

Macroeconometrics: Problem Set 1

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Agenda

Granger-Causality

Diagnostic Tests

Unit Root Tests

Granger-Causality

- ▶ The variable y does not “Granger-cause” x if we cannot anticipate x with values of y . Essentially, we contrast the following two models

$$\begin{bmatrix} x_t \\ y_t \end{bmatrix} = \sum_{\ell=1}^p \begin{bmatrix} \phi_{\ell}^{xx} & 0 \\ \phi_{\ell}^{yx} & \phi_{\ell}^{yy} \end{bmatrix} \begin{bmatrix} x_{t-\ell} \\ y_{t-\ell} \end{bmatrix} + \begin{bmatrix} \pi^x \\ \pi^y \end{bmatrix} D_t + \begin{bmatrix} e_t^x \\ e_t^y \end{bmatrix} \quad (C)$$

$$\begin{bmatrix} x_t \\ y_t \end{bmatrix} = \sum_{\ell=1}^p \begin{bmatrix} \phi_{\ell}^{xx} & \phi_{\ell}^{xy} \\ \phi_{\ell}^{yx} & \phi_{\ell}^{yy} \end{bmatrix} \begin{bmatrix} x_{t-\ell} \\ y_{t-\ell} \end{bmatrix} + \begin{bmatrix} \pi^x \\ \pi^y \end{bmatrix} D_t + \begin{bmatrix} e_t^x \\ e_t^y \end{bmatrix} \quad (F)$$

- ▶ A test is carried out with these hypothesis

$$H_0 : \phi_1^{xy} = 0 \cap \dots \cap \phi_p^{xy} = 0 \text{ (all } \phi_{\ell}^{xy} \text{ are zero)}$$

$$H_A : \phi_1^{xy} \neq 0 \cup \dots \cup \phi_p^{xy} \neq 0 \text{ (at least one } \phi_{\ell}^{xy} \text{ is not zero),}$$

where if H_0 is true, we say that y does not “Granger-cause” x .

- ▶ We can employ either a Wald test (only F is estimated), a LM test (only C is estimated) or a likelihood ratio test (both are estimated).

Granger-Causality

- ▶ In the case of multiple variables, there are two ways on how to test for Granger-causality:
 1. Compare one variable versus the rest as a whole.
 2. Evaluate each variable individually.
- ▶ For example, let's say we have three variables x , y and z and we wish to test if x is "Granger-caused" by the other variables. With the first method, we define the vector $w \equiv (y, z)$ and apply the same logic as in the two variable setup. The constrained model is then

$$\begin{bmatrix} x_t \\ w_t \end{bmatrix} = \sum_{\ell=1}^p \begin{bmatrix} \phi_{\ell}^{xx} & 0 \\ \phi_{\ell}^{wx} & \Phi_{\ell}^{ww} \end{bmatrix} \begin{bmatrix} x_{t-\ell} \\ w_{t-\ell} \end{bmatrix} + \begin{bmatrix} \pi^x \\ \Pi^w \end{bmatrix} D_t + \begin{bmatrix} e_t^x \\ e_t^w \end{bmatrix}$$

whereas with the second method we would have

$$\begin{bmatrix} x_t \\ y_t \\ z_t \end{bmatrix} = \sum_{\ell=1}^p \begin{bmatrix} \phi_{\ell}^{xxx} & \phi_{\ell}^{xyy} & 0 \\ \phi_{\ell}^{yxx} & \phi_{\ell}^{yyy} & \phi_{\ell}^{yzz} \\ \phi_{\ell}^{zxx} & \phi_{\ell}^{zyy} & \phi_{\ell}^{zzz} \end{bmatrix} \begin{bmatrix} x_{t-\ell} \\ y_{t-\ell} \\ z_{t-\ell} \end{bmatrix} + \begin{bmatrix} \pi^x \\ \pi^y \\ \pi^z \end{bmatrix} D_t + \begin{bmatrix} e_t^x \\ e_t^y \\ e_t^z \end{bmatrix},$$

the difference being on how to treat each ϕ_{ℓ}^{xyy} in the first equation.

Granger-Causality

- ▶ The R package `vars` implements Granger-causality as a Wald test that is executed with the command `causality()`.
- ▶ Unfortunately, the second method described previously is not contemplated in the `vars` library and requires manual computation.
- ▶ `vars` also allows to use a bootstrap version instead of an asymptotic approximation.
- ▶ The function `causality()` also checks for “instantaneous causality”. A Wald test is conducted for the following hypothesis

$$H_0 : \text{vec}(\Sigma_e^{xy}) = 0$$

$$H_A : \text{vec}(\Sigma_e^{xy}) \neq 0,$$

involving the (block) covariance of the residuals in question.

Serial Correlation: Portmanteau Test

- ▶ This is a multivariate generalization of the Ljung-Box test. It is used to test that there is no auto- or cross correlation in the residuals. The hypothesis of the test are

$$H_0 : R_1 = 0 \cap \dots \cap R_h = 0 \text{ (all } R_\ell \text{ are zero)}$$

$$H_A : R_1 \neq 0 \cup \dots \cup R_h \neq 0 \text{ (at least one } R_\ell \text{ is not zero),}$$

where each R_ℓ is a matrix of cross-correlations of order ℓ .

- ▶ The test statistic has an approximate asymptotically chi-squared distribution under H_0

$$Q_m(h) = T \sum_{\ell=1}^h \text{tr}(\hat{\Gamma}'_\ell \hat{\Gamma}_0^{-1} \hat{\Gamma}_\ell \hat{\Gamma}_0^{-1}) \xrightarrow{d} \chi^2_{m^2(h-p)}$$

where each $\hat{\Gamma}_\ell$ are the sample matrices of cross-covariances of order ℓ computed according to the following formula

$$\hat{\Gamma}_\ell = \frac{1}{T} \sum_{t=\ell+1}^T (\hat{e}_t - \bar{\hat{e}})(\hat{e}_{t-\ell} - \bar{\hat{e}})'$$

Serial Correlation: Portmanteau Test

- ▶ There is a trade-off between a good approximation to the chi-squared random variable and a loss of power depending on the specified value for h . Large values should be preferred.
- ▶ The test is implemented in the package `vars` via the command `serial.test()` when selecting `type = "PT.asymptotic"` in R.
- ▶ For smaller sample sizes, a “corrected” version is available (`type = "PT.adjusted"`). The adjusted test statistic is

$$Q_m^C(h) = T^2 \sum_{\ell=1}^h \frac{1}{T-\ell} \text{tr}(\hat{\Gamma}'_{\ell} \hat{\Gamma}_0^{-1} \hat{\Gamma}_{\ell} \hat{\Gamma}_0^{-1}) \overset{d}{\approx} \chi_{m^2(h-p)}^2$$

- ▶ This test requires a slight modification when applied to VARs estimated with constrained coefficients. Unluckily, `vars` does not include this modification.

Serial Correlation: Breusch–Godfrey Test

- ▶ A multivariate generalization of the standard Breusch–Godfrey test. We estimate the auxiliary regression

$$\hat{\varepsilon}_t = F_1 y_{t-1} + \dots + F_p y_{t-p} + \Upsilon D_t + G_1 \hat{\varepsilon}_{t-1} + \dots + G_h \hat{\varepsilon}_{t-h} + \varepsilon_t$$

and contrast the following hypothesis to test for auto- or cross correlation in the residuals

$$H_0 : G_1 = 0 \cap \dots \cap G_h = 0 \text{ (all } G_j \text{ are zero)}$$

$$H_A : G_1 \neq 0 \cup \dots \cup G_h \neq 0 \text{ (at least one } G_j \text{ is not zero),}$$

the null hypothesis corresponds to no serial correlation in the residuals.

- ▶ Under H_0 , the test statistic converges asymptotically to a chi-squared random variable

$$BG(h) = T[m - \text{tr}(\hat{\Sigma}_e^{-1} \hat{\Sigma}_\varepsilon)] \xrightarrow{d} \chi_{hm^2}^2.$$

Serial Correlation: Breusch–Godfrey Test

- ▶ Suitable for small h should be used. Degrees of freedom are easily exhausted.
- ▶ The test is implemented in `vars` with the command `serial.test()` when selecting `type = "BG"`.
- ▶ A “corrected” statistic with better properties in small samples is also available with the option `type = "ES"`

$$BG^C(h) = \frac{1 - (1 - \delta)^{\frac{1}{\zeta}}}{(1 - \theta)^{\frac{1}{\zeta}}} \left(\frac{m^2 h}{\kappa \zeta - \eta} \right)^{-1} \xrightarrow{d} F_{\text{int}(\kappa \zeta - \eta)}^{m^2 h}$$

with

$$\begin{aligned} \theta &= 1 - |\hat{\Sigma}_\varepsilon|/|\hat{\Sigma}_e| & \zeta &= \left(\frac{m^4 h^2 - 4}{m^2 + (mh)^2 - 5} \right)^{\frac{1}{2}} \\ \eta &= \frac{1}{2} m^2 h - 1 & \kappa &= T - m - mh - \frac{1}{2}(m - mh - 1), \end{aligned}$$

which converges to an F random variable under H_0 .

Normality: Jarque-Bera Test

- ▶ This a goodness-of-fit test of whether sample residuals have the skewness and kurtosis matching a normal distribution.
- ▶ The test statistic converges in distribution to a chi-squared random variable

$$JB = \lambda_3 + \lambda_4 \xrightarrow{d} \chi_{2m}^2,$$

with

$$\lambda_3 = \frac{T}{6} \hat{b}'_1 \hat{b}_1 \xrightarrow{d} \chi_m^2$$

$$\lambda_4 = \frac{T}{24} (\hat{b}'_2 - 3_m)' (\hat{b}'_2 - 3_m) \xrightarrow{d} \chi_m^2,$$

where \hat{b}_1 and \hat{b}_2 are the third and fourth moment vectors of the standardized residuals, $\hat{e}_t^s = \hat{P}^{-1}(\hat{e}_t - \bar{\hat{e}})$.

- ▶ P is a matrix with positive diagonal elements such that $\hat{P}\hat{P}' = \hat{\Sigma}_e$. The Cholesky decomposition of $\hat{\Sigma}_e$ is an example of \hat{P} .
- ▶ The implementation in `vars` is via `normality.test()`.
- ▶ The test can be inaccurate unless there is a really large sample.

Heteroskedasticity: ARCH Test

- ▶ The ARCH test involves an auxiliary regression

$$\text{vech}(\hat{e}_t \hat{e}_t') = \eta + S_1 \text{vech}(\hat{e}_{t-1} \hat{e}_{t-1}') + \dots + S_q \text{vech}(\hat{e}_{t-q} \hat{e}_{t-q}') + \nu_t$$

where the following hypothesis are tested

$$H_0 : S_1 = 0 \cap \dots \cap S_q = 0 \text{ (all } S_j \text{ are zero)}$$

$$H_A : S_1 \neq 0 \cup \dots \cup S_q \neq 0 \text{ (at least one } S_j \text{ is not zero).}$$

- ▶ If H_0 is true, then the test statistic shown below converges to a chi-squared random variable

$$ARCH(q) = \frac{1}{2} T m(m+1) \theta \xrightarrow{d} \chi_{qm^2(m+1)^2/4}^2$$

with

$$\theta = 1 - \frac{2}{m(m+1)} \text{tr}(\hat{\Omega} \hat{\Omega}_0^{-1}),$$

and we say that there are no ARCH effects (i.e., heteroskedasticity).

- ▶ Using `vars`, it is performed via the command `arch.test()`.

Unit Root Tests: Augmented Dickey–Fuller (ADF) Test

- ▶ The R library `urca` implements various unit root tests.
- ▶ In particular, we'll focus on the ADF test.
- ▶ The command `ur.df` allows for three different specifications

$$\textbf{Case 1} : \Delta y_t = \pi y_{t-1} + \sum_{j=1}^k \gamma_j \Delta y_{t-j} + u_t$$

$$\textbf{Case 2} : \Delta y_t = \beta_1 + \pi y_{t-1} + \sum_{j=1}^k \gamma_j \Delta y_{t-j} + u_t$$

$$\textbf{Case 3} : \Delta y_t = \beta_1 + \beta_2 t + \pi y_{t-1} + \sum_{j=1}^k \gamma_j \Delta y_{t-j} + u_t.$$

- ▶ The number of lags k can be chosen by information criteria.

Unit Root Test: Augmented Dickey–Fuller Test

- ▶ The null hypothesis are as follows depending on each case

Case 1 : $\tau_1 : \pi = 0.$

Case 2 : $\tau_2 : \pi = 0$
 $\phi_1 : \pi = 0 \cap \beta_1 = 0.$

Case 3 : $\tau_3 : \pi = 0$
 $\phi_2 : \pi = 0 \cap \beta_2 = 0$
 $\phi_3 : \pi = 0 \cap \beta_2 = 0 \cap \beta_1 = 0.$

- ▶ The ADF test is a **left-tailed** test. The null hypothesis is rejected if the test statistic is larger than the specified critical value for a given significance level.
- ▶ A general drawback of unit root tests is that they have low power against persistent stationary processes.