# HW2 (Size Performance) 2011/03/28

Generate sequence from GARCH(1,1) process with non-Gaussian innovations. For each case, consider the sample size T=(100, 300, 500, 700, 900). For each sample size, please simulate two tests (Ljung-Box and Q-Star) at least 1000 times and evaluate rejection frequency.

$$y_t = \sqrt{h_t} u_t , u_t \sim t(\nu)$$
 with conditional variance :  $h_t = \alpha_0 + \alpha_1 y_{t-1}^2 + \beta_1 h_{t-1}$ , where  $\alpha_0 = 0.5$ ,  $\beta_1 = (0.95 - \alpha_1)$ 

- 1. Given  $\alpha_1=(0.2,\ 0.5,\ 0.8)$ , simulate GARCH(1,1) process with t(3) innovations. Plot rejection frequency for two tests. Please specify the color or marks of each line. For example, "dot line:  $\alpha_1=0.2$ .; solid line:  $\alpha_1=0.5$ ."
- 2. Given  $\alpha_1 = (0.2, 0.5, 0.8)$ , simulate GARCH(1,1) process with t(5) innovations. Plot rejection frequency for two tests.
- 3. How are the changes in  $\nu$  (degree of freedom) related to rejection frequency? For example, does increasing of degree of freedom cause rejection frequency decrease? Why or why not? Think carefully about the underlying statistical inference.

#### R: Tips & tricks

Download library(fGarch), then you can use garchSpec() and garchSim() to generate GARCH sequences.

Please refer to the following links: <a href="http://help.rmetrics.org/fGarch/html/garchSim.html">http://help.rmetrics.org/fGarch/html/garchSim.html</a> http://help.rmetrics.org/fGarch/html/garchSpec.html

Examples 1 :

Suppose you want to generate a GARCH(1,1) sequence with coefficient  $\alpha_0=0.5$ ,  $\alpha_1=0.2$ ,  $\beta_1=0.75$  and Gaussian innovations.

```
spec = garchSpec(model = list(alpha = 0.2, beta = 0.75, omega = 0.5));
y <- garchSim(spec, n = T); ## T is sample size. For example, T is 100.</pre>
```

Examples 2 :

Suppose you want to generate a GARCH(1,1) sequence with coefficient  $\alpha_0=0.5$ ,  $\alpha_1=0.2$ ,  $\beta_1=0.75$  and t(3) innovations.

```
spec = garchSpec(model = list(alpha = 0.2, beta = 0.75, omega = 0.5, shape = 3), cond.dist =
"std");
```

y <- garchSim(spec, n = T); ## T is sample size. For example, T is 100.

#### HW2 (Bonus)

Similar to HW2, please evaluate size performance of modified Cramer-von Mises test (Deo 2000) and compare to original Cramer-von Mises test (Durlauf 1991). You may try difference GARCH sequence. For each case, consider the sample size T=(100, 300, 500, 700, 900). For each sample size, please simulate at least 1000 times and evaluate rejection frequency.

- 1. GARCH(1,1) with coefficient  $\alpha_1 = (0.2, 0.5, 0.8)$  and Gaussian innovations
- 2. GARCH(1,1) with coefficient  $\alpha_1=(0.2,\ 0.5,\ 0.8)$  and t(3) innovations
- 3. GARCH(1,1) with coefficient  $\alpha_1=(0.2,\ 0.5,\ 0.8)$  and t(5) innovations

### Deo (2000):

$$\begin{split} D_T^c(t) &= \frac{\sqrt{2T}}{\pi} \sum_{j=1}^{m(T)} \frac{\widehat{\rho}(j)}{\sqrt{\widehat{v_{JJ}}}} \frac{\sin(j\pi t)}{j}, \\ \text{where } \sqrt{\widehat{v_{JJ}}} &= \frac{1}{\widehat{\gamma}(0)} (\frac{1}{T-j} \sum_{t=1}^{T-j} (y_t - \bar{y})^2 (y_{t+j} - \bar{y})^2)^{0.5}. \\ \text{When T is large,} \\ \text{CVM} &= \int_0^1 [D_T^c(t)]^2 dt \stackrel{T \to \infty}{\longrightarrow} \int_0^1 [B^0(t)]^2 dt \end{split}$$

## R: Tips & tricks

Programming flow chart (modified Cramer-von Mises test)

