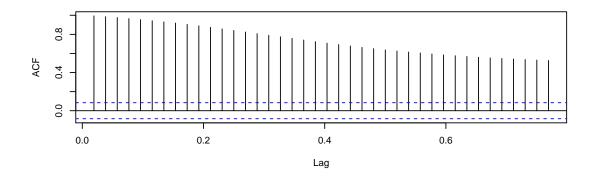
Assignment 3

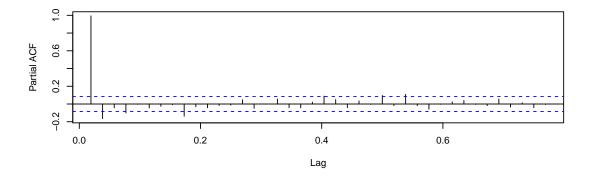
```
plot_helper <- function(data, title) {</pre>
    old <- par(mfrow=c(4, 1))</pre>
    plot(data, main=title)
    acf(data, lag.max=40, main="")
    pacf(data, lag.max=40, main="")
    qqnorm(data, main="", las=1)
    qqline(data)
    par(old)
}
test_helper <- function(data) {</pre>
    print(Box.test(data, lag = 1, type = "Ljung-Box"))
    print(suppressWarnings(adf.test(data)))
    e <- eacf(data)
}
fit_plot <- function(model) {</pre>
    pred <- predict(model, n.ahead=20, se.fit=TRUE)</pre>
    upper_band <- pred$pred + 1.96 * pred$se
    lower_band <- pred$pred - 1.96 * pred$se</pre>
    plot(c(model$x, pred$pred), type="l", xlim=c(500, length(oil) + 20), ylim=c(min(lower_band), max(up)
    lines(length(oil) + 1:20, upper_band, lty=2, col="red")
    lines(length(oil) + 1:20, lower_band, lty=2, col="red")
```

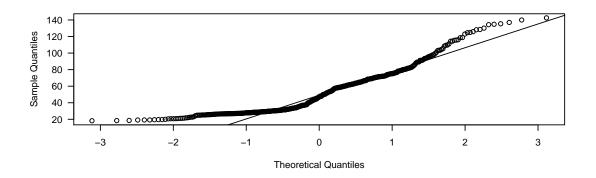
a)

```
loil <- log(oil)
doil <- diff(oil)
ddoil <- diff(oil, 2)
dloil <- diff(loil)
ddloil <- diff(loil, 2)</pre>
```

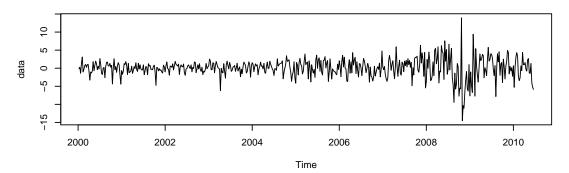


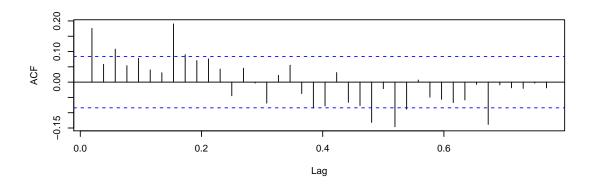


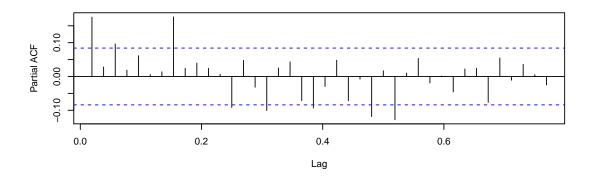


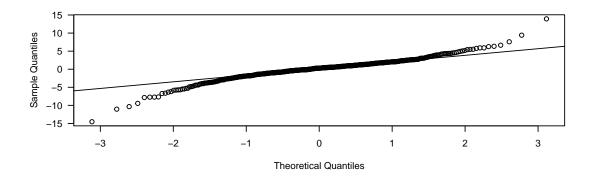


1 Difference Oil

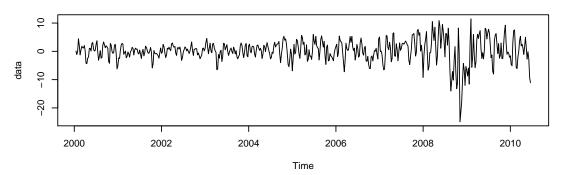


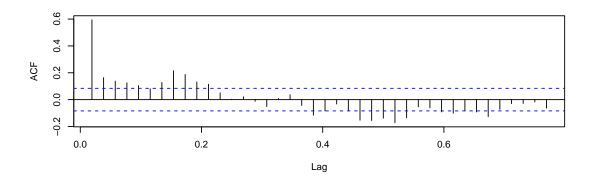


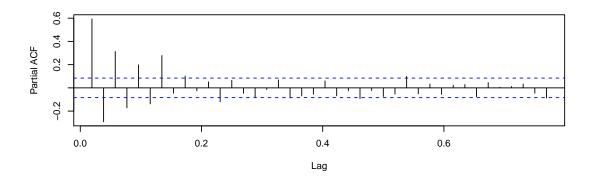


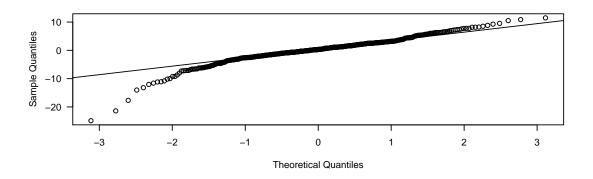


2 Difference Oil

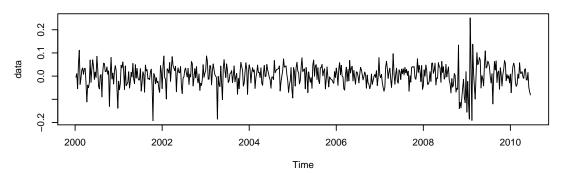


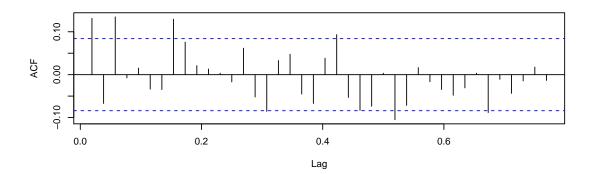


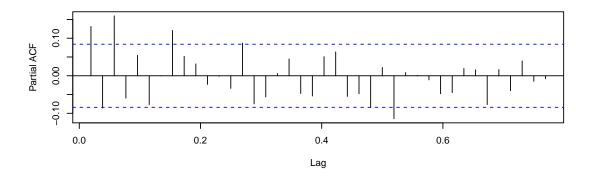


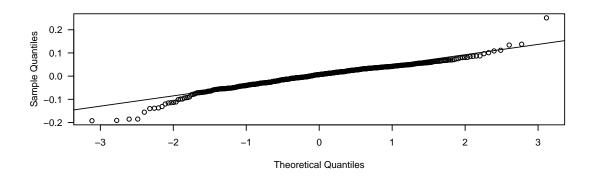


1 Difference log Oil

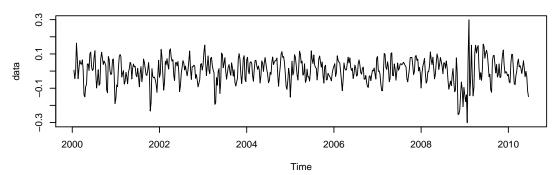


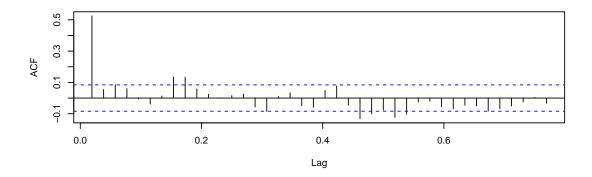


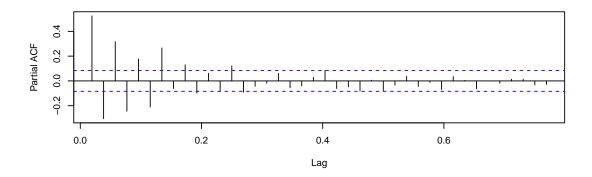


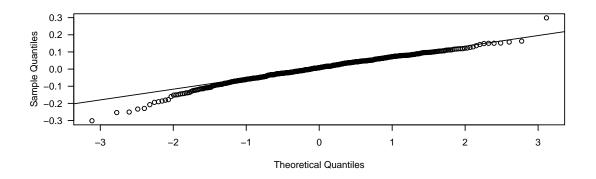


2 Difference log Oil









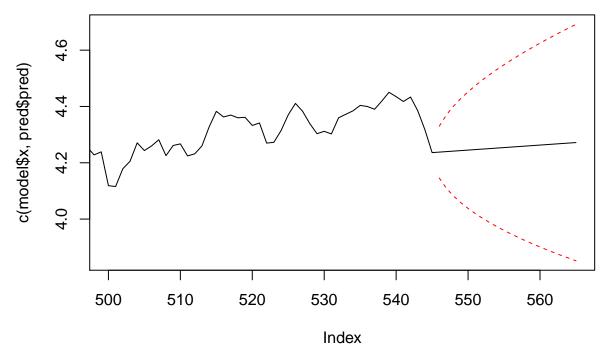
```
test_helper(doil)
##
## Box-Ljung test
##
## data: data
## X-squared = 16.884, df = 1, p-value = 3.974e-05
##
##
## Augmented Dickey-Fuller Test
##
## data: data
## Dickey-Fuller = -5.3269, Lag order = 8, p-value = 0.01
## alternative hypothesis: stationary
##
## AR/MA
## 0 1 2 3 4 5 6 7 8 9 10 11 12 13
## 0 x o x o o o o x x o o o o
## 1 x o o o o o o x o o o o
## 2 x x o o o o o x o o o o
## 3 x x x o o o o x o o o o
## 4 x x x o o o o x o o o o
## 5 x x o o o o o x o o o o
## 6 x o x o x o o x o o o o
## 7 o o x o x o x x o o o o
test_helper(ddoil)
##
## Box-Ljung test
##
## data: data
## X-squared = 192.72, df = 1, p-value < 2.2e-16
##
##
## Augmented Dickey-Fuller Test
##
## data: data
## Dickey-Fuller = -4.7773, Lag order = 8, p-value = 0.01
## alternative hypothesis: stationary
##
## AR/MA
## 0 1 2 3 4 5 6 7 8 9 10 11 12 13
## 0 x x x x x o x x x x x o o
## 1 x x o o o o o x o o o o
## 2 x x x o o o o x x o x o o
## 3 x x x o o o o x o o x o o
## 4 x x o o o o o x x o o o o
## 5 x x o x x x o x o o o o
## 6 x x o x x x x x o o o o o
## 7 x x o x x x o x o x o o o
test_helper(dloil)
```

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##

```
## Box-Ljung test
##
## data: data
## X-squared = 9.4307, df = 1, p-value = 0.002134
##
## Augmented Dickey-Fuller Test
##
## data: data
## Dickey-Fuller = -6.3708, Lag order = 8, p-value = 0.01
## alternative hypothesis: stationary
## AR/MA
## 0 1 2 3 4 5 6 7 8 9 10 11 12 13
## 0 x o x o o o o x o o o o
## 1 x o x o o o o x o o o
## 2 x x x o o o o x o o o o o
## 3 x x x o o o o x o o o o
## 4 x o x o o o o x o o o o
## 5 x x x o x o o x o o o o
## 6 o x x o x x o x o o o o x
## 7 o x x x x x x x o x o o o
test_helper(ddloil)
##
## Box-Ljung test
##
## data: data
## X-squared = 150.51, df = 1, p-value < 2.2e-16
##
##
## Augmented Dickey-Fuller Test
##
## data: data
## Dickey-Fuller = -5.6251, Lag order = 8, p-value = 0.01
## alternative hypothesis: stationary
##
## AR/MA
## 0 1 2 3 4 5 6 7 8 9 10 11 12 13
## 0 x o o o o o o x x o o o o
## 1 x x o o o o o x x o o o o
## 2 x x x o o o o x x o o o o
## 3 x x x x o o o x x o o o o
## 4 x x x x o o o x x o o o o
## 5 x o x x x o o x x o o o o
## 6 x o x x x x x x o x o o o
## 7 x x x x x x x x x o x o o o
```

```
fit1 <- Arima(loil, order=c(0, 2, 1))</pre>
fit1
## Series: loil
## ARIMA(0,2,1)
##
## Coefficients:
##
##
         -1.0000
## s.e.
        0.0061
##
## sigma^2 estimated as 0.002213: log likelihood=886.63
## AIC=-1769.26
                  AICc=-1769.24
                                 BIC=-1760.67
fit_plot(fit1)
```



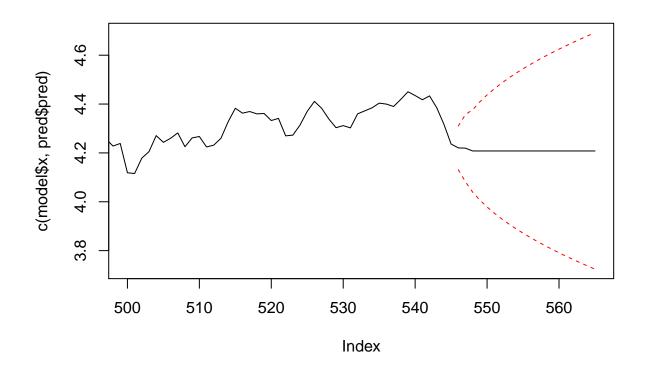
```
fit2 <- Arima(loil, order=c(0, 1, 3))
fit2

## Series: loil
## ARIMA(0,1,3)
##

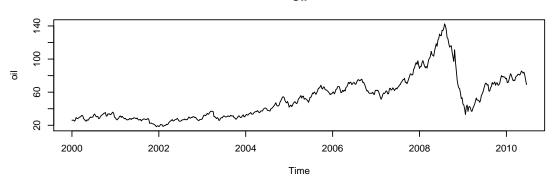
## Coefficients:
## ma1 ma2 ma3
## 0.1696 -0.0886 0.1458
## s.e. 0.0424 0.0424 0.0429
##</pre>
```

```
## sigma^2 estimated as 0.002094: log likelihood=907.41
## AIC=-1806.83 AICc=-1806.75 BIC=-1789.63
```

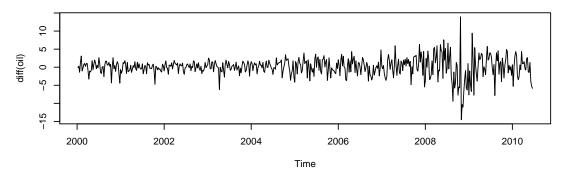
fit_plot(fit2)



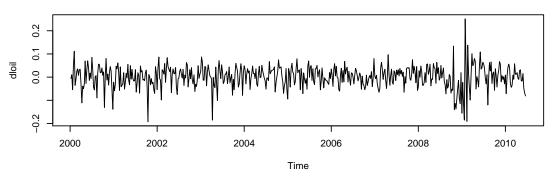




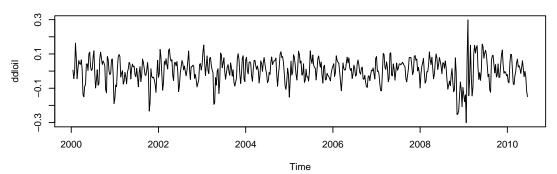
Difference 1 Oil



Difference 1 Log Oil

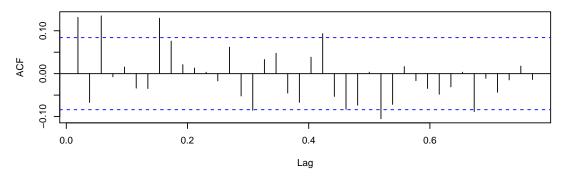


Difference 2 Log Oil

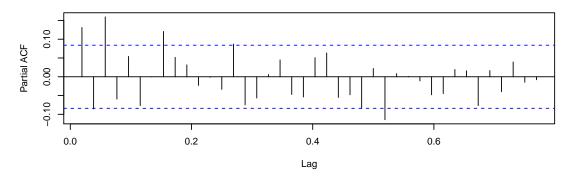


Clearly difference log is the data we should work with. bla, bla, \dots

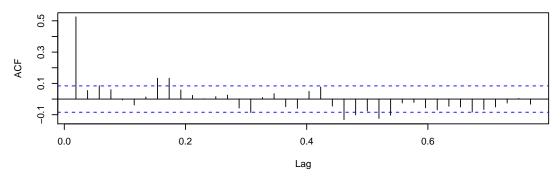
Difference 1 Log Oil ACF



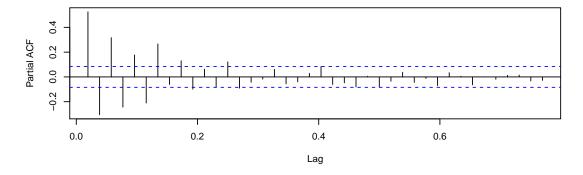
Difference 1 Log Oil PACF



Difference 2 Log Oil ACF



Difference 2 Log Oil PACF

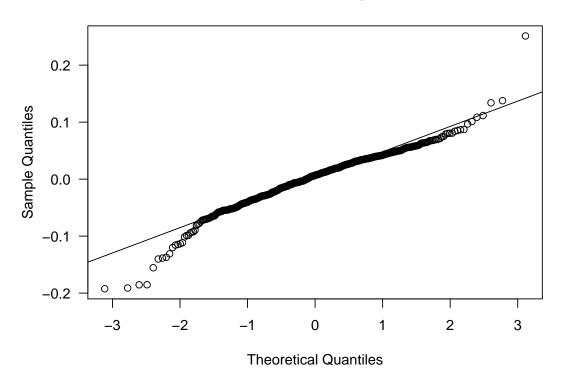


eacf(dloil)

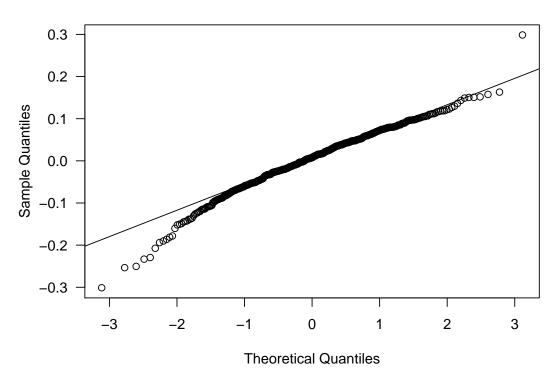
eacf(ddloil)

AR/MA

Difference 1 Log Oil



Difference 2 Log Oil

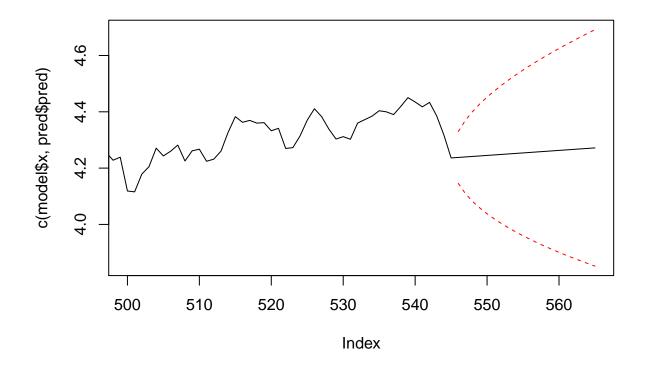


```
fit1 <- Arima(loil, order=c(1, 1, 1))</pre>
fit1
## Series: loil
## ARIMA(1,1,1)
##
## Coefficients:
##
            ar1
##
        -0.5253 0.7142
## s.e. 0.0872 0.0683
##
## sigma^2 estimated as 0.002112: log likelihood=904.58
## AIC=-1803.15 AICc=-1803.11 BIC=-1790.25
fit2 <- Arima(loil, order=c(0, 1, 3))</pre>
## Series: loil
## ARIMA(0,1,3)
## Coefficients:
                            ma3
          ma1
                ma2
        0.1696 -0.0886 0.1458
##
## s.e. 0.0424 0.0424 0.0429
## sigma^2 estimated as 0.002094: log likelihood=907.41
## AIC=-1806.83 AICc=-1806.75 BIC=-1789.63
fit3 <- Arima(loil, order=c(0, 2, 1))</pre>
fit3
## Series: loil
## ARIMA(0,2,1)
##
## Coefficients:
##
            ma1
##
        -1.0000
## s.e. 0.0061
## sigma^2 estimated as 0.002213: log likelihood=886.63
## AIC=-1769.26 AICc=-1769.24 BIC=-1760.67
```

```
complex_dist <- function(x) {
    sqrt(Re(x)^2 + Im(x)^2)
}
sapply(polyroot(c(1, -2, 1)), complex_dist)

## [1] 1 1
sapply(polyroot(c(1, -1)), complex_dist)

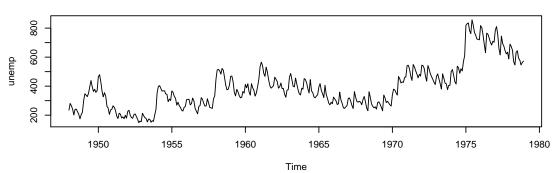
## [1] 1
fit_plot(fit3)</pre>
```



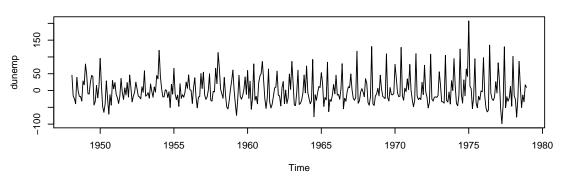
b)

```
lunemp <- log(unemp)
dunemp <- diff(unemp)
ddunemp <- diff(dunemp, 2)
dlunemp <- diff(lunemp)
ddlunemp <- diff(lunemp, 2)</pre>
```

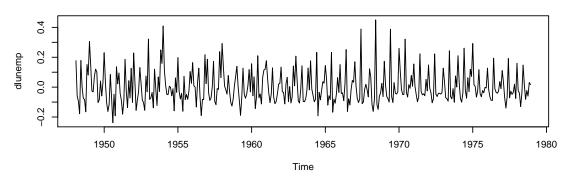




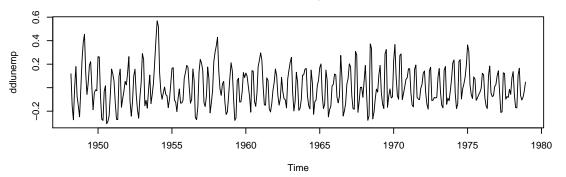
Difference 1 Unemp



Difference 1 Log Unemp

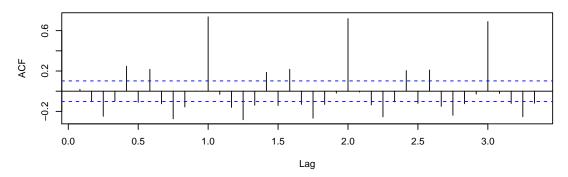


Difference 2 Log Unemp

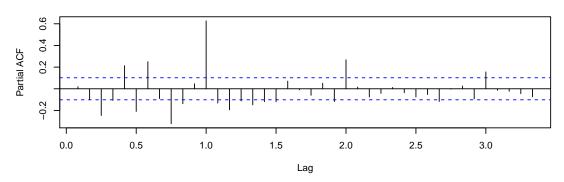


Clearly difference log is the data we should work with. bla, bla, \dots

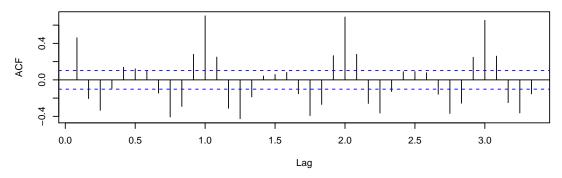
Difference 1 Log Unemp ACF



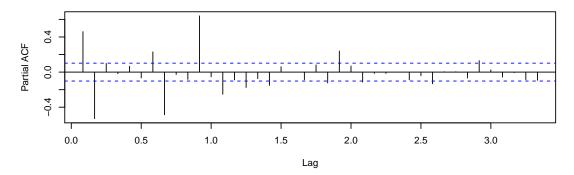
Difference 1 Log Unemp PACF



Difference 2 Log Unemp ACF



Difference 2 Log Unemp PACF

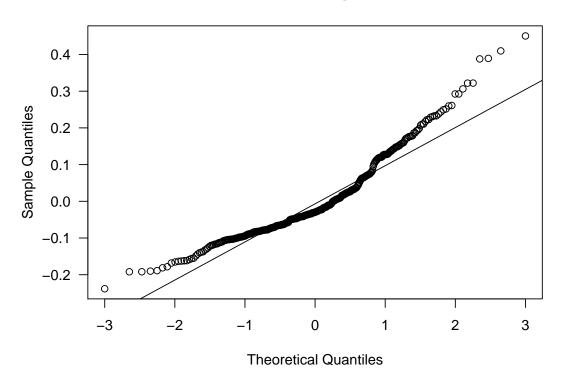


eacf(dlunemp)

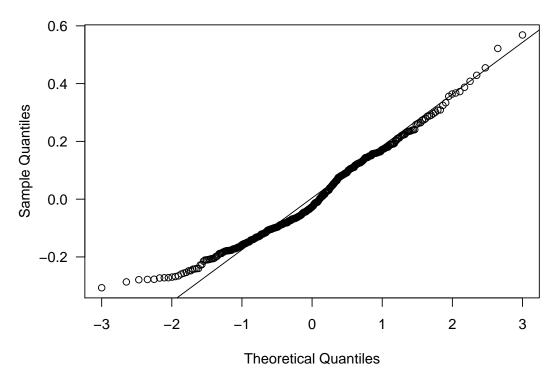
eacf(ddlunemp)

 $\#\#\ 7\ \verb"x" o o o x x o x o o o x o x$

Difference 1 Log Unemp



Difference 2 Log Unemp



```
fit1 <- Arima(lunemp, order=c(1, 1, 1))</pre>
## Series: lunemp
## ARIMA(1,1,1)
##
## Coefficients:
##
##
        -0.7592 0.8157
## s.e. 0.0952 0.0796
##
## sigma^2 estimated as 0.01289: log likelihood=281.71
## AIC=-557.42 AICc=-557.35 BIC=-545.67
fit2 <- Arima(lunemp, order=c(0, 1, 3))</pre>
## Series: lunemp
## ARIMA(0,1,3)
## Coefficients:
           ma1
                  ma2
                             ma3
        -0.0079 0.0277 -0.3629
##
## s.e. 0.0470 0.0506 0.0481
## sigma^2 estimated as 0.01177: log likelihood=298.85
## AIC=-589.7 AICc=-589.59 BIC=-574.03
fit3 <- Arima(lunemp, order=c(0, 2, 1))</pre>
fit3
## Series: lunemp
## ARIMA(0,2,1)
##
## Coefficients:
##
           ma1
##
        -1.000
## s.e. 0.007
## sigma^2 estimated as 0.01298: log likelihood=276.19
## AIC=-548.39 AICc=-548.36 BIC=-540.56
```