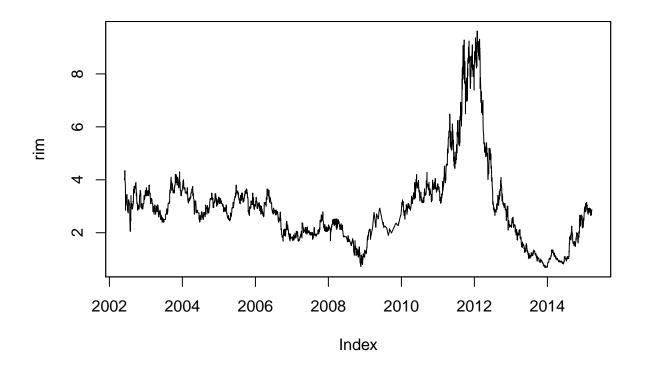
a4q3

```
part (a)
```

```
library("tseries")
rim<-get.hist.quote(instrument="rim",quote="Close")</pre>
```

time series starts 2002-05-31

```
par(mfcol=c(1,1))
plot(rim)
```

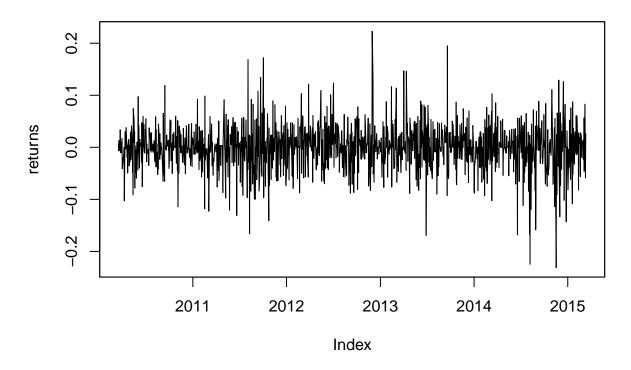


part b

```
rim<-get.hist.quote(instrument="rim",quote="Close",start="2010-03-17",end="2015-03-16")
returns<-log(rim/lag(rim,h=-1))</pre>
```

 $part\ c$

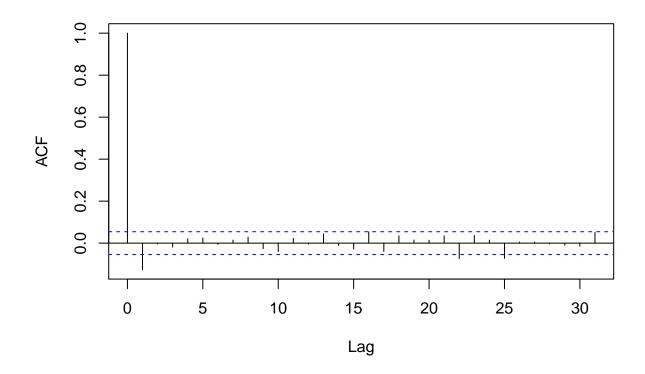
```
plot(returns)
```



part d

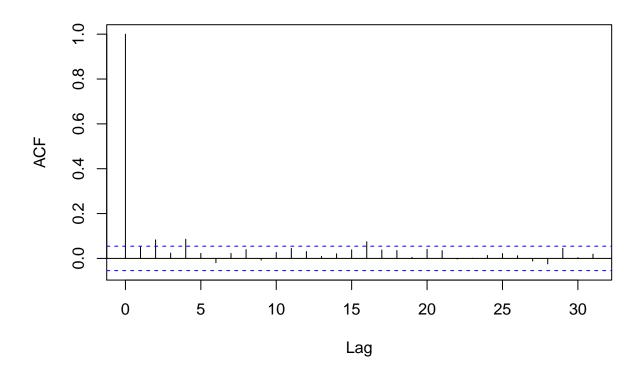
returns<-as.numeric(returns)
acf(returns)</pre>

Series returns



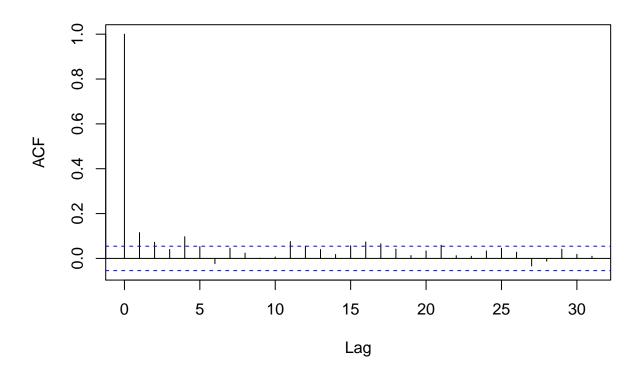
acf(returns^2)

Series returns^2



acf(abs(returns))

Series abs(returns)



```
par(mfcol=c(1,2))

part e

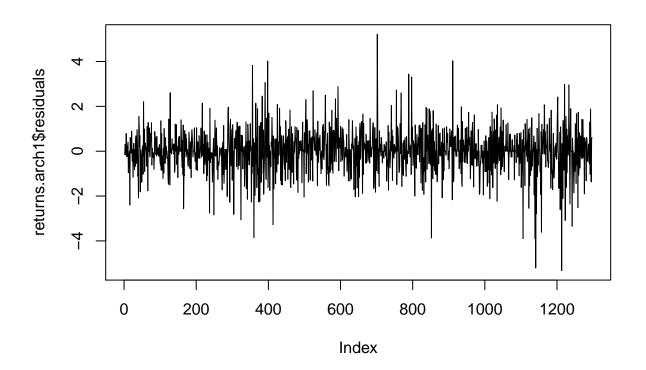
returns.arch1 <- garch(returns, order=c(0,1), trace=F)
summary(returns.arch1)

##
## Gall</pre>
```

```
## garch(x = returns, order = c(0, 1), trace = F)
##
## Model:
## GARCH(0,1)
##
## Residuals:
##
               1Q Median
                                      Max
  -5.3258 -0.5088 0.0000 0.5779
##
                                  5.2184
##
## Coefficient(s):
      Estimate Std. Error
                           t value Pr(>|t|)
## a0 1.828e-03
                 5.332e-05
                             34.290
                                      <2e-16 ***
## a1 5.890e-02
                 2.502e-02
                              2.354
                                      0.0186 *
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
```

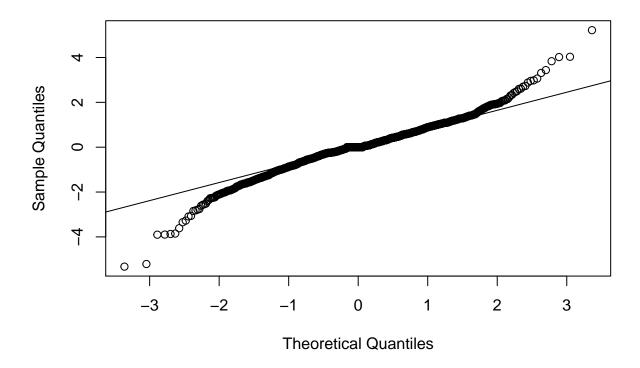
```
## Diagnostic Tests:
## Jarque Bera Test
##
## data: Residuals
## X-squared = 492.8855, df = 2, p-value < 2.2e-16
##
##
## Box-Ljung test
##
## data: Squared.Residuals
## X-squared = 0.0079, df = 1, p-value = 0.9291</pre>
```

plot(returns.arch1\$residuals,type="l")



qqnorm(returns.arch1\$residuals)
qqline(returns.arch1\$residuals)

Normal Q-Q Plot



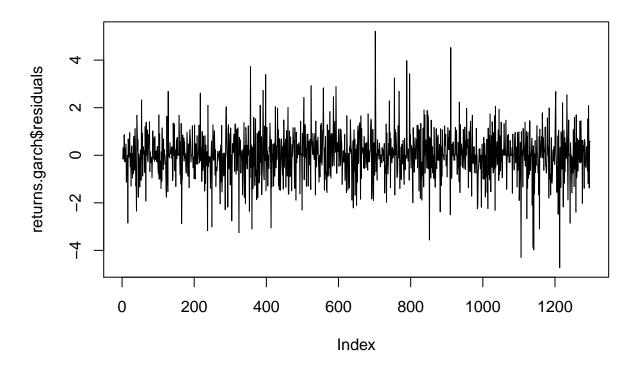
part f

```
returns.garch <- garch(returns, order=c(1,1), trace=F)
summary(returns.garch)</pre>
```

```
##
## garch(x = returns, order = c(1, 1), trace = F)
##
## Model:
## GARCH(1,1)
##
## Residuals:
##
      Min
               1Q Median
                               ЗQ
                                      Max
## -4.7300 -0.5077 0.0000 0.5689
                                  5.2114
##
## Coefficient(s):
      Estimate Std. Error t value Pr(>|t|)
##
## a0 1.328e-04
                 3.352e-05
                              3.962 7.42e-05 ***
## a1 6.097e-02
                              5.473 4.42e-08 ***
                 1.114e-02
## b1 8.724e-01
                 2.424e-02
                             35.996 < 2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Diagnostic Tests:
   Jarque Bera Test
```

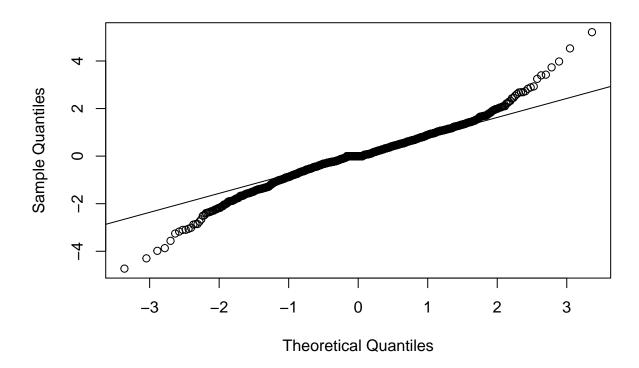
```
##
## data: Residuals
## X-squared = 326.8644, df = 2, p-value < 2.2e-16
##
##
## Box-Ljung test
##
## data: Squared.Residuals
## X-squared = 0.4095, df = 1, p-value = 0.5222</pre>
```

plot(returns.garch\$residuals,type="1")



qqnorm(returns.garch\$residuals)
qqline(returns.garch\$residuals)

Normal Q-Q Plot



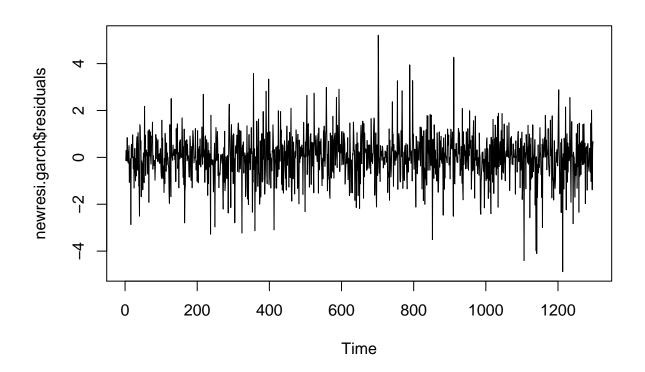
part g

```
new<-arima(returns,order=c(0,0,1),method="ML")
newresi.garch<-garch(new$residuals, order=c(1,1), trace=F)
summary(newresi.garch)</pre>
```

```
##
## Call:
## garch(x = new$residuals, order = c(1, 1), trace = F)
##
## Model:
## GARCH(1,1)
##
## Residuals:
##
       Min
                  1Q
                      Median
                                   ЗQ
  -4.87803 -0.50408 0.01952 0.55300 5.21028
##
##
## Coefficient(s):
##
       Estimate Std. Error t value Pr(>|t|)
                 3.068e-05
                              3.875 0.000107 ***
## a0 1.189e-04
## a1 5.611e-02
                  1.023e-02
                              5.486 4.12e-08 ***
## b1 8.830e-01
                  2.264e-02
                              39.002 < 2e-16 ***
##
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Diagnostic Tests:
```

```
## Jarque Bera Test
##
## data: Residuals
## X-squared = 342.9794, df = 2, p-value < 2.2e-16
##
##
## Box-Ljung test
##
## data: Squared.Residuals
## X-squared = 0.2711, df = 1, p-value = 0.6026

plot(newresi.garch$residuals,type="l")</pre>
```



```
qqnorm(newresi.garch$residuals)
qqline(newresi.garch$residuals)
```

Normal Q-Q Plot

