Homework 4

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Homewor4 (Question1)

Indices in the CRIX family

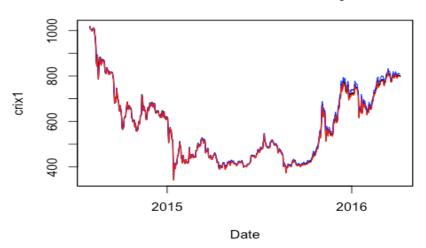


Figure 3: The daily value of indices in the CRIX family

The log return of CRIX index

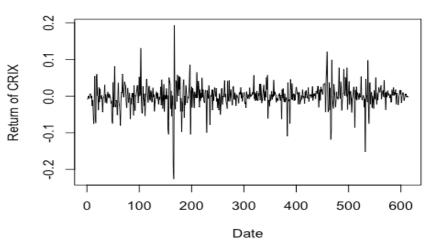


Figure 4: The log returns of CRIX index

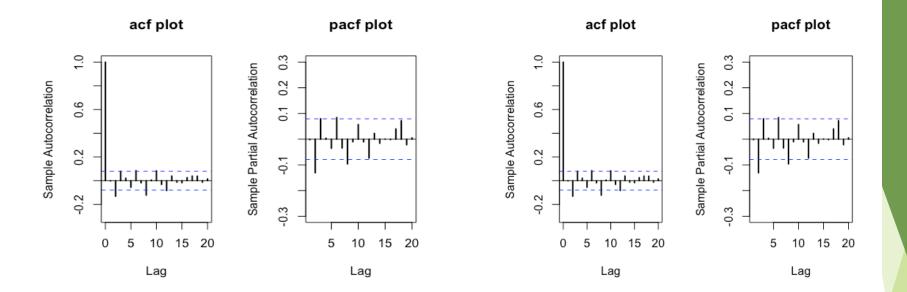


Figure 5: Histogram and QQ plot of CRIX returns

Figure 6: The sample ACF and PACF of CRIX returns

```
rm(list = Is(all = TRUE))
graphics.off()
# install and load packages
libraries = c("zoo", "tseries", "xts","ccgarch")
lapply(libraries, function(x) if (!(x %in% installed.packages())) { install.packages(x)}
```

```
lapply(libraries, library, quietly = TRUE, character.only = TRUE)
# load dataset
load(file.choose())
load(file.choose())
load(file.choose())
# three indices return
ecrix1 = zoo(ecrix, order.by = index(crix1))
efcrix1 = zoo(efcrix, order.by = index(crix1))
# plot with different x-axis scales with zoo
my.panel <- function(x, ...) {
 lines(x, ...)
 lines(ecrix1, col = "blue")
 lines(efcrix1, col = "red")
plot.zoo(crix1, plot.type = "multiple", type = "l", lwd = 1.5, panel = my.panel,
      main = "Indices in the CRIX family", xlab = "Date")
```

```
# plot of crix
# plot(as.xts(crix), type="l", auto.grid=FALSE, main = NA)
plot(crix1, ylab = "Price of CRIX", xlab = "Date")
# plot of crix return
ret = diff(log(crix1))
# plot(as.xts(ret), type="l", auto.grid=FALSE, main = NA)
plot(ret, ylab = "Return of CRIX", xlab = "Date")
# stationary test
adf.test(ret, alternative = "stationary")
kpss.test(ret, null = "Trend")
par(mfrow = c(1, 2))
# histogram of returns
hist(ret, col = "grey", breaks = 20, freq = FALSE, ylim = c(0, 25), xlab = "Return of CRIX")
lines(density(ret), lwd = 2)
mu = mean(ret)
sigma = sd(ret)
x = seq(-4, 4, length = 100)
curve(dnorm(x, mean = mean(ret), sd = sd(ret)), add = TRUE, col = "red",
    lwd = 2
```

Homewor4 (Question2)

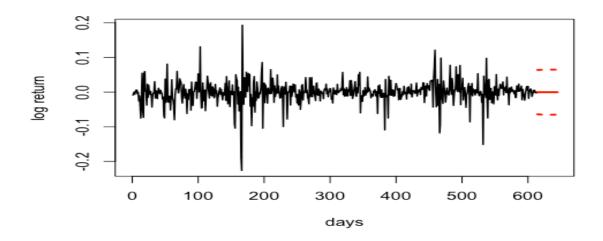


Figure 7: CRIX returns and predicted values.

```
# bic
Codes:
                                             bic = matrix(NA, 6, 6)
                                             for (p in 0:4) {
# arima model
par(mfrow = c(1, 1))
                                              for (q in 0:3) {
fit1 = arima(ret, order = c(1, 0, 1))
                                               b.p.q = arima(ret, order = c(p, 0, q))
                                               bic.p.q = AIC(b.p.q, k = log(length(ret)))
tsdiag(fit1)
Box.test(fit1$residuals, lag = 1)
                                               bic[p + 1, q + 1] = bic.p.q
# aic
aic = matrix(NA, 6, 6)
for (p in 0:4) {
                                             # select p and q order of ARIMA model
 for (q in 0:3) {
                                             fit4 = arima(ret, order = c(2, 0, 3))
  a.p.q = arima(ret, order = c(p, 0, q))
                                             tsdiag(fit4)
                                             Box.test(fit4$residuals, lag = 1)
  aic.p.q = a.p.q$aic
  aic[p + 1, q + 1] = aic.p.q
                                             fitr4 = arima(ret, order = c(2, 1, 3))
                                             tsdiag(fitr4)
                                             Box.test(fitr4$residuals, lag = 1)
```

Homewor4 (Question2)

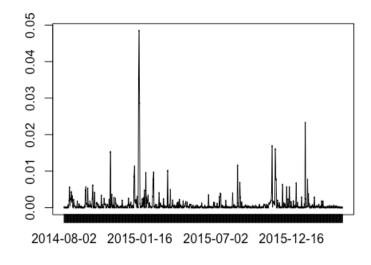


Figure 8: The squared ARIMA(2,0,2) residuals of CRIX returns.

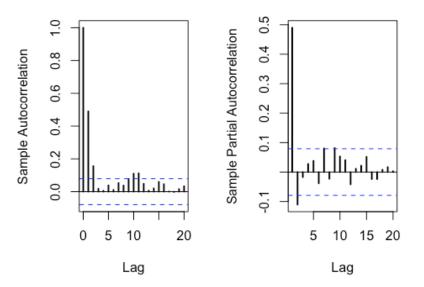


Figure 9: The ACF and PACF of squared ARIMA(2,0,2) residuals

```
Codes:
rm(list = ls(all = TRUE))
graphics.off()
# install and load packages
libraries = c("tseries")
lapply(libraries, function(x) if (!(x %in% installed.packages())) {
 install.packages(x)
lapply(libraries, library, quietly = TRUE, character.only = TRUE)
# please change your working directory
setwd()
load(file.choose())
Pr = as.numeric(crix)
Da = factor(date1)
crx = data.frame(Da, Pr)
# plot of crix return
ret = diff(log(crx$Pr))
Dare = factor(date1[-1])
retts = data.frame(Dare, ret)
# arima202 predict
fit202 = arima(ret, order = c(2, 0, 2))
```

```
# vola cluster
par(mfrow = c(1, 1))
res = fit202$residuals
res2 = fit202$residuals^2
tsres202 = data.frame(Dare,
res2)
plot(tsres202$Dare,
tsres202$res2, type = "o", ylab
= NA)
lines(tsres202$res2)
# plot(res2, ylab='Squared
residuals', main=NA)
par(mfrow = c(1, 2))
acfres2 = acf(res2, main = NA,
lag.max = 20, ylab = "Sample
```

Autocorrelation", lwd = 2)

= NA)

pacfres2 = pacf(res2, lag.max

Autocorrelation", lwd = 2, main

= 20, ylab = "Sample Partial

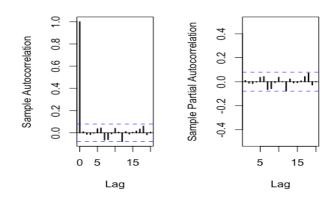


Figure 10: The ACF and PACF of squared ARIMA(2,0,2) residuals

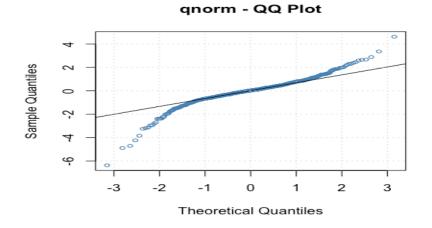


Figure 11: The QQ plots of model residuals of ARIMA-GARCH process.

```
Codes:
rm(list = ls(all = TRUE))
graphics.off()
# install and load packages
libraries = c("forecast", "fGarch")
lapply(libraries, function(x) if (!(x %in%
installed.packages())) {
 install.packages(x)
})
lapply(libraries, library, quietly = TRUE,
character.only = TRUE)
# load dataset
load(file.choose())
ret = diff(log(crix1))
# vol cluster
fit202 = arima(ret, order = c(2, 0, 2))
par(mfrow = c(1, 1))
res = fit202$residuals
res2 = fit202$residuals^2
```

```
# different garch model
fg11 = garchFit(data = res, data
~ garch(1, 1))
summary(fg11)
fg12 = garchFit(data = res, data
~ garch(1, 2))
summary(fg12)
fg21 = garchFit(data = res, data
~ garch(2, 1))
summary(fg21)
fg22 = garchFit(data = res, data
~ garch(2, 2))
summary(fg22)
# residual plot
reszo = zoo(fg11@residuals,
order.by = index(crix1))
plot(reszo, ylab = NA, lwd = 2)
```

```
par(mfrow = c(1, 2))
fg11res2 = fg11@residuals
acfres2 = acf(fg11res2, lag.max = 20, ylab = "Sample Autocorrelation",
         main = NA, Iwd = 2)
pacfres2 = pacf(fg11res2, lag.max = 20, ylab = "Sample Partial Autocorrelation",
         main = NA, lwd = 2, ylim = c(-0.5, 0.5))
fg12res2 = fg12@residuals
acfres2 = acf(fg12res2, lag.max = 20, ylab = "Sample Autocorrelation",
         main = NA, Iwd = 2)
pacfres2 = pacf(fg12res2, lag.max = 20, ylab = "Sample Partial Autocorrelation",
         main = NA, lwd = 2, ylim = c(-0.5, 0.5)
# qq plot
par(mfrow = c(1, 1))
plot(fg11, which = 13) #9,10,11,13
```

ACF of Squared ResiduPACF of Squared Residu

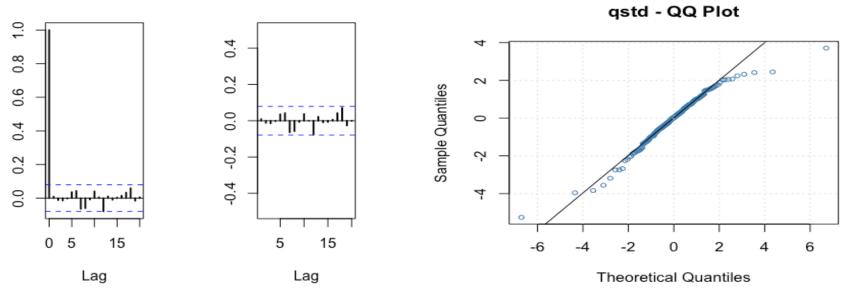


Figure 12: The ACF and PACF plots for model residuals of ARIMA(2,0,2)- t-GARCH(1,1) process.

Figure 13: The QQ plots of model residuals of ARIMA-t-GARCH process.

```
Codes:
fg11stu = garchFit(data = res, data ~ garch(1, 1), cond.dist = "std")
# different forecast with t-garch
# fg11stufore = predict(fg11stu, n.ahead = 30, plot=TRUE, mse='uncond', auto.grid=FALSE)
fg11stufore = predict(fg11stu, n.ahead = 30, plot = TRUE, cond.dist = "QMLE",
             auto.grid = FALSE)
par(mfrow = c(1, 2))
stu.fg11res2 = fg11stu@residuals
# acf and pacf for t-garch
stu.acfres2 = acf(stu.fg11res2, ylab = NA, lag.max = 20, main = "ACF of Squared Residuals",
           lwd = 2)
stu.pacfres2 = pacf(stu.fg11res2, lag.max = 20, main = "PACF of Squared Residuals",
            lwd = 2, ylab = NA, ylim = c(-0.5, 0.5)
# ARIMA-t-GARCH qq plot
par(mfrow = c(1, 1))
plot(fg11stu, which = 13)
```