# Untitled

2024-04-22

## Extract data

Draw the Amazon stock data from 2023-01-01 to 2023-12-31 as train set

```
library(quantmod)
## Loading required package: xts
## Loading required package: zoo
##
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
       as.Date, as.Date.numeric
## Loading required package: TTR
## Registered S3 method overwritten by 'quantmod':
##
     method
     as.zoo.data.frame zoo
##
getSymbols("AMZN",src='yahoo',return.class='ts', from = '2023-01-01', to = '2023-12-31')
## [1] "AMZN"
train <- AMZN[,4]</pre>
length(train)
## [1] 250
```

Draw the Amazon stock data from 2024-01-01 to 2024-01-31 as test set

```
getSymbols("AMZN",src='yahoo',return.class='ts', from = '2024-01-01', to = '2024-01-31')
## [1] "AMZN"
```

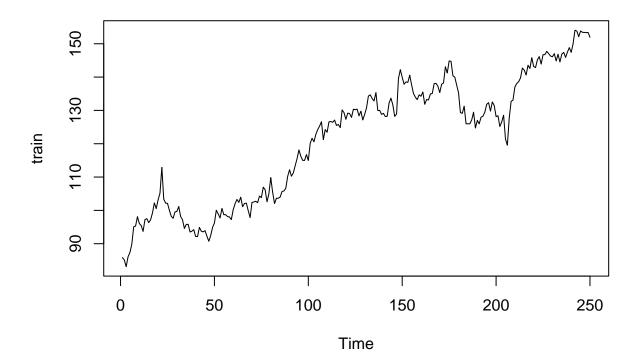
```
test <- AMZN[,4]
length(test)</pre>
```

## [1] 20

# EDA

we can first observe the time series plot of the data. And we can see that the  $scale(10^2)$  is large, and there is trend on the long term.

```
ts.plot(train)
```



So we can reduce the scale by log transformation and fit a trend model for the data.

```
train_set <- log(train)
test_set <- log(test)
tfit <- time(train_set)</pre>
```

## Remove the Deterministic trend

#### linear trend model

```
mlr.lin <- lm(train_set ~ tfit)
summary(mlr.lin)
##
## Call:
## lm(formula = train_set ~ tfit)
## Residuals:
       Min
                 1Q Median
                                      30
## -0.168507 -0.044383 -0.005994 0.049328 0.153878
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 4.527e+00 7.544e-03 600.1 <2e-16 ***
## tfit
             2.064e-03 5.211e-05 39.6 <2e-16 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
## Residual standard error: 0.05946 on 248 degrees of freedom
## Multiple R-squared: 0.8635, Adjusted R-squared: 0.8629
## F-statistic: 1568 on 1 and 248 DF, p-value: < 2.2e-16
```

#### quadratic trend model

```
tsqfit <- tfit^2/factorial(2)
mlr.quad <- lm(train_set ~ tfit + tsqfit)
summary(mlr.quad)</pre>
```

```
## Call:
## lm(formula = train_set ~ tfit + tsqfit)
## Residuals:
##
                  1Q
                        Median
## -0.162385 -0.049980 0.009945 0.043060 0.180370
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 4.477e+00 1.055e-02 424.422 < 2e-16 ***
              3.272e-03 1.940e-04 16.861 < 2e-16 ***
## tfit
## tsqfit
             -9.627e-06 1.497e-06 -6.429 6.59e-10 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
```

```
##
## Residual standard error: 0.05515 on 247 degrees of freedom
## Multiple R-squared: 0.883, Adjusted R-squared: 0.8821
## F-statistic: 932.3 on 2 and 247 DF, p-value: < 2.2e-16</pre>
```

#### cubic trend model

```
tcubfit <- tfit^3/factorial(3)</pre>
mlr.cub <- lm(train_set ~ tfit + tsqfit + tcubfit)</pre>
summary(mlr.cub)
##
## Call:
## lm(formula = train_set ~ tfit + tsqfit + tcubfit)
## Residuals:
##
       Min
                 1Q
                     Median
                                   3Q
## -0.16760 -0.04541 0.01161 0.04416 0.17727
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 4.494e+00 1.409e-02 319.025 < 2e-16 ***
              2.424e-03 4.851e-04 4.997 1.1e-06 ***
## tfit
## tsqfit
              7.221e-06 8.973e-06 0.805
                                            0.4217
## tcubfit
             -1.342e-07 7.050e-08 -1.904
                                            0.0581 .
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.05486 on 246 degrees of freedom
## Multiple R-squared: 0.8847, Adjusted R-squared: 0.8833
## F-statistic: 629.3 on 3 and 246 DF, p-value: < 2.2e-16
```

#### quartic trend model

```
tquarfit <- tfit^4/factorial(4)
mlr.quar <- lm(train_set ~ tfit + tsqfit + tcubfit + tquarfit)
summary(mlr.quar)</pre>
```

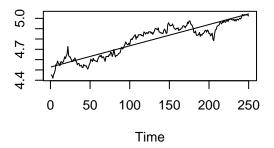
```
##
## Call:
## lm(formula = train_set ~ tfit + tsqfit + tcubfit + tquarfit)
##
## Residuals:
## Min 1Q Median 3Q Max
## -0.156319 -0.025418 0.003902 0.028243 0.190171
##
## Coefficients:
```

```
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept)
               4.591e+00 1.468e-02 312.807 < 2e-16 ***
              -5.109e-03
                         8.067e-04 -6.332 1.14e-09 ***
               2.761e-04
                          2.607e-05 10.591
                                            < 2e-16 ***
## tsqfit
## tcubfit
               -5.126e-06
                          4.677e-07 -10.961
                                             < 2e-16 ***
                         3.697e-09 10.758
## tquarfit
               3.977e-08
                                             < 2e-16 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Residual standard error: 0.0453 on 245 degrees of freedom
## Multiple R-squared: 0.9217, Adjusted R-squared: 0.9204
## F-statistic: 721.1 on 4 and 245 DF, p-value: < 2.2e-16
```

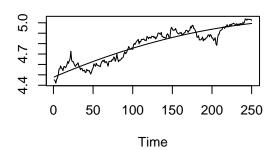
#### trend model selection

```
par(mfrow=c(2,2))
plin=cbind(train_set,mlr.lin$fitted)
ts.plot(plin,main="xfit and fit.linear")
pquad=cbind(train_set,mlr.quad$fitted)
ts.plot(pquad,main="xfit and fit.quadratic")
pcub=cbind(train_set,mlr.cub$fitted)
ts.plot(pcub,main="xfit and fitt.cubic")
pquar=cbind(train_set,mlr.quar$fitted)
ts.plot(pquar,main="xfit and fitt.quartic")
```

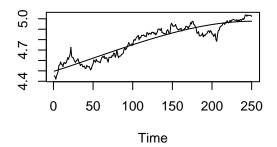
#### xfit and fit.linear



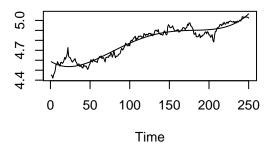
### xfit and fit.quadratic



### xfit and fitt.cubic



xfit and fitt.quartic



#### in-sample metric

we can compare the trend models by AIC

```
nfit <- length(train_set)</pre>
AIC.lin <- AIC(mlr.lin)/nfit
AIC.quad <- AIC(mlr.quad)/nfit
AIC.cub <- AIC(mlr.cub)/nfit
AIC.quar <- AIC(mlr.quar)/nfit
data.frame(
 model = c("lin", "quad", "cub", "quar"),
 AIC = c(AIC.lin, AIC.quad, AIC.cub, AIC.quar)
)
##
    model
                AIC
## 1 lin -2.791029
## 2 quad -2.937741
## 3
     cub -2.944371
## 4 quar -3.323230
```

By AIC, we can see that quartic trend model has the best effect on the combination of fitting and complexity.

#### out of sample metric

we can calculate the MAPE of each model.

```
new <- data.frame(tfit=c(378:397))</pre>
pfore.lin <- predict(mlr.lin,new,se.fit = TRUE)</pre>
efore.lin <- test_set - pfore.lin$fit</pre>
tfit <- c(378:397)
tsqfit <- tfit^2/factorial(2)</pre>
mat <- matrix(c(tfit,tsqfit),nrow=20,ncol=2,dimnames = list(c(),c("tfit","tsqfit")))</pre>
newnq <- data.frame(mat)</pre>
pfore.quad <- predict(mlr.quad,newnq,se.fit = TRUE)</pre>
efore.quad <- test_set - pfore.quad$fit</pre>
tfit <- c(378:397)
tcubfit <- tfit^3/factorial(3)</pre>
mat <- matrix(c(tfit,tsqfit,tcubfit),nrow=20,ncol=3, dimnames = list(c(),c("tfit","tsqfit","tcubfit")))</pre>
newnc <- data.frame(mat)</pre>
pfore.cub <- predict(mlr.cub,newnc,se.fit = TRUE)</pre>
efore.cub <- test_set - pfore.cub$fit
tfit <- c(378:397)
tquarfit <- tfit^4/factorial(4)</pre>
mat <- matrix(c(tfit,tsqfit,tcubfit,tquarfit),nrow=20,ncol=4,</pre>
                dimnames = list(c(),c("tfit","tsqfit","tcubfit","tquarfit")))
newnc <- data.frame(mat)</pre>
pfore.quar <- predict(mlr.quar,newnc,se.fit = TRUE)</pre>
efore.quar <- test_set - pfore.quar$fit</pre>
```

```
mape.lin <- 100*(mean(abs((efore.lin)/test_set)))
mape.quad <- 100*(mean(abs((efore.quad)/test_set)))
mape.cub <- 100*(mean(abs((efore.cub)/test_set)))
mape.quar <- 100*(mean(abs((efore.quar)/test_set)))

data.frame(
  model = c("lin", "quad", "cub", "quar"),
  MAPE = c(mape.lin, mape.quad, mape.cub, mape.quar)</pre>
```

```
## 1 din 5.8314833
## 2 quad 0.5580105
## 3 cub 7.1480535
## 4 quar 118.7957671
```

We can see that the quadratic model has the smallest MAPE, which means that it has the best predictive performance. Although the quartic model do well in AIC, but it is a overfitted model, it has too large MAPE.

Considering the AIC and MAPE, I decide to use quadratic model to remove the deterministic trend for further analyze.

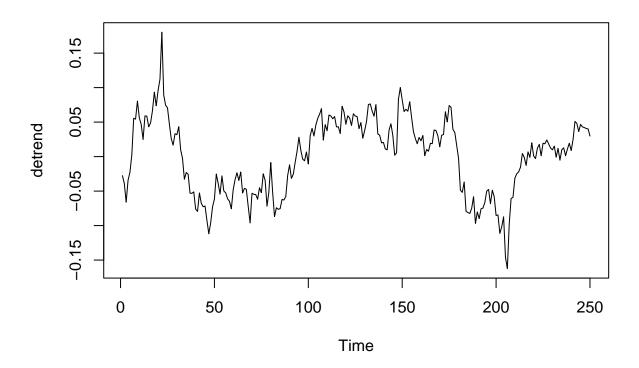
### Stochastic trend

After removing the deterministic trend, we need to explore the stochastic trend of the data to help us better forecast.

```
detrend <- mlr.quad$resid
detrend_test <- efore.quad</pre>
```

let's see the time series plot of the data we need to analyze now.

```
ts.plot(detrend)
```



Well, the mean is around 0, and the values volatile between around -0.15 and 0.15.

## Sationarity check And let's check whether the data is stationary or not.

```
library(fUnitRoots)
adfTest(detrend, lags=10, type = "c")
##
## Title:
##
    Augmented Dickey-Fuller Test
##
##
   Test Results:
     PARAMETER:
##
##
       Lag Order: 10
##
     STATISTIC:
##
       Dickey-Fuller: -2.3845
     P VALUE:
##
##
       0.169
##
## Description:
    Mon Apr 22 23:53:18 2024 by user:
```

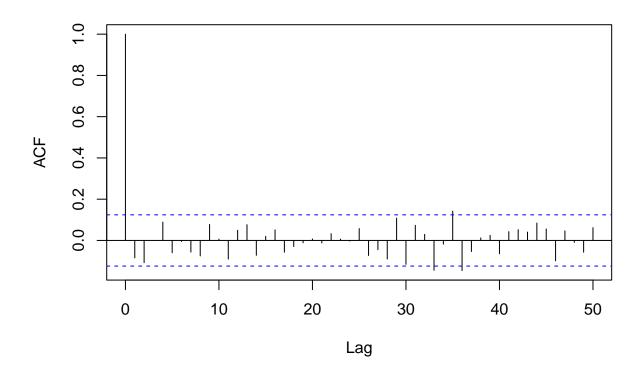
From the Augmented Dickey-Fuller Test, we can know its p-value is significantly large, so we need to difference the data to make it statioanry.

```
diff <- diff(detrend)</pre>
adfTest(diff, lags=10, type = "c")
## Warning in adfTest(diff, lags = 10, type = "c"): p-value smaller than printed
## p-value
##
## Title:
##
    Augmented Dickey-Fuller Test
##
## Test Results:
     PARAMETER:
##
##
       Lag Order: 10
##
     STATISTIC:
##
       Dickey-Fuller: -5.3002
     P VALUE:
##
       0.01
##
##
## Description:
    Mon Apr 22 23:53:18 2024 by user:
```

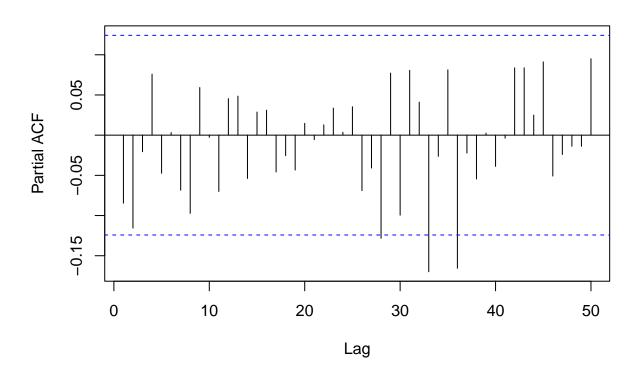
After differencing, we can see that the data is stationary now.

```
acf(diff, lag.max = 50)
```

# Series diff



# Series diff



From the acf and pacf plot, we can barely observe any autocorrelation.

```
ar(x = diff)
```

```
##
## Call:
## ar(x = diff)
##
## Coefficients:
## 1 2
## -0.0944 -0.1156
##
## Order selected 2 sigma^2 estimated as 0.0004243
```

But I still want to fit an arima model.

And the ar() function recommend me to fit AR(2).

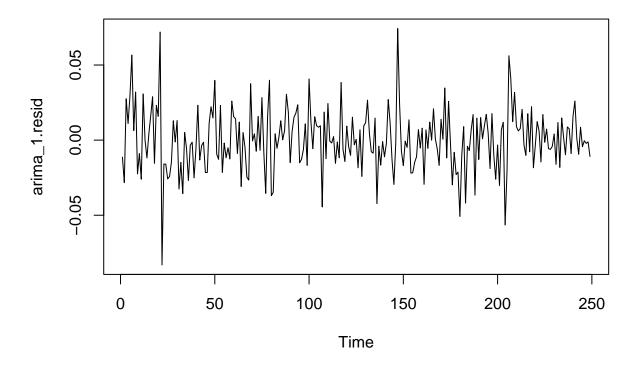
# ARIMA(2,1,0)

```
arima_1 <- arima(diff, order=c(2,0,0), include.mean = TRUE)
arima_1</pre>
```

```
##
## Call:
  arima(x = diff, order = c(2, 0, 0), include.mean = TRUE)
##
##
  Coefficients:
##
##
                           intercept
                              0.0003
##
         -0.0947
                  -0.1161
          0.0629
                   0.0630
                              0.0011
## s.e.
##
## sigma^2 estimated as 0.0004191: log likelihood = 614.97, aic = -1221.93
```

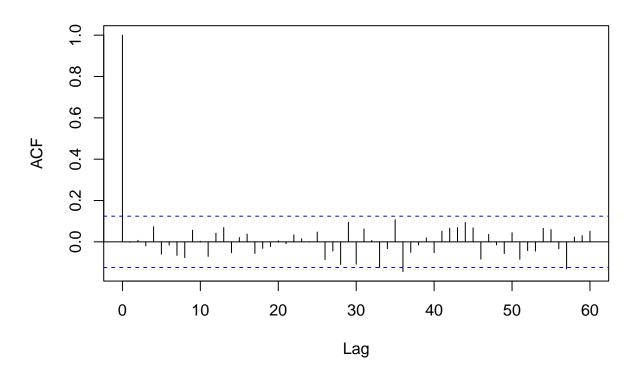
let's see check the residuals of the arima model

```
arima_1.resid <- arima_1$residuals
ts.plot(arima_1.resid)</pre>
```



```
acf(arima_1.resid, lag.max = 60)
```

# Series arima\_1.resid



```
Box.test(arima_1.resid, lag=30, fitdf=1, type = c("Ljung-Box"))
```

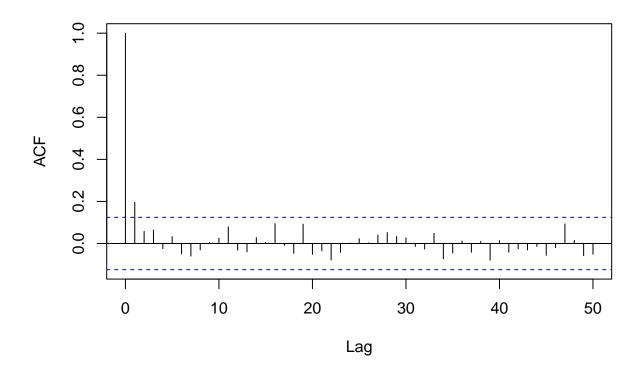
```
##
## Box-Ljung test
##
## data: arima_1.resid
## X-squared = 24.332, df = 29, p-value = 0.7124
```

From the acf plot and the result of Box-Ljung test, we can know that the mean model is adequate. And let's check whether there is some ARCH effect.

## Heteroscedasticity check

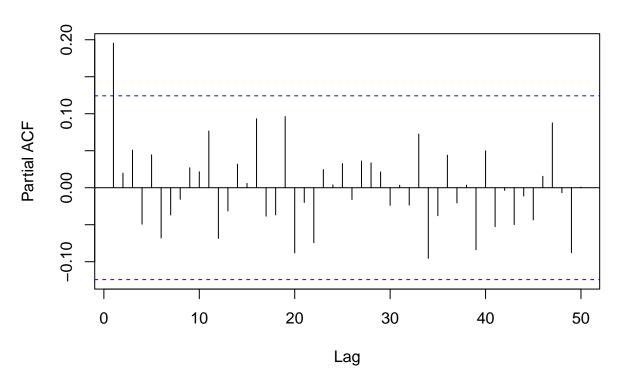
```
acf((arima_1.resid)^2, lag.max = 50)
```

# Series (arima\_1.resid)^2



pacf((arima\_1.resid)^2, lag.max = 50)

# Series (arima\_1.resid)^2



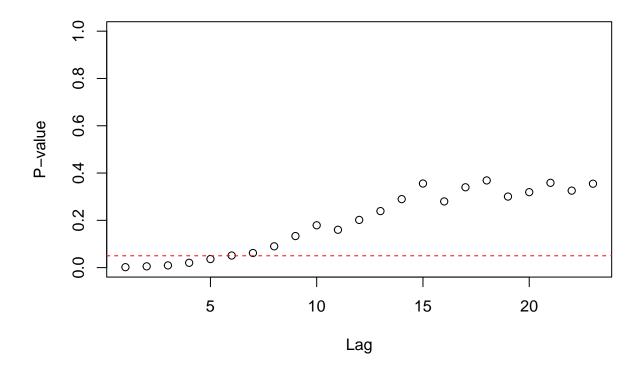
# library(TSA)

```
##
## Attaching package: 'TSA'

## The following objects are masked from 'package:stats':
##
## acf, arima

## The following object is masked from 'package:utils':
##
## tar

McLeod.Li.test(y = arima_1.resid)
```



From the acf and pacf plot of the square of the residuals, we can see there is heteroscedasticity, and we need to fit some ARCH model to improve it.

# Heteroscedastic Model

# ARCH(1)

we can first try ARCH(1) model.

#### library(fGarch)

```
## NOTE: Packages 'fBasics', 'timeDate', and 'timeSeries' are no longer
## attached to the search() path when 'fGarch' is attached.
##
## If needed attach them yourself in your R script by e.g.,
## require("timeSeries")

##
## Attaching package: 'fGarch'

## The following object is masked from 'package:TTR':
##
## volatility
```

```
arch_1 <- garchFit(~garch(1,0), data = arima_1.resid, trace=FALSE,</pre>
                   cond.dist=c("norm"), include.mean=FALSE)
summary(arch_1)
##
## Title:
## GARCH Modelling
## Call:
   garchFit(formula = ~garch(1, 0), data = arima_1.resid, cond.dist = c("norm"),
       include.mean = FALSE, trace = FALSE)
##
## Mean and Variance Equation:
## data ~ garch(1, 0)
## <environment: 0x1145a3dd0>
## [data = arima_1.resid]
## Conditional Distribution:
## norm
##
## Coefficient(s):
##
      omega
                alpha1
## 0.0003667 0.1147555
##
## Std. Errors:
## based on Hessian
## Error Analysis:
          Estimate Std. Error t value Pr(>|t|)
## omega 3.667e-04 3.938e-05
                                9.312
                                          <2e-16 ***
## alpha1 1.148e-01
                    6.808e-02
                                1.686
                                          0.0919 .
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Log Likelihood:
## 618.222
              normalized: 2.482819
##
## Description:
## Mon Apr 22 23:53:19 2024 by user:
##
##
## Standardised Residuals Tests:
##
                                   Statistic
                                                  p-Value
## Jarque-Bera Test
                      R
                           Chi^2 18.3431389 0.0001039532
## Shapiro-Wilk Test R
                                   0.9858868 0.0147014066
                           W
## Ljung-Box Test
                      R
                           Q(10)
                                   4.8716325 0.8995863699
## Ljung-Box Test
                      R
                           Q(15)
                                   9.8557127 0.8287164482
## Ljung-Box Test
                      R
                           Q(20) 11.9929702 0.9163177318
## Ljung-Box Test
                      R^2 Q(10)
                                   4.8915148 0.8983060284
## Ljung-Box Test
                      R^2 Q(15)
                                   7.1105536 0.9545090009
## Ljung-Box Test
                      R<sup>2</sup> Q(20) 12.8840517 0.8823032312
```

6.7969287 0.8707363449

## LM Arch Test

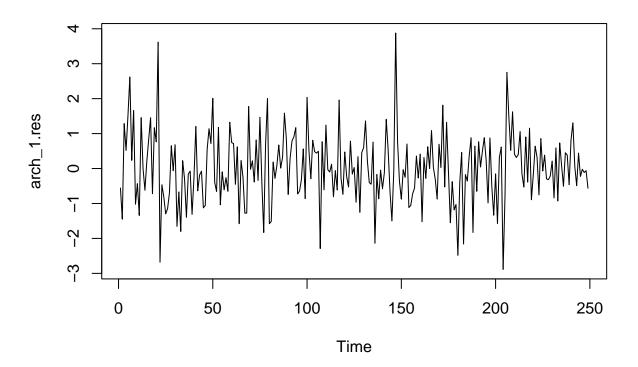
R.

TR^2

```
##
## Information Criterion Statistics:
## AIC BIC SIC HQIC
## -4.949574 -4.921322 -4.949702 -4.938202
```

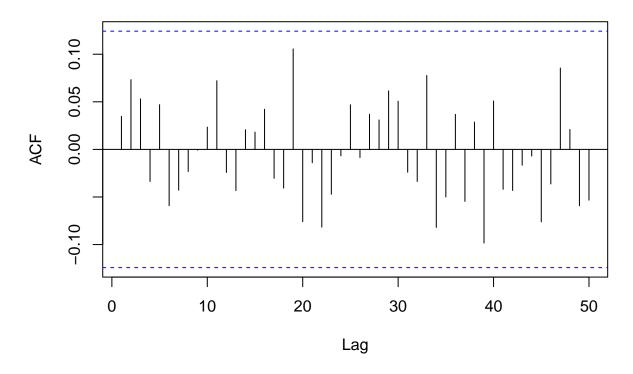
Let's check the residuals of the model.

```
arch_1.res <- residuals(arch_1, standardize=TRUE)
ts.plot(arch_1.res)</pre>
```



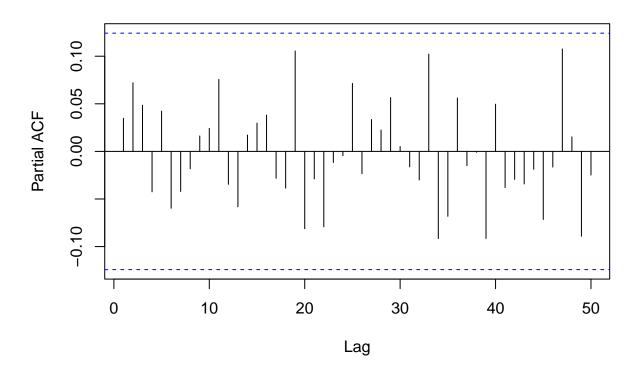
```
acf((arch_1.res)^2, lag.max = 50)
```

# Series (arch\_1.res)^2

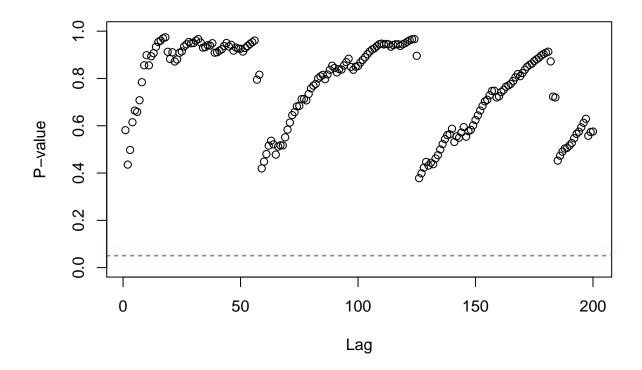


pacf((arch\_1.res)^2, lag.max = 50)

# Series (arch\_1.res)^2



McLeod.Li.test(y = arch\_1.res, gof.lag = 200)



The acf and pacf plot shows there is no further ARCH effect, and McLeod-Li test shows that the model is adequate. But it shows some abnormal pattern that I cannot explain. Let's check the normality of the residuals.

```
shapiro.test(arch_1.res)
```

```
##
## Shapiro-Wilk normality test
##
## data: arch_1.res
## W = 0.98589, p-value = 0.0147
```

The Shapiro-Wilk Normality Test shows that the residuals is not a white noise under the significant level  $\alpha = 0.05$ .

So we need to fit other models or greater orders.

# GARCH(1,1)

```
garch_11 <- garchFit(~arma(2,0)+garch(1,1), data=arima_1.resid, trace=FALSE, cond.dist=c("norm"), inclusion
summary(garch_11)</pre>
```

##

```
## Title:
## GARCH Modelling
##
## Call:
##
   garchFit(formula = ~arma(2, 0) + garch(1, 1), data = arima_1.resid,
##
       cond.dist = c("norm"), include.mean = FALSE, trace = FALSE)
## Mean and Variance Equation:
## data ~ arma(2, 0) + garch(1, 1)
## <environment: 0x113910860>
## [data = arima_1.resid]
## Conditional Distribution:
## norm
##
## Coefficient(s):
##
                                               alpha1
                                                             beta1
          ar1
                        ar2
                                   omega
## -0.06446519
                0.04390722
                              0.00017196
                                           0.16109034
                                                        0.42513185
##
## Std. Errors:
## based on Hessian
## Error Analysis:
           Estimate Std. Error t value Pr(>|t|)
## ar1
                                           0.3859
         -6.447e-02 7.435e-02
                                 -0.867
## ar2
          4.391e-02
                     6.872e-02
                                    0.639
                                            0.5228
                                           0.0271 *
          1.720e-04
                     7.783e-05
                                    2.209
## omega
## alpha1 1.611e-01
                      8.418e-02
                                    1.914
                                           0.0557 .
                                           0.0386 *
## beta1
          4.251e-01
                      2.056e-01
                                    2.068
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Log Likelihood:
## 621.2061
               normalized: 2.494804
## Description:
   Mon Apr 22 23:53:19 2024 by user:
##
##
## Standardised Residuals Tests:
##
                                    Statistic
                                                  p-Value
## Jarque-Bera Test
                           Chi^2 15.4862157 0.0004337215
                      R
## Shapiro-Wilk Test R
                                    0.9881126 0.0376363158
                           W
## Ljung-Box Test
                            Q(10)
                      R
                                    5.6074356 0.8470965898
## Ljung-Box Test
                            Q(15) 10.6279659 0.7784938891
                      R
## Ljung-Box Test
                            Q(20) 12.6848340 0.8904932510
                       R
## Ljung-Box Test
                      R^2 Q(10)
                                    3.3491555 0.9719736875
## Ljung-Box Test
                       R^2 Q(15)
                                    6.9962619 0.9577533939
## Ljung-Box Test
                      R<sup>2</sup> Q(20) 13.4698595 0.8563270952
## LM Arch Test
                       R
                            TR^2
                                    5.9906542 0.9165524389
## Information Criterion Statistics:
        AIC
                  BIC
                            SIC
                                      HQIC
## -4.949447 -4.878815 -4.950232 -4.921017
```

#### shapiro.test(residuals(garch\_11, standardize=TRUE))

```
##
## Shapiro-Wilk normality test
##
## data: residuals(garch_11, standardize = TRUE)
## W = 0.98811, p-value = 0.03764
```

We can see that the data is still not normally distributed under the significant level  $\alpha = 0.05$ .

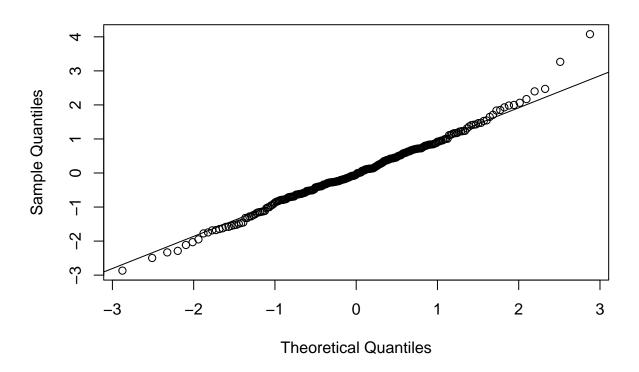
## APARCH(1,1)

Since I observed that in the time series plot, the absolute value of positive peaks(~4) is larger than the absolute value of the negative peaks(~3), so I think that this might be the leverage effect, and I fit a APARCH/TGARCH model.

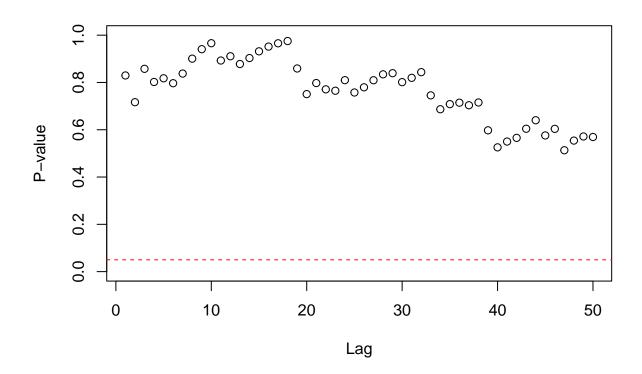
```
##
## Title:
##
    GARCH Modelling
##
## Call:
##
    garchFit(formula = ~arma(2, 0) + aparch(1, 1), data = arima_1.resid,
       delta = 1, cond.dist = c("norm"), include.mean = FALSE, include.delta = F,
##
       trace = FALSE)
##
##
## Mean and Variance Equation:
    data \sim arma(2, 0) + aparch(1, 1)
## <environment: 0x116438a08>
    [data = arima_1.resid]
##
##
## Conditional Distribution:
##
   norm
##
## Coefficient(s):
                                                                       beta1
          ar1
                       ar2
                                 omega
                                             alpha1
                                                         gamma1
## -0.0787227
                0.0327566
                             0.0079085
                                          0.1851578 -0.1844504
                                                                   0.4634344
##
## Std. Errors:
   based on Hessian
##
## Error Analysis:
##
           Estimate
                     Std. Error t value Pr(>|t|)
## ar1
          -0.078723
                        0.076237
                                   -1.033
                                             0.3018
           0.032757
                                    0.495
                                             0.6206
## ar2
                        0.066171
## omega
           0.007909
                        0.004308
                                    1.836
                                             0.0664 .
```

```
## alpha1 0.185158
                    0.076463
                                  2.422
                                          0.0155 *
## gamma1 -0.184450
                      0.255376
                                 -0.722
                                          0.4701
## beta1 0.463434
                      0.226398
                                  2.047
                                          0.0407 *
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
##
## Log Likelihood:
## 622.924
              normalized: 2.501703
##
## Description:
## Mon Apr 22 23:53:19 2024 by user:
##
##
## Standardised Residuals Tests:
##
                                   Statistic
                                                p-Value
## Jarque-Bera Test
                            Chi^2 13.347759 0.001263487
## Shapiro-Wilk Test R
                           W
                                   0.989754 0.075907476
## Ljung-Box Test
                      R
                            Q(10)
                                   5.982691 0.816715344
## Ljung-Box Test
                            Q(15) 11.152999 0.741674999
                      R
## Ljung-Box Test
                            Q(20) 12.893123 0.881922391
                      R
## Ljung-Box Test
                      R<sup>2</sup> Q(10) 3.531179 0.966029079
## Ljung-Box Test
                      R<sup>2</sup> Q(15) 7.796317 0.931688603
## Ljung-Box Test
                      R<sup>2</sup> Q(20) 15.440130 0.750693918
## LM Arch Test
                           TR^2
                                   5.942989 0.918928583
##
## Information Criterion Statistics:
##
        AIC
                  BIC
                            SIC
                                     HQIC
## -4.955213 -4.870455 -4.956338 -4.921097
aparch_11.resid <- residuals(aparch_11, standardize=TRUE)</pre>
qqnorm(aparch_11.resid)
qqline(aparch_11.resid)
```

# Normal Q-Q Plot



McLeod.Li.test(y = aparch\_11.resid, gof.lag = 50)



#### shapiro.test(aparch\_11.resid)

```
##
## Shapiro-Wilk normality test
##
## data: aparch_11.resid
## W = 0.98975, p-value = 0.07591
```

We can see that the residuals is normally distributed now, and the model is adequate.

## Final Model

According to the analyze above, we can get a model with the following expression:

```
\begin{split} X_t &= \mu + 3.272 \times 10^{-3} t - 4.8135 \times 10^{-6} t^2 + Y_t \\ Y_t - Y_{t-1} &= Z_t \\ Z_t &= -0.0787 Z_{t-1} + 0.0328 Z_{t-2} + e_t \\ e_t &= \sigma_t \epsilon_t \\ \sigma_t^2 &= 0.0079 + 0.185 e_{t-1} - 0.184 e_{t-1} I\{e_{t-1} < 0\} + 0.4634 \sigma_{t-1} I\{e_{t-1} < 0\} + 0.463
```

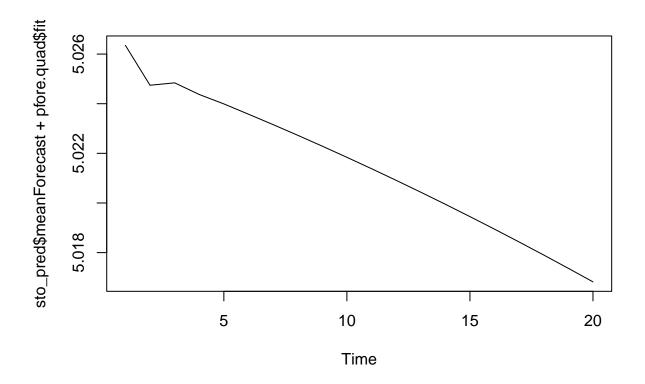
## **Forecast**

We have already forecasted the deterministic trend, so what we need to do now is to forecast the stochastic trend and combine them.

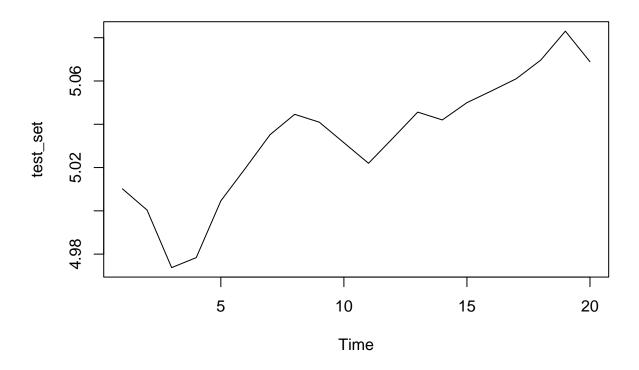
# Forecast the ARIMA(2,1,0)+TGARCH(1,1)

```
sto_pred <- predict(aparch_11, n.ahead=20, plot=FALSE, conf=.95, nx=100)</pre>
```

ts.plot(sto\_pred\$meanForecast + pfore.quad\$fit)



ts.plot(test\_set)



Let's calculate the MAPE of the whole model(deterministic trend + stochastic trend model)

```
mape_log <- 100*(mean(abs((test_set - (sto_pred$meanForecast + pfore.quad$fit))/test_set)))
cat("The MAPE of the log transformation process is:",mape_log)</pre>
```

## The MAPE of the log transformation process is: 0.5584481

```
mape <- 100*(mean(abs((test - exp(sto_pred$meanForecast + pfore.quad$fit))/test)))
cat("The MAPE of the whole model is:",mape)</pre>
```

## The MAPE of the whole model is: 2.789165