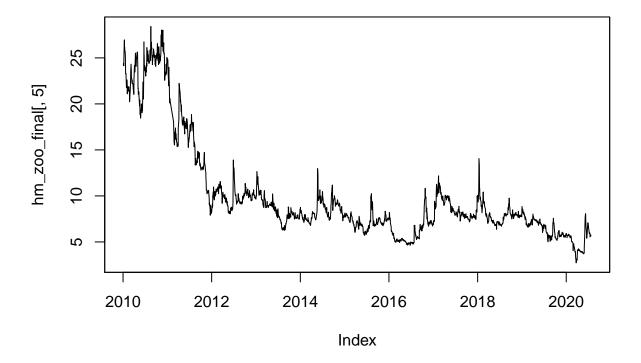
Untitled

me

7/26/2020

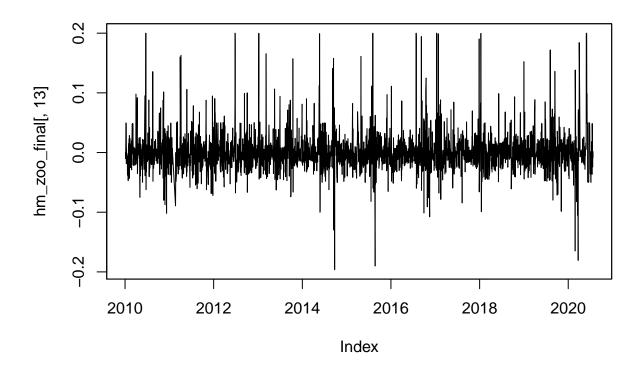
```
5+5
## [1] 10
library(zoo)
##
## Attaching package: 'zoo'
## The following objects are masked from 'package:base':
##
##
       as.Date, as.Date.numeric
library(tseries)
## Warning: package 'tseries' was built under R version 4.0.2
## Registered S3 method overwritten by 'quantmod':
    method
     as.zoo.data.frame zoo
library(FinTS)
## Warning: package 'FinTS' was built under R version 4.0.2
library(rugarch)
## Warning: package 'rugarch' was built under R version 4.0.2
## Loading required package: parallel
##
## Attaching package: 'rugarch'
## The following object is masked from 'package:stats':
##
##
       sigma
```

hm_zoo_final<-read.zoo("hindustan motors.csv",header=TRUE,sep=",",format="%d-%b-%y",FUN = as.Date)
plot(hm_zoo_final[,5])</pre>



there is a trend in the closing price hence i am taking the differnce and considering the log return.

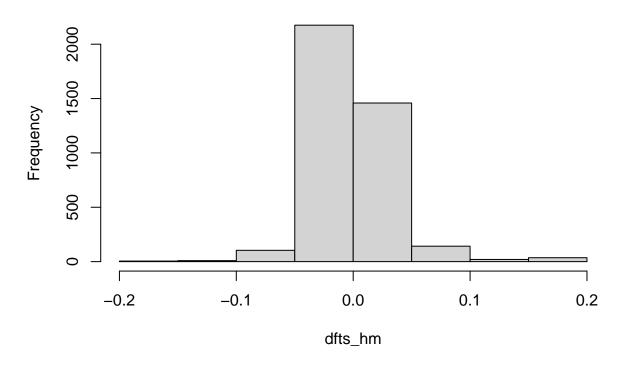
plot(hm_zoo_final[,13])



the return series seems to be constant at mean zero and volatility clustering is visible in graph

```
ret_hm<-hm_zoo_final[-1,13]
dfts_hm<- ts(ret_hm,start=c(2010,1),end=c(2020,300),frequency = 365)
hist(dfts_hm)</pre>
```

Histogram of dfts_hm



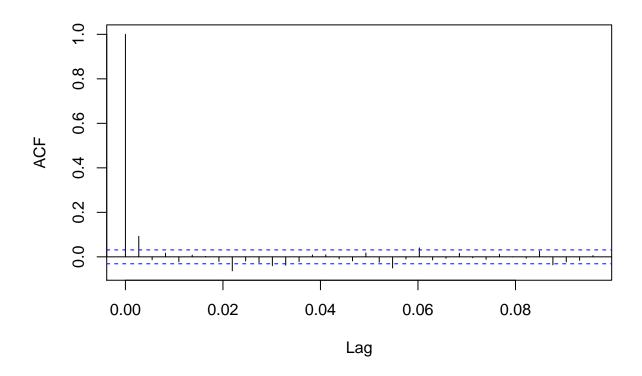
the return appears to be noormally distributed.

```
shapiro.test(dfts_hm)
##
    Shapiro-Wilk normality test
##
## data: dfts_hm
## W = 0.85846, p-value < 2.2e-16
the series is staionary.
mean(dfts_hm)
## [1] -0.0001571083
adf.test(ret_hm)
## Warning in adf.test(ret_hm): p-value smaller than printed p-value
##
    Augmented Dickey-Fuller Test
##
## data: ret_hm
## Dickey-Fuller = -15.391, Lag order = 13, p-value = 0.01
## alternative hypothesis: stationary
```

since p value is smaller than 0.05 hence we are rejecting the null hypothesis hence the series is stationary. also the series is normal around mean 0.

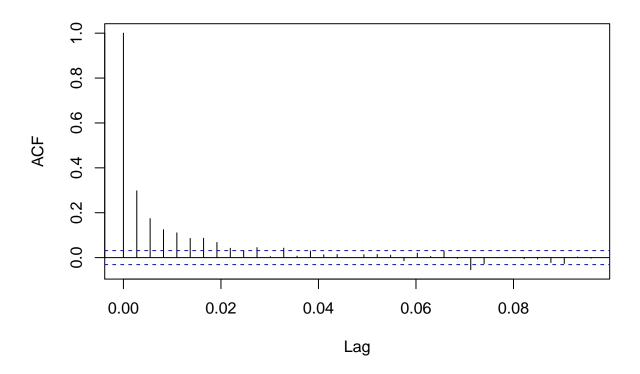
acf(dfts_hm)

Series dfts_hm



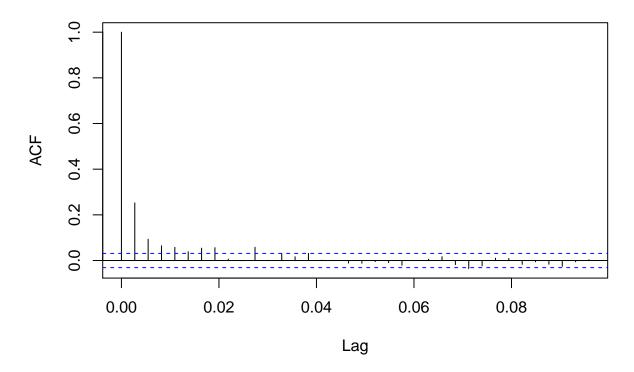
acf(abs(dfts_hm))

Series abs(dfts_hm)



acf(dfts_hm^2)

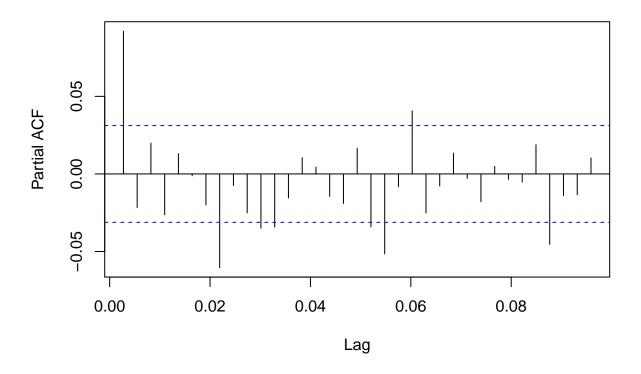
Series dfts_hm^2



by acf function of return we can say there is no auto correlation and there will be no need of MA model also by looking at the acf of absolute return we can say that the large return are followed by large returns regardless of sign

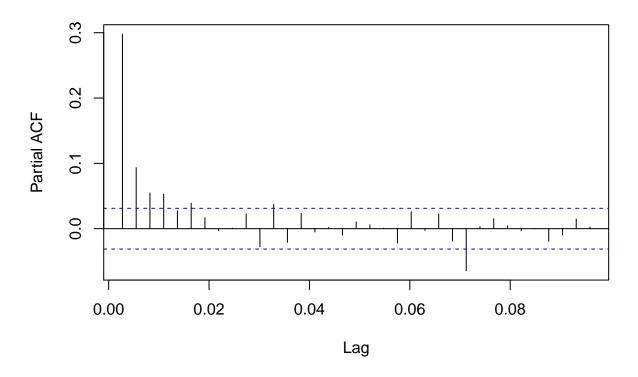
pacf(dfts_hm)

Series dfts_hm



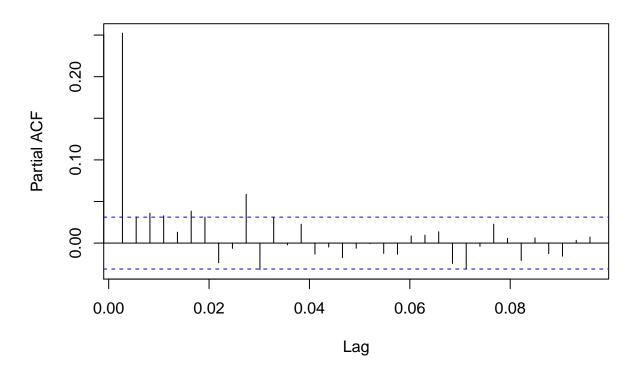
pacf(abs(dfts_hm))

Series abs(dfts_hm)



pacf(dfts_hm^2)

Series dfts_hm^2



there may need arise of an AR model

```
Box.test(ret_hm,lag=12,type="Ljung")

##

## Box-Ljung test

##

## data: ret_hm

## X-squared = 57.374, df = 12, p-value = 6.767e-08

ArchTest(ret_hm)

##

## ARCH LM-test; Null hypothesis: no ARCH effects

##

## data: ret_hm

## Chi-squared = 217.35, df = 12, p-value < 2.2e-16</pre>
```

since p value is less than 0.05 we are rejecting the null hypothesis and thus we can say there is arch effect in the series.

```
arima010=arima(ret_hm,order=c(0,1,0))
AIC(arima010)
```

```
## [1] -8327.755
```

```
arima110=arima(ret_hm,order=c(1,1,0))
AIC(arima110)
## [1] -9025.426
arima011=arima(ret_hm,order=c(0,1,1))
AIC(arima011)
## [1] -10097.98
arima111=arima(ret_hm,order=c(1,1,1))
AIC(arima111)
## [1] -10111.63
arima012=arima(ret_hm,order=c(0,1,2))
AIC(arima012)
## [1] -10111.46
arima210=arima(ret_hm,order=c(2,1,0))
AIC(arima210)
## [1] -9408.992
arima112=arima(ret_hm, order=c(1,1,2))
AIC(arima112)
## [1] -10110.4
arima211=arima(ret_hm,order=c(2,1,1))
AIC(arima211)
## [1] -10109.79
arima212=arima(ret_hm,order=c(2,1,2))
AIC(arima212)
## [1] -10111.82
arima020=arima(ret_hm,order=c(0,2,0))
AIC(arima020)
## [1] -4899.385
```

```
arima120=arima(ret_hm,order=c(1,2,0))
AIC(arima120)
## [1] -6454.954
arima021=arima(ret_hm,order=c(0,2,1))
AIC(arima021)
## [1] -8313.035
arima121=arima(ret_hm,order=c(1,2,1))
AIC(arima121)
## [1] -9009.657
arima022=arima(ret_hm,order=c(0,2,2))
AIC(arima022)
## [1] -9983.073
arima220=arima(ret_hm,order=c(2,2,0))
AIC(arima220)
## [1] -7443.981
arima122=arima(ret_hm, order=c(1,2,2))
AIC(arima122)
## [1] -10056.46
arima221=arima(ret_hm,order=c(2,2,1))
AIC(arima221)
## [1] -9392.478
arima222=arima(ret_hm,order=c(2,2,2))
AIC(arima222)
## [1] -10041.97
arima002=arima(ret_hm,order=c(0,0,2))
AIC(arima002)
## [1] -10122.1
```

```
arima001=arima(ret_hm,order=c(0,0,1))
AIC(arima001)
## [1] -10124.09
arima100=arima(ret_hm,order=c(1,0,0))
AIC(arima100)
## [1] -10124.25
arima001=arima(ret_hm,order=c(1,0,0))
AIC(arima100)
## [1] -10124.25
(2,1,2) and (1,1,1) we may ignore these since the acf and pacf function didnt recommended any also there is
no need to make the model more complex.
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(1,0)),mean.model = list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
            GARCH Model Fit
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(1,0)
## Mean Model : ARFIMA(0,0,0)
## Distribution : norm
##
## Optimal Parameters
  _____
          Estimate Std. Error t value Pr(>|t|)
##
## mu
        ## omega 0.000845 0.000030 27.8692
                                        0.0000
## alpha1 0.314338
                     0.038859
                              8.0892
                                        0.0000
##
## Robust Standard Errors:
##
          Estimate Std. Error t value Pr(>|t|)
         -0.000614
                    0.000643 -0.95542 0.339365
## mu
          0.000845
                     0.000085 9.90138 0.000000
## omega
## alpha1 0.314338
                     0.068704 4.57528 0.000005
## LogLikelihood : 5214.172
##
## Information Criteria
##
```

```
-4.0086
## Akaike
          -4.0018
-4.0086
## Bayes
## Shibata
## Hannan-Quinn -4.0061
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                      statistic p-value
## Lag[1]
                        8.621 0.003323
## Lag[2*(p+q)+(p+q)-1][2]
                        8.715 0.004181
## Lag[4*(p+q)+(p+q)-1][5] 10.018 0.009108
## d.o.f=0
## HO : No serial correlation
##
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                      statistic p-value
## Lag[1]
                       0.1205 0.7285
## Lag[2*(p+q)+(p+q)-1][2] 0.2122 0.8467
## Lag[4*(p+q)+(p+q)-1][5] 2.5254 0.5005
## d.o.f=1
##
## Weighted ARCH LM Tests
## -----
## Statistic Shape Scale P-Value
## ARCH Lag[2] 0.1831 0.500 2.000 0.6687
## ARCH Lag[4] 2.7094 1.397 1.611 0.3058
## ARCH Lag[6] 4.3851 2.222 1.500 0.2582
## Nyblom stability test
## -----
## Joint Statistic: 0.4867
## Individual Statistics:
## mu
     0.07616
## omega 0.09277
## alpha1 0.23295
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 0.846 1.01 1.35
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## -----
             t-value prob sig
##
## Sign Bias
                 2.1093 0.03501 **
## Negative Sign Bias 0.5128 0.60813
## Positive Sign Bias 0.7466 0.45540
## Joint Effect 5.0259 0.16991
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 445.8 1.235e-82
```

```
## 2
      30 460.3
                    4.040e-79
## 3
    40 488.6 4.774e-79
## 4
      50 506.8 2.396e-77
##
## Elapsed time : 0.3822222
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(0,1)),mean.model = list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
     GARCH Model Fit *
## *----*
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(0,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : norm
##
## Optimal Parameters
## -----
        Estimate Std. Error t value Pr(>|t|)
      ## omega 0.000001 0.000000 4.4583e+01
                                      0.00
## beta1 0.999000 0.000012 8.1291e+04
                                      0.00
##
## Robust Standard Errors:
##
       Estimate Std. Error t value Pr(>|t|)
        0.000017 0.000657 2.5841e-02 0.97938
## omega 0.000001 0.000000 1.0805e+01 0.00000
## beta1 0.999000 0.000127 7.8354e+03 0.00000
##
## LogLikelihood : 5058.001
##
## Information Criteria
## -----
##
## Akaike
            -3.8885
## Bayes
            -3.8817
         -3.8885
## Shibata
## Hannan-Quinn -3.8860
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                     statistic p-value
## Lag[1]
                      25.29 4.941e-07
## Lag[2*(p+q)+(p+q)-1][2] 25.34 1.766e-07
## Lag[4*(p+q)+(p+q)-1][5] 26.80 2.505e-07
## d.o.f=0
## HO : No serial correlation
##
```

```
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                         statistic p-value
## Lag[1]
                           181.9
## Lag[2*(p+q)+(p+q)-1][2] 191.2
## Lag[4*(p+q)+(p+q)-1][5] 205.9
                                        0
## d.o.f=1
##
## Weighted ARCH LM Tests
## -----
             Statistic Shape Scale P-Value
## ARCH Lag[2] 18.57 0.500 2.000 1.641e-05
## ARCH Lag[4] 29.41 1.397 1.611 4.307e-08
## ARCH Lag[6] 34.41 2.222 1.500 4.748e-09
## Nyblom stability test
## -----
## Joint Statistic: 6.1311
## Individual Statistics:
## mu 0.06183
## omega 0.75330
## beta1 0.17709
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 0.846 1.01 1.35
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
                    t-value prob sig
## Sign Bias
                     0.2677 7.890e-01
## Negative Sign Bias 4.9577 7.594e-07 ***
## Positive Sign Bias 11.4503 1.192e-29 ***
## Joint Effect 171.4625 6.151e-37 ***
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
   group statistic p-value(g-1)
## 1 20 586.7 3.190e-112
## 2 30 639.7 3.575e-116
## 3 40 657.5 2.302e-113
## 4 50 679.6 7.012e-112
##
## Elapsed time : 0.4278071
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(1,1)),mean.model = list(armaOrder=c(0,0)</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
```

*

GARCH Model Fit

```
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(1,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : norm
## Optimal Parameters
         Estimate Std. Error t value Pr(>|t|)
## mu -0.000552 0.000603 -0.91485 0.36027
## omega 0.000225 0.000030 7.44504 0.00000
## alpha1 0.159197 0.022238 7.15875 0.00000
## beta1 0.657757 0.038145 17.24349 0.00000
##
## Robust Standard Errors:
        Estimate Std. Error t value Pr(>|t|)
       ## omega 0.000225 0.000080 2.81468 0.004883
## alpha1 0.159197 0.040695 3.91197 0.000092
## beta1 0.657757 0.092723 7.09380 0.000000
##
## LogLikelihood : 5242.235
##
## Information Criteria
## Akaike
            -4.0294
## Bayes
            -4.0204
## Shibata -4.0294
## Hannan-Quinn -4.0261
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                      statistic p-value
## Lag[1]
                        8.024 0.004617
## Lag[2*(p+q)+(p+q)-1][2] 8.037 0.006326
## Lag[4*(p+q)+(p+q)-1][5] 8.839 0.018171
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
                       statistic p-value
                          1.296 0.2549
## Lag[1]
## Lag[2*(p+q)+(p+q)-1][5] 2.259 0.5586
## Lag[4*(p+q)+(p+q)-1][9] 2.648 0.8154
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
## Statistic Shape Scale P-Value
## ARCH Lag[3] 0.3664 0.500 2.000 0.5450
```

```
## ARCH Lag[5]
               0.5075 1.440 1.667 0.8814
## ARCH Lag[7] 0.5443 2.315 1.543 0.9742
## Nyblom stability test
## -----
## Joint Statistic: 0.3636
## Individual Statistics:
## mu
        0.01862
## omega 0.04597
## alpha1 0.12434
## beta1 0.08184
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.07 1.24 1.6
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
## -----
                  t-value prob sig
##
## Sign Bias
                   2.0499 0.04047 **
## Negative Sign Bias 0.6458 0.51846
## Positive Sign Bias 0.1415 0.88746
## Joint Effect 6.1153 0.10613
##
##
## Adjusted Pearson Goodness-of-Fit Test:
   group statistic p-value(g-1)
##
## 1 20 422.6 8.701e-78
## 2 30 460.6 3.359e-79
    40 485.5 1.924e-78
50 497.2 1.842e-75
## 3
## 4
##
##
## Elapsed time : 0.5473549
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(2,0)),mean.model = list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
    GARCH Model Fit *
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(2,0)
## Mean Model : ARFIMA(0,0,0)
## Distribution : norm
##
## Optimal Parameters
        Estimate Std. Error t value Pr(>|t|)
##
```

```
## mu 2.1e-05 6.9e-05 0.31123 0.755626
## omega 1.0e-06 0.0e+00 4.66329 0.000003
## alpha1 2.5e-02 2.7e-05 930.66806 0.000000
## alpha2 2.5e-02 1.2e-05 2133.70221 0.000000
## Robust Standard Errors:
## Estimate Std. Error t value Pr(>|t|)
## mu 2.1e-05 0.016853 0.001273 0.998984
## omega 1.0e-06 0.000072 0.016574 0.986777
## alpha1 2.5e-02 0.007329 3.411008 0.000647
## alpha2 2.5e-02 0.004466 5.598294 0.000000
## LogLikelihood: -53282.89
##
## Information Criteria
## -----
##
## Akaike
             40.990
## Bayes
              40.999
## Shibata 40.990
## Hannan-Quinn 40.993
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                        statistic p-value
## Lag[1]
                          2.555 0.1099
## Lag[2*(p+q)+(p+q)-1][2] 2.624 0.1775
## Lag[4*(p+q)+(p+q)-1][5] 3.292 0.3563
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                         statistic p-value
## Lag[1]
                          1.140 0.2856
## Lag[2*(p+q)+(p+q)-1][5] 2.534 0.4987
## Lag[4*(p+q)+(p+q)-1][9] 2.829 0.7868
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
              Statistic Shape Scale P-Value
## ARCH Lag[3] 0.3599 0.500 2.000 0.5485
## ARCH Lag[5] 0.4041 1.440 1.667 0.9115
## ARCH Lag[7] 0.4616 2.315 1.543 0.9817
##
## Nyblom stability test
## -----
## Joint Statistic: 123.5262
## Individual Statistics:
## mu 0.1919
## omega 53.7989
## alpha1 76.3994
## alpha2 80.6063
```

```
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.07 1.24 1.6
## Individual Statistic:
                     0.35 0.47 0.75
## Sign Bias Test
## -----
                 t-value prob sig
             1.705 8.836e-02
## Sign Bias
## Negative Sign Bias 2.695 7.079e-03 ***
## Positive Sign Bias 3.561 3.763e-04 ***
## Joint Effect 23.583 3.052e-05 ***
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
  group statistic p-value(g-1)
## 1 20 11685
## 2
      30 17157
                        0
    40 22056
## 3
                         0
## 4 50 27016
                         Ω
##
##
## Elapsed time : 0.08400297
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(0,2)),mean.model = list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
         GARCH Model Fit
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(0,2)
## Mean Model : ARFIMA(0,0,0)
## Distribution : norm
##
## Optimal Parameters
## -----
       Estimate Std. Error t value Pr(>|t|)
## mu
      ## omega 0.000001 0.000000 7.6185e+01 0.00000
## beta1 0.000404 0.000060 6.7675e+00 0.00000
```

beta2 0.998596 0.000005 1.9383e+05 0.00000

omega 0.000001 0.000000 2.0857e+01 0.000000 ## beta1 0.000404 0.000179 2.2602e+00 0.023812 ## beta2 0.998596 0.000055 1.8307e+04 0.000000

Estimate Std. Error t value Pr(>|t|)

##

##

mu

Robust Standard Errors:

```
##
## LogLikelihood: 5058.314
## Information Criteria
## -----
##
## Akaike -3.8879
## Bayes -3.8789
## Shibata -3.8879
## Hannan-Quinn -3.8847
## Weighted Ljung-Box Test on Standardized Residuals
## -----
         statistic p-value
##
## Lag[1]
                            25.18 5.235e-07
## Lag[2*(p+q)+(p+q)-1][2] 25.23 1.884e-07
## Lag[4*(p+q)+(p+q)-1][5] 26.68 2.708e-07
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
           statistic p-value
## Lag[1] 181.2 0
## Lag[2*(p+q)+(p+q)-1][5] 205.1 0
## Lag[4*(p+q)+(p+q)-1][9] 217.0 0
## d.o.f=2
## Weighted ARCH LM Tests
   Statistic Shape Scale P-Value
## ARCH Lag[3] 10.19 0.500 2.000 1.415e-03
## ARCH Lag[5] 16.54 1.440 1.667 1.580e-04
## ARCH Lag[7] 24.75 2.315 1.543 3.819e-06
## Nyblom stability test
## -----
## Joint Statistic: 135.2244
## Individual Statistics:
## mu 0.06419
## omega 3.17938
## beta1 0.18254
## beta2 0.18254
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.07 1.24 1.6
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
## -----
                    t-value prob sig
##
## Sign Bias 0.2662 7.901e-01
## Negative Sign Bias 4.9517 7.831e-07 ***
## Positive Sign Bias 11.4431 1.290e-29 ***
```

```
## Joint Effect 171.1982 7.014e-37 ***
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 592.4 2.025e-113
       30 639.3 4.460e-116
## 2
## 3
    40 665.0 6.745e-115
## 4 50 678.5 1.178e-111
##
##
## Elapsed time : 0.4849632
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(2,2)),mean.model = list(armaOrder=c(0,0)</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my model hm
## *----*
      GARCH Model Fit *
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(2,2)
## Mean Model : ARFIMA(0,0,0)
## Distribution : norm
##
## Optimal Parameters
         Estimate Std. Error t value Pr(>|t|)
## mu
       -0.000484 0.000602 -0.805285 0.420655
## omega 0.000290 0.000061 4.791537 0.000002
## alpha1 0.202246 0.029480 6.860471 0.000000 ## alpha2 0.000000 0.047878 0.000002 0.999999
## beta1 0.212046 0.142375 1.489344 0.136397
## beta2  0.345850  0.077568  4.458657  0.000008
## Robust Standard Errors:
        Estimate Std. Error t value Pr(>|t|)
## mu -0.000484 0.000582 -0.833061 0.404810
## omega 0.000290 0.000132 2.203845 0.027535
## alpha1 0.202246 0.049369 4.096613 0.000042
## alpha2 0.000000 0.088477 0.000001 0.999999
## beta1 0.212046 0.249739 0.849071 0.395842
## beta2  0.345850  0.109771  3.150636  0.001629
##
## LogLikelihood : 5249.53
## Information Criteria
## -----
```

Akaike -4.0335

```
## Bayes -4.0200
## Shibata -4.0335
## Hannan-Quinn -4.0286
## Weighted Ljung-Box Test on Standardized Residuals
## -----
         statistic p-value
##
                        7.463 0.006297
## Lag[1]
## Lag[2*(p+q)+(p+q)-1][2] 7.464 0.008977
## Lag[4*(p+q)+(p+q)-1][5] 8.318 0.024555
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                       statistic p-value
## Lag[1]
                        0.3072 0.5794
## Lag[2*(p+q)+(p+q)-1][11] 2.4052 0.9300
## Lag[4*(p+q)+(p+q)-1][19] 6.4889 0.8322
## d.o.f=4
##
## Weighted ARCH LM Tests
## -----
    Statistic Shape Scale P-Value
## ARCH Lag[5] 0.1947 0.500 2.000 0.6590
## ARCH Lag[7] 0.2191 1.473 1.746 0.9665
## ARCH Lag[9] 0.8876 2.402 1.619 0.9460
## Nyblom stability test
## -----
## Joint Statistic: 1.5067
## Individual Statistics:
## mu
     0.01951
## omega 0.04316
## alpha1 0.13032
## alpha2 0.13782
## beta1 0.05696
## beta2 0.08128
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.49 1.68 2.12
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
                  t-value prob sig
##
## Sign Bias
                  2.0756 0.03803 **
## Negative Sign Bias 0.5143 0.60712
## Positive Sign Bias 0.2657 0.79051
## Joint Effect 5.5250 0.13715
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
```

```
group statistic p-value(g-1)
           422.6
## 1
      20
                   8.765e-78
           457.4
## 2
      30
                   1.520e-78
## 3
      40 478.8 4.321e-77
## 4
      50
            498.5
                 1.055e-75
##
## Elapsed time : 1.055024
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(1,2)),mean.model = list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
## *----*
         GARCH Model Fit
## *----*
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(1,2)
## Mean Model : ARFIMA(0,0,0)
## Distribution : norm
## Optimal Parameters
## -----
        Estimate Std. Error t value Pr(>|t|)
## mu
                0.000601 -0.80606 0.420211
        -0.000484
## omega 0.000290 0.000037 7.79368 0.000000
## alpha1 0.202245 0.027417 7.37666 0.000000
## beta1 0.212046 0.066862 3.17138 0.001517
                 0.057166 6.04995 0.000000
## beta2 0.345850
##
## Robust Standard Errors:
       Estimate Std. Error t value Pr(>|t|)
##
## mu
      ## omega 0.000290 0.000089 3.25101 0.001150
## alpha1 0.202245 0.048266 4.19022 0.000028
## beta1 0.212046 0.102512 2.06850 0.038593
## beta2  0.345850  0.089089  3.88209  0.000104
##
## LogLikelihood : 5249.53
##
## Information Criteria
## -----
##
## Akaike
           -4.0343
            -4.0230
## Bayes
## Shibata -4.0343
## Hannan-Quinn -4.0302
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                      statistic p-value
```

```
## Lag[1]
                          7.463 0.006296
## Lag[2*(p+q)+(p+q)-1][2] 7.464 0.008977
## Lag[4*(p+q)+(p+q)-1][5] 8.318 0.024554
## d.o.f=0
## HO : No serial correlation
##
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                       statistic p-value
## Lag[1] 0.3073 0.5794
## Lag[2*(p+q)+(p+q)-1][8] 0.9251 0.9789
## Lag [4*(p+q)+(p+q)-1] [14] 4.0162 0.8787
## d.o.f=3
##
## Weighted ARCH LM Tests
## -----
##
            Statistic Shape Scale P-Value
## ARCH Lag[4] 0.003164 0.500 2.000 0.9551
## ARCH Lag[6] 0.184123 1.461 1.711 0.9721
## ARCH Lag[8] 0.384186 2.368 1.583 0.9898
##
## Nyblom stability test
## -----
## Joint Statistic: 0.665
## Individual Statistics:
## mu 0.01951
## omega 0.04316
## alpha1 0.13032
## beta1 0.05696
## beta2 0.08128
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.28 1.47 1.88
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## -----
##
                  t-value prob sig
## Sign Bias 2.0756 0.03803 **
## Negative Sign Bias 0.5143 0.60711
## Positive Sign Bias 0.2657 0.79052
## Joint Effect 5.5250 0.13715
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 422.6 8.765e-78
## 2 30 457.4 1.520e-78
## 3 40 478.8 4.321e-77
## 4 50 498.5 1.055e-75
##
##
## Elapsed time : 0.563163
```

```
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(2,1)),mean.model =list(armaOrder=c(0,0
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)
my_model_hm</pre>
```

```
##
      GARCH Model Fit *
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(2,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : norm
##
## Optimal Parameters
## -----
         Estimate Std. Error t value Pr(>|t|)
## mu -0.000553 0.000607 -0.91202 0.361758
## omega 0.000224 0.000049 4.61934 0.000004
## alpha1 0.159147 0.023637 6.73302 0.000000
## alpha2 0.000000 0.042884 0.00000 1.000000
## beta1 0.657837 0.070170 9.37494 0.000000
##
## Robust Standard Errors:
        Estimate Std. Error t value Pr(>|t|)
## mu -0.000553 0.000624 -0.88601 0.375614
## omega 0.000224 0.000129 1.74167 0.081566
## alpha1 0.159147 0.037289 4.26798 0.000020 ## alpha2 0.000000 0.092549 0.00000 1.000000
## beta1 0.657837 0.186151 3.53389 0.000409
## LogLikelihood : 5242.176
##
## Information Criteria
##
## Akaike
          -4.0286
## Bayes
            -4.0173
## Shibata
            -4.0286
## Hannan-Quinn -4.0245
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                         8.005 0.004666
## Lag[2*(p+q)+(p+q)-1][2] 8.018 0.006398
## Lag[4*(p+q)+(p+q)-1][5] 8.822 0.018352
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
```

```
##
## Weighted ARCH LM Tests
## -----
   Statistic Shape Scale P-Value
## ARCH Lag[4] 0.06946 0.500 2.000 0.7921
## ARCH Lag[6] 0.19900 1.461 1.711 0.9689
## ARCH Lag[8] 0.42808 2.368 1.583 0.9870
## Nyblom stability test
## -----
## Joint Statistic: 1.9671
## Individual Statistics:
## mu
       0.01857
## omega 0.04597
## alpha1 0.12457
## alpha2 0.41416
## beta1 0.08186
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.28 1.47 1.88
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
                   t-value prob sig
## Sign Bias
                   2.0482 0.04064 **
## Negative Sign Bias 0.6462 0.51819
## Positive Sign Bias 0.1426 0.88665
## Joint Effect 6.1066 0.10654
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## group statistic p-value(g-1)
## 1 20 422.6 8.701e-78
## 2 30 460.6 3.359e-79
## 3 40 485.1 2.314e-78
## 4
     50 497.2 1.874e-75
##
## Elapsed time : 0.449388
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(0,1)),mean.model = list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
## *
          GARCH Model Fit
```

statistic p-value 1.298 0.2546

2.499 0.7761

##

Lag[1]

d.o.f=3

Lag[2*(p+q)+(p+q)-1][8]

Lag[4*(p+q)+(p+q)-1][14] 6.104 0.6248

```
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(0,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
##
## Optimal Parameters
        Estimate Std. Error t value Pr(>|t|)
## mu -0.000456 0.000646 -0.70624 0.48004
## omega 0.000027 0.000005 5.86996 0.00000
## beta1 0.984486 0.000343 2869.34191 0.00000
## skew 1.189873 0.030812 38.61744 0.00000
## shape 2.511569 0.125716 19.97815 0.00000
##
## Robust Standard Errors:
##
       Estimate Std. Error t value Pr(>|t|)
## mu
       -0.000456 0.000635 -0.71866 0.47235
## omega 0.000027 0.000005 5.05107 0.00000
## beta1 0.984486 0.000119 8298.71590 0.00000
## skew 1.189873 0.028916 41.14864 0.00000
## shape 2.511569 0.131867 19.04617 0.00000
##
## LogLikelihood : 5594.417
##
## Information Criteria
## Akaike
            -4.2996
           -4.2883
-4.2996
## Bayes
## Shibata
## Hannan-Quinn -4.2955
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                          25.51 4.393e-07
## Lag[2*(p+q)+(p+q)-1][2]
                       25.57 1.542e-07
## Lag[4*(p+q)+(p+q)-1][5]
                         27.07 2.102e-07
## d.o.f=0
## HO : No serial correlation
##
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                       statistic p-value
## Lag[1]
                         181.6
                       190.8
## Lag[2*(p+q)+(p+q)-1][2]
                                     0
## Lag[4*(p+q)+(p+q)-1][5]
                                     0
## d.o.f=1
##
## Weighted ARCH LM Tests
## -----
```

```
Statistic Shape Scale P-Value
## ARCH Lag[2] 18.28 0.500 2.000 1.910e-05
              28.92 1.397 1.611 5.800e-08
## ARCH Lag[4]
## ARCH Lag[6] 33.89 2.222 1.500 6.598e-09
## Nyblom stability test
## -----
## Joint Statistic: 0.5297
## Individual Statistics:
## mu
      0.1209
## omega 0.1891
## beta1 0.1917
## skew 0.1009
## shape 0.1836
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.28 1.47 1.88
## Individual Statistic:
                      0.35 0.47 0.75
## Sign Bias Test
## -----
                  t-value prob sig
                  0.2807 7.790e-01
## Sign Bias
## Negative Sign Bias 4.8843 1.101e-06 ***
## Positive Sign Bias 11.6008 2.261e-30 ***
## Joint Effect 170.8591 8.302e-37 ***
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
   group statistic p-value(g-1)
## 1
      20
           8.646 9.789e-01
      30 72.585 1.332e-05
## 2
    40 45.231 2.279e-01
## 3
      50 66.346
                  4.994e-02
## 4
##
##
## Elapsed time : 1.407672
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(1,0)),mean.model = list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
## *----*
          GARCH Model Fit
## *----*
##
## Conditional Variance Dynamics
```

GARCH Model : sGARCH(1,0)
Mean Model : ARFIMA(0,0,0)

Distribution : sstd

##

```
## Optimal Parameters
## -----
        Estimate Std. Error t value Pr(>|t|)
## mu
       ## omega 0.000851 0.000114 7.45631 0.00000
## alpha1 0.690368 0.121578 5.67841 0.00000
## skew 1.252926 0.034898 35.90250 0.00000
## shape 2.831086 0.178200 15.88715 0.00000
##
## Robust Standard Errors:
        Estimate Std. Error t value Pr(>|t|)
        -0.000239 0.000592 -0.40448 0.68586
## mu
## omega 0.000851 0.000124 6.87921 0.00000
## alpha1 0.690368 0.125707 5.49188 0.00000
## skew
        1.252926 0.039121 32.02688 0.00000
        ## shape
##
## LogLikelihood: 5716.244
##
## Information Criteria
## -----
           -4.3933
## Akaike
## Bayes -4.3820
## Shibata -4.3933
## Hannan-Quinn -4.3892
## Weighted Ljung-Box Test on Standardized Residuals
##
                     statistic p-value
## Lag[1]
                        6.501 0.01078
## Lag[2*(p+q)+(p+q)-1][2] 6.531 0.01588
## Lag[4*(p+q)+(p+q)-1][5] 7.969 0.02999
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                     statistic p-value
## Lag[1]
                       1.789 0.1811
## Lag[2*(p+q)+(p+q)-1][2] 1.800 0.2982
## Lag[4*(p+q)+(p+q)-1][5] 3.070 0.3945
## d.o.f=1
##
## Weighted ARCH LM Tests
## -----
            Statistic Shape Scale P-Value
## ARCH Lag[2] 0.02226 0.500 2.000 0.8814
## ARCH Lag[4] 1.44103 1.397 1.611 0.5779
## ARCH Lag[6] 2.60748 2.222 1.500 0.5486
##
## Nyblom stability test
## -----
## Joint Statistic: 1.0512
```

```
## Individual Statistics:
## mu
     0.07334
## omega 0.19854
## alpha1 0.33958
## skew 0.15166
## shape 0.10358
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.28 1.47 1.88
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## -----
                 t-value prob sig
##
                 2.0596 0.03954 **
## Sign Bias
## Negative Sign Bias 0.6744 0.50009
## Positive Sign Bias 2.1711 0.03001 **
## Joint Effect 9.2816 0.02577 **
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 19.51 0.4247298
## 2
      30 59.43 0.0007267
## 3 40 42.00 0.3422059
## 4 50 53.69 0.2992545
##
##
## Elapsed time : 1.450274
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(1,1)),mean.model = list(armaOrder=c(0,0)</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
## *----*
         GARCH Model Fit *
## *----*
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(1,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
##
## Optimal Parameters
## -----
##
        Estimate Std. Error t value Pr(>|t|)
## mu 0.000020 0.000540 0.037577 0.970025
## omega 0.000226 0.000049 4.597995 0.000004
## alpha1 0.357891 0.071134 5.031214 0.000000
## beta1 0.582941 0.056504 10.316793 0.000000
## skew 1.298620 0.036983 35.113558 0.000000
```

```
## shape 2.983342 0.192455 15.501508 0.000000
##
## Robust Standard Errors:
         Estimate Std. Error t value Pr(>|t|)
         0.000020 0.000548 0.037023 0.970466
## mu
## omega 0.000226 0.000067 3.361324 0.000776
## alpha1 0.357891 0.080518 4.444866 0.000009
## beta1 0.582941 0.083813 6.955246 0.000000 ## skew 1.298620 0.043492 29.858543 0.000000
## shape 2.983342 0.194355 15.349939 0.000000
## LogLikelihood: 5754.14
## Information Criteria
## -----
##
             -4.4216
## Akaike
## Bayes
             -4.4081
## Shibata
             -4.4217
## Hannan-Quinn -4.4167
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
                         statistic p-value
## Lag[1]
                           6.300 0.01208
## Lag[2*(p+q)+(p+q)-1][2] 6.399 0.01722
## Lag[4*(p+q)+(p+q)-1][5] 7.336 0.04296
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                         statistic p-value
                           0.111 0.7390
## Lag[1]
## Lag[2*(p+q)+(p+q)-1][5] 2.205 0.5709
## Lag[4*(p+q)+(p+q)-1][9] 2.926 0.7712
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
             Statistic Shape Scale P-Value
## ARCH Lag[3] 0.9329 0.500 2.000 0.3341
## ARCH Lag[5] 1.4712 1.440 1.667 0.5999
## ARCH Lag[7] 1.5614 2.315 1.543 0.8090
## Nyblom stability test
## -----
## Joint Statistic: 0.6731
## Individual Statistics:
## mu
       0.05038
## omega 0.14038
## alpha1 0.08650
## beta1 0.07352
## skew 0.16519
```

```
## shape 0.08780
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.49 1.68 2.12
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
## -----
##
                  t-value prob sig
## Sign Bias
                   2.5001 0.01248 **
## Negative Sign Bias 0.4062 0.68461
## Positive Sign Bias 1.8615 0.06279
## Joint Effect 9.8380 0.01999 **
##
##
## Adjusted Pearson Goodness-of-Fit Test:
   group statistic p-value(g-1)
## 1 20 24.86 0.1651505
          63.79
## 2 30
                     0.0002037
                   0.1979377
## 3 40 46.25
## 4 50 60.77 0.1207402
##
## Elapsed time : 1.618748
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(2,0)),mean.model = list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
       GARCH Model Fit
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(2,0)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
##
         Estimate Std. Error t value Pr(>|t|)
       -0.000115 0.000560 -0.20589 0.836876
## omega 0.000742 0.000100 7.43893 0.000000
## alpha1 0.560907 0.104208 5.38260 0.000000
## alpha2 0.218335 0.060340 3.61842 0.000296
## skew 1.270936 0.036083 35.22265 0.000000
## shape 2.860891 0.180668 15.83509 0.000000
##
## Robust Standard Errors:
        Estimate Std. Error t value Pr(>|t|)
## mu
       -0.000115 0.000590 -0.19522 0.845224
```

```
## omega 0.000742 0.000114 6.50996 0.000000 ## alpha1 0.560907 0.100828 5.56302 0.000000
## alpha2 0.218335 0.053950 4.04702 0.000052
          1.270936 0.043216 29.40865 0.000000
## skew
## shape 2.860891 0.190669 15.00450 0.000000
##
## LogLikelihood: 5735.569
##
## Information Criteria
            -4.4074
-4.3938
## Akaike
## Bayes
## Shibata -4.4074
## Hannan-Quinn -4.4025
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                        statistic p-value
## Lag[1]
                           6.377 0.01156
## Lag[2*(p+q)+(p+q)-1][2] 6.446 0.01673
## Lag[4*(p+q)+(p+q)-1][5] 7.614 0.03671
## d.o.f=0
## HO : No serial correlation
##
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
                      statistic p-value
##
## Lag[1]
                          0.9859 0.3207
## Lag[2*(p+q)+(p+q)-1][5] 3.1535 0.3797
## Lag[4*(p+q)+(p+q)-1][9] 4.9587 0.4355
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
     Statistic Shape Scale P-Value
## ARCH Lag[3] 0.02221 0.500 2.000 0.8815
## ARCH Lag[5] 1.04637 1.440 1.667 0.7192
## ARCH Lag[7] 2.63340 2.315 1.543 0.5853
##
## Nyblom stability test
## -----
## Joint Statistic: 0.8797
## Individual Statistics:
## mu
        0.04586
## omega 0.16980
## alpha1 0.19041
## alpha2 0.09935
## skew 0.18499
## shape 0.09122
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.49 1.68 2.12
## Individual Statistic: 0.35 0.47 0.75
```

```
##
## Sign Bias Test
## -----
                 t-value prob sig
##
## Sign Bias
                 1.9766 0.04819 **
## Negative Sign Bias 0.7867 0.43155
## Positive Sign Bias 2.0537 0.04011 **
## Joint Effect
              9.1317 0.02759 **
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 21.23 0.324191
## 2
      30 58.05
                 0.001074
    40 48.37
                 0.144504
## 3
## 4
    50 55.42
                 0.245358
##
##
## Elapsed time : 1.710994
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(0,2)),mean.model = list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
      GARCH Model Fit *
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(0,2)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
## -----
       Estimate Std. Error t value Pr(>|t|)
     -0.000452 0.000648 -0.69759 0.485433
## mu
## omega 0.000026 0.000005
                           5.00256 0.000001
## beta1 0.985380 0.000059 16606.95416 0.000000
## shape 2.512737 0.125611 20.00413 0.000000
## Robust Standard Errors:
       Estimate Std. Error
                            t value Pr(>|t|)
     -0.000452 0.000642 -7.0327e-01 0.48189
## omega 0.000026 0.000004 5.9465e+00 0.00000
## beta1 0.985380 0.000038 2.5657e+04 0.00000
## beta2 0.000000 0.000996 1.0000e-06 1.00000
## skew 1.190096 0.029176 4.0790e+01 0.00000
## shape 2.512737 0.133282 1.8853e+01 0.00000
```

```
##
## LogLikelihood: 5594.406
## Information Criteria
## -----
##
## Akaike -4.2988
## Bayes -4.2852
## Shibata -4.2988
## Hannan-Quinn -4.2939
## Weighted Ljung-Box Test on Standardized Residuals
## -----
          statistic p-value
##
## Lag[1]
                            25.50 4.416e-07
## Lag[2*(p+q)+(p+q)-1][2] 25.56 1.550e-07
## Lag[4*(p+q)+(p+q)-1][5] 27.07 2.109e-07
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
           statistic p-value
## Lag[1] 181.4

## Lag[2*(p+q)+(p+q)-1][5] 205.0

## Lag[4*(p+q)+(p+q)-1][9] 216.7
## d.o.f=2
## Weighted ARCH LM Tests
   Statistic Shape Scale P-Value
## ARCH Lag[3] 10.14 0.500 2.000 1.453e-03
## ARCH Lag[5] 16.32 1.440 1.667 1.792e-04
## ARCH Lag[7] 24.46 2.315 1.543 4.539e-06
## Nyblom stability test
## -----
## Joint Statistic: 0.4894
## Individual Statistics:
## mu
      0.1242
## omega 0.1925
## beta1 0.1957
## beta2 0.1958
## skew 0.1006
## shape 0.1866
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.49 1.68 2.12
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## -----
                    t-value prob sig
                     0.2797 7.797e-01
## Sign Bias
```

```
## Negative Sign Bias 4.8817 1.116e-06 ***
## Positive Sign Bias 11.5980 2.333e-30 ***
## Joint Effect 170.7463 8.781e-37 ***
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
    group statistic p-value(g-1)
## 1
      20
           8.323
                   9.831e-01
      30
## 2
           74.846
                   6.405e-06
## 3
      40 44.985
                   2.356e-01
## 4
      50 64.731
                   6.541e-02
##
## Elapsed time : 1.533
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(2,2)),mean.model = list(armaOrder=c(0,0)</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
## *----*
         GARCH Model Fit
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(2,2)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
## ------
         Estimate Std. Error t value Pr(>|t|)
         0.000075 0.000541 0.138332 0.889978
## mu
         ## omega
## alpha1 0.394337 0.073812 5.342425 0.000000
## alpha2 0.000000 0.239444 0.000002 0.999999
## beta1 0.287611 0.510094 0.563840 0.572863
## beta2 0.255109 0.239515 1.065106 0.286828
## skew 1.305804 0.037387 34.926337 0.000000
## shape 2.993976 0.183234 16.339636 0.000000
##
## Robust Standard Errors:
##
         Estimate Std. Error t value Pr(>|t|)
         0.000075 0.000548 0.136469 0.89145
## mu
## omega
         0.000239
                 0.000307 0.777875 0.43664
## alpha1 0.394337 0.069533 5.671253 0.00000
## alpha2 0.000000 0.523479 0.000001 1.00000
## beta1
         0.287611 1.135802 0.253223 0.80010
        ## beta2
## skew 1.305804 0.045896 28.451198 0.00000
## shape 2.993976 0.238480 12.554424 0.00000
##
```

```
## LogLikelihood : 5757.812
##
## Information Criteria
## -----
## Akaike
             -4.4229
## Bayes
            -4.4049
## Shibata -4.4230
## Hannan-Quinn -4.4164
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                      statistic p-value
## Lag[1]
                         5.799 0.01604
## Lag[2*(p+q)+(p+q)-1][2] 5.865 0.02390
## Lag[4*(p+q)+(p+q)-1][5] 6.722 0.06054
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                        statistic p-value
                           0.231 0.6308
## Lag[1]
## Lag[2*(p+q)+(p+q)-1][11] 3.305 0.8308
## Lag[4*(p+q)+(p+q)-1][19] 5.588 0.9041
## d.o.f=4
##
## Weighted ARCH LM Tests
## Statistic Shape Scale P-Value
## ARCH Lag[5] 0.5054 0.500 2.000 0.4772
## ARCH Lag[7] 0.6870 1.473 1.746 0.8456
## ARCH Lag[9] 1.4805 2.402 1.619 0.8555
##
## Nyblom stability test
## Joint Statistic: 1.5524
## Individual Statistics:
## mu 0.05152
## omega 0.14094
## alpha1 0.07646
## alpha2 0.12909
## beta1 0.07355
## beta2 0.07702
## skew 0.15394
## shape 0.08681
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.89 2.11 2.59
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
## -----
##
                   t-value prob sig
```

```
## Sign Bias
                    2.5278 0.01154 **
## Negative Sign Bias 0.4456 0.65596
## Positive Sign Bias 1.9873 0.04699
## Joint Effect
                   10.3140 0.01608 **
##
## Adjusted Pearson Goodness-of-Fit Test:
    group statistic p-value(g-1)
## 1
      20 17.22
                    0.5752786
## 2
      30
             60.08
                     0.0006042
## 3
      40
            49.05
                     0.1300010
## 4
      50
             61.96
                     0.1011570
##
##
## Elapsed time : 2.716993
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(1,2)),mean.model =list(armaOrder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
## *----*
           GARCH Model Fit
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(1,2)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
##
         Estimate Std. Error t value Pr(>|t|)
## mu
         0.000075 0.000541 0.13812 0.890149
## omega 0.000239 0.000049 4.84768 0.000001
## alpha1 0.394397 0.074605 5.28649 0.000000
## beta1
         ## beta2 0.255060 0.081022 3.14802 0.001644
## skew 1.305789 0.037766 34.57558 0.000000
## shape 2.993875 0.194228 15.41426 0.000000
##
## Robust Standard Errors:
##
         Estimate Std. Error t value Pr(>|t|)
## mu
                  0.000549 0.13618 0.891678
         0.000075
## omega
         0.000239
                    0.000063 3.78107 0.000156
## alpha1 0.394397
                 0.078003 5.05619 0.000000
## beta1
         0.287609 0.067919 4.23455 0.000023
## beta2
         0.044963 29.04169 0.000000
## skew
         1.305789
## shape 2.993875 0.193670 15.45863 0.000000
## LogLikelihood : 5757.812
```

```
##
## Information Criteria
## -----
##
## Akaike -4.4237
## Bayes -4.4079
## Shibata -4.4237
## Hannan-Quinn -4.4180
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                          5.799 0.01604
## Lag[2*(p+q)+(p+q)-1][2] 5.865 0.02390
## Lag[4*(p+q)+(p+q)-1][5] 6.722 0.06053
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
                     statistic p-value
##
## Lag[1]
                         0.2311 0.6307
## Lag[2*(p+q)+(p+q)-1][8] 2.3471 0.8024
## Lag[4*(p+q)+(p+q)-1][14] 4.1872 0.8618
## d.o.f=3
## Weighted ARCH LM Tests
    Statistic Shape Scale P-Value
## ARCH Lag[4] 0.2874 0.500 2.000 0.5919
## ARCH Lag[6] 0.7433 1.461 1.711 0.8222
## ARCH Lag[8] 1.0528 2.368 1.583 0.9164
## Nyblom stability test
## -----
## Joint Statistic: 0.7623
## Individual Statistics:
## mu
        0.05153
## omega 0.14094
## alpha1 0.07645
## beta1 0.07352
## beta2 0.07697
## skew 0.15404
## shape 0.08679
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.69 1.9 2.35
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## t-value prob sig
## Sign Bias 2.5278 0.01154 **
## Negative Sign Bias 0.4456 0.65589
```

```
## Positive Sign Bias 1.9875 0.04698 **
## Joint Effect 10.3144 0.01607 **
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
  group statistic p-value(g-1)
      20 17.22
## 1
                    0.5752786
           60.08
## 2
      30
                    0.0006042
## 3 40 49.05
                    0.1300010
## 4
      50 61.85
                    0.1029340
##
##
## Elapsed time : 1.811992
spec_of_garch_hm<- ugarchspec(variance.model = list(garchOrder=c(2,1)),mean.model = list(armaOrder=c(0,0)</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
## *----*
           GARCH Model Fit
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(2,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
##
## Optimal Parameters
##
         Estimate Std. Error t value Pr(>|t|)
         0.000019 0.000540 0.035509 0.971674
## omega 0.000226 0.000224 1.009946 0.312521
## alpha1 0.357251 0.056287 6.346931 0.000000
## alpha2 0.000000 0.274977 0.000000 1.000000
## beta1 0.583942 0.322237 1.812154 0.069962
         1.298275 0.030926 41.979635 0.000000
## skew
## shape 2.981740 0.213429 13.970636 0.000000
## Robust Standard Errors:
         Estimate Std. Error t value Pr(>|t|)
##
## mu
         0.000019 \qquad 0.000552 \quad 0.034766 \ 0.972266
## omega
         ## alpha1 0.357251 0.465623 0.767254 0.442931
                  1.568333 0.000000 1.000000
## alpha2 0.000000
## beta1 0.583942 1.800795 0.324269 0.745734
## skew 1.298275 0.119602 10.854983 0.000000
## shape 2.981740 1.638870 1.819387 0.068852
##
## LogLikelihood : 5754.062
## Information Criteria
```

```
##
## Akaike
            -4.4208
## Bayes
            -4.4050
## Shibata
            -4.4208
## Hannan-Quinn -4.4151
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                         6.278 0.01223
## Lag[2*(p+q)+(p+q)-1][2] 6.378 0.01745
## Lag[4*(p+q)+(p+q)-1][5] 7.316 0.04345
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                       statistic p-value
## Lag[1]
                         0.1083 0.7421
## Lag[2*(p+q)+(p+q)-1][8]
                         2.7159 0.7373
## Lag[4*(p+q)+(p+q)-1][14] 4.8828 0.7839
## d.o.f=3
## Weighted ARCH LM Tests
## -----
            Statistic Shape Scale P-Value
## ARCH Lag[4] 0.3924 0.500 2.000 0.5311
## ARCH Lag[6] 0.7082 1.461 1.711 0.8324
## ARCH Lag[8] 0.9146 2.368 1.583 0.9365
## Nyblom stability test
## -----
## Joint Statistic: 3.8548
## Individual Statistics:
## mu
       0.05012
## omega 0.14029
## alpha1 0.08666
## alpha2 0.58290
## beta1 0.07364
## skew 0.16354
## shape 0.08786
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.69 1.9 2.35
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## -----
                  t-value prob sig
## Sign Bias
                  2.4981 0.01255 **
## Negative Sign Bias 0.4035 0.68664
## Positive Sign Bias 1.8563 0.06352
## Joint Effect 9.8093 0.02026 **
```

```
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
  group statistic p-value(g-1)
## 1
      20 25.00 0.1605422
## 2
      30 65.11 0.0001373
    40 46.49 0.1911055
## 3
                  0.1241588
## 4
    50 60.58
##
##
## Elapsed time : 1.552002
spec_of_garch_hm<-ugarchspec(variance.model =list(model="eGARCH",garchOrder=c(1,1)),mean.model = list(a</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
      GARCH Model Fit *
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : eGARCH(1,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
##
## Optimal Parameters
##
        Estimate Std. Error t value Pr(>|t|)
       ## omega -1.109089 0.238498 -4.65030 0.000003
## beta1 0.837270 0.034864 24.01507 0.000000
## gamma1 0.472441 0.056943 8.29681 0.000000
## skew 1.289988 0.036823 35.03198 0.000000
## shape 3.045425 0.196872 15.46904 0.000000
##
## Robust Standard Errors:
       Estimate Std. Error t value Pr(>|t|)
## mu
       ## omega -1.109089 0.361619 -3.06700 0.002162
## beta1 0.837270 0.052974 15.80527 0.000000
## gamma1 0.472441 0.070278 6.72245 0.000000
                0.043528 29.63564 0.000000
## skew
        1.289988
## shape 3.045425 0.199407 15.27240 0.000000
## LogLikelihood : 5756.141
##
## Information Criteria
##
```

```
-4.4224
## Akaike
          -4.4066
-4.4224
## Bayes
## Shibata
## Hannan-Quinn -4.4167
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                      statistic p-value
                        6.203 0.01275
## Lag[1]
## Lag[2*(p+q)+(p+q)-1][2] 6.286 0.01846
## Lag[4*(p+q)+(p+q)-1][5] 7.179 0.04691
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                      statistic p-value
## Lag[1]
                       0.0919 0.7618
## Lag[2*(p+q)+(p+q)-1][5] 1.4895 0.7425
## Lag[4*(p+q)+(p+q)-1][9] 1.9581 0.9100
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
## Statistic Shape Scale P-Value
## ARCH Lag[3] 0.6805 0.500 2.000 0.4094
## ARCH Lag[5] 0.9045 1.440 1.667 0.7615
## ARCH Lag[7] 0.9434 2.315 1.543 0.9226
## Nyblom stability test
## -----
## Joint Statistic: 1.0447
## Individual Statistics:
## mu
      0.02739
## omega 0.06910
## alpha1 0.30928
## beta1 0.07473
## gamma1 0.11247
## skew 0.13624
## shape 0.11201
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.69 1.9 2.35
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## -----
     t-value prob sig
##
## Sign Bias
                 2.2309 0.02577 **
## Negative Sign Bias 0.4485 0.65384
## Positive Sign Bias 1.4438 0.14892
## Joint Effect 8.5611 0.03573 **
##
##
```

```
## Adjusted Pearson Goodness-of-Fit Test:
## -----
   group statistic p-value(g-1)
## 1
      20
           17.48
                    0.557591
## 2
      30
           52.30
                    0.005046
## 3
      40 46.89
                 0.180364
## 4
      50
           69.54
                    0.028370
##
##
## Elapsed time : 1.609997
this model is rejected since there is no leverage effect.
spec_of_garch_hm<-ugarchspec(variance.model =list(model="gjrGARCH",garchOrder=c(1,1)),mean.model = list</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
## *----*
           GARCH Model Fit
##
## Conditional Variance Dynamics
## -----
## GARCH Model : gjrGARCH(1,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
## -----
##
        Estimate Std. Error t value Pr(>|t|)
       ## mu
                  0.000048 4.5905 0.000004
## omega 0.000220
## alpha1 0.325089 0.068840 4.7224 0.000002
## beta1
        ## gamma1 0.086331
                0.073188
                          1.1796 0.238170
                0.037178 34.9366 0.000000
## skew
        1.298861
## shape
        2.992478
                0.192316 15.5602 0.000000
##
## Robust Standard Errors:
        Estimate Std. Error t value Pr(>|t|)
       ## mu
## omega 0.000220 0.000064 3.45118 0.000558
## alpha1 0.325089 0.075644 4.29764 0.000017
                0.080180 7.26199 0.000000
## beta1
        0.582269
## gamma1 0.086331 0.077849 1.10895 0.267451
## skew
        1.298861 0.043929 29.56746 0.000000
        ## shape
## LogLikelihood: 5754.87
##
```

Information Criteria

```
##
## Akaike
            -4.4214
## Bayes
            -4.4057
## Shibata
            -4.4215
## Hannan-Quinn -4.4157
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                      statistic p-value
                        6.408 0.01136
## Lag[1]
## Lag[2*(p+q)+(p+q)-1][2] 6.505 0.01614
## Lag[4*(p+q)+(p+q)-1][5] 7.403 0.04135
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
             statistic p-value
##
                        0.05588 0.8131
## Lag[1]
## Lag[2*(p+q)+(p+q)-1][5] 2.01184 0.6156
## Lag[4*(p+q)+(p+q)-1][9] 2.69260 0.8084
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
            Statistic Shape Scale P-Value
## ARCH Lag[3] 0.8841 0.500 2.000 0.3471
## ARCH Lag[5] 1.3333 1.440 1.667 0.6372
## ARCH Lag[7] 1.4243 2.315 1.543 0.8364
## Nyblom stability test
## -----
## Joint Statistic: 0.941
## Individual Statistics:
## mu
     0.05304
## omega 0.12890
## alpha1 0.07097
## beta1 0.06696
## gamma1 0.22608
## skew 0.16372
## shape 0.08189
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.69 1.9 2.35
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
## -----
                  t-value prob sig
## Sign Bias
                   2.1343 0.03291 **
## Negative Sign Bias 0.6246 0.53226
## Positive Sign Bias 1.4962 0.13472
## Joint Effect 8.7885 0.03224 **
##
```

```
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
   group statistic p-value(g-1)
## 1 20 23.40 0.2201825
## 2
      30 58.65
                   0.0009075
## 3 40 55.08 0.0454688
    50 63.77 0.0763960
## 4
##
##
## Elapsed time : 3.777005
spec of garch hm<-ugarchspec(variance.model =list(model="gjrGARCH",garchOrder=c(1,2)),mean.model = list</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
## *----*
      GARCH Model Fit
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : gjrGARCH(1,2)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
         Estimate Std. Error t value Pr(>|t|)
       ## omega 0.000232 0.000048 4.802533 0.000002
## alpha1 0.362014 0.074397 4.865945 0.000001
## beta1 0.300193 0.092835 3.233627 0.001222
## beta2  0.243230  0.082327  2.954441  0.003132  ## gamma1  0.082775  0.082140  1.007736  0.313581
## skew 1.305833 0.037683 34.653328 0.000000
## shape 3.002980 0.194327 15.453204 0.000000
## Robust Standard Errors:
       Estimate Std. Error t value Pr(>|t|)
## mu
       ## omega 0.000232 0.000060 3.852563 0.000117
## alpha1 0.362014 0.076143 4.754405 0.000002
## beta1 0.300193 0.070311 4.269483 0.000020
## beta2
        ## gamma1 0.082775 0.089137 0.928634 0.353079
        1.305833 0.044764 29.171484 0.000000
## skew
## shape 3.002980 0.192387 15.609059 0.000000
##
## LogLikelihood : 5758.332
##
## Information Criteria
## -----
```

```
##
## Akaike
           -4.4233
## Bayes
            -4.4053
## Shibata
            -4.4234
## Hannan-Quinn -4.4168
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                      statistic p-value
## Lag[1]
                        5.920 0.01497
## Lag[2*(p+q)+(p+q)-1][2] 5.991 0.02212
## Lag[4*(p+q)+(p+q)-1][5] 6.823 0.05724
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
##
                       statistic p-value
## Lag[1]
                         0.1583 0.6907
                          2.1524 0.8348
\# \text{Lag}[2*(p+q)+(p+q)-1][8]
## Lag[4*(p+q)+(p+q)-1][14] 4.0486 0.8756
## d.o.f=3
##
## Weighted ARCH LM Tests
## -----
            Statistic Shape Scale P-Value
## ARCH Lag[4] 0.2544 0.500 2.000 0.6140
## ARCH Lag[6] 0.6405 1.461 1.711 0.8520
## ARCH Lag[8] 0.9668 2.368 1.583 0.9292
## Nyblom stability test
## -----
## Joint Statistic: 0.9656
## Individual Statistics:
## mu
      0.05254
## omega 0.13075
## alpha1 0.06490
## beta1 0.06861
## beta2 0.06955
## gamma1 0.19658
## skew 0.15424
## shape 0.08244
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.89 2.11 2.59
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## -----
                  t-value prob sig
              2.1647 0.03050 **
## Sign Bias
## Negative Sign Bias 0.6426 0.52055
## Positive Sign Bias 1.6422 0.10066
## Joint Effect 9.1608 0.02723 **
```

```
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
    group statistic p-value(g-1)
## 1
      20 18.52 0.487791
## 2
      30 55.23 0.002328
      40 52.15
                 0.077459
## 3
                  0.203005
## 4
      50
         56.96
##
##
## Elapsed time : 3.951006
spec_of_garch_hm<-ugarchspec(variance.model =list(model="gjrGARCH",garchOrder=c(2,1)),mean.model = list</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
       GARCH Model Fit *
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : gjrGARCH(2,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
##
## Optimal Parameters
##
        Estimate Std. Error t value Pr(>|t|)
       -0.000094 0.000539 -0.174208 0.861702
## mu
## omega 0.000219 0.000154 1.427241 0.153510
## alpha1 0.323861 0.030435 10.640952 0.000000
## alpha2 0.000000 0.191102 0.000000 1.000000
## beta1 0.584135 0.217373 2.687253 0.007204
## gamma1 0.088490 0.105166 0.841436 0.400104
## gamma2 -0.002779 0.095071 -0.029235 0.976677
## skew
         ## shape 2.991178 0.110955 26.958394 0.000000
##
## Robust Standard Errors:
        Estimate Std. Error t value Pr(>|t|)
##
## mu
        -0.000094 0.000656 -0.143249 0.886093
## omega 0.000219 0.000595 0.368175 0.712743
## alpha1 0.323861 0.204884 1.580707 0.113945
## alpha2 0.000000 0.754824 0.000000 1.000000
## beta1
         ## gamma1 0.088490 0.110012 0.804368 0.421184
1.298573 0.067969 19.105427 0.000000
## skew
        ## shape
##
```

LogLikelihood: 5754.794

```
##
## Information Criteria
## -----
##
## Akaike -4.4198
## Bayes -4.3995
## Shibata -4.4199
## Hannan-Quinn -4.4125
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                         6.384 0.01151
## Lag[2*(p+q)+(p+q)-1][2] 6.482 0.01637
## Lag[4*(p+q)+(p+q)-1][5] 7.382 0.04185
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                       statistic p-value
## Lag[1]
                         0.05326 0.8175
## Lag[2*(p+q)+(p+q)-1][8] 2.48372 0.7787
## Lag[4*(p+q)+(p+q)-1][14] 4.72712 0.8025
## d.o.f=3
## Weighted ARCH LM Tests
    Statistic Shape Scale P-Value
## ARCH Lag[4] 0.3297 0.500 2.000 0.5658
## ARCH Lag[6] 0.5928 1.461 1.711 0.8657
## ARCH Lag[8] 0.8223 2.368 1.583 0.9486
## Nyblom stability test
## -----
## Joint Statistic: 4.8984
## Individual Statistics:
## mu
        0.05268
## omega 0.12865
## alpha1 0.07112
## alpha2 0.60819
## beta1 0.06712
## gamma1 0.22618
## gamma2 0.20112
## skew 0.16180
## shape 0.08188
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 2.1 2.32 2.82
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
## -----
##
                   t-value prob sig
```

```
## Sign Bias
                    2.133 0.03301 **
## Negative Sign Bias 0.625 0.53204
## Positive Sign Bias 1.489 0.13673
## Joint Effect
                     8.780 0.03236 **
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
##
    group statistic p-value(g-1)
## 1
      20 23.58
                  0.21257
## 2
      30
             56.45
                      0.00167
            56.28
## 3
      40
                      0.03610
## 4
      50
             62.73
                      0.08993
##
##
## Elapsed time : 4.890011
spec_of_garch_hm<-ugarchspec(variance.model =list(model="gjrGARCH",garchOrder=c(2,2)),mean.model = list</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
## *----*
          GARCH Model Fit
##
## Conditional Variance Dynamics
## -----
## GARCH Model : gjrGARCH(2,2)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
##
         Estimate Std. Error t value Pr(>|t|)
## mu
        -0.000062 0.000544 -0.113278 0.909811
## omega 0.000250 0.000075 3.326406 0.000880
## alpha1 0.392820 0.080214 4.897134 0.000001
## alpha2 0.000000 0.136761 0.000001 0.999999
## beta1 0.217187 0.321488 0.675567 0.499316
## beta2
         ## gamma1 -0.010421 0.091185 -0.114284 0.909013
## gamma2 0.123906 0.050189 2.468772 0.013558
                 0.037370 34.877867 0.000000
## skew
         1.303377
## shape
         3.006382
                  0.192534 15.614837 0.000000
##
## Robust Standard Errors:
##
         Estimate Std. Error t value Pr(>|t|)
        -0.000062 0.000574 -0.107387 0.91448
## mu
## omega 0.000250 0.000144 1.738685 0.08209
## alpha1 0.392820 0.074009 5.307725 0.00000
## alpha2 0.000000 0.228499 0.000001 1.00000
## beta1 0.217187 0.525521 0.413279 0.67940
## beta2 0.283618 0.292396 0.969978 0.33206
```

```
## gamma1 -0.010421 0.092780 -0.112319 0.91057
## gamma2 0.123906 0.098028 1.263983 0.20624
## skew 1.303377 0.044862 29.053180 0.00000
          3.006382 0.201190 14.943004 0.00000
## shape
## LogLikelihood : 5759.305
## Information Criteria
## -----
##
## Akaike
             -4.4225
## Bayes
             -4.4000
## Shibata -4.4226
## Hannan-Quinn -4.4144
## Weighted Ljung-Box Test on Standardized Residuals
##
                        statistic p-value
## Lag[1]
                           5.880 0.01531
## Lag[2*(p+q)+(p+q)-1][2] 5.925 0.02303
## Lag[4*(p+q)+(p+q)-1][5] 6.777 0.05873
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                          statistic p-value
## Lag[1]
                            0.215 0.6428
## Lag[2*(p+q)+(p+q)-1][11] 3.112 0.8552
## Lag[4*(p+q)+(p+q)-1][19] 5.378 0.9178
## d.o.f=4
##
## Weighted ARCH LM Tests
             Statistic Shape Scale P-Value
## ARCH Lag[5] 0.4797 0.500 2.000 0.4886
## ARCH Lag[7] 0.6932 1.473 1.746 0.8439
## ARCH Lag[9] 1.4579 2.402 1.619 0.8594
##
## Nyblom stability test
## -----
## Joint Statistic: 1.1057
## Individual Statistics:
## mu 0.06388
## omega 0.12788
## alpha1 0.05695
## alpha2 0.06136
## beta1 0.06087
## beta2 0.06195
## gamma1 0.18352
## gamma2 0.14391
## skew 0.16535
## shape 0.08022
##
```

```
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 2.29 2.54 3.05
## Individual Statistic:
                       0.35 0.47 0.75
##
## Sign Bias Test
## -----
                   t-value prob sig
## Sign Bias
                   2.0963 0.03615 **
## Negative Sign Bias 0.5505 0.58204
## Positive Sign Bias 1.7315 0.08349
## Joint Effect 8.3630 0.03908 **
##
##
## Adjusted Pearson Goodness-of-Fit Test:
    group statistic p-value(g-1)
      20 15.63 0.68175
## 1
## 2
      30
            48.17
                      0.01411
## 3
      40
            44.55
                      0.24953
      50
## 4
            54.04
                      0.28796
##
## Elapsed time: 10.728
```

sGARCH(0,2) Akaike -4.2988 sstd sGARCH(0,1) Akaike -4.2996 sstd sGARCH(1,0) Akaike -4.3933 sstd sGARCH(2,0) Akaike -4.4074 sstd gjrGARCH(2,1) Akaike -4.4198 sstd sGARCH(2,1) Akaike -4.4208 sstd gjrGARCH(1,1) Akaike -4.4214 sstd sGARCH(1,1) Akaike -4.4216 sstd gjrGARCH(2,2) Akaike -4.4225 sstd sGARCH(2,2) Akaike -4.4229 sstd gjrGARCH(1,2) Akaike -4.4233 sstd sGARCH(1,2) Akaike -4.4237 sstd

interestingly a simple garch model is the best choice also to make the model simple i am taking sGARCH (1,2) with no ARMA component.

```
spec_of_garch_hm<-ugarchspec(variance.model =list(model="gjrGARCH",garchOrder=c(1,3)),mean.model = list
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)
my_model_hm</pre>
```

```
##
## *----*
        GARCH Model Fit
## *----*
##
## Conditional Variance Dynamics
## -----
## GARCH Model : gjrGARCH(1,3)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
## -----
##
        Estimate Std. Error t value Pr(>|t|)
       0.000025 0.000547 0.045533 0.963683
## mu
## omega 0.000237 0.000049 4.810035 0.000002
## alpha1 0.367888 0.074968 4.907253 0.000001
## beta1 0.288182 0.107093 2.690956 0.007125
```

```
## beta2 0.124810 0.109319 1.141698 0.253580
## beta3 0.117955 0.072893 1.618187 0.105622
## gamma1 0.100251 0.086270 1.162063 0.245210
## skew 1.309344 0.037506 34.910587 0.000000
## shape 2.994535 0.193704 15.459313 0.000000
##
## Robust Standard Errors:
         Estimate Std. Error t value Pr(>|t|)
##
## mu
         0.000025 0.000573 0.043518 0.965289
## omega 0.000237 0.000059 4.010179 0.000061
## alpha1 0.367888 0.073876 4.979831 0.000001
## beta1 0.288182 0.092425 3.118009 0.001821 ## beta2 0.124810 0.106565 1.171208 0.241515
## beta3 0.117955 0.074213 1.589413 0.111967
## shape 2.994535 0.192015 15.595303 0.000000
##
## LogLikelihood : 5759.558
## Information Criteria
## -----
##
            -4.4235
## Akaike
## Bayes
            -4.4032
## Shibata
            -4.4235
## Hannan-Quinn -4.4162
## Weighted Ljung-Box Test on Standardized Residuals
##
                       statistic p-value
## Lag[1]
                          5.773 0.01627
## Lag[2*(p+q)+(p+q)-1][2]
                        5.837 0.02432
## Lag[4*(p+q)+(p+q)-1][5]
                        6.623 0.06393
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                        statistic p-value
## Lag[1]
                         0.1674 0.6824
## Lag[2*(p+q)+(p+q)-1][11] 2.8791 0.8826
## Lag[4*(p+q)+(p+q)-1][19] 5.1483 0.9315
## d.o.f=4
## Weighted ARCH LM Tests
             Statistic Shape Scale P-Value
## ARCH Lag[5] 0.3158 0.500 2.000 0.5741
             0.5154 1.473 1.746 0.8931
## ARCH Lag[7]
## ARCH Lag[9]
             1.3908 2.402 1.619 0.8710
## Nyblom stability test
## -----
```

```
## Joint Statistic: 1.2466
## Individual Statistics:
       0.04389
## omega 0.13528
## alpha1 0.05930
## beta1 0.07445
## beta2 0.07391
## beta3 0.06655
## gamma1 0.18270
## skew 0.15168
## shape 0.08080
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 2.1 2.32 2.82
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
## -----
                 t-value prob sig
##
## Sign Bias
                  2.6117 0.009061 ***
## Negative Sign Bias 0.6038 0.546006
## Positive Sign Bias 1.8867 0.059317
## Joint Effect 11.5143 0.009246 ***
##
##
## Adjusted Pearson Goodness-of-Fit Test:
##
   group statistic p-value(g-1)
## 1 20 16.06 0.653168
    30 51.61 0.006027
## 2
     40 40.98
                  0.383511
## 3
## 4
    50 60.85
                  0.119393
##
##
## Elapsed time : 3.780011
spec_of_garch_hm<-ugarchspec(variance.model =list(model="gjrGARCH",garchOrder=c(3,1)),mean.model = list</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
           GARCH Model Fit *
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : gjrGARCH(3,1)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
##
## Optimal Parameters
## -----
##
        Estimate Std. Error t value Pr(>|t|)
```

```
## mu
        -0.000094 0.000545 -0.172581 0.862981
## omega 0.000223 0.000074 3.001972 0.002682
## alpha1 0.328450 0.056066 5.858312 0.000000
## alpha2 0.000000 0.052268 0.000000 1.000000
## alpha3 0.000000 0.029344 0.000001 0.999999
## beta1 0.576413 0.117849 4.891113 0.000001
## gamma1 0.087838 0.106428 0.825326 0.409186
## skew
         1.298689 0.036790 35.300223 0.000000
## shape
          2.991832 0.175448 17.052511 0.000000
##
## Robust Standard Errors:
## Estimate Std. Error t value Pr(>|t|)
## mu -0.000094 0.000565 -0.16647 0.867787
## omega 0.000223 0.000125 1.77463 0.075959
## alpha1 0.328450 0.061530 5.33809 0.000000
## alpha2 0.000000 0.104541 0.00000 1.000000
## alpha3 0.000000 0.094022 0.00000 1.000000
## beta1 0.576413 0.181975 3.16753 0.001537
## gamma1 0.087838 0.099288 0.88467 0.376332
## gamma2 -0.023570 0.120729 -0.19523 0.845215
## gamma3 0.028463 0.089868 0.31672 0.751453
         1.298689 0.045521 28.52944 0.000000
## skew
## shape 2.991832 0.241797 12.37331 0.000000
## LogLikelihood : 5754.926
## Information Criteria
##
## Akaike
            -4.4184
## Bayes
             -4.3936
## Shibata
            -4.4184
## Hannan-Quinn -4.4094
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                         6.354 0.01171
## Lag[2*(p+q)+(p+q)-1][2] 6.457 0.01662
## Lag[4*(p+q)+(p+q)-1][5] 7.344 0.04277
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
##
                        statistic p-value
## Lag[1]
                        0.0622 0.8030
## Lag[2*(p+q)+(p+q)-1][11] 3.5799 0.7936
## Lag[4*(p+q)+(p+q)-1][19] 6.2946 0.8494
## d.o.f=4
##
## Weighted ARCH LM Tests
```

```
## Statistic Shape Scale P-Value
## ARCH Lag[5] 0.3065 0.500 2.000 0.5799
## ARCH Lag[7] 0.3700 1.473 1.746 0.9311
## ARCH Lag[9] 1.0197 2.402 1.619 0.9286
##
## Nyblom stability test
## -----
## Joint Statistic: 5.4518
## Individual Statistics:
       0.05276
## omega 0.12722
## alpha1 0.07353
## alpha2 0.59598
## alpha3 0.30636
## beta1 0.06683
## gamma1 0.22349
## gamma2 0.20180
## gamma3 0.17450
## skew 0.16074
## shape 0.08257
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 2.49 2.75 3.27
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
                  t-value prob sig
## Sign Bias
                  2.1249 0.03369 **
## Negative Sign Bias 0.6319 0.52749
## Positive Sign Bias 1.5047 0.13252
## Joint Effect 8.7621 0.03263 **
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 20.20 0.382635
      30 50.78 0.007439
## 2
## 3 40 52.77 0.069489
## 4 50 61.88 0.102339
##
## Elapsed time : 6.719008
spec_of_garch_hm<-ugarchspec(variance.model =list(model="gjrGARCH",garchOrder=c(2,3)),mean.model = list</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
## *----*
```

* GARCH Model Fit * ## *-----*

```
##
## Conditional Variance Dynamics
## -----
## GARCH Model : gjrGARCH(2,3)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
## -----
##
         Estimate Std. Error t value Pr(>|t|)
## mu
        -0.000011 0.000548 -0.02041 0.983716
                   0.000043 5.84800 0.000000
## omega 0.000250
## alpha1 0.392020 0.080671 4.85950 0.000001
## alpha2 0.000000 0.099375 0.00000 1.000000
## beta1
        ## beta2
## beta3
## gamma1 0.026369 0.107491 0.24531 0.806216
## gamma2 0.095793 0.074995 1.27733 0.201486
## skew
         1.306845 0.037593 34.76264 0.000000
## shape
         3.000046 0.194163 15.45120 0.000000
##
## Robust Standard Errors:
         Estimate Std. Error
##
                            t value Pr(>|t|)
## mu
        -0.000011 0.000580 -0.019313 0.984592
## omega 0.000250 0.000076 3.278399 0.001044
## alpha1 0.392020 0.076777 5.105936 0.000000
## alpha2 0.000000 0.091554 0.000000 1.000000
## beta1 0.216412 0.204553 1.057976 0.290066
## beta2 0.183339 0.086443 2.120927 0.033928
## beta3  0.097993  0.113170  0.865893  0.386549
## gamma1  0.026369  0.111585  0.236312  0.813191
## gamma2 0.095793 0.069281 1.382668 0.166767
         1.306845 0.044321 29.485989 0.000000
## skew
## shape
         3.000046
                  0.193384 15.513390 0.000000
##
## LogLikelihood : 5760.064
##
## Information Criteria
## -----
            -4.4224
## Akaike
## Baves
             -4.3976
## Shibata
            -4.4224
## Hannan-Quinn -4.4134
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                          5.740 0.01658
                       5.786 0.02508
## Lag[2*(p+q)+(p+q)-1][2]
## Lag[4*(p+q)+(p+q)-1][5]
                          6.600 0.06476
## d.o.f=0
## HO : No serial correlation
```

```
##
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                        statistic p-value
## Lag[1]
                          0.2185 0.6402
## Lag[2*(p+q)+(p+q)-1][14] 3.7329 0.9043
## Lag[4*(p+q)+(p+q)-1][24] 6.0974 0.9662
## d.o.f=5
##
## Weighted ARCH LM Tests
## -----
             Statistic Shape Scale P-Value
## ARCH Lag[6] 0.06996 0.500 2.000 0.7914
## ARCH Lag[8] 0.53803 1.480 1.774 0.8907
## ARCH Lag[10] 2.23710 2.424 1.650 0.7260
##
## Nyblom stability test
## -----
## Joint Statistic: 1.5146
## Individual Statistics:
## mu
       0.05340
## omega 0.13186
## alpha1 0.05610
## alpha2 0.05066
## beta1 0.06674
## beta2 0.06798
## beta3 0.05911
## gamma1 0.17166
## gamma2 0.15653
## skew 0.15770
## shape 0.07998
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 2.49 2.75 3.27
## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## -----
## t-value prob sig
## Sign Bias 2.1676 0.03028 **
## Negative Sign Bias 0.5882 0.55644
## Positive Sign Bias 1.7249 0.08467
## Joint Effect 8.9867 0.02947 **
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 10.

## 2 30 46.46 0.0211

## 3 40 41.85 0.3483

50 48.50 0.4933
##
##
```

```
## Elapsed time : 5.529996
spec_of_garch_hm<-ugarchspec(variance.model =list(model="gjrGARCH",garchOrder=c(3,2)),mean.model = list</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
## *----*
       GARCH Model Fit *
## *----*
##
## Conditional Variance Dynamics
## -----
## GARCH Model : gjrGARCH(3,2)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
##
## Convergence Problem:
## Solver Message:
spec_of_garch_hm<-ugarchspec(variance.model =list(model="gjrGARCH",garchOrder=c(3,3)),mean.model = list</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm)</pre>
my_model_hm
##
     GARCH Model Fit *
##
## Conditional Variance Dynamics
## -----
## GARCH Model : gjrGARCH(3,3)
## Mean Model : ARFIMA(0,0,0)
## Distribution : sstd
## Optimal Parameters
## -----
         Estimate Std. Error t value Pr(>|t|)
       -0.000169 0.000001 -193.470
## mu
## omega 0.000137 0.000001 180.895
## alpha1 0.209953 0.001009 208.170
## alpha2 0.001999 0.000024 84.275
## alpha3 0.001229 0.000020 60.930
                                          0
                                          0
## beta1 0.166145 0.000819 202.961
                                          0
## beta2 0.537078 0.002369 226.753
                                          0
## beta3 0.018319 0.000192 95.198
                                          0
## gamma1 0.469651 0.002180 215.426
                                          0
## gamma2 0.096881 0.000536 180.905
## gamma3 -0.457302 0.002027 -225.660
                                          0
         1.334417 0.032719 40.784
## skew
                                          0
## shape 2.913417 0.060530 48.132
```

Robust Standard Errors:

```
Estimate Std. Error t value Pr(>|t|)
## mu
        -0.000169 0.000888 -0.190842 0.84865
## omega 0.000137 0.000089 1.535929 0.12456
## alpha1 0.209953 0.440330 0.476807 0.63350
## alpha2 0.001999 0.015057 0.132773 0.89437
## alpha3 0.001229 0.015425 0.079688 0.93649
## beta1 0.166145 0.319364 0.520238 0.60290
## beta2  0.537078  1.535627  0.349745  0.72653  ## beta3  0.018319  0.110220  0.166202  0.86800
## gamma1 0.469651 1.068786 0.439425 0.66035
## gamma2 0.096881 0.067141 1.442950 0.14904
## gamma3 -0.457302 2.665251 -0.171579 0.86377
## skew
          1.334417 0.171809 7.766846 0.00000
## shape
        2.913417 19.175016 0.151938 0.87924
##
## LogLikelihood: 5748.73
##
## Information Criteria
## Akaike -4.4121
## Bayes
             -4.3828
## Shibata -4.4121
## Hannan-Quinn -4.4015
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                          6.415 0.01132
## Lag[2*(p+q)+(p+q)-1][2] 6.422 0.01698
## Lag[4*(p+q)+(p+q)-1][5] 7.172 0.04710
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                       statistic p-value
## Lag[1]
                           0.2099 0.6468
## Lag[2*(p+q)+(p+q)-1][17] 4.2598 0.9414
## Lag[4*(p+q)+(p+q)-1][29] 8.2184 0.9567
## d.o.f=6
##
## Weighted ARCH LM Tests
## -----
             Statistic Shape Scale P-Value
             0.03311 0.500 2.000 0.8556
## ARCH Lag[7]
## ARCH Lag[9] 1.58863 1.485 1.796 0.6160
## ARCH Lag[11] 3.72096 2.440 1.677 0.4717
## Nyblom stability test
## -----
## Joint Statistic: 13.7408
## Individual Statistics:
## mu 0.05238
```

```
## omega 0.14063
## alpha1 0.03677
## alpha2 0.07893
## alpha3 0.09772
## beta1 0.03707
## beta2 0.03719
## beta3 0.10439
## gamma1 0.03688
## gamma2 0.18498
## gamma3 0.03735
## skew 0.26005
## shape 0.06294
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 2.89 3.15 3.69
## Individual Statistic: 0.35 0.47 0.75
##
## Sign Bias Test
## -----
                  t-value prob sig
## Sign Bias
                  2.2360 0.025436 **
## Negative Sign Bias 0.8967 0.369944
## Positive Sign Bias 0.5444 0.586206
## Joint Effect 12.2257 0.006649 ***
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 18.66 0.4787337
## 2
      30 28.35
                    0.4994676
## 3
    40 50.40 0.1044144
## 4
      50 88.88 0.0004258
##
## Elapsed time : 6.875004
going further is not leading any good results
spec_of_garch_hm<-ugarchspec(variance.model = list(garch0rder=c(1,2)),mean.model = list(arma0rder=c(0,0))</pre>
my_model_hm<-ugarchfit(spec=spec_of_garch_hm,data=ret_hm,out.sample = 500)</pre>
my_model_hm
          GARCH Model Fit
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(1,2)
```

Mean Model : ARFIMA(0,0,0)

Distribution : sstd

```
##
## Optimal Parameters
## -----
         Estimate Std. Error t value Pr(>|t|)
         0.000017 0.000599 0.027762 0.977852
## omega 0.000247 0.000055 4.448606 0.000009
## alpha1 0.396450 0.080932 4.898560 0.000001
## beta1 0.297301 0.103122 2.883017 0.003939
## beta2 0.234754 0.088927 2.639842 0.008294
## skew 1.321476 0.042175 31.332971 0.000000
## shape 2.999525 0.218506 13.727418 0.000000
## Robust Standard Errors:
##
        Estimate Std. Error t value Pr(>|t|)
## mu
       0.000017 0.000615 0.02702 0.978444
## omega 0.000247 0.000065 3.82094 0.000133
## alpha1 0.396450 0.080241 4.94076 0.000001
## beta1 0.297301 0.079405 3.74413 0.000181
## beta2 0.234754 0.086551 2.71233 0.006681
## skew 1.321476 0.049427 26.73572 0.000000
## shape 2.999525 0.207501 14.45549 0.000000
## LogLikelihood: 4672.518
## Information Criteria
## -----
##
## Akaike -4.4434
## Bayes
            -4.4245
## Shibata -4.4434
## Hannan-Quinn -4.4365
##
## Weighted Ljung-Box Test on Standardized Residuals
## -----
                      statistic p-value
##
## Lag[1]
                         5.621 0.01775
## Lag[2*(p+q)+(p+q)-1][2] 5.628 0.02764
## Lag[4*(p+q)+(p+q)-1][5] 5.960 0.09191
## d.o.f=0
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
          statistic p-value
##
                          0.2961 0.5864
## Lag[1]
## Lag[2*(p+q)+(p+q)-1][8] 1.9989 0.8591
## Lag[4*(p+q)+(p+q)-1][14] 4.1402 0.8666
## d.o.f=3
## Weighted ARCH LM Tests
      Statistic Shape Scale P-Value
## ARCH Lag[4] 0.1754 0.500 2.000 0.6753
## ARCH Lag[6] 0.3865 1.461 1.711 0.9230
```

```
## -----
## Joint Statistic: 0.8594
## Individual Statistics:
       0.02994
## omega 0.21873
## alpha1 0.05295
## beta1 0.08089
## beta2 0.07362
## skew 0.07846
## shape 0.09488
##
## Asymptotic Critical Values (10% 5% 1%)
## Joint Statistic: 1.69 1.9 2.35
## Individual Statistic:
                         0.35 0.47 0.75
##
## Sign Bias Test
## -----
##
                    t-value
                              prob sig
                    2.3633 0.01821 **
## Sign Bias
## Negative Sign Bias 0.3329 0.73924
## Positive Sign Bias 1.6455 0.10001
## Joint Effect 8.5895 0.03528 **
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## group statistic p-value(g-1)
## 1
      20 15.68
                      0.6788
## 2
       30 35.23
                        0.1971
     40 48.30
## 3
                        0.1459
## 4
       50 47.67
                         0.5273
##
##
## Elapsed time : 1.310992
back_testing<-ugarchroll(spec_of_garch_hm,ret_hm,n.ahead=1,n.start = 2000,refit.every =30,refit.window
##
## Iter: 1 fn: -4438.3025 Pars: 0.0003191 0.0002506 0.4141641 0.2941073 0.2506736 1.3396645 2.90325
## Iter: 2 fn: -4438.3025 Pars: 0.0003192 0.0002506 0.4141935 0.2940687 0.2506947 1.3396678 2.90321
## Iter: 3 fn: -4438.3025 Pars: 0.0003192 0.0002506 0.4141935 0.2940687 0.2506947 1.3396678 2.90321
## solnp--> Completed in 3 iterations
## Iter: 1 fn: -4432.0539
                          Pars: 0.0002559 0.0002456 0.4115616 0.3170721 0.2328361 1.3382652 2.91266
                          Pars: 0.0002559 0.0002456 0.4115616 0.3170721 0.2328361 1.3382652 2.91266
## Iter: 2 fn: -4432.0539
## solnp--> Completed in 2 iterations
## Iter: 1 fn: -4430.1153
                          Pars: 0.0001412 0.0002383 0.3948317 0.3170265 0.2283056 1.3342451 3.00282
## Iter: 2 fn: -4430.1153 Pars: 0.0001412 0.0002383 0.3948368 0.3170189 0.2283117 1.3342451 3.00280
## Iter: 3 fn: -4430.1153 Pars: 0.0001412 0.0002383 0.3948368 0.3170189 0.2283117 1.3342451 3.00280
## solnp--> Completed in 3 iterations
```

ARCH Lag[8]

Nyblom stability test

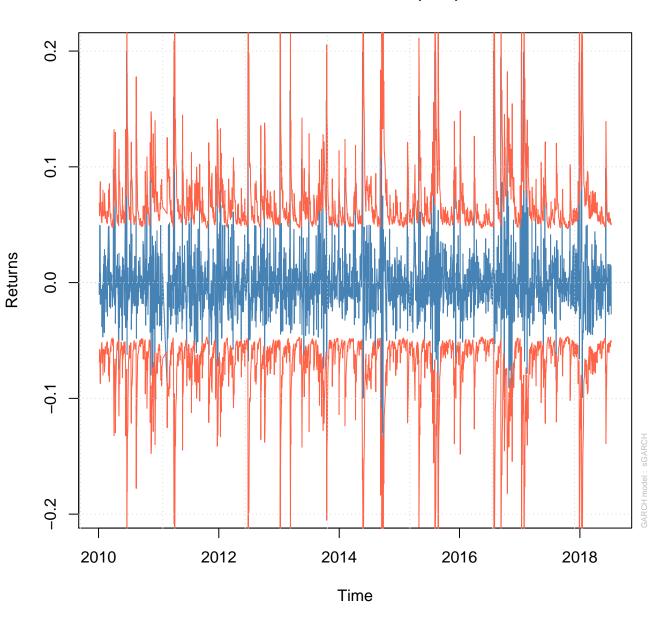
##

0.9404 2.368 1.583 0.9329

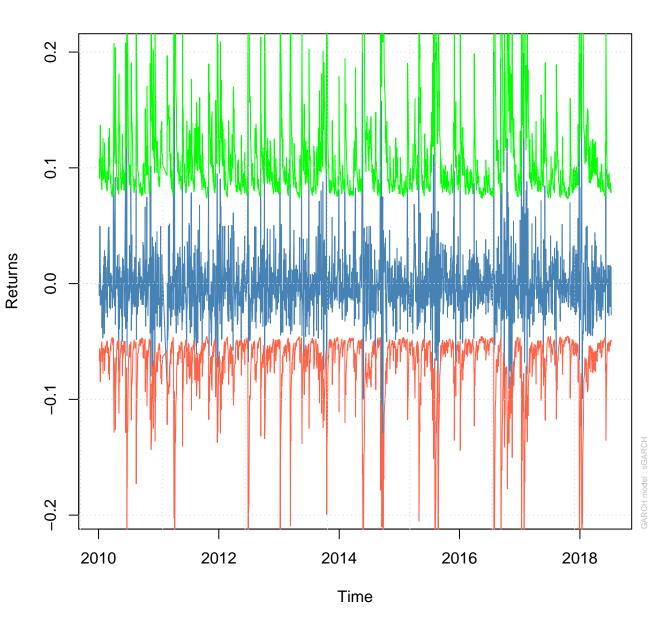
```
##
## Iter: 1 fn: -4437.5357 Pars: 0.00009238 0.00022411 0.37991764 0.32819760 0.22662345 1.32913901 3
## Iter: 2 fn: -4437.5357 Pars: 0.00009235 0.00022411 0.37992346 0.32821076 0.22661947 1.32913287 3
                          Pars: 0.00009235 0.00022411 0.37992346 0.32821076 0.22661947 1.32913287 3
## Iter: 3 fn: -4437.5357
## solnp--> Completed in 3 iterations
##
                            Pars: 0.0001463 0.0002006 0.3642601 0.3219088 0.2617950 1.3257079 3.03613
## Iter: 1 fn: -4457.7985
## Iter: 2 fn: -4457.7985
                          Pars: 0.0001463 0.0002006 0.3642601 0.3219088 0.2617950 1.3257079 3.03613
## solnp--> Completed in 2 iterations
## Iter: 1 fn: -4455.5486
                            Pars: 0.0001491 0.0001870 0.3518530 0.3235934 0.2680945 1.3183685 3.09579
                          Pars: 0.0001491 0.0001870 0.3518530 0.3235934 0.2680945 1.3183685 3.09579
## Iter: 2 fn: -4455.5486
## solnp--> Completed in 2 iterations
##
                            Pars: 0.0002003 0.0001919 0.3571111 0.3028881 0.2830643 1.3167092 3.09720
## Iter: 1 fn: -4449.8214
## Iter: 2 fn: -4449.8214
                            Pars: 0.0002003 0.0001919 0.3571111 0.3028881 0.2830643 1.3167092 3.09720
## solnp--> Completed in 2 iterations
##
                            Pars: 0.00008679 0.00018749 0.35402371 0.31096276 0.27289886 1.30938385 3
## Iter: 1 fn: -4456.0332
## Iter: 2 fn: -4456.0332
                            Pars: 0.00008679 0.00018749 0.35402371 0.31096276 0.27289886 1.30938385 3
## solnp--> Completed in 2 iterations
                            Pars: 0.0001112 0.0001987 0.3567669 0.3358767 0.2341392 1.3304676 3.14201
## Iter: 1 fn: -4476.3422
                            Pars: 0.0001112 0.0001987 0.3567669 0.3358767 0.2341392 1.3304676 3.14201
## Iter: 2 fn: -4476.3422
## solnp--> Completed in 2 iterations
## Iter: 1 fn: -4485.5811
                            Pars: 0.00009406 0.00019011 0.32871442 0.35971664 0.22233564 1.31916175 3
                            Pars: 0.00009406 0.00019011 0.32871441 0.35971664 0.22233564 1.31916175 3
## Iter: 2 fn: -4485.5811
## solnp--> Completed in 2 iterations
##
## Iter: 1 fn: -4497.8384
                            Pars: 0.00007323 0.00019600 0.33984839 0.37879152 0.18772301 1.32552939 3
## Iter: 2 fn: -4497.8384
                            Pars: 0.00007323 0.00019600 0.33984839 0.37879152 0.18772301 1.32552939 3
## solnp--> Completed in 2 iterations
## Iter: 1 fn: -4507.9647
                            Pars: 0.0000911 0.0001967 0.3442811 0.3753093 0.1898310 1.3237932 3.18727
## Iter: 2 fn: -4507.9647
                            Pars: 0.00009114 0.00019667 0.34427479 0.37531025 0.18983020 1.32379872 3
## Iter: 3 fn: -4507.9647
                            Pars: 0.00009114 0.00019667 0.34427478 0.37531025 0.18983021 1.32379872 3
## solnp--> Completed in 3 iterations
##
## Iter: 1 fn: -4501.7633
                            Pars: -0.00002738 0.00019099 0.34118478 0.36788207 0.20101396 1.3171
## Iter: 2 fn: -4501.7633
                            Pars: -0.00002732 0.00019099 0.34119375 0.36788494 0.20101041 1.3171
## Iter: 3 fn: -4501.7633
                            Pars: -0.00002732 0.00019099 0.34119375 0.36788494 0.20101041 1.3171
## solnp--> Completed in 3 iterations
##
                            Pars: 0.00008735 0.00018178 0.33171290 0.32929802 0.25581101 1.32701297 3
## Iter: 1 fn: -4484.4871
                            Pars: 0.00008734 0.00018178 0.33170903 0.32930009 0.25581051 1.32701197 3
## Iter: 2 fn: -4484.4871
## solnp--> Completed in 2 iterations
##
## Iter: 1 fn: -4478.6381
                            Pars: 0.0002059 0.0001860 0.3518316 0.3314664 0.2456153 1.3332225 3.19314
## Iter: 2 fn: -4478.6381
                            Pars: 0.000206 0.000186 0.351842 0.331468 0.245609 1.333226 3.193112
## Iter: 3 fn: -4478.6381
                            Pars: 0.000206 0.000186 0.351842 0.331468 0.245609 1.333226 3.193112
## solnp--> Completed in 3 iterations
##
                            Pars: 0.00009406 0.00018897 0.33836745 0.33116154 0.23848076 1.32168473 3
## Iter: 1 fn: -4473.5635
```

```
## Iter: 2 fn: -4473.5635
                           Pars: 0.00009406 0.00018897 0.33836745 0.33116154 0.23848076 1.32168473 3
## solnp--> Completed in 2 iterations
                         Pars: 0.0002717 0.0001870 0.3349683 0.3328525 0.2444990 1.3331944 3.21822
## Iter: 1 fn: -4494.9272
## Iter: 2 fn: -4494.9272
                           Pars: 0.0002717 0.0001870 0.3349683 0.3328525 0.2444990 1.3331944 3.21822
## solnp--> Completed in 2 iterations
                           Pars: 0.0001051 0.0002015 0.3512651 0.3209563 0.2488804 1.3197470 3.10757
## Iter: 1 fn: -4492.4002
## Iter: 2 fn: -4492.4002
                           Pars: 0.0001051 0.0002015 0.3512651 0.3209563 0.2488804 1.3197470 3.10757
## solnp--> Completed in 2 iterations
## Iter: 1 fn: -4473.8841
                           Pars: -0.0000255 0.0002111 0.3595951 0.3127036 0.2391634 1.3072296
## Iter: 2 fn: -4473.8841
                           Pars: -0.0000255 0.0002111 0.3595951 0.3127036 0.2391634 1.3072296
## solnp--> Completed in 2 iterations
## Iter: 1 fn: -4463.7398
                           Pars: 0.0001854 0.0002160 0.3791622 0.3051964 0.2561402 1.3172979 2.98605
                           Pars: 0.0001854 0.0002160 0.3791619 0.3051971 0.2561402 1.3172983 2.98605
## Iter: 2 fn: -4463.7398
## solnp--> Completed in 2 iterations
report(back_testing)
## VaR Backtest Report
## Model:
                       sGARCH-sstd
## Backtest Length: 600
## Data:
##
## ===========
## alpha:
## Expected Exceed: 6
## Actual VaR Exceed:
                      7
## Actual %:
                       1.2%
##
## Unconditional Coverage (Kupiec)
## Null-Hypothesis: Correct Exceedances
## LR.uc Statistic: 0.16
## LR.uc Critical:
                       3.841
## LR.uc p-value:
                       0.689
## Reject Null:
                   NO
##
## Conditional Coverage (Christoffersen)
## Null-Hypothesis: Correct Exceedances and
##
                   Independence of Failures
## LR.cc Statistic: 0.325
## LR.cc Critical:
                       5.991
## LR.cc p-value:
                       0.85
## Reject Null:
                   NO
```

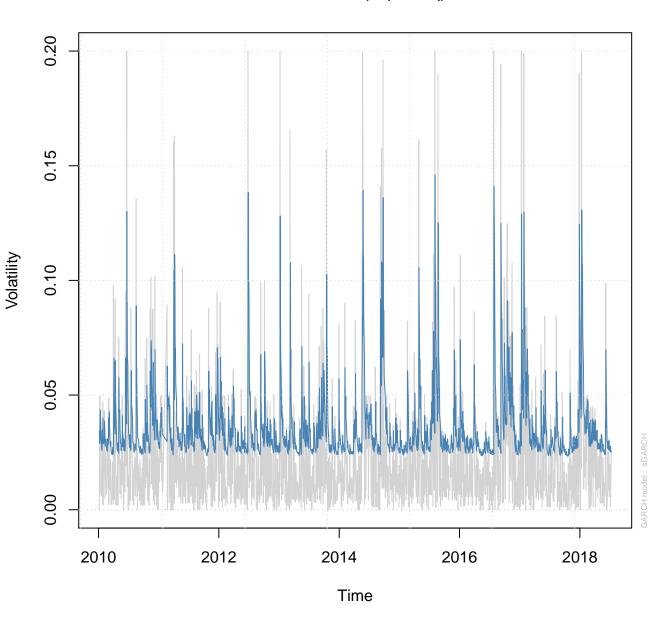
Series with 2 Conditional SD Superimposed



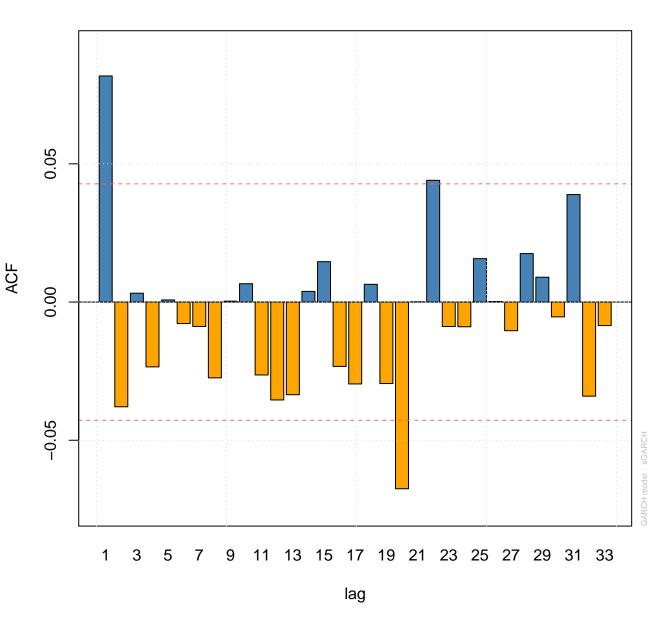
Series with with 1% VaR Limits



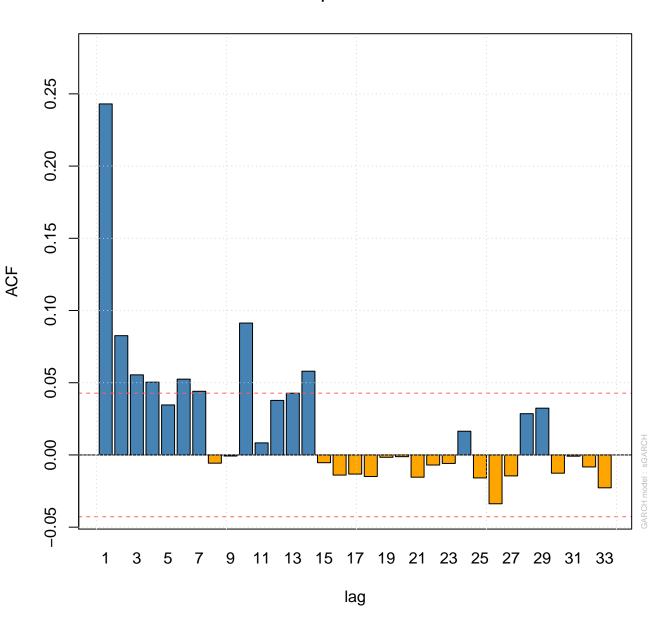
Conditional SD (vs |returns|)



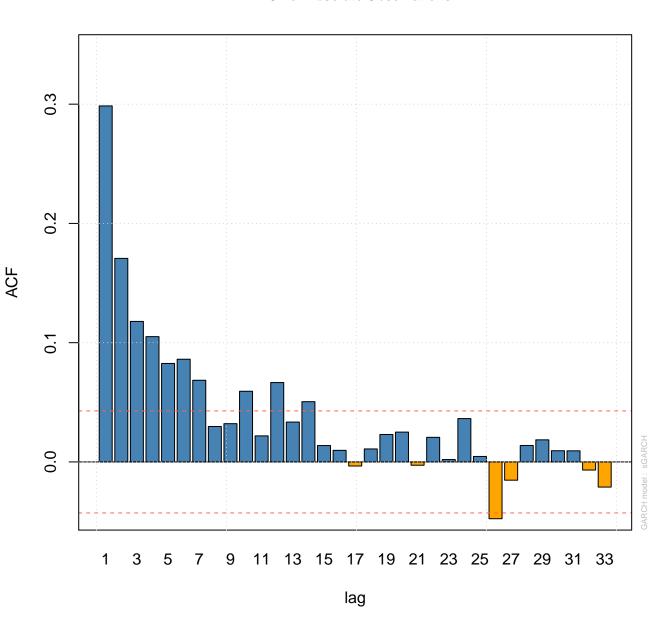
ACF of Observations



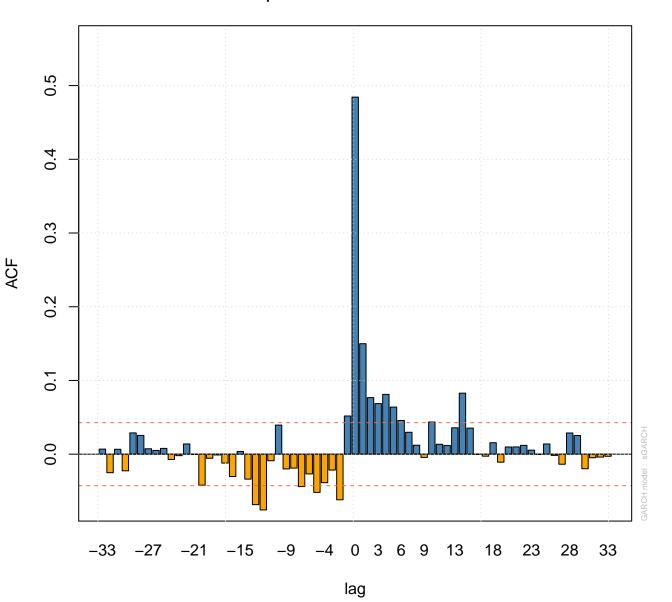
ACF of Squared Observations



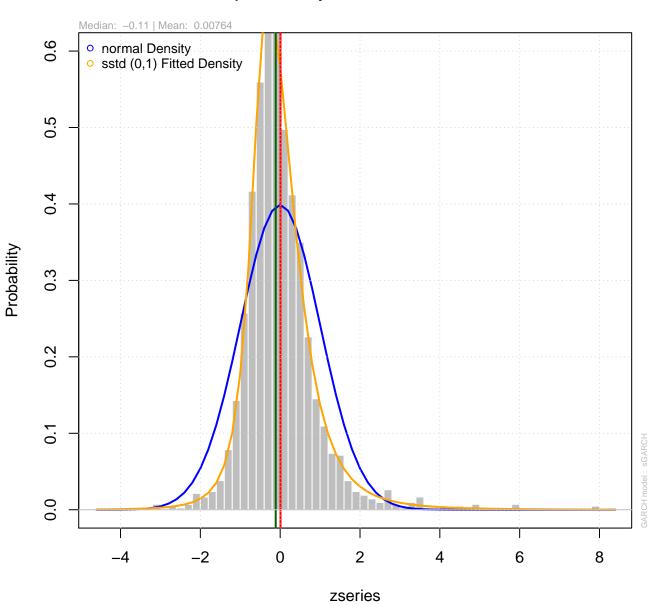
ACF of Absolute Observations



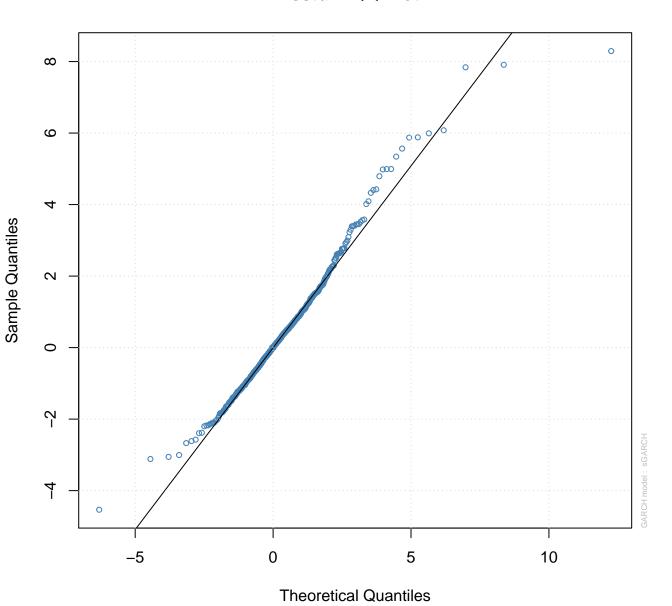
Cross-Correlations of Squared vs Actual Observations



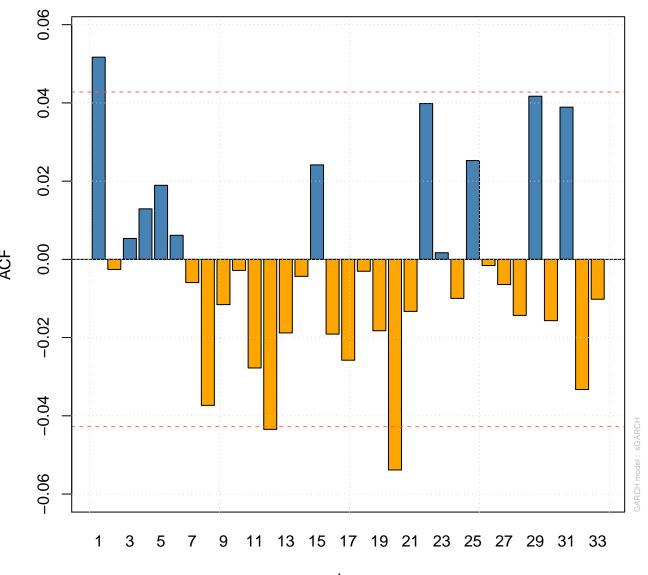
Empirical Density of Standardized Residuals



sstd - QQ Plot

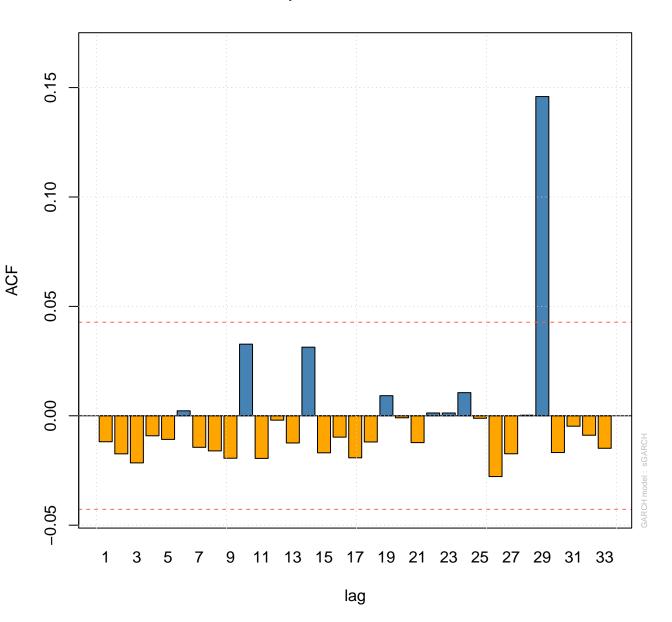


ACF of Standardized Residuals

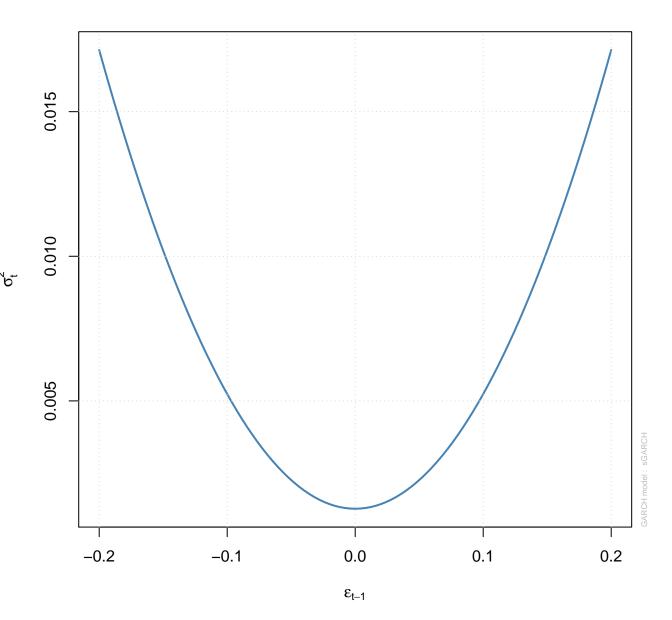


lag

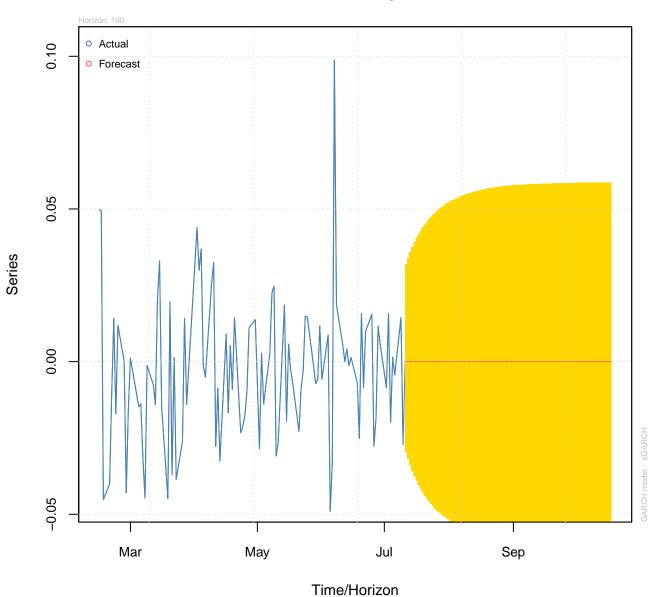
ACF of Squared Standardized Residuals



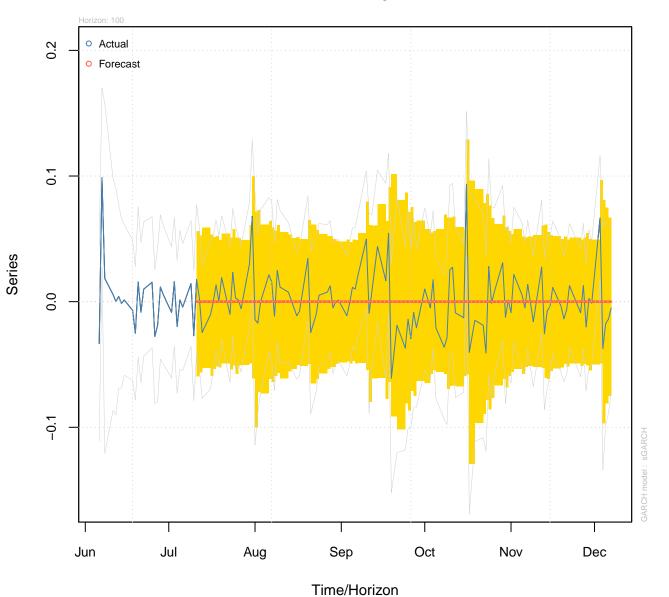
News Impact Curve



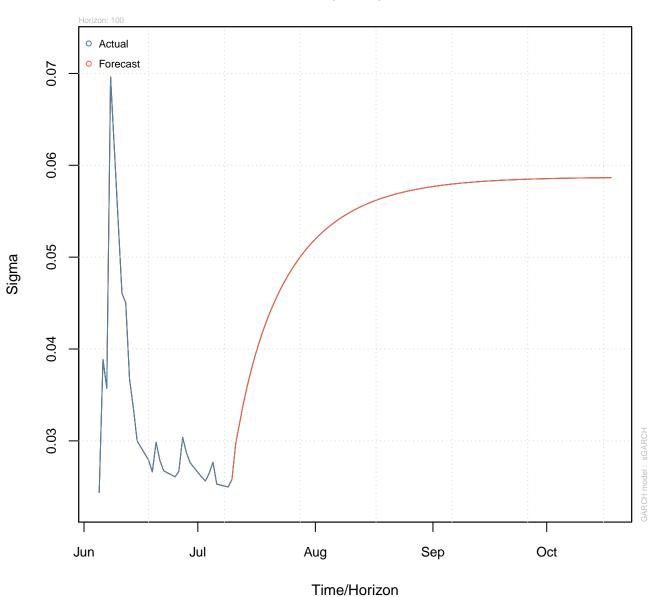
Forecast Series w/th unconditional 1–Sigma bands



Rolling Forecast vs Actual Series w/th conditional 2-Sigma bands



Forecast Unconditional Sigma (n.roll = 0)



Forecast Rolling Sigma vs |Series|

