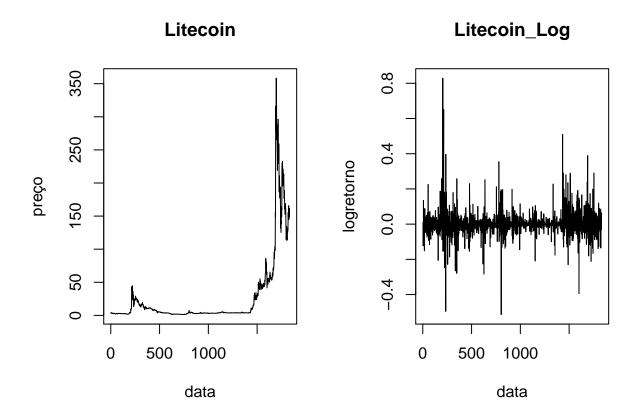
Criptomoeda

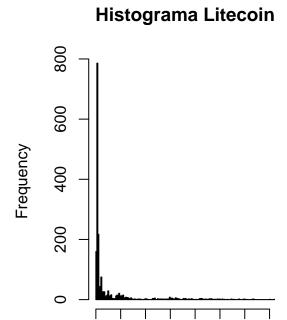
Daniel Ryba Zanardini de Oliveira

```
# inserindo os dados:
library(readxl)
litecoin <- read_excel("C:\\Users\\Daniel\\Documents\\MESTRADO\\II SEMESTRE\\FINANÇAS\\EXERCÍCIOS\\TRAB
Visualização das séries
# transformando em uma série de tempo:
library(timeSeries)
L <- timeSeries(litecoin)
L \leftarrow rev(L)
tail (L)
##
##
        Date
                     Open
                            High
                                  Low
                                           Close Volume
                                                               Market Cap
## [1,] Apr 28, 2018 145.74 153.33 145.74 152.30 337,117,000 8,202,220,000
## [2,] Apr 29, 2018 152.17 155.62 148.90 153.63 388,082,000 8,566,100,000
## [3,] Apr 30, 2018 153.65 154.08 147.87 148.48 341,397,000 8,651,810,000
## [4,] May 01, 2018 148.34 148.55 143.94 148.49 342,152,000 8,355,630,000
## [5,] May 02, 2018 148.45 152.89 146.98 151.58 354,657,000 8,363,530,000
## [6,] May 03, 2018 151.86 164.00 151.01 161.46 617,984,000 8,557,950,000
# para avaliar o logretorno, trabalharemos com o preço de fechamento:
preço <- as.numeric(L[,5])</pre>
# calculando o log retorno:
L.ret <- returns(preço)</pre>
head(L.ret)
##
                 [,1]
## [1,]
                  NA
## [2,] 0.006872879
## [3,] -0.018433702
## [4,] -0.123613956
## [5,] -0.120088322
## [6,] -0.103055229
# visualização gráfica:
par(mfrow=c(1,2))
```

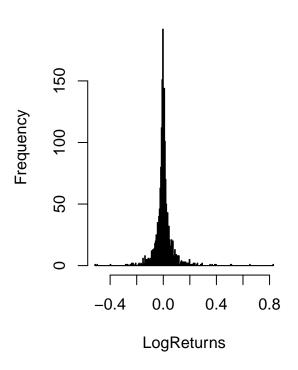
plot(preço, main="Litecoin", type="l", ylab="preço", xlab="data")

plot(L.ret, main="Litecoin_Log", type="l", ylab="logretorno", xlab="data")





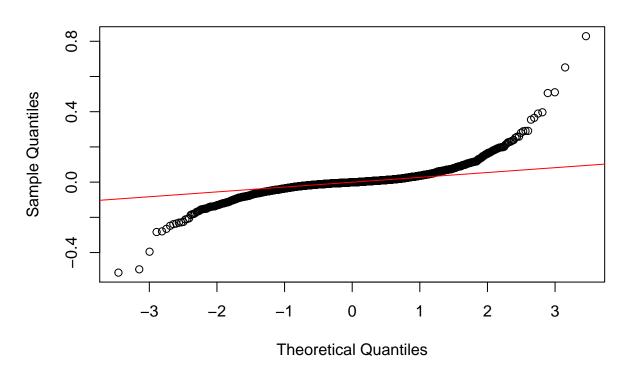
Histograma Litecoin



#qqplot
qqnorm(L.ret); qqline(L.ret, col="red")

Preços

Normal Q-Q Plot



```
\#\#Estatísticas das séries
```

```
# tabela com as principais estaísticas das séries de preços e retornos: média, min, max, desvio, variân
library(tseries)
library(fBasics)
options(digits=4)
#série de preços:
StatsPreço=as.matrix(basicStats(preço), 2,1)
#série de log retornos:
L.ret.NA <- removeNA(L.ret) #removendo o NA
StatsRet=as.matrix(basicStats(L.ret.NA), 2,1)
STATS <- cbind(StatsRet, StatsPreço)</pre>
STATS
```

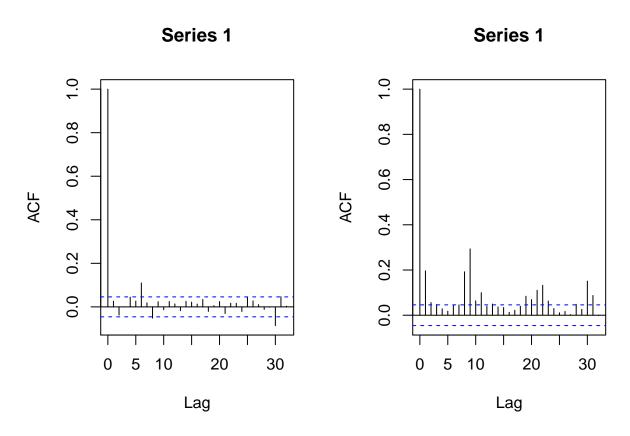
##		L.ret.NA	preço
##	nobs	1.831e+03	1832.000
##	NAs	0.000e+00	0.000
##	Minimum	-5.139e-01	1.160
##	Maximum	8.290e-01	358.340
##	1. Quartile	-1.904e-02	3.147
##	3. Quartile	1.809e-02	16.277
##	Mean	1.974e-03	25.924
##	Median	0.000e+00	3.960

```
## Sum
               3.614e+00 47493.390
              1.619e-03
## SE Mean
                            1.258
## LCL Mean -1.201e-03
                            23.458
## UCL Mean
              5.148e-03
                            28.391
               4.797e-03 2897.497
## Variance
## Stdev
               6.926e-02
                          53.828
## Skewness
              1.774e+00
                            3.158
## Kurtosis
               2.495e+01
                            10.171
# estacionarieade série de preços
adf.test(preço)
##
  Augmented Dickey-Fuller Test
##
## data: preço
## Dickey-Fuller = -2.3, Lag order = 12, p-value = 0.4
## alternative hypothesis: stationary
kpss.test(preço)
## Warning in kpss.test(preço): p-value smaller than printed p-value
## KPSS Test for Level Stationarity
##
## data: preço
## KPSS Level = 7, Truncation lag parameter = 8, p-value = 0.01
# estacionarieade série de retorno
adf.test(L.ret.NA)
## Warning in adf.test(L.ret.NA): p-value smaller than printed p-value
## Augmented Dickey-Fuller Test
##
## data: L.ret.NA
## Dickey-Fuller = -11, Lag order = 12, p-value = 0.01
## alternative hypothesis: stationary
kpss.test(L.ret.NA, null="Trend")
## Warning in kpss.test(L.ret.NA, null = "Trend"): p-value greater than printed p-
## value
##
## KPSS Test for Trend Stationarity
##
## data: L.ret.NA
## KPSS Trend = 0.076, Truncation lag parameter = 8, p-value = 0.1
# teste de normalidade série de preços
jarque.bera.test(preço)
##
## Jarque Bera Test
##
## data: preço
```

```
## X-squared = 10968, df = 2, p-value <2e-16
# teste de normalidade série de retorno
jarque.bera.test(L.ret.NA)
##
   Jarque Bera Test
##
##
## data: L.ret.NA
## X-squared = 48565, df = 2, p-value <2e-16
# testes Box Pierce Ljung Box na série de retornos
Box.test(L.ret.NA, lag = 5)
## Box-Pierce test
##
## data: L.ret.NA
## X-squared = 8.8, df = 5, p-value = 0.1
Box.test(L.ret.NA, lag = 10)
##
## Box-Pierce test
##
## data: L.ret.NA
## X-squared = 38, df = 10, p-value = 4e-05
Box.test(L.ret.NA, lag = 30)
##
## Box-Pierce test
##
## data: L.ret.NA
## X-squared = 69, df = 30, p-value = 7e-05
# testes Box Pierce Ljung Box na série do quadrado dos retornos
Box.test(L.ret.NA^2, lag = 5)
##
## Box-Pierce test
##
## data: L.ret.NA^2
## X-squared = 82, df = 5, p-value = 2e-16
Box.test(L.ret.NA^2, lag = 10)
##
## Box-Pierce test
##
## data: L.ret.NA^2
## X-squared = 323, df = 10, p-value <2e-16
Box.test(L.ret.NA^2, lag = 30)
##
## Box-Pierce test
##
## data: L.ret.NA^2
```

```
## X-squared = 490, df = 30, p-value <2e-16
# teste ARCH-LM na série de retorno
library(FinTS)
ArchTest(L.ret.NA)

##
## ARCH LM-test; Null hypothesis: no ARCH effects
##
## data: L.ret.NA
## Chi-squared = 243, df = 12, p-value <2e-16
# ACF
par(mfrow=c(1,2))
acf(L.ret.NA)
acf(L.ret.NA^2)</pre>
```



```
## Ajustando o modelor ARIMA
```

```
# ajustando o melhor modelo "familia GARCH"
library(forecast)
library(rugarch)
library(fGarch)
library(rmgarch)
library(quantmod)

# primeiro fazeos um FIT para ter uma idéia
fit <- auto.arima(L.ret.NA, seasonal=FALSE, stepwise=FALSE, approximation=FALSE)</pre>
```

```
summary(fit)
## Series: L.ret.NA
## ARIMA(2,0,3) with zero mean
## Coefficients:
##
         ar1 ar2 ma1 ma2
                                   ma3
       1.475 -0.854 -1.456 0.789 0.075
##
## s.e. 0.050 0.046 0.055 0.064 0.026
## sigma^2 estimated as 0.00474: log likelihood=2303
## AIC=-4594 AICc=-4594 BIC=-4561
##
## Training set error measures:
##
                  ME
                               MAE MPE MAPE
                                            MASE
                                                     ACF1
                        RMSE
## Training set 0.001839 0.06878 0.03807 NaN Inf 0.6776 -0.001263
Ajustando o modelo GARCH
# modelos GARCH
garch_model <- ugarchspec(mean.model = list(armaOrder=c(2,3)),</pre>
                       variance.model = list(garchOrder=c(1,1), model = "sGARCH"),
                       distribution.model = "sstd")
garch_model_1 <- ugarchspec(mean.model = list(armaOrder=c(1,1)),</pre>
                       variance.model = list(garchOrder=c(1,1), model = "sGARCH"),
                       distribution.model = "std")
garch <- ugarchfit(data = L.ret.NA, spec = garch model)</pre>
print(garch)
## *----*
          GARCH Model Fit
## *----*
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(1,1)
## Mean Model : ARFIMA(2,0,3)
## Distribution : sstd
##
## Optimal Parameters
## -----
        Estimate Std. Error t value Pr(>|t|)
        -0.000112 0.000567 -0.19695 0.843870
## mu
        1.493662 0.010543 141.66717 0.000000
## ar1
## ar2
        -0.929453 0.007649 -121.52063 0.000000
      -1.587060 0.009860 -160.95606 0.000000
## ma1
        ## ma2
## ma3
## omega 0.000018 0.000008 2.27105 0.023144
## alpha1 0.110106 0.015566 7.07350 0.000000
## beta1 0.888894 0.015734 56.49401 0.000000
```

```
1.034550 0.025977 39.82616 0.000000
## skew
## shape 2.791510 0.114966 24.28124 0.000000
##
## Robust Standard Errors:
        Estimate Std. Error t value Pr(>|t|)
## mu
        -0.000112 0.000535 -0.20871 0.83467
## ar1
        1.493662 0.007830 190.75218 0.00000
## ma3 -0.063834 0.000622 -102.63230 0.00000
## omega 0.000018 0.000015 1.23280 0.21765
## alpha1 0.110106 0.021476 5.12690 0.00000
## beta1 0.888894 0.028964 30.68949 0.00000
## skew 1.034550 0.024750 41.79958 0.00000
## shape 2.791510 0.114771 24.32238 0.00000
##
## LogLikelihood : 3255
##
## Information Criteria
## -----
## Akaike -3.5434
## Bayes -3.5103
## Shibata -3.5435
## Hannan-Quinn -3.5312
## Weighted Ljung-Box Test on Standardized Residuals
                        statistic p-value
##
## Lag[1]
                          6.623 0.010068
## Lag[2*(p+q)+(p+q)-1][14] 14.449 0.000000
## Lag[4*(p+q)+(p+q)-1][24] 20.097 0.006191
## d.o.f=5
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
                      statistic p-value
##
## Lag[1]
                       0.07444 0.7850
## Lag[2*(p+q)+(p+q)-1][5] 0.31551 0.9821
## Lag[4*(p+q)+(p+q)-1][9] 0.47468 0.9987
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
             Statistic Shape Scale P-Value
## ARCH Lag[3] 0.1247 0.500 2.000 0.7240
## ARCH Lag[5] 0.3142 1.440 1.667 0.9365
             0.3800 2.315 1.543 0.9879
## ARCH Lag[7]
##
## Nyblom stability test
## -----
## Joint Statistic: 10.76
```

```
## Individual Statistics:
## mu
       0.45477
## ar1
       0.02664
      0.04446
## ar2
## ma1
       0.01984
## ma2
      0.08710
## ma3
      0.23755
## omega 0.57550
## alpha1 1.05199
## beta1 0.76915
## skew 0.09177
## shape 0.91898
## 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
                  1.2492 0.2118
## Negative Sign Bias 0.1391 0.8894
## Positive Sign Bias 0.8991 0.3687
## Joint Effect
              2.2038 0.5312
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 44.51 0.0008022
## 2
      30 48.02
                    0.0146195
## 3 40 66.21 0.0042043
## 4
    50 70.94 0.0218557
##
## Elapsed time : 0.9485
garch_1 <- ugarchfit(data = L.ret.NA, spec = garch_model_1)</pre>
print(garch_1)
       GARCH Model Fit *
##
## Conditional Variance Dynamics
## -----
## GARCH Model : sGARCH(1,1)
## Mean Model : ARFIMA(1,0,1)
## Distribution : std
## Optimal Parameters
        Estimate Std. Error t value Pr(>|t|)
## mu -0.000598 0.000426 -1.4035 0.160465
```

```
0.417623 0.109821 3.8027 0.000143
-0.512906 0.102203 -5.0185 0.000001
## ar1
## ma1
## omega 0.000019 0.000009 2.1820 0.029109
## alpha1 0.110623 0.016192 6.8320 0.000000
## beta1 0.888377 0.016957 52.3901 0.000000
## shape 2.779880 0.114359 24.3084 0.000000
## Robust Standard Errors:
##
        Estimate Std. Error t value Pr(>|t|)
## mu
        ## ar1
        0.417623 0.085329 4.8942 0.000001
## ma1 -0.512906 0.079305 -6.4675 0.000000
## omega 0.000019 0.000017 1.1264 0.260008
## alpha1 0.110623 0.024222 4.5671 0.000005
## beta1 0.888377 0.033231 26.7336 0.000000
         ## shape
##
## LogLikelihood : 3248
##
## Information Criteria
## -----
## Akaike -3.5399
## Bayes -3.5189
## Shibata -3.5400
## Hannan-Quinn -3.5322
## Weighted Ljung-Box Test on Standardized Residuals
                      statistic p-value
##
## Lag[1]
                         8.618 3.329e-03
## Lag[2*(p+q)+(p+q)-1][5] 12.127 0.000e+00
## Lag[4*(p+q)+(p+q)-1][9] 17.745 1.325e-06
## d.o.f=2
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                      statistic p-value
## Lag[1]
                       0.06936 0.7923
## Lag[2*(p+q)+(p+q)-1][5] 0.29922 0.9838
## Lag[4*(p+q)+(p+q)-1][9] 0.45860 0.9988
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
             Statistic Shape Scale P-Value
## ARCH Lag[3] 0.1244 0.500 2.000 0.7243
## ARCH Lag[5] 0.3084 1.440 1.667 0.9380
             0.3760 2.315 1.543 0.9882
## ARCH Lag[7]
##
## Nyblom stability test
## -----
## Joint Statistic: 10.15
```

```
## Individual Statistics:
## m11
       0.6220
       0.1787
## ar1
## ma1
      0.1499
## omega 0.5854
## alpha1 1.1371
## beta1 0.8363
## shape 0.9461
##
## 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
                  1.1221 0.2620
## Sign Bias
## Negative Sign Bias 0.1417 0.8873
## Positive Sign Bias 0.7739 0.4391
## Joint Effect
                   1.7904 0.6170
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 42.74 0.0014062
## 2
      30
           52.64 0.0046179
    40 73.12 0.0007608
## 3
## 4 50 78.48 0.0047431
##
##
## Elapsed time : 0.3381
```

Exponencial GARCH

```
## GARCH Model : eGARCH(1,1)
## Mean Model : ARFIMA(2,0,3)
## Distribution : sstd
##
## Optimal Parameters
## -----
        Estimate Std. Error t value Pr(>|t|)
        1.093229 0.052514 20.81800 0.000000
## ar1
        ## ar2
## ma1
      -1.190092 0.055136 -21.58470 0.000000
## ma2
       0.646880 0.055009 11.75952 0.000000
## ma3 -0.022182 0.009438 -2.35026 0.018760
## omega -0.050616 0.012757 -3.96778 0.000073
## alpha1 0.021231 0.030537 0.69526 0.486895
## beta1
         0.989755 0.001749 565.92523 0.000000
## gamma1 0.477235 0.144280 3.30769 0.000941
## skew
         ## shape
         2.133296 0.083319 25.60391 0.000000
## Robust Standard Errors:
##
       Estimate Std. Error t value Pr(>|t|)
        1.093229 0.015147 72.17511 0.000000
## ar1
      ## ar2
      -1.190092 0.021365 -55.70217 0.000000
## ma1
## ma2
       0.646880 0.025863 25.01211 0.000000
## ma3
        -0.022182 0.003112 -7.12724 0.000000
## omega -0.050616 0.011039 -4.58536 0.000005
## alpha1 0.021231 0.035074 0.60533 0.544962
        0.989755 0.000927 1067.22699 0.000000
## beta1
## gamma1 0.477235 0.153064 3.11787 0.001822
                 0.015416 67.34845 0.000000
## skew
         1.038213
## shape
         2.133296
                 0.089188 23.91911 0.000000
##
## LogLikelihood : 3276
## Information Criteria
## -----
##
## Akaike
            -3.5665
## Bayes
            -3.5334
## Shibata
            -3.5666
## Hannan-Quinn -3.5543
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                         9.945 0.0016127
## Lag[2*(p+q)+(p+q)-1][14]
                       17.765 0.0000000
## Lag[4*(p+q)+(p+q)-1][24]
                       22.843 0.0006195
## d.o.f=5
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
```

```
##
                       statistic p-value
## Lag[1]
                       0.05165 0.8202
## Lag[2*(p+q)+(p+q)-1][5] 0.34069 0.9793
## Lag[4*(p+q)+(p+q)-1][9] 0.56118 0.9978
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
  Statistic Shape Scale P-Value
## ARCH Lag[3] 0.1586 0.500 2.000 0.6905
## ARCH Lag[5] 0.3912 1.440 1.667 0.9152
## ARCH Lag[7] 0.5003 2.315 1.543 0.9784
## Nyblom stability test
## -----
## Joint Statistic: 2.822
## Individual Statistics:
## ar1
       0.03724
## ar2
      0.10641
      0.03997
## ma1
## ma2
      0.07425
## ma3
      0.05495
## omega 0.49198
## alpha1 0.44894
## beta1 0.46660
## gamma1 0.17413
## skew 0.02746
## shape 0.29907
## 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
           1.40751 0.1594
## Sign Bias
## Negative Sign Bias 0.07262 0.9421
## Positive Sign Bias 0.94794 0.3433
## Joint Effect 2.61246 0.4553
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 22.03
                     0.2827
## 2
      30.03
                      0.4124
## 3 40 34.58
                     0.6716
## 4 50 45.76
                     0.6052
##
## Elapsed time : 2.235
```

```
egarch_1 <- ugarchfit(data = L.ret.NA, spec = egarch_model_1)</pre>
print(egarch_1)
##
           GARCH Model Fit
## *----*
##
## Conditional Variance Dynamics
## -----
## GARCH Model : eGARCH(1,1)
## Mean Model : ARFIMA(1,0,1)
## Distribution : std
##
## Optimal Parameters
## -----
##
         Estimate Std. Error t value Pr(>|t|)
        ## ar1
## ma1 -0.511154 0.044402 -11.51198 0.000000
## omega -0.049404 0.013878 -3.55976 0.000371
## alpha1 0.024709 0.031375 0.78754 0.430968
## beta1 0.989647 0.001771 558.78039 0.000000
## gamma1 0.501016 0.166799 3.00371 0.002667
## shape
         2.122821 0.084405 25.15041 0.000000
##
## Robust Standard Errors:
    Estimate Std. Error t value Pr(>|t|)
## ar1
        -0.511154 0.019834 -25.7716 0.000000
## ma1
## omega -0.049404 0.012059 -4.0969 0.000042
## alpha1 0.024709 0.035862 0.6890 0.490822
## beta1
         0.989647 0.000946 1046.1061 0.000000
## gamma1 0.501016 0.167521 2.9908 0.002783
## shape
         2.122821 0.084519 25.1164 0.000000
## LogLikelihood : 3271
##
## Information Criteria
##
## Akaike
             -3.5655
## Bayes
             -3.5444
            -3.5655
## Shibata
## Hannan-Quinn -3.5577
##
## Weighted Ljung-Box Test on Standardized Residuals
##
                        statistic
                                  p-value
## Lag[1]
                         11.22 8.076e-04
## Lag[2*(p+q)+(p+q)-1][5] 14.12 0.000e+00
## Lag[4*(p+q)+(p+q)-1][9] 18.47 5.318e-07
## d.o.f=2
## HO : No serial correlation
##
```

```
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                      statistic p-value
                        0.04778 0.8270
## Lag[1]
## Lag[2*(p+q)+(p+q)-1][5] 0.32993 0.9805
## Lag[4*(p+q)+(p+q)-1][9] 0.54598 0.9980
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
            Statistic Shape Scale P-Value
## ARCH Lag[3] 0.1580 0.500 2.000 0.6910
             0.3792 1.440 1.667 0.9186
## ARCH Lag[5]
## ARCH Lag[7] 0.4877 2.315 1.543 0.9795
## Nyblom stability test
## Joint Statistic: 2.237
## Individual Statistics:
## ar1
      0.1651
## ma1
      0.1429
## omega 0.4991
## alpha1 0.4149
## beta1 0.4704
## gamma1 0.1772
## shape 0.2939
## 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
                  1.31103 0.1900
## Negative Sign Bias 0.08813 0.9298
## Positive Sign Bias 0.90581 0.3652
## Joint Effect 2.30012 0.5125
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1 20 23.67 0.2091
## 2 30 28.10
                     0.5126
                    0.6869
## 3 40 34.23
## 4 50 46.47
                      0.5762
##
## Elapsed time: 0.8089
```

APARCH

skew

1.036420

0.015650

```
# modelos APARCH
aparch_model <- ugarchspec(mean.model = list(armaOrder=c(2,3), include.mean=FALSE),</pre>
                        variance.model = list(garchOrder=c(1,1), model = "apARCH"),
                        distribution.model = "sstd")
aparch_model_1 <- ugarchspec(mean.model = list(armaOrder=c(1,1), include.mean=FALSE),</pre>
                        variance.model = list(garchOrder=c(1,1), model = "apARCH"),
                        distribution.model = "std")
aparch <- ugarchfit(data = L.ret.NA, spec = aparch_model)</pre>
print(aparch)
##
## *----*
           GARCH Model Fit
## *----*
##
## Conditional Variance Dynamics
## -----
## GARCH Model : apARCH(1,1)
## Mean Model : ARFIMA(2,0,3)
## Distribution : sstd
##
## Optimal Parameters
## -----
         Estimate Std. Error t value Pr(>|t|)
                  0.010041 148.4021 0.000000
## ar1
         1.490045
        ## ar2
## ma1
        -1.587411 0.009702 -163.6122 0.000000
         1.050190 0.000903 1162.8049 0.000000
## ma2
        -0.068835 0.000471 -146.1374 0.000000
## ma3
## omega 0.000810 0.000625 1.2946 0.195443
## alpha1 0.369318 0.079327
                             4.6556 0.000003
                  0.017085 50.9396 0.000000
## beta1
         0.870284
                  0.070295
## gamma1 -0.086774
                             -1.2344 0.217046
## delta
                             6.0126 0.000000
         1.064638
                  0.177068
                    0.018746 55.2872 0.000000
## skew
         1.036420
         2.155017
                    0.055971
                             38.5023 0.000000
## shape
##
## Robust Standard Errors:
##
         Estimate Std. Error
                             t value Pr(>|t|)
         1.490045
                  0.006674 223.2668 0.000000
## ar1
## ar2
        -0.925236
                    0.004219 -219.3119 0.000000
## ma1
        -1.587411 0.006180 -256.8725 0.000000
         1.050190 0.000532 1975.6355 0.000000
## ma2
                  0.000493 -139.7105 0.000000
## ma3
        -0.068835
        0.000810 0.000787
                              1.0289 0.303549
## omega
## alpha1 0.369318
                 0.062446
                             5.9142 0.000000
                  0.021934
## beta1
         0.870284
                              39.6779 0.000000
## gamma1 -0.086774
                    0.078401
                              -1.1068 0.268381
## delta
         1.064638
                    0.236897
                             4.4941 0.000007
```

66.2265 0.000000

```
## shape 2.155017 0.044922 47.9723 0.000000
##
## LogLikelihood : 3281
##
## Information Criteria
## -----
## Akaike
            -3.5703
## Bayes -3.5342
## Shibata -3.5704
## Hannan-Quinn -3.5570
## Weighted Ljung-Box Test on Standardized Residuals
## -----
##
                       statistic p-value
## Lag[1]
                          9.921 0.001634
## Lag[2*(p+q)+(p+q)-1][14] 17.280 0.000000
## Lag[4*(p+q)+(p+q)-1][24] 21.551 0.001900
## d.o.f=5
## HO : No serial correlation
##
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                       statistic p-value
## Lag[1]
                       0.04334 0.8351
## Lag[2*(p+q)+(p+q)-1][5] 0.30455 0.9832
## Lag[4*(p+q)+(p+q)-1][9] 0.46369 0.9988
## d.o.f=2
##
## Weighted ARCH LM Tests
## -----
   Statistic Shape Scale P-Value
##
## ARCH Lag[3] 0.1194 0.500 2.000 0.7297
## ARCH Lag[5] 0.3273 1.440 1.667 0.9329
## ARCH Lag[7] 0.3872 2.315 1.543 0.9874
##
## Nyblom stability test
## -----
## Joint Statistic: 3.413
## Individual Statistics:
## ar1
      0.03886
## ar2 0.07093
      0.02641
## ma1
## ma2 0.14153
      0.35497
## ma3
## omega 0.35903
## alpha1 0.30880
## beta1 0.35009
## gamma1 0.83875
## delta 0.36986
## skew 0.03924
## shape 0.39200
##
## Asymptotic Critical Values (10% 5% 1%)
```

```
## Joint Statistic: 2.69 2.96 3.51 ## Individual Statistic: 0.35 0.47 0.75
## Sign Bias Test
## -----
##
               t-value prob sig
## Sign Bias
            1.2350 0.2170
## Negative Sign Bias 0.3101 0.7565
## Positive Sign Bias 1.1607 0.2459
## Joint Effect 2.7027 0.4398
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1
    20
          17.88
                  0.5304
## 2
     30 28.33
                    0.5004
## 3 40 33.18
                   0.7317
## 4 50 42.59
                   0.7289
##
##
## Elapsed time : 38.45
aparch_1 <- ugarchfit(data = L.ret.NA, spec = aparch_model_1)</pre>
print(aparch_1)
##
## *----*
    GARCH Model Fit *
## *----*
##
## Conditional Variance Dynamics
## -----
## GARCH Model : apARCH(1,1)
## Mean Model : ARFIMA(1,0,1)
## Distribution : std
##
## Optimal Parameters
## -----
       Estimate Std. Error t value Pr(>|t|)
## ar1
       ## ma1
     -0.519937 0.031749 -16.3766 0.00000
## omega 0.001214 0.000814 1.4918 0.13576
## alpha1 0.461968 0.077530 5.9586 0.00000
       0.864629 0.018410 46.9644 0.00000
## beta1
## delta 1.033309 0.163287 6.3282 0.00000
        2.100000 0.005459 384.6814 0.00000
## shape
## Robust Standard Errors:
      Estimate Std. Error t value Pr(>|t|)
       ## ar1
## ma1 -0.519937 0.011978 -43.4064 0.000000
## omega 0.001214 0.000937 1.2961 0.194951
## alpha1 0.461968 0.092072 5.0175 0.000001
```

```
## beta1 0.864629 0.024993 34.5955 0.000000
## gamma1 -0.080628 0.075025 -1.0747 0.282518
## delta 1.033309 0.187213 5.5194 0.000000
## shape 2.100000 0.000781 2687.5415 0.000000
## LogLikelihood : 3274
## Information Criteria
## -----
##
## Akaike
             -3.5677
## Bayes
             -3.5436
## Shibata -3.5678
## Hannan-Quinn -3.5588
## Weighted Ljung-Box Test on Standardized Residuals
##
                       statistic p-value
## Lag[1]
                          12.17 4.863e-04
## Lag[2*(p+q)+(p+q)-1][5] 15.46 0.000e+00
## Lag[4*(p+q)+(p+q)-1][9] 20.60 3.423e-08
## d.o.f=2
## HO : No serial correlation
## Weighted Ljung-Box Test on Standardized Squared Residuals
## -----
##
                        statistic p-value
## Lag[1]
                        0.03741 0.8466
## Lag[2*(p+q)+(p+q)-1][5] 0.30139 0.9836
## Lag[4*(p+q)+(p+q)-1][9] 0.47158 0.9987
## d.o.f=2
##
## Weighted ARCH LM Tests
            Statistic Shape Scale P-Value
## ARCH Lag[3] 0.1264 0.500 2.000 0.7222
## ARCH Lag[5] 0.3432 1.440 1.667 0.9286
## ARCH Lag[7] 0.4099 2.315 1.543 0.9858
##
## Nyblom stability test
## -----
## Joint Statistic: 2.903
## Individual Statistics:
## ar1 0.09294
      0.08483
## ma1
## omega 0.38196
## alpha1 0.31939
## beta1 0.34788
## gamma1 0.80636
## delta 0.28197
## shape 0.38697
## 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
                 1.2368 0.2163
## Negative Sign Bias 0.2973 0.7663
## Positive Sign Bias 1.1315 0.2580
## Joint Effect
             2.6342 0.4515
##
##
## Adjusted Pearson Goodness-of-Fit Test:
## -----
## group statistic p-value(g-1)
## 1
      20
           30.44
                    0.04644
## 2
           35.28
      30
                    0.19564
## 3
     40 56.12
                    0.03721
## 4
     50 51.33
                     0.38244
##
##
## Elapsed time : 15.06
```

Melhor modelo pelo critério AKAIKE ARMA (2,3) APARCH (1,1)

Avaliando o modelo

```
FR=residuals(aparch, standardize=TRUE)

# ACF
par(mfrow=c(1,2))
acf(FR)
acf(FR^2)
```

ACF 0.2 0.4 0.6 0.8 1.0

1000000

Lag

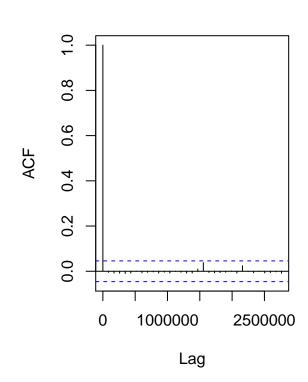
2500000

0.0

0

Series FR

Series FR²



```
# estacionáridade dos resíduos
adf.test(FR)
## Warning in adf.test(FR): p-value smaller than printed p-value
##
    Augmented Dickey-Fuller Test
##
##
## data: FR
## Dickey-Fuller = -11, Lag order = 12, p-value = 0.01
## alternative hypothesis: stationary
kpss.test(FR, null="Trend")
## Warning in kpss.test(FR, null = "Trend"): p-value greater than printed p-value
##
   KPSS Test for Trend Stationarity
##
##
## KPSS Trend = 0.078, Truncation lag parameter = 8, p-value = 0.1
# normalidade dos resíduos
jarque.bera.test(FR)
##
##
    Jarque Bera Test
##
## data: FR
```

```
## X-squared = 329372, df = 2, p-value <2e-16
#testes Box Pierce Ljung Box nos resíduos
Box.test(FR, lag = 5)
##
   Box-Pierce test
##
##
## data: FR
## X-squared = 12, df = 5, p-value = 0.03
Box.test(FR, lag = 10)
##
##
   Box-Pierce test
##
## data: FR
## X-squared = 21, df = 10, p-value = 0.02
Box.test(FR, lag = 50)
##
##
   Box-Pierce test
##
## data: FR
## X-squared = 53, df = 50, p-value = 0.3
# qq plot dos resíduos
qqnorm(FR); qqline(FR, col="red")
```

Normal Q-Q Plot

