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The Fractionally Cointegrated Vector Autoregression Model in R

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Abstract

This article illustrates how to estimate the fractionally cointegrated vector autoregression model in R.

Keywords: cofractional process, cointegration rank, fractional autoregressive model, fractional cointegration, fractional unit root, VAR model, Matlab, R.

1. Introduction: Cointegration and fractional integration in R

> The introduction is in principle "as usual". However, it should usually embed both the implemented *methods* and the *software* into the respective relevant literature. For the latter both competing and complementary software should be discussed (within the same software environment and beyond), bringing out relative (dis)advantages. All software mentioned should be properly `\cite{}`d.

The fractionally cointegrated vector autoregression model is an excellent model...

Let's stick to a description with words here. We'll get into the ugly details in the next section. For now, talk about CVAR as a tool for testing equilibrium relationships. Then the fractional part is to accommodate a potenially long-lived deviation from equilibrium.

An exhaustive listing of the R packages available for time series analysis were compiled in *CRAN Task View: Time Series Analysis* by Rob J Hyndman, available at `https://CRAN.R-project.org/vie`

1.1. R packages for estimating the CVAR model

In R (R Core Team 2017), contegration analysis can be conducted using a variety of packages.

- One such package is **aTSA** for *Alternative Time Series Analysis* (Qiu 2015). In this package, the `coint.test()` function performs Engle-Granger tests, as in Engle and Granger (1987), for the null hypothesis that two or more time series, each of which is I(1), are not cointegrated.
  – This package is designed for bivariate analysis.

- Another package following the Engle and Granger (1987) approach is the **egcm** package in (Clegg 2017). – This package is also designed for bivariate analysis.

- The `po.test()` from the **tseries** package Trapletti, Hornik, and LeBaron (2019) implements the tests in Phillips and Ouliaris (1990).

- Another option is the **cointReg** in Aschersleben and Wagner (2016). This implements three types of regressions designed for integrated variables[1].

- An option without the limitation to bivariate series is the **urca** package to perform unit root tests and cointegration analysis (Pfaff, Zivot, and Stigler 2016)[2].
  – They have the function `ca.po()` implementing the tests in Phillips and Ouliaris (1990).
  – This package also follows the approach in Johansen (1995) in the function `ca.jo()`.
  – Within this framework, there is a function `cajo.test()` to test restrictions in the CVAR model.

- The `VECM()` function in the **tsDyn** package allows for the application of either the Engle and Granger (1987) or the Johansen (1995) MLE method.
  – This package is designed primarily with nonlinear time series models in mind.

1.2. Segue from CVAR to $I(d)$

These packages allow for only a discrete form of cointegration between the series. For example, the series are all integrated, i.e. $I(1)$, and the residuals from a regression are stationary and $I(0)$. The fractionally cointegrated VAR model allows for the posibility that variables can be integrated of order d and cointegrated of order $d-b$, where d and $b>0$ can be real numbers.

Analysis using the above packages typically involves a preliminary alanlysis of the form of non-stationarity of the variables, using a number of unit root tests, i.e. to test whether the series are $I(1)$. With fractionally integrated variables, the first stage of the analysis is to determine the order of fractional integration, i.e. the parameter d.

[1] I will describe this later, if we decide to keep it in (the above and below are sufficient, in my view) but here is a quote from their description: "These methods include the fully modified OLS (FM-OLS) approach of Phillips and Hansen (1990) <doi:10.2307/2297545>, the dynamic OLS (D-OLS) approach of Phillips and Loretan (1991) <doi:10.2307/2298004>, Saikkonen (1991) <doi:10.1017/S0266466600004217> and Stock and Watson (1993) <doi:10.2307/2951763> and the new estimation approach called integrated modified OLS (IM-OLS) of Vogelsang and Wagner (2014) <doi:10.1016/j.jeconom.2013.10.015>. The latter is based on an augmented partial sum (integration) transformation of the regression model".

[2] They also use the Danish data, so it would be worthwhile to include in the documentation an example with and without the restriction $d=b=1$ to compare

1.3. R packages for fractional integration

Quoted from *CRAN Task View: Time Series Analysis* by Rob J Hyndman, referenced above:
"Some facilities for fractional differenced ARFIMA models are provided in the **fracdiff** package. The **arfima** package has more advanced and general facilities for ARFIMA and ARIMA models, including dynamic regression (transfer function) models. Additional methods for fitting and simulating non-stationary ARFIMA models are in **nsarfima**. **LongMemoryTS** provides a collection of functions for analysing long memory time series."

- The **fracdiff** package (Maechler, Fraley, Leisch, Reisen, Lemonte, and Hyndman (2020)) includes functions for fitting $ARFIMA(p, d, q)$ models, including the step of estimating the long memory parameter d. The namesake function `fracdiff()` calculates the maximum likelihood estimators of the parameters of a fractionally-differenced $ARIMA(p, d, q)$ model. A few notable functions in this package estimate the long memory parameter d within this model. [3]
- **arfima** Veenstra and McLeod (2018) fits a wider variety of arfima models.
- **nsarfima** Groebe (2019) is more innovative in terms of the types of optimization problems built on the $ARFIMA$ model, including both maximum likelihood (as in Beran (1995)) and minimum distance (as in Mayoral, L. (2007)) estimators.
- **LongMemoryTS** this model uses a wide variety of methods methods to investigate both cointegrating relationships and fractional integration. However, it seems to me that this package (Leschinski, Voges, and Wenger (2019)) chains together several separate issues, such as, estimating the fractional integration parameter and then plugging that value into the rank test as if it were known. Another example is testing cointegration by estimating the fractional parameter from the residuals. Morten, you have a lot of citations in this package. It implements Nielsen (JoE, 2010) with a given d, where apparently you recommend `d = 0.1`. It also implements Nielsen and Shimotsu (JoE, 2007) for rank estimation in fractionally cointegrated systems. Also: Hausman-type test for fractional cointegration (Robinson (2008)). Basically a who's who of analyzing the cointegration of fractional systems, except for Johansen's MLE framework, which is what we implement. Ours is more (what's the word?) thorough, in that all the parameters are estimated jointly, aside from the rank and lag selection. It captures all of these in one maximum likelihood framework.

This package is closest to a cross between the Johansen cointegration model in **urca** and the fractionally differencing models in **a**. In fact[4], the model estimated in the **urca** packages is the special case in which the fractional integration parameters d and b are both equal to one.

1.4. The Matlab program

This R package is based on a companion package **FCVARmodel.m**, written in Matlab. The Matlab package is documented in Nielsen and Popiel (2016) and Nielsen and Morin (2014).

[3] Note that the `diffseries()` function in **fracdiff** is based on the same algorithm in Jensen and Nielsen (2014) as `FracDiff()` in **FCVAR**, except that `diffseries()` demeans the data first. Specifically, `all(fracdiff::diffseries(x, d) == FCVAR::FracDiff(x - mean(x), d))` returns `TRUE`.

[4] Maybe we should add this example back, since we had it in the first version.

The latest version of the package can be downloaded from the first author's website at Queen's University:

http://www.econ.queensu.ca/faculty/mon/software/

It is freely available for non-commercial, academic use. The use of this program requires a functioning installation of Matlab. Any recent version should work. (more precisely: which version? And this part should go at the end)

Unzip the contents of the zip file into any directory that you plan to use as the working directory of the program.

1.5. Roadmap

The next section describes the FCVAR model and the restricted models that can be estimated with this program. Section 3 describes the functioning of the main program, which is a replication of one of the tables of results in Jones, Nielsen, and Popiel (2014). Section 4 describes another example program, which demonstrates some additional functionality of the software. Importantly, these are the only two files that would need to be changed to apply the program for other empirical analyses.

2. The fractionally cointegrated VAR model

The fractionally cointegrated vector autoregressive (FCVAR) model was proposed in Johansen (2008) and analyzed by, e.g., Johansen and Nielsen (2010, 2012). For a time series X_t of dimension p, the fractionally cointegrated VAR model is given in error correction form as

$$\Delta^d X_t = \alpha\beta' \Delta^{d-b} L_b X_t + \sum_{i=1}^{k} \Gamma_i \Delta^d \ L_b^i X_t + \varepsilon_t, \tag{1}$$

where ε_t is p-dimensional $i.i.d.(0, \Omega)$, Δ^d is the fractional difference operator, and $L_b = 1 - \Delta^b$ is the fractional lag operator.[5] Johansen and Nielsen (2012) imposed two restrictions on the parameter space, $d \geq b$ and $d - b < 1/2$, in their asymptotic analysis. However, these restrictions were relaxed in Johansen and Nielsen (2018a,b).

Model (1) includes the Johansen (1995) CVAR model as the special case $d = b = 1$; see Johansen and Nielsen (2018b). Some of the parameters are well-known from the CVAR model and these have the usual interpretations also in the FCVAR model. The most important of these are the long-run parameters α and β, which are $p \times r$ matrices with $0 \leq r \leq p$. The rank r is termed the cointegration, or cofractional, rank. The columns of β constitute the r cointegration (cofractional) vectors such that $\beta' X_t$ are the cointegrating combinations of the variables in the system, i.e. the long-run equilibrium relations. The parameters in α are the adjustment or loading coefficients which represent the speed of adjustment towards equilibrium for each of the variables. The short-run dynamics of the variables are governed by the parameters $\Gamma = (\Gamma_1, \ldots, \Gamma_k)$ in the autoregressive augmentation.

[5] Both the fractional difference and fractional lag operators are defined in terms of their binomial expansion in the lag operator, L. Note that the expansion of L_b has no term in L^0 and thus only lagged disequilibrium errors appear in (1).

The FCVAR model has two additional parameters compared with the CVAR model, namely the fractional parameters d and b. Here, d denotes the fractional integration order of the observable time series and b determines the degree of fractional cointegration, i.e. the reduction in fractional integration order of $\beta' X_t$ compared to X_t itself. These parameters are estimated jointly with the remaining parameters. This model thus has the same main structure as in the standard CVAR model in that it allows for modeling of both cointegration and adjustment towards equilibrium, but is more general since it accommodates fractional integration and cointegration.

In the next four subsections we briefly describe the accommodation of deterministic terms as well as estimation and testing in the FCVAR model.

2.1. Deterministic terms

There are several ways to accommodate deterministic terms in the FCVAR model (1). The inclusion of the so-called restricted constant was considered in Johansen and Nielsen (2012), and the so-called unrestricted constant term was considered in Dolatabadi, Nielsen, and Xu (2016). A general formulation that encompasses both models is[6]

$$\Delta^d X_t = \alpha \Delta^{d-b} L_b(\beta' X_t + \rho') + \sum_{i=1}^{k} \Gamma_i \Delta^d \; L_b^i X_t + \xi + \varepsilon_t. \tag{2}$$

The parameter ρ is the so-called restricted constant term (since the constant term in the model is restricted to be of the form $\alpha\rho'$), which is interpreted as the mean level of the long-run equilibria when these are stationary, i.e. $E\beta' X_t + \rho' = 0$. The parameter ξ is the unrestricted constant term, which gives rise to a deterministic trend in the levels of the variables. When $d = 1$ this trend is linear. Thus, the model (2) contains both a restricted constant and an unrestricted constant. In the usual CVAR model, i.e. with $d = b = 1$, the former would be absorbed in the latter, but in the fractional model they can both be present and are interpreted differently. For the representation theory related to (2), and in particular for additional interpretation of the two types of constant terms, see Dolatabadi *et al.* (2016).

An alternative formulation of deterministic terms was suggested by Johansen and Nielsen (2016), albeit in a simpler model, with the aim of reducing the impact of pre-sample observations of the process. This model is

$$\Delta^d (X_t - \mu) = \alpha \beta' \Delta^{d-b} L_b (X_t - \mu) + \sum_{i=1}^{k} \Gamma_i \Delta^d \; L_b^i (X_t - \mu) + \varepsilon_t, \tag{3}$$

which can be derived easily from the unobserved components formulation

$$X_t = \mu + X_t^0, \quad \Delta^d X_t^0 = L_b \alpha \beta' X_t^0 + \sum_{i=1}^{k} \Gamma_i \Delta^d \; L_b^i X_t^0 + \varepsilon_t. \tag{4}$$

The formulation (3), or equivalently (4), includes the restricted constant, which may be obtained as $\rho' = \beta' \mu$. More generally, the level parameter μ is meant to accommodate a

[6] In Dolatabadi *et al.* (2016) the constants are included as $\rho' \pi_t(1)$ and $\xi \pi_t(1)$, where $\pi_t(u)$ denotes coefficients in the binomial expansion of $(1 - z)^{-u}$. This is mathematically convenient, but makes no difference in terms of the practical implementation.

non-zero starting point for the first observation on the process, i.e., for X_1. It has the added advantage of reducing the bias arising due to pre-sample behavior of X_t, at least in simple models, even when conditioning on no initial values (see below). For details, see Johansen and Nielsen (2016).

2.2. Maximum likelihood estimation

It is assumed that a sample of length $T+N$ is available on X_t, where N denotes the number of observations used for conditioning, for details see Johansen and Nielsen (2016). The models (1), (2), and (3) are estimated by conditional maximum likelihood, conditional on N initial values, by maximizing the function

$$\log L_T(\lambda) = -\frac{Tp}{2}(\log(2\pi)+1) - \frac{T}{2}\log\det\left\{T^{-1}\sum_{t=N+1}^{T+N}\varepsilon_t(\lambda)\varepsilon_t(\lambda)'\right\}, \tag{5}$$

where the residuals are defined as

$$\varepsilon_t(\lambda) = \Delta^d X_t - \alpha\Delta^{d-b}L_b(\beta' X_t + \rho') - \sum_{i=1}^{k}\Gamma_i\Delta^d\ L_b^i X_t - \xi, \quad \lambda = (d,b,\alpha,\beta,\Gamma,\rho,\xi), \tag{6}$$

for model (2), and hence also for submodels of model (2), such as (1), with the appropriate restrictions imposed on ρ and ξ. For model (3) the residuals are

$$\varepsilon_t(\lambda) = \Delta^d(X_t-\mu) - \alpha\beta'\Delta^{d-b}L_b(X_t-\mu) - \sum_{i=1}^{k}\Gamma_i\Delta^d\ L_b^i(X_t-\mu), \quad \lambda = (d,b,\alpha,\beta,\Gamma,\mu). \tag{7}$$

It is shown in Johansen and Nielsen (2012) and Dolatabadi *et al.* (2016) how, for fixed (d,b), the estimation of model (2) reduces to regression and reduced rank regression as in Johansen (1995). In this way the parameters $(\alpha,\beta,\Gamma,\rho,\xi)$ can be concentrated out of the likelihood function, and numerical optimization is only needed to optimize the profile likelihood function over the two fractional parameters, d and b. In model (3) we can similarly concentrate the parameters (α,β,Γ) out of the likelihood function resulting in numerical optimization over (d,b,μ), making the estimation of model (3) slightly more involved numerically than that of model (2).

For model (2) with $\xi = 0$, Johansen and Nielsen (2012) shows that asymptotic theory is standard when $b < 0.5$, and for the case $b > 0.5$ asymptotic theory is non-standard and involves fractional Brownian motion of type II. Specifically, when $b > 0.5$, Johansen and Nielsen (2012) shows that under i.i.d. errors with suitable moment conditions, the conditional maximum likelihood parameter estimates $(\hat{d},\hat{b},\hat{\alpha},\hat{\Gamma}_1,\ldots,\hat{\Gamma}_k)$ are asymptotically Gaussian, while $(\hat{\beta},\hat{\rho})$ are locally asymptotically mixed normal. These results allow asymptotically standard (chi-squared) inference on all parameters of the model, including the cointegrating relations and orders of fractionality, using quasi-likelihood ratio tests. As in the CVAR model, see Johansen (1995), the same results hold for the same parameters in the full models (2) and (3), whereas the asymptotic distribution theory for the remaining parameters, ξ and μ, is currently unknown.

2.3. Cointegration rank tests

Letting $\Pi = \alpha\beta'$, the likelihood ratio (LR) test statistic of the hypothesis $\mathcal{H}_r : \text{rank}(\Pi) = r$ against $\mathcal{H}_p : \text{rank}(\Pi) = p$ is of particular interest because it deals with an important empirical

question. This statistic is often denoted the "trace" statistic. Let $\theta = (d, b)$ for model (2) and $\theta = (d, b, \mu)$ for model (3) denote the parameters for which the likelihood is numerically maximized. Then let $L(\theta, r)$ be the profile likelihood function given rank r, where (α, β, Γ), and possibly (ρ, ξ) if appropriate, have been concentrated out by regression and reduced rank regression; see Johansen and Nielsen (2012) and Dolatabadi *et al.* (2016) for details.

The profile likelihood function is maximized both under the hypothesis $\mathcal{H}_r$ and under $\mathcal{H}_p$ and the LR test statistic is then $\mathrm{LR}_T(q) = 2\log(L(\hat{\theta}_p, p)/L(\hat{\theta}_r, r))$, where

$$L(\hat{\theta}_p, p) = \max_{\theta} L(\theta, p), \quad L(\hat{\theta}_r, r) = \max_{\theta} L(\theta, r),$$

and $q = p - r$. This problem is qualitatively different from that in Johansen (1995) since the asymptotic distribution of $\mathrm{LR}_T(q)$ depends qualitatively (and quantitatively) on the parameter b. In the case with $0 < b < 1/2$ (sometimes known as "weak cointegration"), $\mathrm{LR}_T(q)$ has a standard asymptotic distribution, see Johansen and Nielsen (2012, Theorem 11(ii)), namely

$$\mathrm{LR}_T(q) \xrightarrow{D} \chi^2(q^2), \quad 0 < b < 1/2. \tag{8}$$

On the other hand, when $1/2 < b \leq d$ ("strong cointegration"), asymptotic theory is non-standard and

$$\mathrm{LR}_T(q) \xrightarrow{D} \mathrm{Tr}\left\{ \int_0^1 \mathsf{d}W(s) F(s)' \left(\int_0^1 F(s)F(s)'\mathsf{d}s \right)^{-1} \int_0^1 F(s)\mathsf{d}W(s)' \right\}, \quad b > 1/2, \tag{9}$$

where the vector process $\mathsf{d}W$ is the increment of ordinary (non-fractional) vector standard Brownian motion of dimension $q = p - r$. The vector process F depends on the deterministics in a similar way as in the CVAR model in Johansen (1995), although the fractional orders complicate matters. The following cases have been derived in the literature:

1. When no deterministic term is in the model, $F(u) = W_b(u)$, where $W_b(u) = \Gamma(b)^{-1} \int_0^u (u-s)^{b-1}\mathsf{d}W(s)$ is vector fractional Brownian motion of type II, see Johansen and Nielsen (2012, Theorem 11(i)).

2. When only the restricted constant term is included in model (2), $F(u) = (W_b(u)', u^{-(d-b)})'$, see Johansen and Nielsen (2012, Theorem 11(iv)) for the result with $d = b$ and an earlier working paper version for the general result.

3. In model (3) the same result as in bullet 2. holds because $\beta'\mu = \rho'$ is the restricted constant and $\beta_\perp'\mu$ has no influence on the asymptotic distribution (in a similar way to X_0 in a random walk).

4. When both the restricted and unrestricted constants are included in model (2) with $d = 1$,

$$F_i(u) = W_{b,i}(u) - \int_0^1 W_{b,i}(u)\mathsf{d}u, \ i = 1, ..., q-1,$$
$$F_q(u) = u^b - \int_0^1 u^b \mathsf{d}u = u^b - 1/(b+1),$$
$$F_{q+1}(u) = u^{b-1} - \int_0^1 u^{b-1}\mathsf{d}u = u^{b-1} - 1/b,$$

see Dolatabadi *et al.* (2016).

Importantly, the asymptotic distribution (9) of the test statistic $\mathrm{LR}_T(q)$ depends on both b and $q = p - r$. The dependence on the unknown (true value of the) scalar parameter b complicates empirical analysis compared to the CVAR model. Generally, the distribution (9) would need to be simulated on a case-by-case basis. However, for model (1) and for model (2) with $d = b$ and $\xi = 0$, and hence also for model (3) with $d = b$ in light of bullet 3. above, computer programs for computing asymptotic critical values and asymptotic P values for the LR cointegration rank tests based on numerical distribution functions, are made available by MacKinnon and Nielsen (2014). Their computer programs are incorporated in the present program for the relevant cases/models as discussed and illustrated below.

2.4. Restricted models

Note that a reduced rank restriction has already been imposed on models (1)–(3), where the coefficient matrix $\Pi = \alpha\beta'$ has been restricted to rank $r \leq p$. Other restrictions on the model parameters can be considered as in Johansen (1995). The most interesting restrictions from an economic theory point of view would likely be restrictions on the adjustment parameters α and cointegration vectors β.

We formulate hypotheses as

$$R_\psi \psi = r_\psi, \tag{10}$$
$$R_\alpha \mathrm{vec}(\alpha) = 0, \tag{11}$$
$$R_\beta \mathrm{vec}(\beta^*) = r_\beta, \tag{12}$$

with $\beta^* = (\beta', \rho')'$, and use the switching algorithm in (Boswijk and Doornik 2004, p. 455) to optimize the likelihood numerically subject to the restrictions. The switching algorithm can be improved by adding a line search, see Doornik (2018). This is done by setting the option `opt.LineSearch = 1`, which is the default setting.

The only limitation on the linear restrictions that can be imposed on (d, b, α, β^*) in (10)–(12) is that only homogenous restrictions can be imposed on $\mathrm{vec}(\alpha)$ in (11). Otherwise, any combination of linear restrictions can be imposed on these parameters. For now, the remaining parameters cannot be restricted.

Note that, when the restricted constant term ρ is included in the model, restrictions on β and ρ must be written in the form given by (12). This is without loss of generality.

The restrictions in (10)–(12) above can be implemented individually or simultaneously in the Matlab program. The next section provides an example session illustrating the use of the program with a step-by-step description of a typical empirical analysis, including several restricted models in Section 3.6.

2.5. Forecasting from the FCVAR model

Because the FCVAR model is autoregressive, the best linear predictor takes a simple form and is relatively straightforward to calculate. Consider, for example, the model with level parameter in (3). We first note that

$$\Delta^d(X_{t+1} - \mu) = X_{t+1} - \mu - (X_{t+1} - \mu) + \Delta^d(X_{t+1} - \mu) = X_{t+1} - \mu - L_d(X_{t+1} - \mu)$$

and then rearrange (3) as

$$X_{t+1} = \mu + L_d(X_{t+1} - \mu) + \alpha\beta'\Delta^{d-b}L_b(X_{t+1} - \mu) + \sum_{i=1}^{k}\Gamma_i\Delta^d L_b^i(X_{t+1} - \mu) + \varepsilon_{t+1}. \quad (13)$$

Since $L_b = 1 - \Delta^b$ is a lag operator, so that $L_b^i X_{t+1}$ is known at time t for $i \geq 1$, this equation can be used as the basis to calculate forecasts from the model.

We let conditional expectation given the information set at time t be denoted $E_t(\cdot)$, and the best linear predictor forecast of any variable Z_{t+1} given information available at time t be denoted $\hat{Z}_{t+1|t} = E_t(Z_{t+1})$. Clearly, we then have that the forecast of the innovation for period $t+1$ at time t is $\hat{\varepsilon}_{t+1|t} = E_t(\varepsilon_{t+1}) = 0$, and $\hat{X}_{t+1|t}$ is then easily found from (13). Inserting also coefficient estimates based on data available up to time t, denoted[7] $(\hat{d}, \hat{b}, \hat{\mu}, \hat{\alpha}, \hat{\beta}, \hat{\Gamma}_1, \ldots, \hat{\Gamma}_k)$, we have that

$$\hat{X}_{t+1|t} = \hat{\mu} + L_{\hat{d}}(X_{t+1} - \hat{\mu}) + \hat{\alpha}\hat{\beta}'\Delta^{\hat{d}-\hat{b}}L_{\hat{b}}(X_{t+1} - \hat{\mu}) + \sum_{i=1}^{k}\hat{\Gamma}_i\Delta^{\hat{d}}L_{\hat{b}}^i(X_{t+1} - \hat{\mu}). \quad (14)$$

This defines the one-step ahead forecast of X_{t+1} given information at time t.

Multi-period ahead forecasts can be generated recursively. That is, to calculate the h-step ahead forecast, we first generalize (14) as

$$\hat{X}_{t+j|t} = \hat{\mu} + L_{\hat{d}}(\hat{X}_{t+j|t} - \hat{\mu}) + \hat{\alpha}\hat{\beta}'\Delta^{\hat{d}-\hat{b}}L_{\hat{b}}(\hat{X}_{t+j|t} - \hat{\mu}) + \sum_{i=1}^{k}\hat{\Gamma}_i\Delta^{\hat{d}}L_{\hat{b}}^i(\hat{X}_{t+j|t} - \hat{\mu}), \quad (15)$$

where $\hat{X}_{s|t} = X_s$ for $s \leq t$. Then forecasts are calculated recursively from (15) for $j = 1, 2, \ldots, h$ to generate h-step ahead forecasts, $\hat{X}_{t+h|t}$.

Clearly, one-step ahead and h-step ahead forecasts for the model (2) with a restricted constant term, and possibly also an unrestricted constant term, instead of the level parameter can be calculated entirely analogously.

3. Main Program

Estimating the fractionally cointegrated vector autoregression model works like this...

Example session: replicationJNP2014.m

The main file is `fcvar_replication_JNP2014.R` and it serves as an example of what a typical session of estimation, testing, and forecasting can include. This code replicates "Table 4: FCVAR results for Model 1" from Jones *et al.* (2014) and follows the empirical procedure developed in that paper. This procedure includes the following steps:

1. Importing data
2. Choosing estimation options

[7] To emphasize that these estimates are based on data available at time t, they could be denoted by a subscript t. However, to avoid cluttering the notation we omit this subscript and let it be understood in the sequel.

3. Lag selection
4. Cointegration rank selection
5. Model estimation
6. Hypothesis testing

3.1. Importing data

The first step is importing the data. Executing the code shown below, assigns the data from the file `data_JNP2014.csv` to a matrix called `data`.

> Better to use names of variables for clarity. Are we actually using these matrices? Maybe we don't need numbers. I think we only `x1`, so we should just call it `x`. Finally, the actual command for a user will be from a data folder in the package.

```
R> x1 <- votingJNP2014[, c("lib", "ir_can", "un_can")]
```

The columns contain the following variables: (1) aggregate support for the Liberal party, (2) aggregate support for the Conservative party, (3) Canadian 3-month T-bill rates, (4) US 3-month T-bill rates, (5) Canadian unemployment rate, and (6) US unemployment rate. Since each of the models in JNP (2014) contain different combinations of these variables, the relevant columns of `data` for each model are assigned to different matrices of variables named `x1` through `x6`.

3.2. Choosing options

Once the data is imported, the user sets the program options. The script contains two sets of options: variables set for function arguments in the script itself and model/estimation related options. The first of set of options is as follows.

```
p              <- ncol(x1)
kmax           <- 3
order          <- 12
printWNtest    <-  1
```

The variable `kmax` determines the highest lag order for the sequential testing that is performed in the lag selection, whereas `p` is the dimension of the system. The `order` specifies the number of lags used for the white noise test in lag selection, while `printWNtest` indicates whether to print results of white noise tests post-estimation.

The next set of initialization commands assign values to the variables contained in object `opt` defined by the function `FCVARoptions`.

> I wonder if we should leave some of this to the package documentation. Maybe for this article, we should concentrate on the options that are necessary for the examples below. Besides, JSS style prefers code blocks do not have comments, i.e. state it in the text. We could say that "the rest of the options are described in the package documentation".

```
opt <- EstOptions() # Define variable to store Estimation Options (object).

opt$dbMin        <- c(0.01, 0.01) # lower bound for d,b.
opt$dbMax        <- c(2.00, 2.00) # upper bound for d,b.
opt$unrConstant  <- 0 # include an unrestricted constant? 1 <- yes, 0 <- no.
opt$rConstant    <- 0 # include a restricted constant? 1 <- yes, 0 <- no.
opt$levelParam   <- 1 # include level parameter? 1 <- yes, 0 <- no.
opt$constrained  <- 0 # impose restriction dbMax >= d >= b >= dbMin ? 1 <- yes, 0 <- no.
opt$restrictDB   <- 1 # impose restriction d=b ? 1 <- yes, 0 <- no.
opt$db0          <- c(0.8, 0.8) # set starting values for optimization algorithm.
opt$N            <- 0 # number of initial values to condition upon.
opt$print2screen <- 1 # print output.
opt$printRoots   <- 1 # do not print roots of characteristic polynomial.
opt$plotRoots    <- 1 # do not plot roots of characteristic polynomial.
opt$gridSearch   <- 1 # For more accurate estimation, perform the grid search.
                                    # This will make estimation take longer.
opt$plotLike     <- 0 # Plot the likelihood (if gridSearch <- 1).
opt$progress           <- 0 # Show grid search progress indicator waitbar.
opt$updateTime   <- 0.5 # How often progress is updated (seconds).

# Linux example:
opt$progLoc <- '"/usr/bin/fdpval"'  # location path with program name
                                    # of fracdist program, if installed
                                    # Note: use both single (outside) and double
                                    # quotes (inside). This is especially important
                                    # if path name has spaces.
# Windows example:
# opt$progLoc <- '".\fdpval\fdpval"'  # program located in folder fdpval in current direct

# There are many other options (see EstOptions.m for
# everything else. These can be accessed/modified as in, for example:
# opt$dbFminOptions$Algorithm <- 'interior-point'

DefaultOpt <- opt # Store the options for restoring them in between hypothesis tests.
```

The first line initializes the object `opt` and assigns all of the default options set in `FCVARoptions`. The user can see the full set of options by typing `DefaultOpt` (or `opt` after initialization) in the command line. The code block above shows how to easily change any of the default options. Defining the program options in this way allows the user to create and store several option objects with different attributes. This can be very convenient when, for example, performing the same hypothesis tests on different data sets.

The set of available options can be broken into several categories: numerical optimization, model deterministics and restrictions, output, grid search, and P-values for the rank test. We recommend that only advanced users make changes to the numerical optimization options. Adding deterministics requires setting the variable corresponding to the type of deterministic component to 1. For instance, in the present example, a model estimated with options `opt`

will include the level parameter μ but no restricted or unrestricted constant. Output variables refer to either printing or plotting various information post-estimation and usually take values 1 or 0 (on or off). For example, if the user is not interested in the estimates of Γ, they can be suppressed by setting `opt$printGammas <- 0`.

The bounds on the parameter space for d and b are specified in `opt$dbMin` and `opt$dbMax`. In this example, these are both specified as 2-dimensional column vectors, in which case the first element specifies the bound on d and the second element the bound on b. Alternatively, one can set `opt$dbMin` and `opt$dbMax` as scalars, which imposes the same bounds on d and b.

An important feature in this package is the ability to pre-estimate by using a grid search. If the user selects this option, they can view progress by setting `opt$progress` to 1 (wait-bar) or 2 (output in command line). The minimum frequency of these updates is set by `opt$updateTime`. The user also has the option (`opt$plotLike`) to view a plot of the likelihood over d and/or b after the grid search completes. The output of the grid search is a preliminary estimate of the fractional parameters. These are used as starting values in the subsequent numerical optimization, and the bounds on d and b are set to these starting values plus/minus 0.1 but still within the original `dbMin` and `dbMax` settings.

As of v.1.4.0, the new option `opt$LocalMax` allows more control over the grid search. If `opt$LocalMax <- 0`, the function `FCVARlikeGrid()` returns the parameter values corresponding to the global maximum of the likelihood on the grid. If `opt$LocalMax <- 1`, then `FCVARlikeGrid()` returns the parameter values for the local maximum corresponding to the highest value of b. This is meant to alleviate the identification problem discussed in Johansen and Nielsen (2010, Section 2.3) and Carlini and de Magistris (2017). As of v.1.4.0, the default setting is `opt$LocalMax <- 1`.

Another new option in v.1.4.0 is the addition of a line search to the switching algorithm for estimation of models with restrictions on α and/or β. This is added via the option `opt$LineSearch <- 1` and is the default. See Doornik (2018, Section 2.2) for details.

> We might have to rewrite this to reflect our updated strategy for the P-values for cointegration rank tests – once we figure out what that strategy is. Maybe it will be as simple as calling the **Rcpp** package.

In order to automatically obtain P-values for cointegration rank tests when $b > 0.5$, the user needs to download and install the necessary program (see Section 5.1). The last option, `opt$progLoc`, identifies the location of that program.

After all options have been set, the last line stores them in `DefaultOpt` so that the user can recall them at any point in the estimation. This is particularly useful if the user wants to change only a few options in between estimations.

3.3. Lag-order selection

Once the options are set, the user moves to the next step, which involves choosing the appropriate lag order. The relevant information is obtained with a call to `FCVARlagSelect()` which performs estimation of models with lag-orders from 0 to `kmax`. The program performs lag selection on the full-rank unrestricted model.

Our Matlab output is wider than 80 characters. This might sometimes be a problem. In particular, in the JSS draft, 80 characters in the `CodeChunk` environment will fit well within the margins. This is the revised format of the output.

```
R> LagSelectStats <- FCVARlagSelect(x1, kmax, p, order, opt)

--------------------------------------------------------------------------------
                        Lag Selection Results
--------------------------------------------------------------------------------
Dimension of system:       3     Number of observations in sample:          316
Order for WN tests:       12     Number of observations for estimation:     316
Restricted constant:      No     Initial values:                              0
Unrestricted constant:    No     Level parameter:                           Yes
--------------------------------------------------------------------------------
Parameter Estimates and Information Criteria:
--------------------------------------------------------------------------------
 k  r     d     b      LogL     LR    pv    AIC       BIC
 3  3 0.676 0.676  456.42   7.31 0.605  -832.85   -682.62
 2  3 0.581 0.581  452.77  20.59 0.015  -843.53*  -727.11
 1  3 1.043 1.043  442.47  56.99 0.000  -840.94   -758.31*
 0  3 1.036 1.036  413.97   0.00 0.000  -801.95   -753.12
--------------------------------------------------------------------------------
Tests for Serial Correlation of Residuals:
--------------------------------------------------------------------------------
 k   pmvQ  pQ1   pLM1  pQ2   pLM2  pQ3   pLM3
 3   0.94  0.72  0.46  0.49  0.89  0.51  0.47
 2   0.82  0.69  0.45  0.29  0.75  0.54  0.40
 1   0.34  0.75  0.52  0.15  0.58  0.34  0.18
 0   0.00  0.01  0.01  0.00  0.08  0.37  0.17
--------------------------------------------------------------------------------
```

Estimates of d and b are reported for each lag (k) with rank (r) set to the number of variables in the system. Note that in this example the restriction $d = b$ has been imposed. The log-likelihood for each lag is shown in column `LogL`. The likelihood ratio test-statistic `LR` is for the null hypothesis $\Gamma_k = 0$ with P-value reported in column `pv`. This is followed by AIC and BIC information criteria. The next set of columns provides P-values for white noise tests on the residuals. The first P-value, `pmvQ`, is for the multivariate Q-test followed by univarite Q-tests as well as LM tests on the p individual residuals; that is, `pQ1` and `pLM1` are the P-values for the residuals in the first equation, `pQ2` and `pLM2` are for the residuals in the second equation, and so on.

3.4. Cointegration rank testing

The user now chooses the lag-order based on the information provided above and can move to the next step, which is cointegration rank testing. In the next code block, the user first assigns the lag augmentation, $k = 2$ in this case, and then calls the function `RankTests()`.

```
R> k <- 2
R> rankTestStats <- RankTests(x1, k, opt)
```

```
--------------------------------------------------------------------------------
             Likelihood Ratio Tests for Cointegrating Rank
--------------------------------------------------------------------------------
Dimension of system:       3     Number of observations in sample:          316
Number of lags:            2     Number of observations for estimation:     316
Restricted constant:      No     Initial values:                              0
Unestricted constant:     No     Level parameter:                           Yes
--------------------------------------------------------------------------------
Rank     d      b     Log-likelihood   LR statistic   P-value
 0     0.643  0.643          440.040         25.454      ----
 1     0.569  0.569          451.174          3.186      ----
 2     0.576  0.576          452.707          0.120      ----
 3     0.581  0.581          452.767           ----      ----
--------------------------------------------------------------------------------
```

The first block of output provides a summary of the model specification. The second block provides the test results relevant for selecting the appropriate rank. The table is meant to be read sequentially from lowest to highest rank, i.e. from top to bottom. Since we can reject the null of rank 0 against the alternative of rank 3 we move to the test of rank 1 against rank 3. This test fails to reject with a P-value of 0.820, so this is the appropriate choice in this case.

3.5. Unrestricted model estimation

With the rank and lag selected, the user can now move to the next code section. Here the user first specifies the choice for the rank based on the previously performed cointegrating rank tests (thus setting $r = 1$ in this example). Next, the default options set in the initialization, see Section 3.2, are assigned to `opt1`, which is used as an argument in the call to the function `FCVARestn()`. This function is the main part of the program since it performs the estimation of the parameters, obtains model residuals and standard errors, and calculates many other relevant components such as the number of free parameters and the roots of the characteristic polynomial. If `opt1$print2screen <- 1` then, in addition to storing all of these results in the list `m1`, the function outputs the estimation results to the command window. To see a list of variables stored in `m1`, the user can type `m1` in the command line.

The program output is shown below. It begins with a table summarizing relevant model specifications and then the coefficients and their standard errors. The roots of the characteristic polynomial are displayed at the bottom.

I think we should trim this output to 80 characters wide. There are no numbers in that range, only comments.

```
R> r <- 1
R> opt1 <- DefaultOpt
R> m1 <- FCVARestn(x1, k, r, opt1)
```

```
------------------------------------------------------------------------------
                    Fractionally Cointegrated VAR: Estimation Results
------------------------------------------------------------------------------
Dimension of system:       3      Number of observations in sample:          316
Number of lags:            2      Number of observations for estimation:     316
Restricted constant:      No      Initial values:                              0
Unrestricted constant:    No      Level parameter:                           Yes
Starting value for d:    0.800    Parameter space for d: (0.010 , 2.000)
Starting value for b:    0.800    Parameter space for b: (0.010 , 2.000)
------------------------------------------------------------------------------
Cointegrating rank:            1  AIC:              -848.348
Log-likelihood:          451.174  BIC:              -746.943
log(det(Omega_hat)):     -11.369  Free parameters:        27
------------------------------------------------------------------------------
    Fractional parameters:
------------------------------------------------------------------------------
    Coefficient               Estimate                Standard error
------------------------------------------------------------------------------
         d                       0.569                     0.049
------------------------------------------------------------------------------
------------------------------------------------------------------------------
    Cointegrating equations (beta):
------------------------------------------------------------------------------
      Variable        CI equation 1
------------------------------------------------------------------------------
        Var1              1.000
        Var2              0.111
        Var3             -0.240
------------------------------------------------------------------------------
Note: Identifying restriction imposed.
------------------------------------------------------------------------------
    Adjustment matrix (alpha):
------------------------------------------------------------------------------
      Variable        CI equation 1
------------------------------------------------------------------------------
        Var 1            -0.180
         SE 1         (   0.064  )
        Var 2             0.167
         SE 2         (   0.194  )
        Var 3             0.037
         SE 3         (   0.014  )
------------------------------------------------------------------------------
Note: Standard errors in parenthesis.
------------------------------------------------------------------------------
    Long-run matrix (Pi):
------------------------------------------------------------------------------
      Variable         Var 1          Var 2          Var 3
```

```
--------------------------------------------------------------------------------
      Var 1           -0.180         -0.020          0.043
      Var 2            0.167          0.019         -0.040
      Var 3            0.037          0.004         -0.009
--------------------------------------------------------------------------------

--------------------------------------------------------------------------------
    Level parameter (mu):
--------------------------------------------------------------------------------
        Var 1            -0.345
         SE 1         (   0.069  )
        Var 2            11.481
         SE 2         (   0.548  )
        Var 3            -2.873
         SE 3         (   0.033  )
--------------------------------------------------------------------------------
Note: Standard errors in parenthesis (from numerical Hessian)
      but asymptotic distribution is unknown.
--------------------------------------------------------------------------------
    Lag matrix 1 (Gamma_1):
--------------------------------------------------------------------------------
      Variable         Var 1          Var 2          Var 3
--------------------------------------------------------------------------------
      Var 1            0.276         -0.032         -0.510
       SE 1        (   0.160  )   (   0.026  )   (   0.513  )
      Var 2           -0.148          1.126         -3.289
       SE 2        (   0.378  )   (   0.196  )   (   1.975  )
      Var 3           -0.052          0.008          0.711
       SE 3        (   0.022  )   (   0.005  )   (   0.170  )
--------------------------------------------------------------------------------
Note: Standard errors in parentheses.
--------------------------------------------------------------------------------
    Lag matrix 2 (Gamma_2):
--------------------------------------------------------------------------------
      Variable         Var 1          Var 2          Var 3
--------------------------------------------------------------------------------
      Var 1            0.566          0.106          0.608
       SE 1        (   0.182  )   (   0.045  )   (   0.612  )
      Var 2            0.493         -0.462          0.457
       SE 2        (   0.562  )   (   0.198  )   (   2.628  )
      Var 3           -0.039         -0.020          0.318
       SE 3        (   0.033  )   (   0.008  )   (   0.143  )
--------------------------------------------------------------------------------
Note: Standard errors in parentheses.
--------------------------------------------------------------------------------
--------------------------------------------------------------------------------
    Roots of the characteristic polynomial
```

```
------------------------------------------------------------------------------
    Number     Real part    Imaginary part       Modulus
------------------------------------------------------------------------------
       1         -2.893         -0.000            2.893
       2         -1.522         -0.000            1.522
       3          1.010         -0.927            1.371
       4          1.010          0.927            1.371
       5          1.107          0.000            1.107
       6          1.000          0.000            1.000
       7          1.000          0.000            1.000
       8          0.944         -0.261            0.980
       9          0.944          0.261            0.980
------------------------------------------------------------------------------

------------------------------------------------------------------------------
Restrictions imposed on the following parameters:
- Psi. For details see "options$R_psi"
------------------------------------------------------------------------------
```

At the end of the output, a notice is printed to remind the user that restrictions were imposed on `Psi`, i.e. on (d, b). In this case, this is the restriction $d = b$ imposed via `opt$restrictDB <- 1`.

In addition to the coefficient estimates, we are also interested in testing the model residuals for serial correlation. After the unrestricted model has been estimated, this code section concludes with a call to `MVWNtest()`, which performs a series of white noise tests on the residuals and prints the output in the command window. The results of the white noise tests are shown below. For each residual both the Q- and LM-test statistics and their P-values are reported, in addition to the multivariate Q-test and P-value in the first line of the table. From the output of this table we can conclude that there does not appear to be any problems with serial correlation in the residuals.

```
R> MVWNtest_m1 <- MVWNtest(m1$Residuals, order, printWNtest)

       White Noise Test Results (lag = 12)
-------------------------------------------
Variable |       Q  P-val |      LM  P-val  |
-------------------------------------------
Multivar |  97.879  0.747 |     ----  ----  |
Var1     |   9.300  0.677 |  11.238  0.509  |
Var2     |  14.450  0.273 |   8.568  0.739  |
Var3     |  10.591  0.564 |  12.265  0.425  |
-------------------------------------------
```

Because `opt$plotRoots <- 1` in the options, the roots of the characteristic polynomial is also plotted along with the unit circle and the transformed unit circle, $\mathbb{C}_{\hat{b}}$, see Johansen (2008). The plot is shown in Figure 1.

Example: A basic frequency distribution of the response variable is displayed in

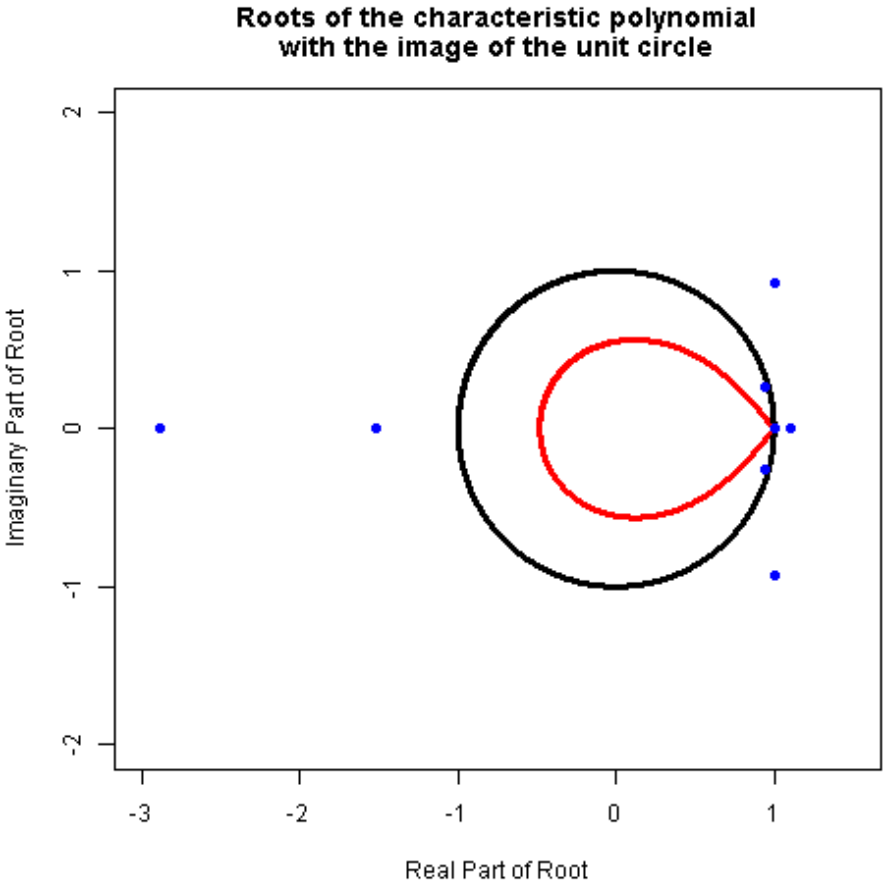


Figure 1: Roots of characteristic polynomial

Furthermore, the estimation was performed with the grid search and the plot option selected, i.e. with `opt$gridSearch <- 1` and `opt$plotLike <- 1`, which produces a plot of the log-likelihood. The plot for this model is shown in Figure 2.

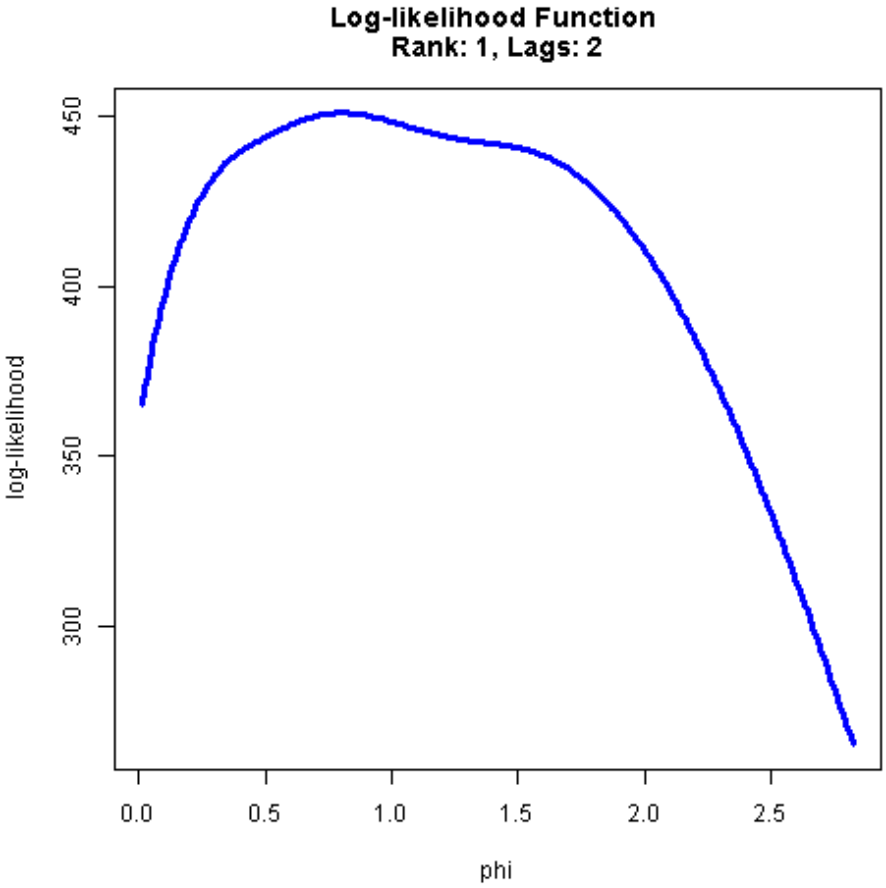


Figure 2: Plot of log-likelihood

The complete results for the unrestricted model are stored in the list `m1` and can be accessed anytime. For instance, if the user would like to perform a more careful analysis of the residuals they are stored in `m1$Residuals`.

3.6. Hypothesis testing

We now move into the hypothesis testing section of the code where we can test several restricted models and perform inference. For restricted model estimation the grid search option is switched off because computation can be very slow, especially in the presence of the level parameter. However, if the user wishes to verify the accuracy of the results or if estimates are close to the upper or lower bound, the grid search option can resolve these issues and give the user additional insight about the behaviour of the likelihood.

All hypotheses are defined as shown in (10)–(12). The first hypothesis test is $\mathscr{H}_d^1$ (for precise definitions of each hypothesis, please see Jones *et al.* (2014)), shown below.

```
R> opt1 <- DefaultOpt
R> opt1$R_psi <- matrix(c(1, 0), nrow = 1, ncol = 2)
R> opt1$r_psi <- 1
```

```
R> m1r1 <- FCVARestn(x1, k, r, opt1)
R> MVWNtest_m1r1 <- MVWNtest(m1r1$Residuals, order, printWNtest)
R> Hdb <- HypoTest(m1, m1r1)
```

Here we test the CVAR model (null hypothesis $d = b = 1$) against the FCVAR model (alternative hypothesis $d = b \neq 1$). Since `opt1$restrictDB <- 1` was selected in the choice of options, the restriction that $d = b$ is already imposed. Thus, the user needs to only impose an additional restriction that either d or b is equal to one. In this example, the restriction that $d = 1$ is imposed by setting `opt1.R_psi = [1 0]` and `opt1.r_psi = 1`, but the result would be the same if $b = 1$ were imposed instead. The restricted model is then estimated and the results are stored in the Matlab structure `m1r1`. As before, the user can perform a series of white noise tests on the residuals by calling the `mv_wntest.m` function. The next step is to perform the actual test. With the results structures from the restricted and unrestricted models, the user can call the function `HypoTest.m` and perform an LR test. This function takes the two model result structures as inputs, automatically compares the number of free parameters to obtain the degrees of freedom, computes the LR test statistic, and displays the output. The results of this test are then stored in the list `Hdb` and can be accessed at any time.

Since the output of the estimated model and the white noise tests are similar to the previous example, we only show the output from the hypothesis test.

```
Unrestricted log-likelihood: 451.174
Restricted log-likelihood:   442.027
Test results (df = 1):
LR statistic:   18.295
P-value:   0.000
```

The log-likelihoods from both models are reported, along with the degrees of freedom, the LR test statistic, and its P-value. In this case the test clearly rejects the null hypothesis that the model is a CVAR. For more significant digits, or to access any of these values from the command window, the user can type `Hdb`.

The next hypothesis of interest is $\mathscr{H}_\beta^1$, which is a zero restriction on the first element of the cointegration vector.

```
R> opt1 <- DefaultOpt
R> opt1$R_Beta <- matrix(c(1, 0, 0), nrow = 1, ncol = 3)
R> m1r2 <- FCVARestn(x1, k, r, opt1)
R> MVWNtest_m1r2 <- MVWNtest(m1r2$Residuals, order, printWNtest)
R> Hbeta1 <- HypoTest(m1, m1r2)
```

Since the object `opt1` has the restriction $d = b = 1$ stored, the first step is to reset the options to default. The restriction on β is then specified as in (12). There are two things to note here. First, the column length of R_β must equal $p_1 r$, where $p_1 = p + 1$ if a restricted constant is present and $p_1 = p$ otherwise; recall that p is the number of variables in the system and r is the number of cointegrating vectors. Second, zero restrictions are the default and automatically imposed when r_β is empty. Therefore, the user only needs to specify r_β

if it includes non-zero elements. Recall that for restrictions on α only $r_\alpha = 0$ is allowed so that there is no need to specify r_α. As before, the restricted model is estimated with results stored in `m1r2`, the residuals are tested for white noise, and the model under the null is tested against the unrestricted model `m1` with results stored in `Hbeta1`.

Again, since the estimation output is similar to the first example, we only show the results of the hypothesis test here. With a P-value close to zero, this hypothesis is also strongly rejected.

```
Unrestricted log-likelihood: 451.174
Restricted log-likelihood:   444.395
Test results (df = 1):
LR statistic:   13.557
P-value:   0.000
```

Next, we move to tests on α. In this case, we test restriction that political variable is long-run exogenous.

```
R> opt1 <- DefaultOpt
R> opt1$R_Alpha <- matrix(c(1, 0, 0), nrow = 1, ncol = 3)
R> opt1$gridSearch <- 0
R> m1r3 <- FCVARestn(x1, k, r, opt1)
R> MVWNtest_m1r3 <- MVWNtest(m1r3$Residuals, order, printWNtest)
R> Halpha1 <- HypoTest(m1, m1r3)
```

Again we first reset `opt1` to the default options to clear previously imposed restrictions. Note that, if it were the case that we failed to reject $\mathscr{H}_\beta^1$ and wanted to leave it imposed while adding a restriction on α, we could either omit the first line `opt1 <- DefaultOpt`, or we could replace it with `opt1 <- m1r2$options`. The latter assignment is preferred in this case because it is explicit about which model options we are leaving imposed.

The hypothesis $\mathscr{H}_\alpha^1$ is tested in the exact same way as before, only now we are changing the variable R_α instead of R_β. The results are shown below and we can see that this hypothesis is also rejected.

```
Unrestricted log-likelihood: 451.174
Restricted log-likelihood:   446.086
Test results (df = 1):
LR statistic:   10.176
P-value:   0.001
```

We next move to the remaining long-run exogeneity tests, $\mathscr{H}_\alpha^2$ and $\mathscr{H}_\alpha^3$, shown the examples below. The hypothesis $\mathscr{H}_\alpha^2$ tests that the interest-rate is long-run exogenous.

```
R> opt1 <- DefaultOpt
R> opt1$R_Alpha <- matrix(c(0, 1, 0), nrow = 1, ncol = 3)
R> opt1$gridSearch <- 0
R> m1r4 <- FCVARestn(x1, k, r, opt1)
R> MVWNtest_m1r4 <- MVWNtest(m1r4$Residuals, order, printWNtest)
R> Halpha2 <- HypoTest(m1, m1r4)
```

```
Unrestricted log-likelihood: 451.174
Restricted log-likelihood:   450.857
Test results (df = 1):
LR statistic:          0.633
P-value:           0.426
```

Next, we test the hypothesis $\mathscr{H}_\alpha^3$ that unemployment is long-run exogenous.

```
R> opt1 <- DefaultOpt
R> opt1$gridSearch <- 0
R> opt1$R_Alpha <- matrix(c(0, 0, 1), nrow = 1, ncol = 3)
R> m1r5 <- FCVARestn(x1, k, r, opt1)
R> MVWNtest_m1r5 <- MVWNtest(m1r5$Residuals, order, printWNtest)
R> Halpha3 <- HypoTest(m1, m1r5)

Unrestricted log-likelihood: 451.174
Restricted log-likelihood:   446.184
Test results (df = 1):
LR statistic:          9.979
P-value:           0.002
```

The only hypothesis that we fail to reject is $\mathscr{H}_\alpha^2$, under which interest rates are long-run exogenous. After having estimated all of the restricted models of interest, we provide the full estimation output for the model `m1r4`, with the restriction imposed for $\mathscr{H}_\alpha^2$. Note from the output that $\alpha_2 = 0$ as imposed by the restriction.

```
--------------------------------------------------------------------------------
                     Fractionally Cointegrated VAR: Estimation Results
--------------------------------------------------------------------------------
Dimension of system:       3      Number of observations in sample:          316
Number of lags:            2      Number of observations for estimation:     316
Restricted constant:      No      Initial values:                              0
Unrestricted constant:    No      Level parameter:                           Yes
Starting value for d:    0.800    Parameter space for d: (0.010 , 2.000)
Starting value for b:    0.800    Parameter space for b: (0.010 , 2.000)
--------------------------------------------------------------------------------
Cointegrating rank:            1  AIC:              -849.715
Log-likelihood:          450.857  BIC:              -752.065
log(det(Omega_hat)):     -11.367  Free parameters:        26
--------------------------------------------------------------------------------
    Fractional parameters:
--------------------------------------------------------------------------------
    Coefficient               Estimate                Standard error
--------------------------------------------------------------------------------
         d                       0.575                     0.048
--------------------------------------------------------------------------------
```

```
------------------------------------------------------------------------------
    Cointegrating equations (beta):
------------------------------------------------------------------------------
      Variable        CI equation 1
------------------------------------------------------------------------------
        Var1              0.994
        Var2              0.105
        Var3             -0.181
------------------------------------------------------------------------------
    Adjustment matrix (alpha):
------------------------------------------------------------------------------
      Variable        CI equation 1
------------------------------------------------------------------------------
        Var 1            -0.189
         SE 1         (   0.065  )
        Var 2             0.000
         SE 2         (   0.000  )
        Var 3             0.039
         SE 3         (   0.014  )
------------------------------------------------------------------------------
Note: Standard errors in parenthesis.
------------------------------------------------------------------------------
    Long-run matrix (Pi):
------------------------------------------------------------------------------
      Variable         Var 1          Var 2          Var 3
------------------------------------------------------------------------------
      Var 1           -0.188         -0.020          0.034
      Var 2            0.000          0.000          0.000
      Var 3            0.039          0.004         -0.007
------------------------------------------------------------------------------

------------------------------------------------------------------------------
    Level parameter (mu):
------------------------------------------------------------------------------
        Var 1            -0.310
         SE 1         (   0.067  )
        Var 2            11.538
         SE 2         (   0.553  )
        Var 3            -2.874
         SE 3         (   0.033  )
------------------------------------------------------------------------------
Note: Standard errors in parenthesis (from numerical Hessian)
      but asymptotic distribution is unknown.
------------------------------------------------------------------------------
    Lag matrix 1 (Gamma_1):
------------------------------------------------------------------------------
      Variable         Var 1          Var 2          Var 3
```

```
------------------------------------------------------------------------------
      Var 1            0.269          -0.032          -0.511
       SE 1        (   0.157  )   (   0.026  )   (   0.507  )
      Var 2           -0.013           1.115          -3.004
       SE 2        (   0.345  )   (   0.189  )   (   1.909  )
      Var 3           -0.053           0.008           0.694
       SE 3        (   0.022  )   (   0.005  )   (   0.164  )
------------------------------------------------------------------------------
Note: Standard errors in parentheses.
------------------------------------------------------------------------------
    Lag matrix 2 (Gamma_2):
------------------------------------------------------------------------------
      Variable         Var 1           Var 2           Var 3
------------------------------------------------------------------------------
      Var 1            0.570           0.104           0.585
       SE 1        (   0.184  )   (   0.044  )   (   0.606  )
      Var 2            0.685          -0.371           0.229
       SE 2        (   0.508  )   (   0.159  )   (   2.509  )
      Var 3           -0.043          -0.020           0.330
       SE 3        (   0.032  )   (   0.008  )   (   0.138  )
------------------------------------------------------------------------------
Note: Standard errors in parentheses.
------------------------------------------------------------------------------
------------------------------------------------------------------------------
    Roots of the characteristic polynomial
------------------------------------------------------------------------------
    Number     Real part    Imaginary part       Modulus
------------------------------------------------------------------------------
       1         -2.710          -0.000           2.710
       2         -1.498          -0.000           1.498
       3          1.130          -0.939           1.469
       4          1.130           0.939           1.469
       5          1.098           0.000           1.098
       6          1.000           0.000           1.000
       7          1.000           0.000           1.000
       8          0.934          -0.281           0.976
       9          0.934           0.281           0.976
------------------------------------------------------------------------------

------------------------------------------------------------------------------
Restrictions imposed on the following parameters:
- Psi. For details see "options$R_psi"
- Alpha. For details see "options$R_Alpha"
------------------------------------------------------------------------------


       White Noise Test Results (lag = 12)
```

```
-------------------------------------------
Variable |       Q  P-val |      LM  P-val  |
-------------------------------------------
Multivar |  97.674  0.752 |     ----  ----  |
Var1     |   9.083  0.696 |  11.268  0.506  |
Var2     |  14.937  0.245 |   9.339  0.674  |
Var3     |  10.725  0.553 |  12.237  0.427  |
-------------------------------------------
```

Sometimes it is the case that the model output is not normalized with respect to the user's variable of interest, for example when restrictions are imposed on α or β. For this reason, we also include a code section that normalizes the output, i.e. imposes an identity matrix in the first $r \times r$ block of β. That is, $\hat{\alpha}$ is post multiplied by G^{-1} so that $\pi = \hat{\alpha}(G^{-1})G\hat{\beta}' = ab'$[8]. Of course, this code section should only be executed if it does not interfere with any restrictions imposed on the model.

```
R> modelRstrct <- m1r4
R> G <- solve(modelRstrct$coeffs$betaHat[1:r, 1:r])
R> betaHatR <- modelRstrct$coeffs$betaHat %*% G
R> alphaHatR <- modelRstrct$coeffs$alphaHat %*% t(solve(G))
R> print('betaHatR = ')
R> print(betaHatR)
R> print('alphaHatR = ')
R> print(alphaHatR)

[1] "betaHatR = "
           [,1]
[1,]  1.0000000
[2,]  0.1056815
[3,] -0.1822645
[1] "alphaHatR = "
            [,1]
[1,] -0.18773922
[2,]  0.00000000
[3,]  0.03855402
```

As an example of when this feature can be useful, consider model $\mathscr{H}_\alpha^2$. In the output above, we notice that the cointegrating vector has not been normalized (because restrictions are imposed). The user assigns the model of interest to the variable `modelRstrct`, in this case `m1r4`, and executes the cell. The output is shown below.

4. Additional examples: `MoreExamples.R`

To show some additional functionality of the FCVAR software package, this section contains several other examples, which are based on Jones *et al.* (2014), but are not part of that paper.

[8]Let's revise this sentence to account for the changing definition of $\hat{\alpha}$ and $\hat{\beta}$

4.1. Forecasting

This code block performs recursive one-step ahead forecasts for each of the variables as well as the equilibrium relation.

```
R> NumPeriods <- 12
R> modelF <- m1r4
R> xf <- FCVARforecast(x1, modelF, NumPeriods)
R> seriesF <- rbind(x1, xf)
R> equilF <- seriesF %*% modelF$coeffs$betaHat
```

The user specifies the forecast horizon (`NumPeriods`) as well as the model (in this case, `modelF <- m1r4`). These two inputs, along with the data, are used in the call to the function `FCVARforecast()`. This function returns `xf`, a `NumPeriods` by p matrix of forecasted values of X, which are forecasted to take place after `x1`. Figure 3 plots the original series and Figure 4 plots the equilibrium relation $X\hat{\beta}$, stored in `equilF`, along with the forecasts.

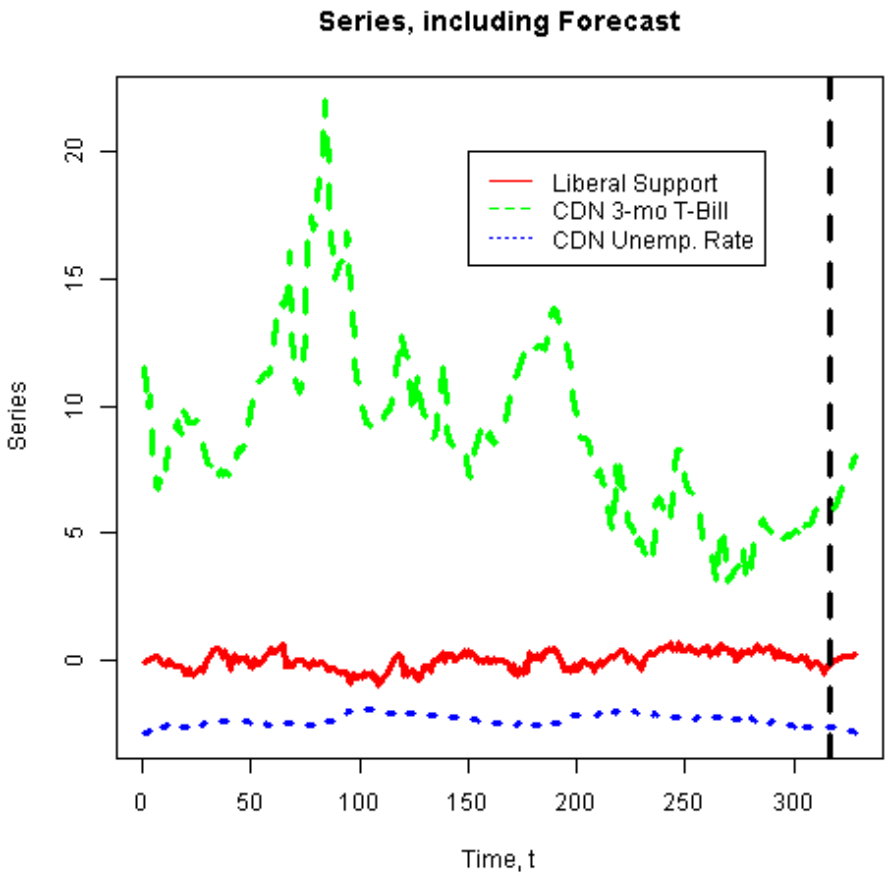


Figure 3: Forecast of final model 12 steps ahead

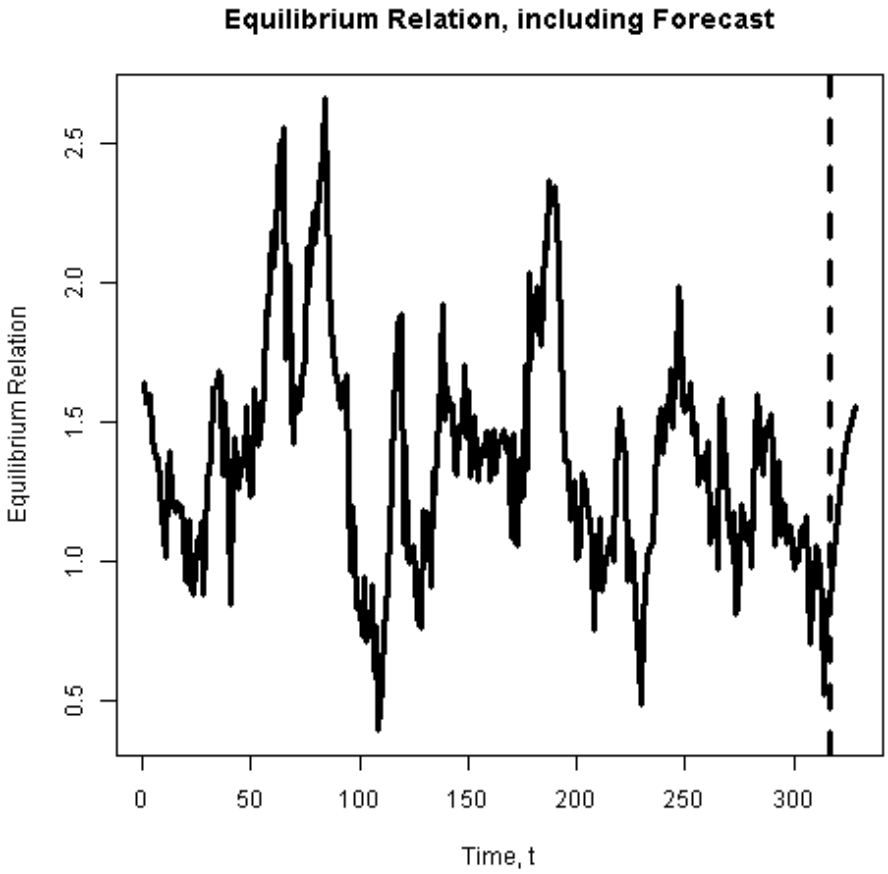


Figure 4: Forecast equilibrium relationship of final model 12 steps ahead

4.2. Bootstrap hypothesis test

This code block demonstrates the use of the wild bootstrap for hypothesis tests on the parameters, as developed by Boswijk, Cavaliere, Rahbek, and Taylor (2016) for the CVAR model. The function `FCVARboot()` returns the results of the wild bootstrap. The user specifies two sets of options corresponding to two different nested models, the restricted model with `optRES` and the unrestricted model with `optUNR`. This particlar example tests the restriction that political variables do not enter the cointegrating relation(s).

```
R> DefaultOpt$plotRoots <- 0
R> optUNR <- DefaultOpt
R> optRES <- DefaultOpt
R> optRES$R_Beta <- matrix(c(1, 0, 0), nrow = 1, ncol = 3)
R> set.seed(42)
R> FCVARboot_stats <- FCVARboot(x1, k, r, optRES, optUNR, B = 999)
R> LRbs_density <- density(FCVARboot_stats$LRbs)
```

An example of the output is

```
Bootstrap results:
Unrestricted log-likelihood: 451.174
Restricted log-likelihood:   444.395
Test results (df <- 1):
LR statistic:   13.557
P-value:   0.000
P-value (BS):   0.021
```

The user might also be interested in comparing the bootstrap likelihood ratio test statistic distribution to the asymptotic one, a χ-squared distribution with `H.df` degrees of freedom. These objects can be used to produce a plot of the two distributions, shown in Figure 5.

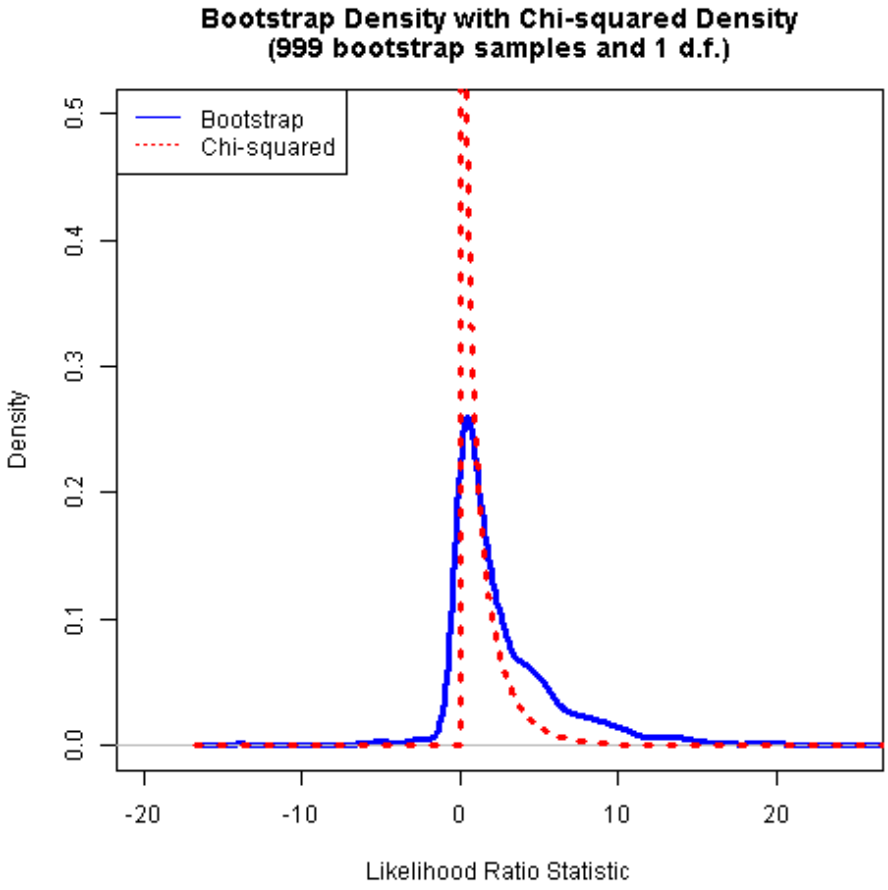


Figure 5: Density of bootstrap LR test statistic

4.3. Bootstrap rank test

This code block shows how to perform a wild bootstrap rank test, following the methodology of Cavaliere, Rahbek, and Taylor (2010) for the CVAR model. This procedure works in much the same way as the bootstrap hypothesis test described in Section 4.2. The difference is that, instead of providing two sets of estimation options, the user specifies two different ranks

for comparison.

```
R> r1 <- 0
R> r2 <- 1
R> FCVARbootRank_stats <- FCVARbootRank(x1, k, DefaultOpt, r1, r2, B = 999)
R> cat(sprintf('P-value: \t %1.3f\n', rankTestStats$pv[1]))
```

The results are printed as

```
Bootstrap results:
Unrestricted log-likelihood: 451.174
Restricted log-likelihood:   440.040
Test results:
LR statistic:   22.268
P-value (BS):   0.031
# Need to modify program:
P-value:   0.043
```

in which the last P-value is printed to compare the bootstrap P-value to that based on the asymptotic distribution.

4.4. Simulation

Finally, this example shows how to simulate an FCVAR model for a given set of parameters. The user provides data for starting values and a list containing model parameters for simulation as well as the number of periods to simulate. The simulated data are generated using Gaussian errors.

```
R> T_sim <- 100
R> xSim <- FCVARsim(x1, modelF, T_sim)
```

For the example above, using the same data as for the forecasting example above, the generated data is shown in Figure 6.

5. Extensions

5.1. Extension for P-values

Although the Matlab program can run standalone, one of the functions, `RankTests.m`, makes an external system call to a separately installed program, `fdpval`. This external program is the C++ implementation of a Fortran program used to obtain simulated P-values from MacKinnon and Nielsen (2014). If the user would like P-values for the cointegration rank tests to be automatically calculated, we recommend obtaining this companion program, which is made available by Jason Rhinelander and can be downloaded from:

https://github.com/jagerman/fracdist/releases

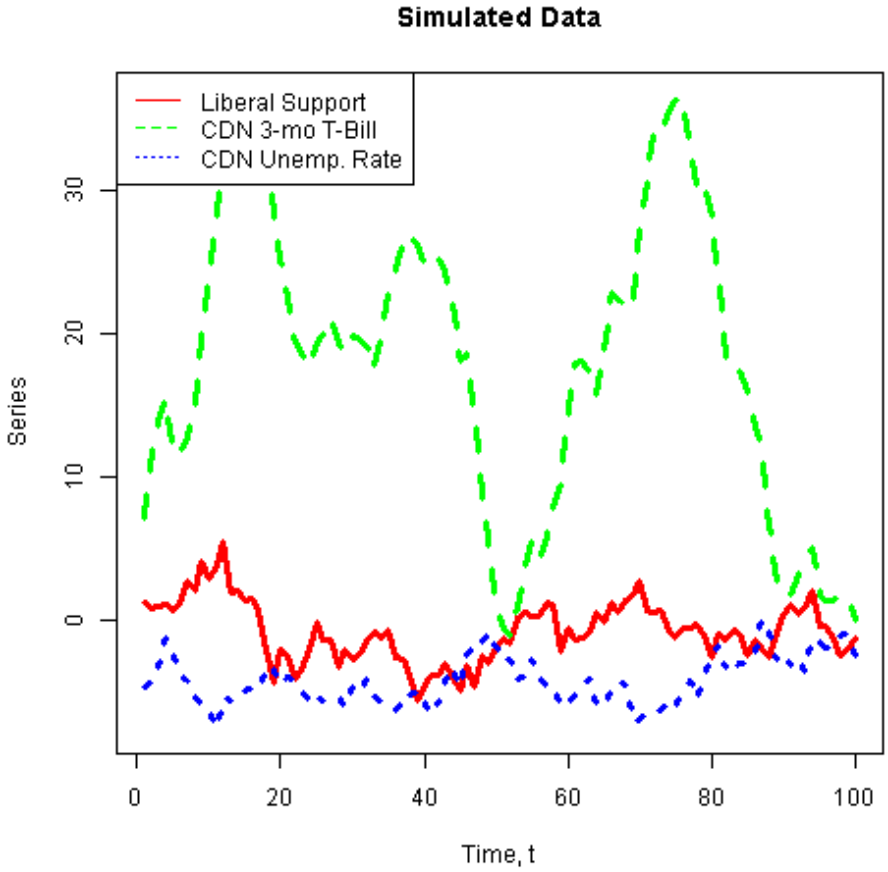


Figure 6: Simulated data

It can be either installed or downloaded in a compressed folder. It is important to note where the program is stored or installed, because the Matlab program requires the program location as an input in the estimation options. For example, if the program is stored in the folder `/usr/bin/` on a Linux system, the location variable is defined as follows, `progLoc = '"/usr/bin/fdpval"'`. For details see Sections ?? and ??.

5.2. Badly behaved objective function

We also make use of the excellent `extrema.m` and `extrema2.m` functions, which are written by Carlos Adrián Vargas Aguilera and are freely available from the Mathworks website. For simplicity these are included in the Auxiliary subfolder.

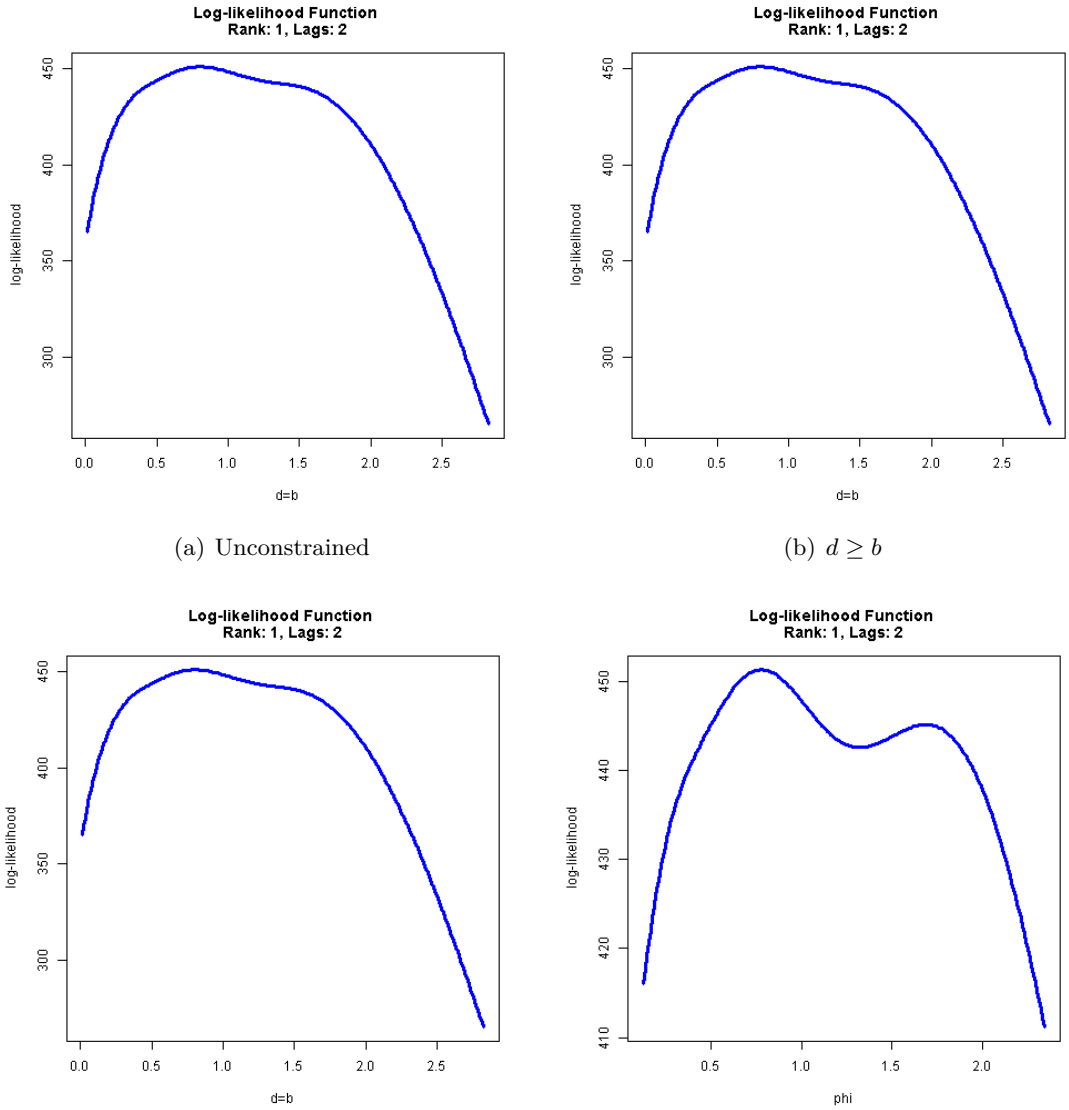
> The above functions are currently not used in the R package. Can we provide a use case in which the functions correct a problem or improve efficiency in the optimization?

These functions are used in `LikeGrid.m()`, which allows the user to pre-estimate to obtain starting values by using a grid search. There are four types of estimation that the grid search can perform. If d and b are completely unconstrained, the grid search is over two

dimensions within the bounds specified by `opt.dbMin` and `opt.dbMax`. An example of the likelihood obtained in an unconstrained grid search is shown in Figure 7(a). Next, if $d \geq b$ is imposed via `opt.constrained = 1` (imposed in Johansen and Nielsen (2012) but relaxed in Johansen and Nielsen (2018b)) the computation can be cut in half. An example of this likelihood is shown in Figure 7(b). If the restriction $d = b$ is imposed, then the grid search is one-dimensional as shown in Figure 7(c). Finally, if a restriction is imposed on either d or b via R_ψ and r_ψ in (10), then the grid search is also one-dimensional. An example of this situation is shown in Figure 7(d). Note that the x-axis is over the parameter ϕ and the fractional parameters are found from

$$\begin{bmatrix} d \\ b \end{bmatrix} = H\phi + h, \tag{16}$$

where $H = (R'_\psi)_\perp$ and $h = R'_\psi(R_\psi R'_\psi)^{-1} r_\psi$. The bounds on ϕ are derived from `opt.dbMin` and `opt.dbMax` in a similar way.



(a) Unconstrained

(b) $d \geq b$

(c) $d = b$

(d) Restrictions imposed

Figure 7: Grid search

6. Summary and discussion

Summary goes here.

Computational details

The results in this paper were obtained using R 3.4.1 with the **MASS** 7.3.47 package. R itself and all packages used are available from the Comprehensive R Archive Network (CRAN) at https://CRAN.R-project.org/.

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A. More technical details

Technical details go here.

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