

CH5350: Applied **Time Series Analysis**

Project Report

By:

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Declaration:

I state that the following work has been done by me and only me without the assistance of any of the other students in this course (CH5350: Applied Time Series Analysis).

Sincerely,
Vishal Subbiah
(MM12B035)

Vishal Subbiah

1.R

```
library(TSA)
```

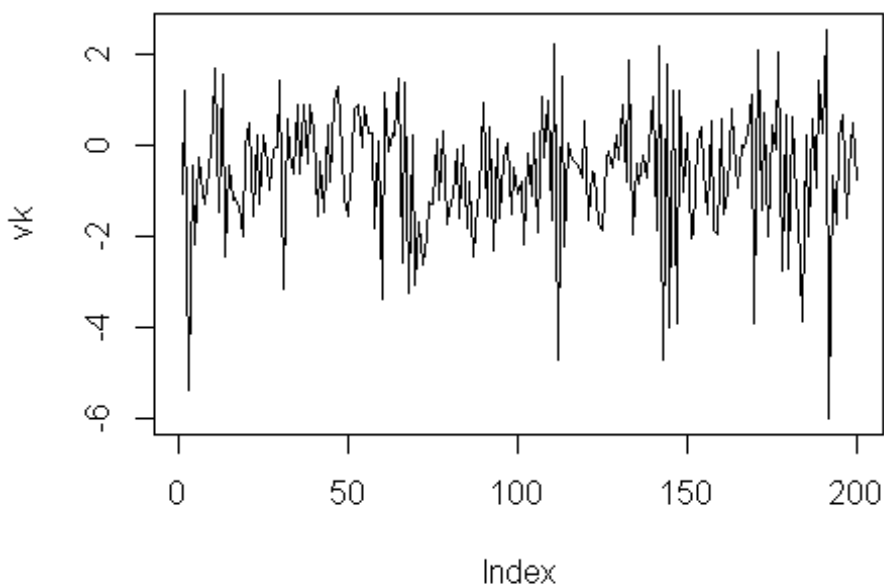
```
## Loading required package: leaps
## Loading required package: locfit
## locfit 1.5-9.1 2013-03-22
## Loading required package: mgcv
## Loading required package: nlme
## This is mgcv 1.8-0. For overview type 'help("mgcv-package")'.
## Loading required package: tseries
##
## Attaching package: 'TSA'
##
## The following objects are masked from 'package:stats':
##
##   acf, arima
##
## The following object is masked from 'package:utils':
##
##   tar
```

```
library(tsDyn)
```

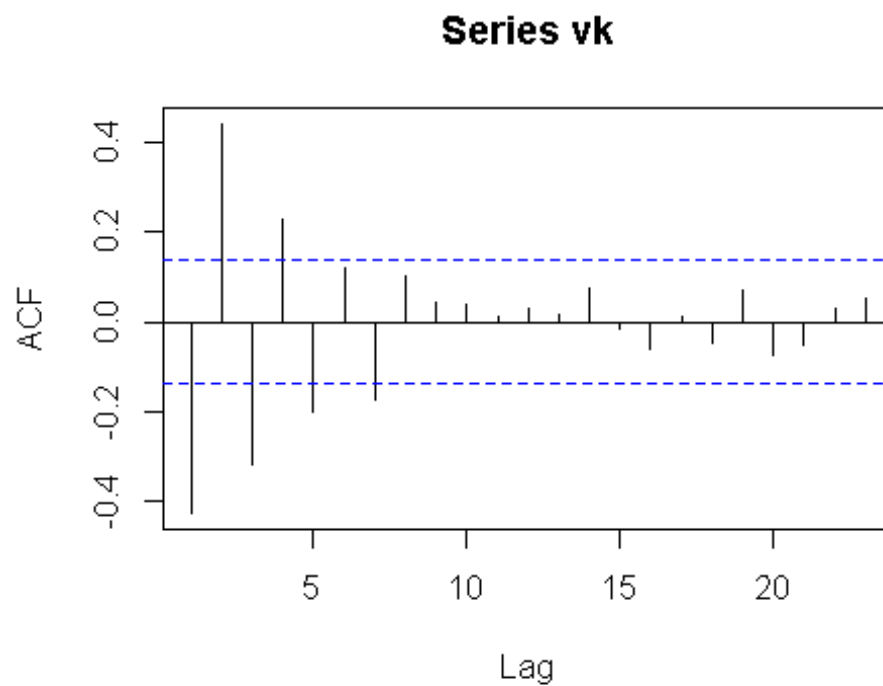
```
## Warning: package 'tsDyn' was built under R version 3.1.2
```

```
setwd("C:/Users/Toshiba/Desktop/vishal iit/5th sem/Applied time series
analysis/assignments/project")
load("projq1a.Rdata")
```

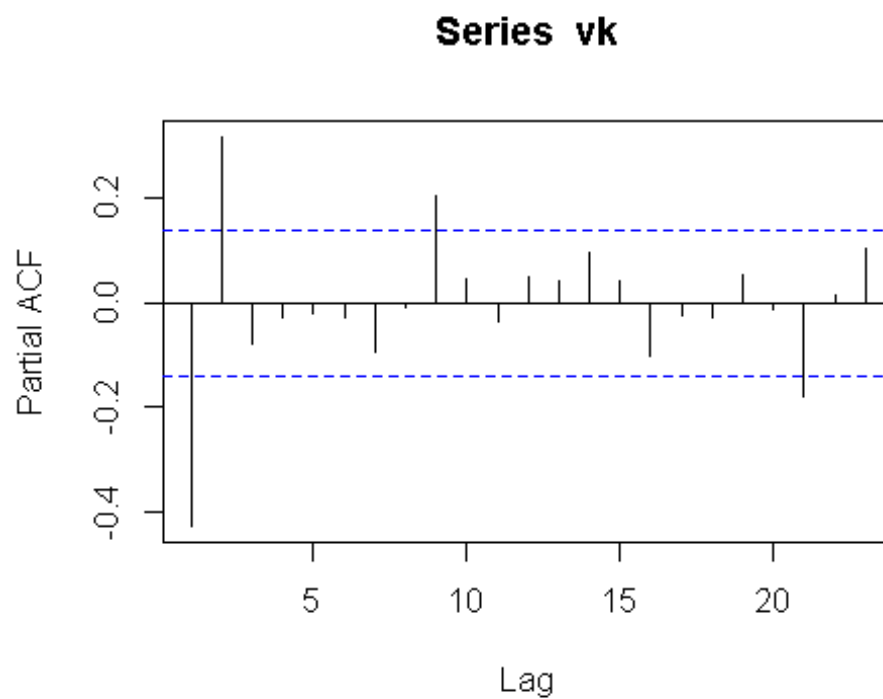
```
plot(vk,type='l')
```



```
acf(vk)
```



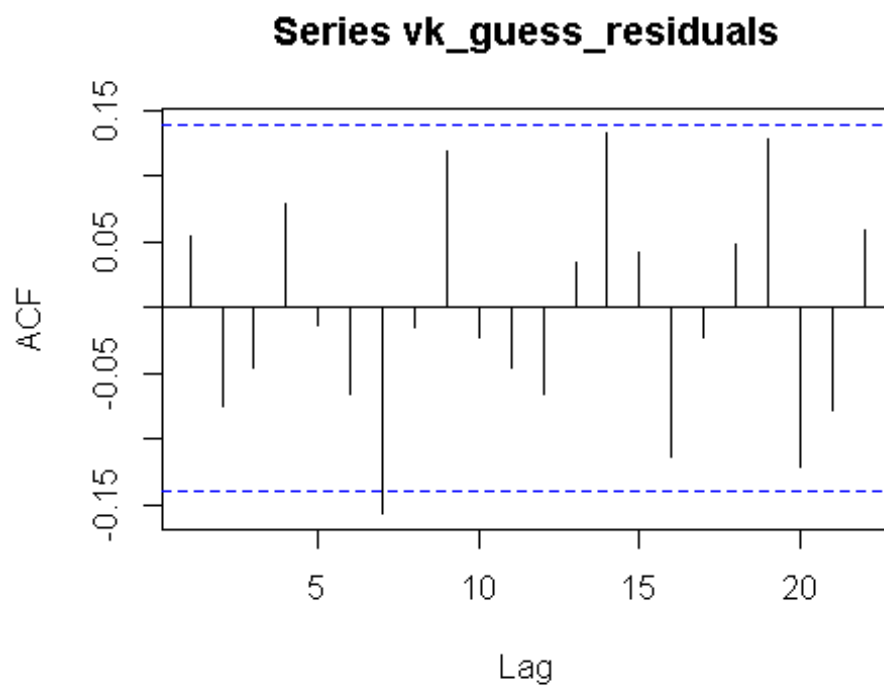
```
pacf(vk)
```



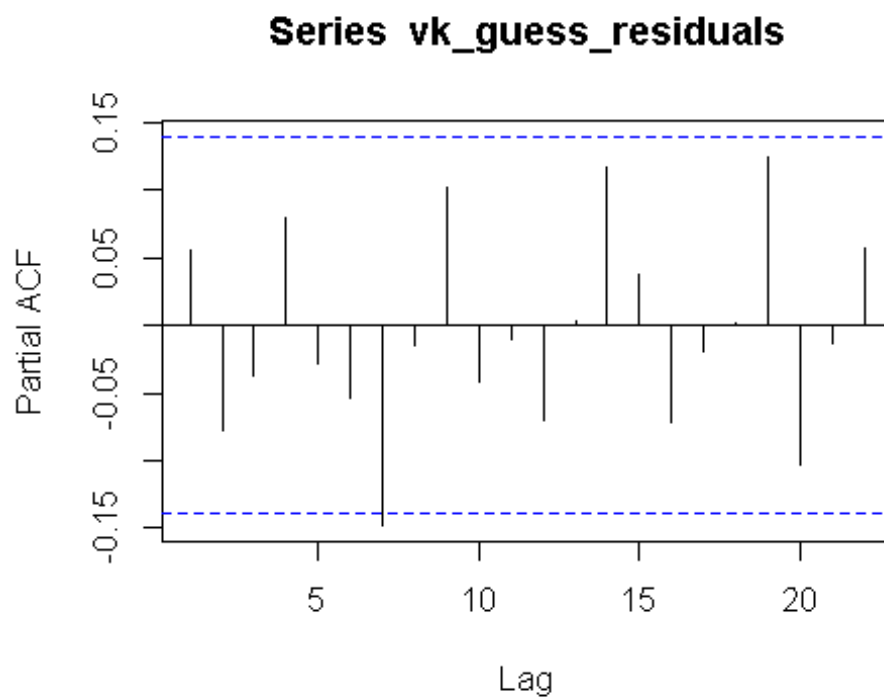
```
nonlinear_test=Tsay.test(vk,order=2)# non linear
```

```
vk_guess=tar(vk,2,1,1,method="MAIC",estimate.thd=TRUE)
```

```
vk_guess_residuals=vk_guess$residuals  
acf(vk_guess_residuals)
```



```
pacf(vk_guess_residuals)
```



```
Threshold=vk_guess$n1 # Threshold  
vk_guess_1_coeff=vk_guess$qr1$coefficients # coefficients for first regime  
vk_guess_2_coeff=vk_guess$qr2$coefficients # coefficients for second regime
```

```

phi_intercept_1=vk_guess_1_coeff[1]
phi_1_1=vk_guess_1_coeff[2]
phi_2_1=vk_guess_1_coeff[3]

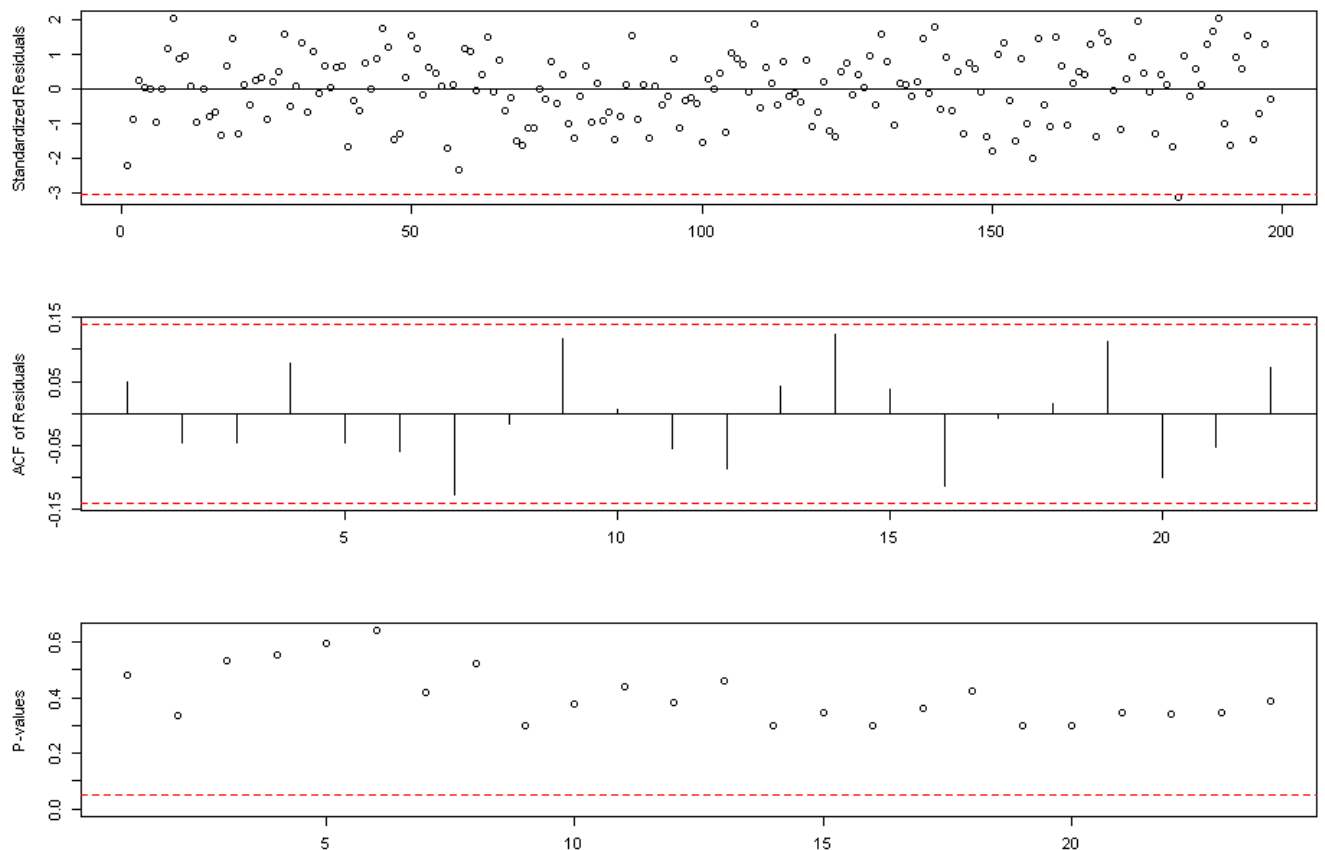
```

```

phi_intercept_2=vk_guess_2_coeff[1]
phi_1_2=vk_guess_2_coeff[2]

```

```
tsdiag.TAR(vk_guess)
```



part B

```
epsk=vk_guess_residuals
```

```
vk_guess_y=vk_guess$y
```

```

Threshold_mod_store=0
phi_mod_intercept_1_store=0
phi_mod_1_1_store=0
phi_mod_2_1_store=0

```

```

phi_mod_intercept_2_store=0
phi_mod_1_2_store=0
j=1
k=1

```

```

l=1
m=1
n=1
o=1
for (i in 1:200)
{

epskr1<-sample(epsk,size=200,replace=T)
vk_mod=vk_guess_y+epskr1
vk_guess_mod=tar(vk_mod,2,1,1,method="MAIC",estimate.thd=TRUE)

if(!is.na(vk_guess_mod$n1))
{
Threshold_mod_store[j]=vk_guess_mod$n1
j=j+1
}

vk_guess_mod_1_coeff=vk_guess_mod$qr1$coefficients # coefficients for first regime
vk_guess_mod_2_coeff=vk_guess_mod$qr2$coefficients # coefficients for second regime

if(!is.na(vk_guess_mod_1_coeff[1]))
{

phi_mod_intercept_1_store[k]=vk_guess_mod_1_coeff[1]
k=k+1
}

if(!is.na(vk_guess_mod_1_coeff[2]))
{
phi_mod_1_1_store[l]=vk_guess_mod_1_coeff[2]
l=l+1
}

if(!is.na(vk_guess_mod_1_coeff[3]))
{
phi_mod_2_1_store[m]=vk_guess_mod_1_coeff[3]
m=m+1
}

if(!is.na(vk_guess_mod_2_coeff[1]))
{
phi_mod_intercept_2_store[n]=vk_guess_mod_2_coeff[1]
n=n+1
}

if(!is.na(vk_guess_mod_2_coeff[2]))
{
phi_mod_1_2_store[o]=vk_guess_mod_2_coeff[2]
o=o+1
}

}

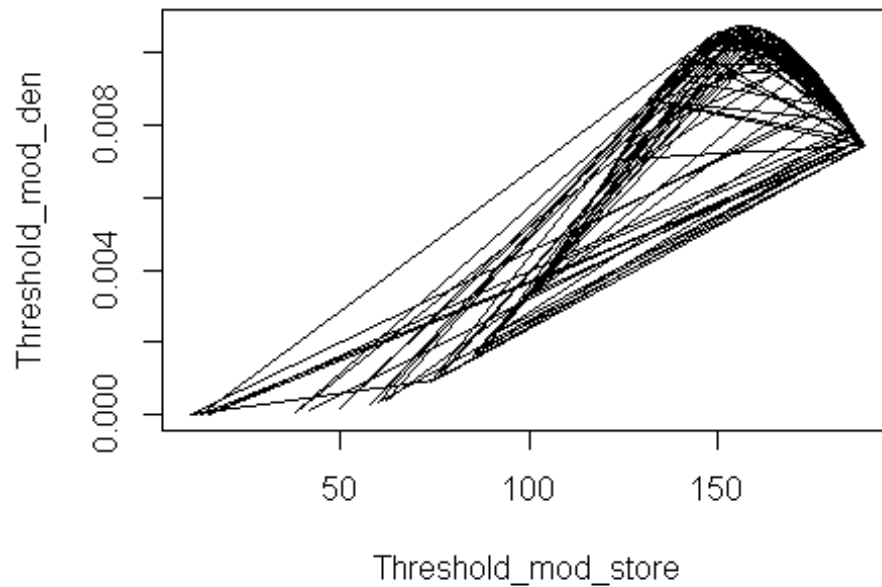
Threshold_mod_mean=mean(Threshold_mod_store)

```

```

Threshold_mod_var=var(Threshold_mod_store)
Threshold_mod_den=dnorm(Threshold_mod_store,mean=Threshold_mod_mean,sd=sqrt(Threshold_mod_var))
plot(Threshold_mod_store,Threshold_mod_den,type="l")

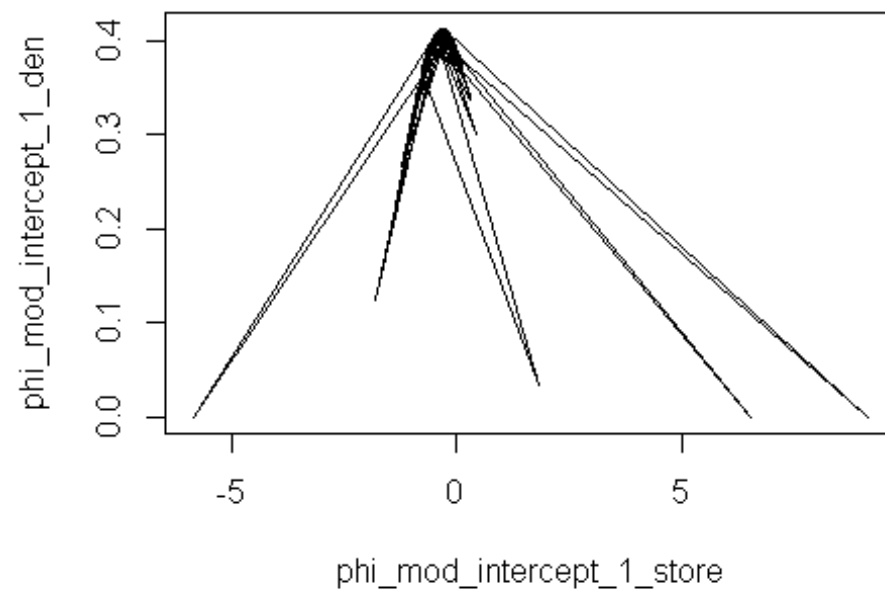
```



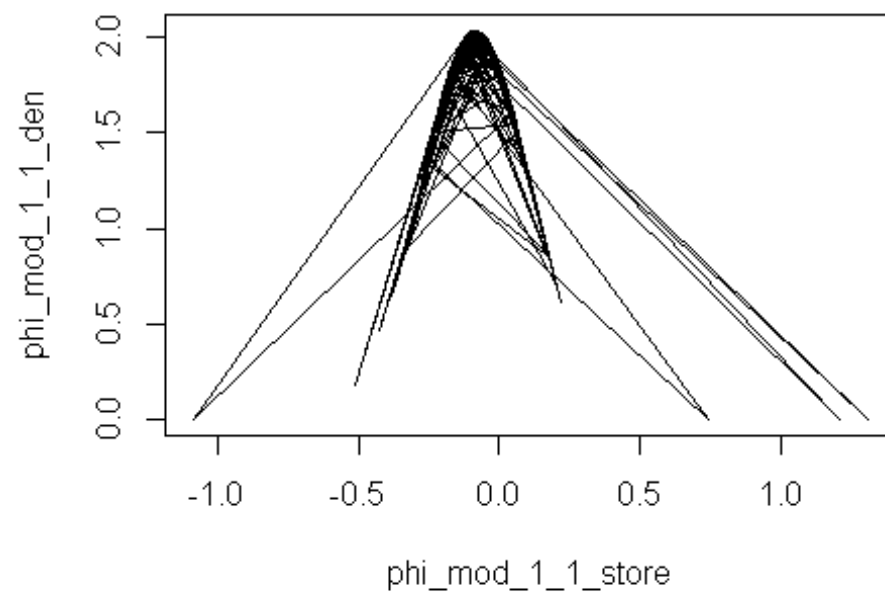
```

phi_mod_intercept_1_mean=mean(phi_mod_intercept_1_store)
phi_mod_intercept_1_var=var(phi_mod_intercept_1_store)
phi_mod_intercept_1_den=dnorm(phi_mod_intercept_1_store,mean=phi_mod_intercept_1_mean,sd
=sqrt(phi_mod_intercept_1_var))
plot(phi_mod_intercept_1_store,phi_mod_intercept_1_den,type="l")

```

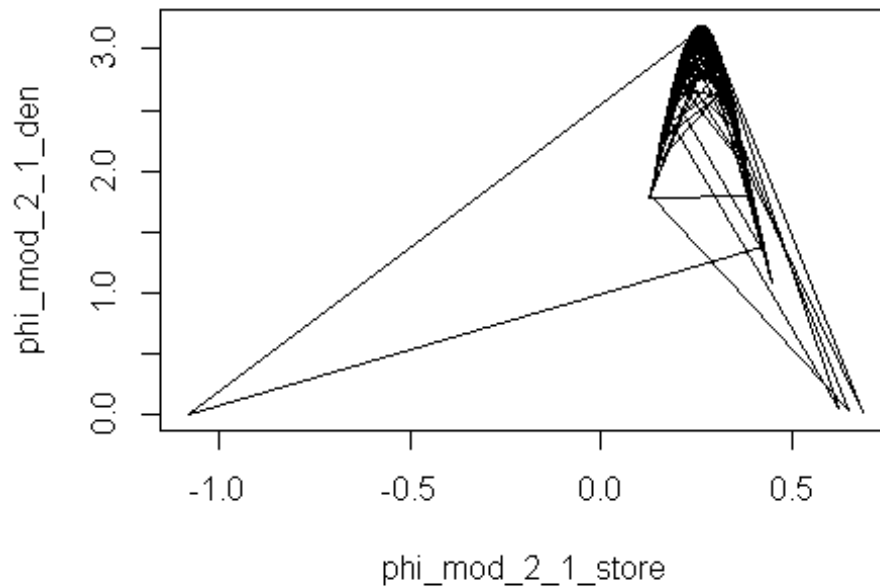



```
phi_mod_1_1_mean=mean(phi_mod_1_1_store)
phi_mod_1_1_var=var(phi_mod_1_1_store)
phi_mod_1_1_den=dnorm(phi_mod_1_1_store,mean=phi_mod_1_1_mean,sd=sqrt(phi_mod_1_1_var))
plot(phi_mod_1_1_store,phi_mod_1_1_den,type='l')
```

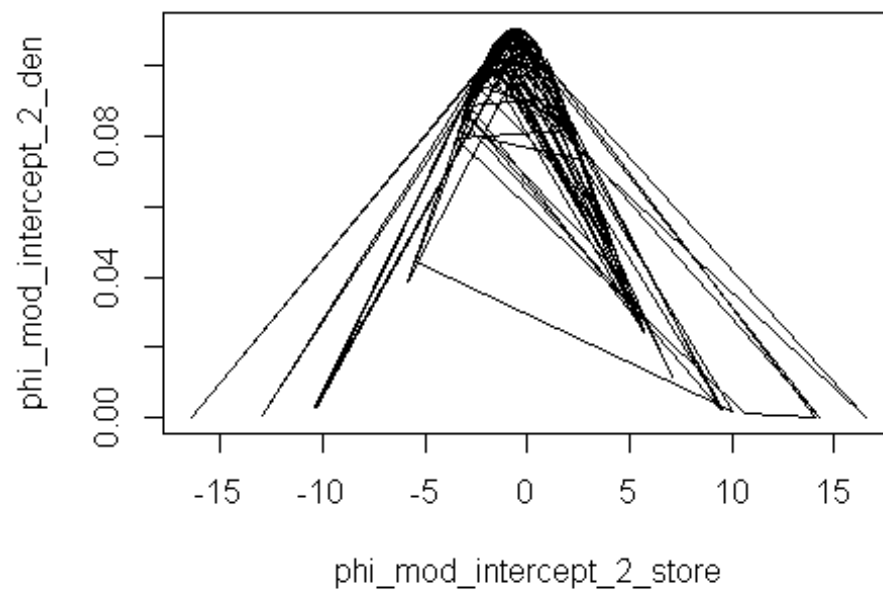


```
phi_mod_2_1_mean=mean(phi_mod_2_1_store)
phi_mod_2_1_var=var(phi_mod_2_1_store)
```

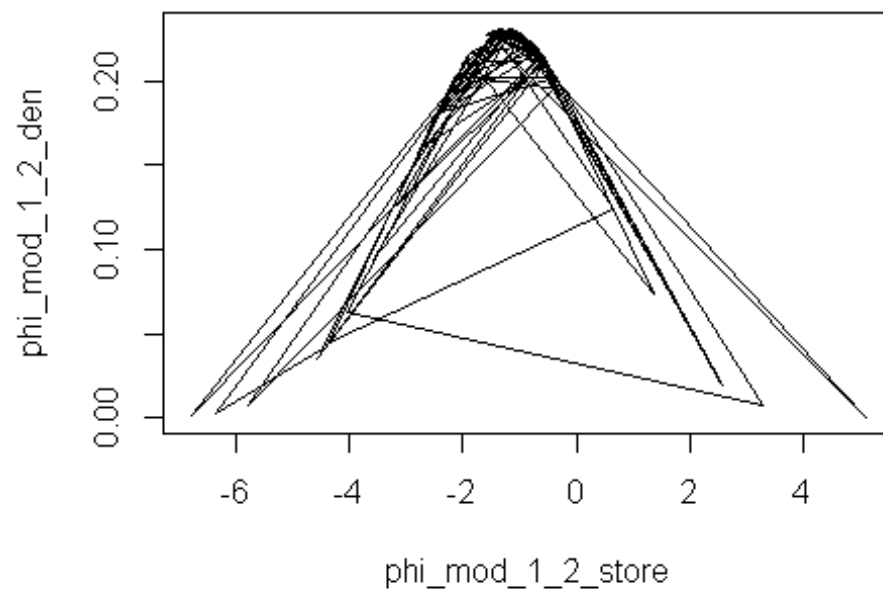
```
phi_mod_2_1_den=dnorm(phi_mod_2_1_store,mean=phi_mod_2_1_mean,sd=sqrt(phi_mod_2_1_v
ar))
plot(phi_mod_2_1_store,phi_mod_2_1_den,type='l')
```



```
phi_mod_intercept_2_mean=mean(phi_mod_intercept_2_store)
phi_mod_intercept_2_var=var(phi_mod_intercept_2_store)
phi_mod_intercept_2_den=dnorm(phi_mod_intercept_2_store,mean=phi_mod_intercept_2_mean,sd
=sqrt(phi_mod_intercept_2_var))
plot(phi_mod_intercept_2_store,phi_mod_intercept_2_den,type='l')
```



```
phi_mod_1_2_mean=mean(phi_mod_1_2_store)
phi_mod_1_2_var=var(phi_mod_1_2_store)
phi_mod_1_2_den=dnorm(phi_mod_1_2_store,mean=phi_mod_1_2_mean,sd=sqrt(phi_mod_1_2_v
ar))
plot(phi_mod_1_2_store,phi_mod_1_2_den,type='l')
```



Variable	From tar routine	Mean	Variance
Threshold	175	156.66	1290.74
Intercept ¹	-0.165	-0.419	0.3845
-d ₁ ¹	-0.0248	-0.107	0.0262
-d ₂ ¹	0.422	0.272	0.0042
Intercept ²	0.366	-0.13	6.9465
-d ₁ ²	-1.88	-1.74	1.5018

The delay is 1.

This model satisfies since the residuals are white based on looking at the ACF and PACF and by running Box.test on the residuals.

2.R

```
library(TSA)
```

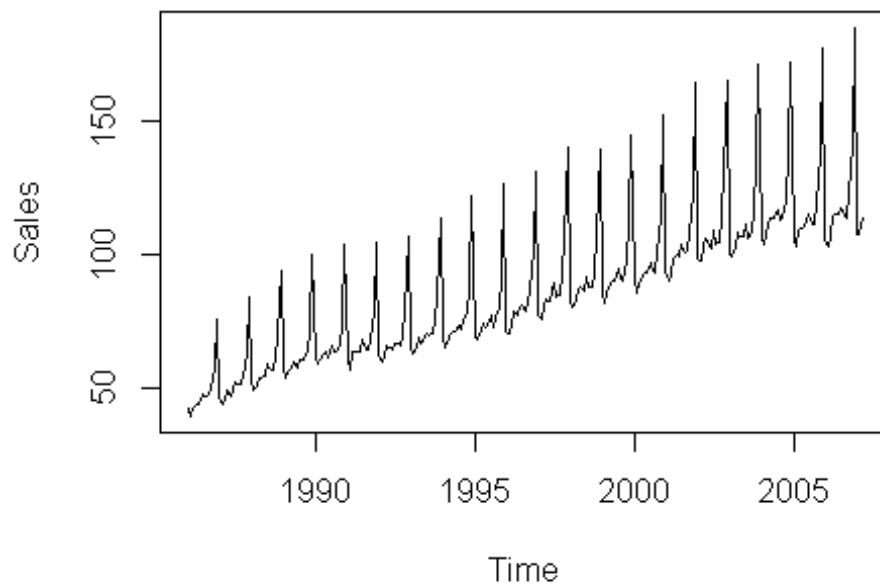
```
## Loading required package: leaps
## Loading required package: locfit
## locfit 1.5-9.1 2013-03-22
## Loading required package: mgcv
## Loading required package: nlme
## This is mgcv 1.8-0. For overview type 'help("mgcv-package")'.
## Loading required package: tseries
##
## Attaching package: 'TSA'
##
## The following objects are masked from 'package:stats':
##
##   acf, arima
##
## The following object is masked from 'package:utils':
##
##   tar
```

```
library(astsa)
```

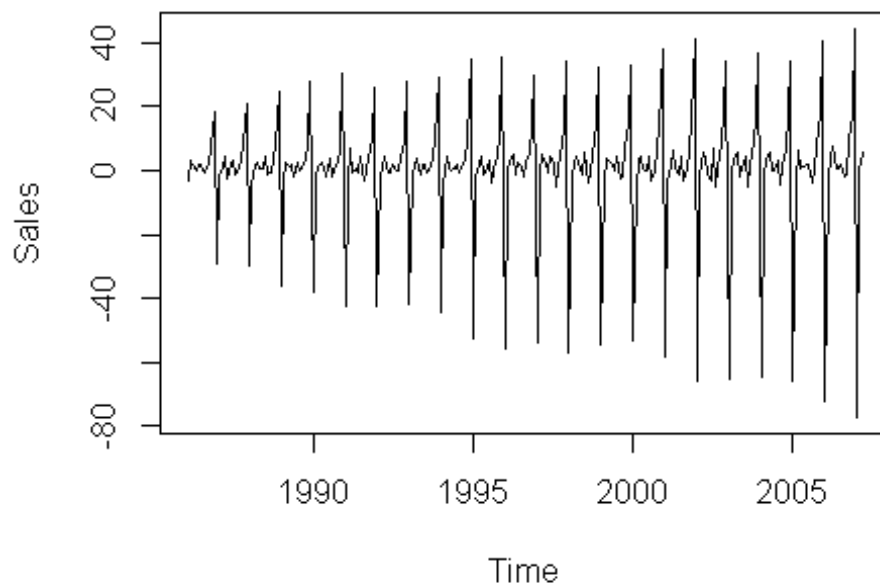
```
library(forecast)
```

```
## Loading required package: zoo
##
## Attaching package: 'zoo'
##
## The following objects are masked from 'package:base':
##
##   as.Date, as.Date.numeric
##
## Loading required package: timeDate
##
## Attaching package: 'timeDate'
##
## The following objects are masked from 'package:TSA':
##
##   kurtosis, skewness
##
## This is forecast 5.6
##
## Attaching package: 'forecast'
##
## The following object is masked from 'package:astsa':
##
##   gas
##
## The following object is masked from 'package:TSA':
##
```

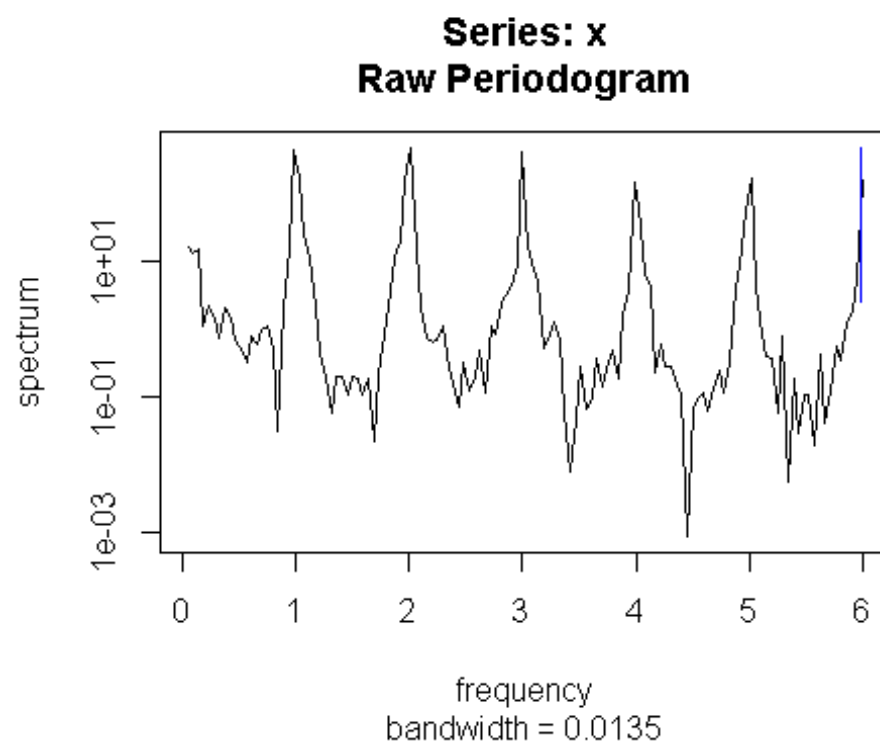
```
## fitted.Arima
##
## The following object is masked from 'package:nlme':
##
## getResponse
data(retail)
#retail_new=retail[1:255]
plot(retail,type='l')
```



```
retail_diff=diff(retail)
plot(retail_diff)# d=1, D=1
```

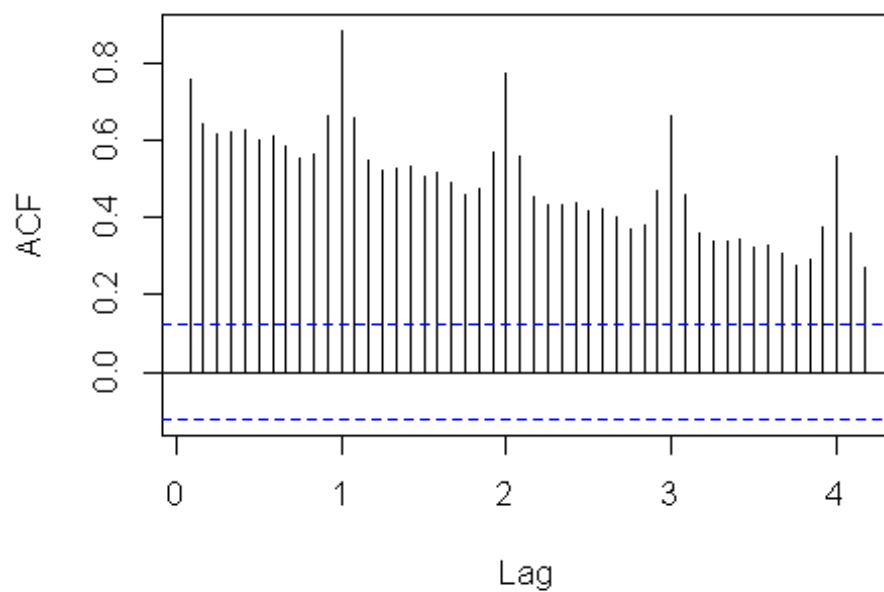


```
spectrum(retail)# S=12
```



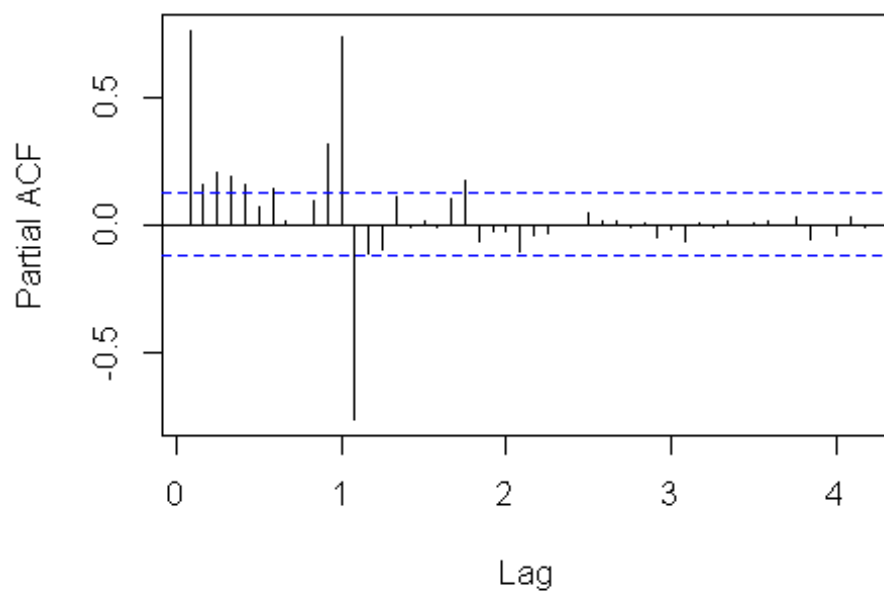
```
retail_acf=acf(retail,lag.max=50)#q=2,Q=1
```

Series retail



```
retail_pacf=pacf(retail,lag.max=50)#p=2,P=0
```

Series retail

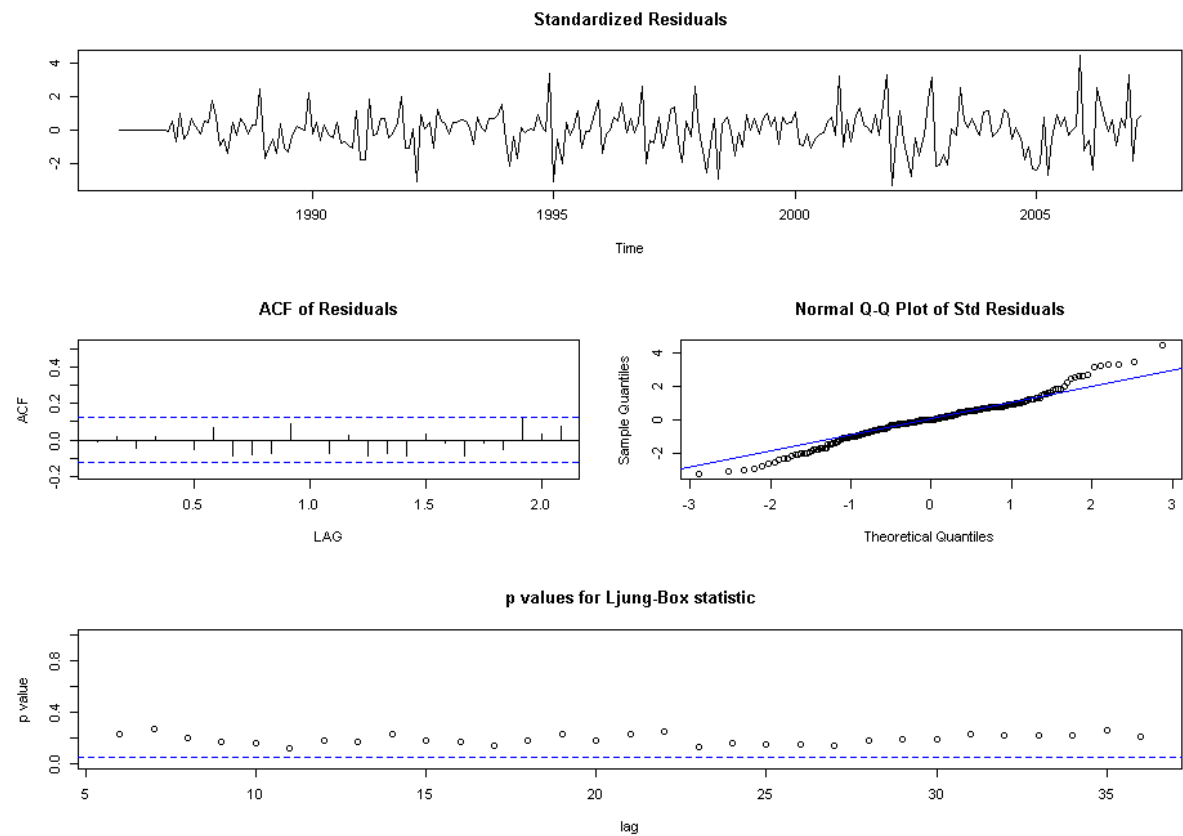


```
retail_test=sarima(retail,2,1,2,D=1,Q=1,S=12)
```

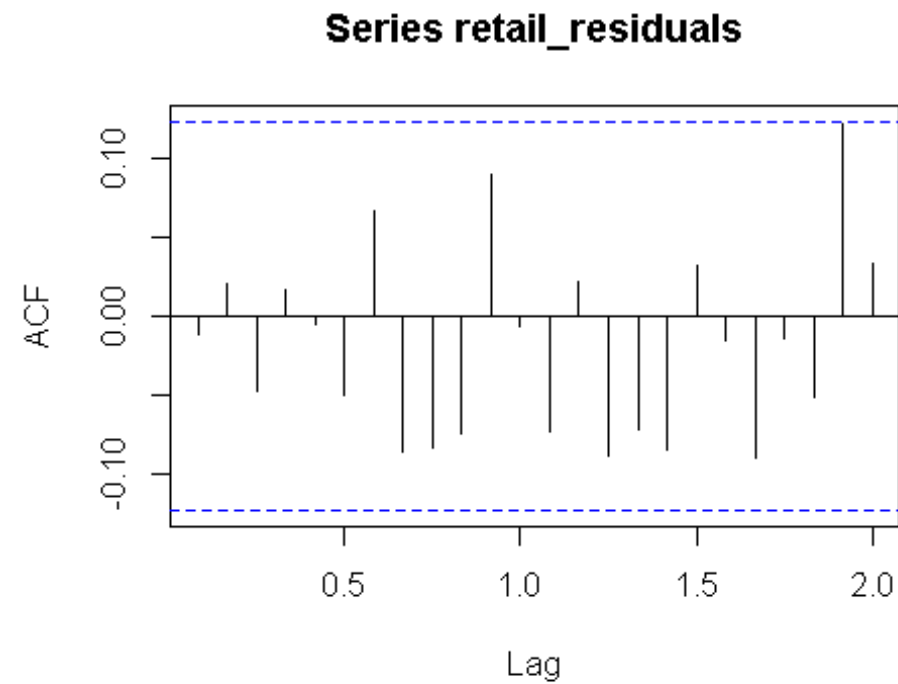
```
## initial value 0.840627  
## iter 2 value 0.685920  
## iter 3 value 0.645495  
## iter 4 value 0.645147
```



```
## iter 5 value 0.644846
## iter 6 value 0.644707
## iter 7 value 0.644628
## iter 8 value 0.644592
## iter 9 value 0.644499
## iter 10 value 0.644275
## iter 11 value 0.643740
## iter 12 value 0.642994
## iter 13 value 0.642226
## iter 14 value 0.641474
## iter 15 value 0.641163
## iter 16 value 0.641076
## iter 17 value 0.641016
## iter 18 value 0.640740
## iter 19 value 0.640226
## iter 20 value 0.639682
## iter 21 value 0.639121
## iter 22 value 0.638806
## iter 23 value 0.638802
## iter 24 value 0.638802
## iter 25 value 0.638802
## iter 26 value 0.638802
## iter 27 value 0.638802
## iter 28 value 0.638802
## iter 29 value 0.638802
## iter 30 value 0.638802
## iter 31 value 0.638801
## iter 32 value 0.638801
## iter 32 value 0.638801
## iter 32 value 0.638801
## final value 0.638801
## converged
## initial value 0.632797
## iter 2 value 0.632772
## iter 3 value 0.632748
## iter 4 value 0.632689
## iter 5 value 0.632626
## iter 6 value 0.632520
## iter 7 value 0.632494
## iter 8 value 0.632480
## iter 9 value 0.632477
## iter 10 value 0.632475
## iter 11 value 0.632475
## iter 11 value 0.632475
## iter 11 value 0.632475
## final value 0.632475
## converged
```

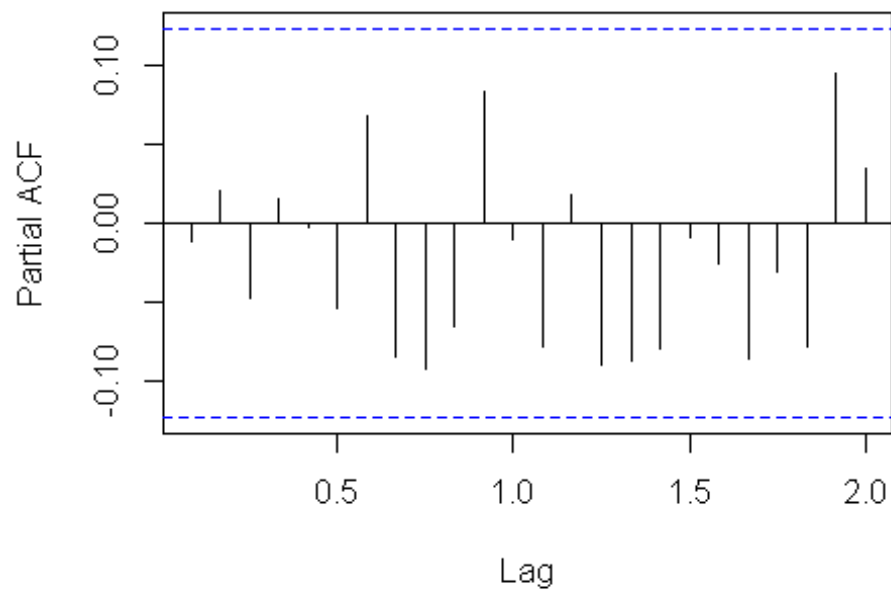


```
retail_residuals=retail_test$fit$residuals
acf(retail_residuals)
```



```
pacf(retail_residuals)
```

Series retail_residuals



The Model is $ARIMA(2,1,2) \times (0,1,1)_{12}$ as seen based on the values and plots above.

This model satisfies since the residuals are white based on looking at the ACF and PACF and by running Box.test on the residuals.