GARCH parameters and quantiles estimation

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Input

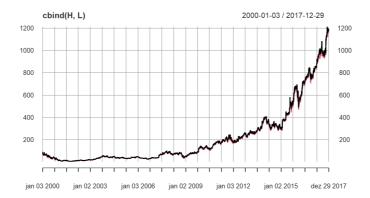
```
symbol = "AMZN"
from=as.Date('2000-01-01')
to=as.Date('2017-12-31')
C_Trend = 0.95
C_Reaction = 0.50
```

Data download

```
x <- getSymbols.yahoo(symbol,auto.assign = FALSE, from=from, to=to)
```

High and Low

```
H <- Hi(x)
L <- Lo(x)
C <- Cl(x)
plot(cbind(H,L))</pre>
```

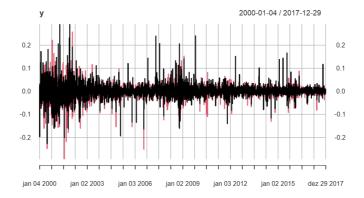


Returns

```
y <- cbind( diff(log(H)),  diff(log(L)) )
y <- na.omit(y)
y %>% cor() # Returns correlation
```

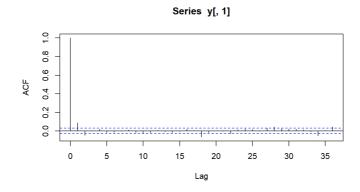
```
## AMZN.High AMZN.Low
## AMZN.High 1.000000 0.732942
## AMZN.Low 0.732942 1.000000
```

plot(y)

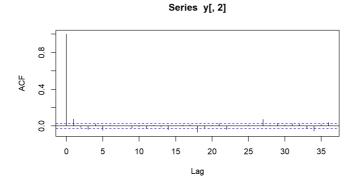


Autocorrelation

acf(y[,1])

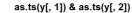


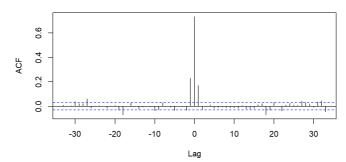
acf(y[,2])



Cross correlation

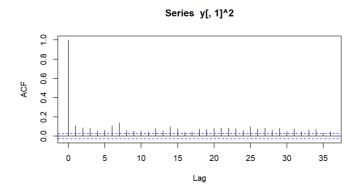
ccf(as.ts(y[,1]),as.ts(y[,2]))



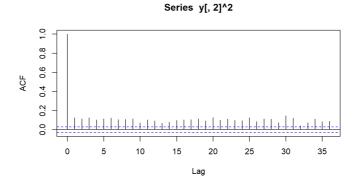


Volatility verification

```
acf(y[,1]^2)
```



acf(y[,2]^2)



Bivariate DCC-GARCH

We will consider the DCC-GARCH to model the volatility of $y=(r_H,r_L)'$, where r_H and r_L denote the $100\times$ log-returns from hight's and low's observations.

```
# returns
mY <- 100*y

# generates the Markov Chain
start <- Sys.time()

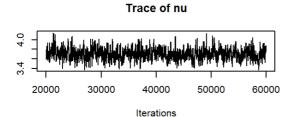
out <- bayesDccGarch(mY, control=list(print=FALSE, nPilotSim=3000))</pre>
```

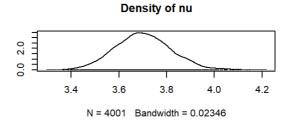
```
## Maximizing the log-posterior density function.
## Done.
## One approximation for covariance matrix of parameters cannot be directly computed through
the hessian matrix.
## Calibrating the standard deviations for simulation:
## Accept Rate:
## phi_1 phi_2 phi_3 phi_4 phi_5 phi_6 phi_7 phi_8 phi_9 phi_10 phi_11
##
   0.26
          0.09
                 0.21
                       0.05
                               0.08
                                     0.09
                                            0.18
                                                   0.05
                                                         0.05
                                                                 0.19
## Accept Rate:
## phi_1 phi_2 phi_3 phi_4 phi_5 phi_6 phi_7 phi_8 phi_9 phi_10 phi_11
   0.27
          0.17
                 0.22
                       0.08
                              0.10
                                     0.16
                                            0.19
                                                   0.05
                                                         0.07
                                                                 0.20
## Accept Rate:
## phi_1 phi_2 phi_3 phi_4 phi_5 phi_6 phi_7 phi_8 phi_9 phi_10 phi_11
   0.26
           0.16
                0.20 0.11 0.16
                                    0.16 0.20
                                                   0.09
                                                        0.14
                                                                 0.21
##
## Accept Rate:
## phi_1 phi_2 phi_3 phi_4 phi_5 phi_6 phi_7
                                                  phi_8 phi_9 phi_10 phi_11
  0.28
                                     0.18
                                            0.20
         0.16
                0.21
                       0.17
                               0.16
                                                   0.13
                                                         0.20
                                                                 0.21
## Accept Rate:
## phi_1 phi_2 phi_3 phi_4 phi_5 phi_6 phi_7
                                                  phi_8 phi_9 phi_10 phi_11
                 0.22
                               0.17
                                     0.14
                                            0.20
   0.27
          0.16
                       0.21
                                                   0.23
                                                         0.23
                                                                 0.20
## Accept Rate:
## phi_1 phi_2 phi_3 phi_4 phi_5 phi_6 phi_7 phi_8 phi_9 phi_10 phi_11
   0.25
          0.17
                  0.21
                       0.20
                              0.17
                                     0.26
                                            0.22
                                                   0.22
                                                         0.21
                                                                 0.18
## Computing the covariance matrix of pilot sample.
## Warning in if (class(control$cholCov) != "try-error") {: a condição tem
## comprimento > 1 e somente o primeiro elemento será usado
## Done.
## Calibrating the Lambda coefficient:
## lambda: 0.4
## Accept Rate: 0.26
## Done.
## Starting the simulation by one-block random walk Metropolis-Hasting algorithm.
## Done.
out2 <- increaseSim(out, nSim=50000)</pre>
## Calibrating the Lambda coefficient:
## lambda: 0.4
## Accept Rate: 0.26
## Done.
## Starting the simulation by one-block random walk Metropolis-Hasting algorithm.
## Done.
out <- window(out2, start=20000, thin=10)
rm(out2)
end <- Sys.time()</pre>
# elapsed time
end-start
```

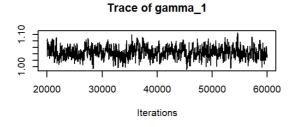
Time difference of 21.81299 mins

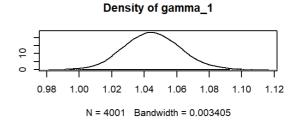
Estimative of parameters
parEst <- summary(out)\$statistics[,'Mean']</pre>

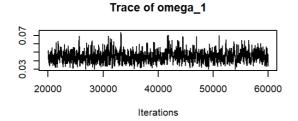
plot Markov Chain
plot(out\$MC)

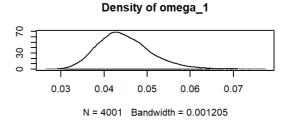


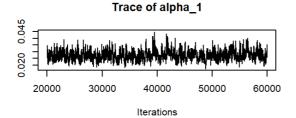


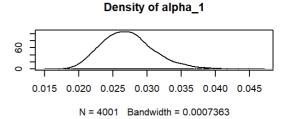


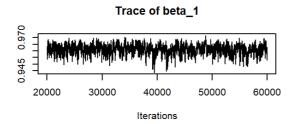


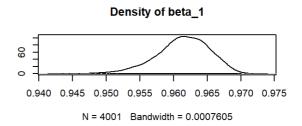


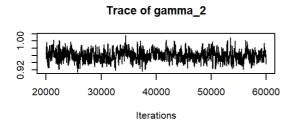


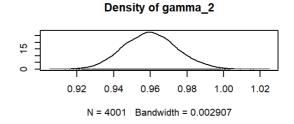


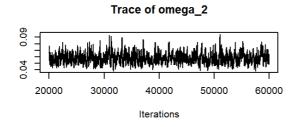


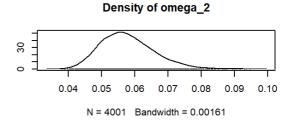


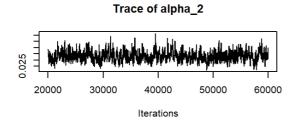


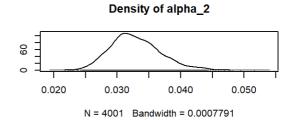


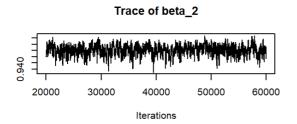


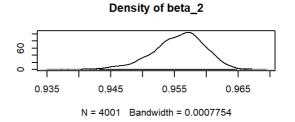


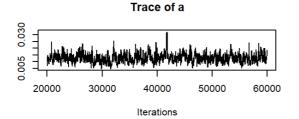


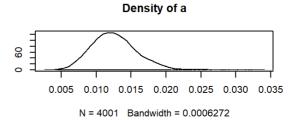


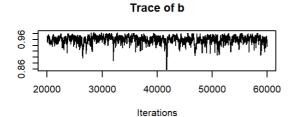


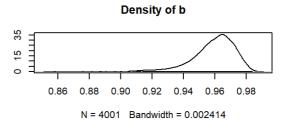












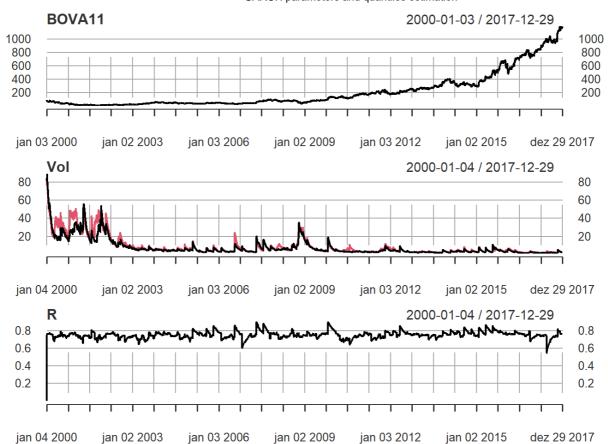
Estimative of parameters
out\$MC %>% summary()

```
##
## Iterations = 20000:60000
## Thinning interval = 10
## Number of chains = 1
## Sample size per chain = 4001
##
## 1. Empirical mean and standard deviation for each variable,
##
      plus standard error of the mean:
##
##
                         SD Naive SE Time-series SE
## nu
           3.70159 0.116252 1.838e-03
                                           0.0055738
## gamma 1 1.04483 0.016877 2.668e-04
                                           0.0009341
## omega_1 0.04475 0.006231 9.851e-05
                                         0.0002925
## alpha_1 0.02717 0.003732 5.901e-05
                                         0.0001918
## beta 1 0.96139 0.003841 6.073e-05
                                         0.0001858
## gamma_2 0.96002 0.014408 2.278e-04
                                          0.0007164
                                          0.0004043
## omega 2 0.05782 0.008086 1.278e-04
## alpha_2 0.03289 0.003945 6.236e-05
                                         0.0002168
## beta_2 0.95566 0.004030 6.372e-05
                                         0.0002118
          0.01261 0.003284 5.191e-05
                                          0.0001990
## b
          0.95930 0.013532 2.139e-04
                                           0.0007718
##
## 2. Quantiles for each variable:
##
##
                        25%
                                50%
               2.5%
                                        75%
                                              97.5%
## nu
           3.482750 3.62182 3.69779 3.77763 3.93316
## gamma 1 1.012682 1.03326 1.04460 1.05610 1.07901
## omega_1 0.034289 0.04039 0.04405 0.04839 0.05890
## alpha 1 0.020897 0.02450 0.02693 0.02939 0.03543
## beta 1 0.952966 0.95908 0.96161 0.96413 0.96787
## gamma 2 0.932804 0.94976 0.95980 0.96953 0.98960
## omega_2 0.044498 0.05200 0.05703 0.06269 0.07565
## alpha_2 0.026166 0.03016 0.03253 0.03534 0.04184
## beta_2 0.946535 0.95327 0.95606 0.95842 0.96255
## a
          0.007031 0.01037 0.01236 0.01454 0.01953
## b
           0.925185 0.95263 0.96167 0.96866 0.97895
## Conditional Correlation
```

```
## Conditional Correlation
R <- xts(out$R[,2], order.by=index(y))

## Volatility
Vol <- xts(out$H[,c("H_1,1","H_2,2")], order.by=index(y))

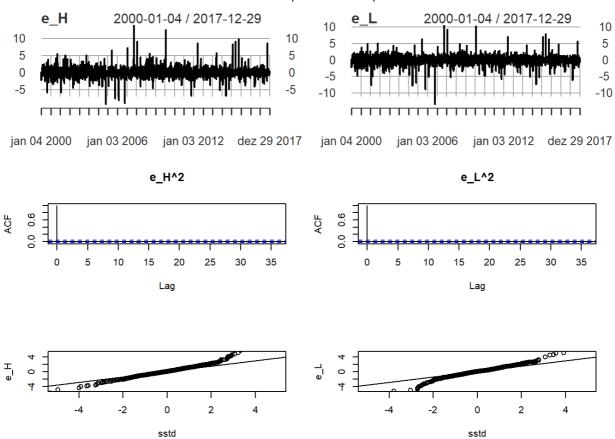
par(mfrow=c(3,1))
plot(C, main="BOVA11")
plot(Vol)
plot(R, main="R")</pre>
```



```
## Standard Residuals
r <- mY / sqrt(Vol)

par(mfrow=c(3,2))

plot(r[,1], main="e_H")
plot(r[,2], main="e_L")
acf(r[,1]^2, main="e_H^2")
acf(r[,2]^2, main="e_L^2")
r1 <- as.numeric(r[,1])
x <- rsstd(2000, mean = 0, sd = 1, nu = parEst['nu'], xi =parEst['gamma_1'])
qqplot(x=x, y=r1, xlim=c(-5, 5), ylim=c(-5, 5), ylab="e_H",xlab="sstd")
qqline(r1)
r2 <- as.numeric(r[,2])
x <- rsstd(2000, mean = 0, sd = 1, nu = parEst['nu'], xi =parEst['gamma_2'])
qqplot(x=x, y=r2, xlim=c(-5, 5), ylim=c(-5, 5), ylab="e_L",xlab="sstd")
qqline(r2)</pre>
```



```
# Prepare input for the expert advisor
## High
#HBOP
High UB HBOP = qsstd(p=1-(1-C Trend)/2, mean = 0, sd = 1, nu = parEst['nu'], xi = parEst['gam
ma_1'])
#S1
High_UB_S1 = qsstd(p=1-(1-C_Reaction)/2, mean = 0, sd = 1, nu = parEst['nu'], xi = parEst['ga
mma 1'])
## Low
#B1
Low_LB_B1 = qsstd(p=(1-C_Reaction)/2, mean = 0, sd = 1, nu = parEst['nu'], xi = parEst['gamma
_2'])
#LBOP
Low_LB_LBOP = qsstd(p=(1-C_Trend)/2, mean = 0, sd = 1, nu = parEst['nu'], xi = parEst['gamma_
2'])
pH <- c(0.6, 0.65, 0.7, 0.75, 0.8, 0.85, 0.9, 0.95, 0.975, 0.99, 0.995)
qH \leftarrow round(qsstd(p=pH, mean = 0, sd = 1, nu = parEst['nu'], xi = parEst['gamma 1']),3)
names(qH) <- paste0(100*pH,"%")</pre>
pL <- 1 - pH
qL <- round(qsstd(p=pL, mean = 0, sd = 1, nu = parEst['nu'], xi = parEst['gamma_2']),3)
names(qL) <- paste0(100*pL,"%")
qC <- rbind(qH, qL)
rownames(qC) <- c("High_UB", "Low_LB")</pre>
colnames(qC) <- paste0(100*pL,"%")</pre>
m = matrix(NA,nrow=10,ncol=1)
rownames(m) = c("High_UB_HBOP","High_UB_S1","Low_LB_B1","Low_LB_LBOP",
               "High_omega", "High_alpha", "High_beta",
                      "Low_omega", "Low_alpha", "Low_beta")
colnames(m) = 'Value'
m["High UB HBOP",1] = High UB HBOP
m["High UB S1",1] = High UB S1
m["Low_LB_B1",1] = Low_LB_B1
m["Low_LB_LBOP",1] = Low_LB_LBOP
m["High omega",1] = parEst["omega 1"]
m["High_alpha",1] = parEst["alpha_1"]
m["High_beta",1] = parEst["beta_1"]
m["Low omega",1] = parEst["omega 2"]
m["Low alpha",1] = parEst["alpha 2"]
m["Low beta",1] = parEst["beta 2"]
# Input for expert advisor
print(qC)
```

```
40% 35% 30% 25% 20% 15% 10% 5% 2.5% 1%

High_UB 0.165 0.265 0.374 0.495 0.638 0.817 1.067 1.511 1.998 2.749

Low_LB -0.167 -0.266 -0.375 -0.496 -0.638 -0.817 -1.066 -1.508 -1.994 -2.743

0.5%

High_UB 3.426

Low_LB -3.417
```

print(round(m,3))

```
Value
High_UB_HBOP 1.998
High_UB_S1
             0.495
Low_LB_B1
             -0.496
Low_LB_LBOP -1.994
High_omega
             0.045
High_alpha
             0.027
High_beta
             0.961
Low_omega
             0.058
Low_alpha
             0.033
Low_beta
              0.956
```