

$$\text{ECM: } \nabla \begin{bmatrix} X_t \\ Y_t \end{bmatrix} = \begin{bmatrix} C_1 \\ C_2 \end{bmatrix} + \begin{bmatrix} -2\rho_1 & \rho_2 \\ -2\rho_2 & \rho_2 \end{bmatrix} \begin{bmatrix} X_{t-1} \\ Y_{t-1} \end{bmatrix} + \sum_i^n \begin{bmatrix} \beta_{x,i} & \beta_{y,i} \\ r_{x,i} & r_{y,i} \end{bmatrix} \nabla \begin{bmatrix} X_{t-1} \\ Y_{t-1} \end{bmatrix} + \begin{bmatrix} \epsilon_{x,t} \\ \epsilon_{y,t} \end{bmatrix}$$

$\begin{bmatrix} -\hat{\alpha} \\ -\hat{\alpha} \end{bmatrix} [\rho_1, \rho_2]$

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## \* State-space model (SSM)

State-space models are useful tools for expressing dynamic systems that involve w/ unobserved state variables.

Two eqns:

(1) Measurement eqn (observation eqn.)

→ describes the relation b/w observed variables (charts) and unobserved state variable.

(2) Transition eqn. (state eqn.)

→ describes the dynamics of the state variables.

The transition equation has the form of

a first-order difference equation in the state vector.

$$(1) \quad y_t = F \tilde{x}_t + \epsilon_t$$

$$(2) \quad \tilde{x}_t = G \tilde{x}_{t-1} + w_t$$

$y_t$  = univariate time series;  $t=1, \dots, T$ .

$\tilde{x}_{t-1}$  = state vector,  $\tilde{x}_t \sim \text{VAR}(1)$ .

$F_{p \times p}, G_{p \times p}$  = coefficient matrices

$\epsilon_t \sim N(0, \sigma^2)$

$w_t \sim N_p(0, W)$ .

e.g.  $y_t \sim AR(2)$ .

$$\Phi_2(\beta)y_t = a_t, \quad a_t \sim N(0, \sigma^2).$$

$$(1) \tilde{y}_t = F \tilde{x}_t + \varepsilon_t \quad \varepsilon_t \sim N(0, \sigma^2) \quad 81$$

$$(2) \tilde{x}_t = G \tilde{x}_{t-1} + w_t \quad w_t \sim N_p(\vec{0}, W)$$

~~$\tilde{x}_t \sim \text{VAR}(1)$~~   $\tilde{x}_t \sim \text{VAR}(1)$   $W = E(W_t W_t')$

$$y_t = \phi_1 y_{t-1} + \phi_2 y_{t-2} + a_t. \Rightarrow (y_{t-1} = \phi_1 y_{t-2} + \phi_2 y_{t-3} + a_{t-1})$$

$$= [\phi_1 \phi_2] [y_{t-1}] + a_t$$

$$\begin{bmatrix} y_{t-1} \\ y_{t-2} \end{bmatrix} = [\phi_1 \phi_2] \begin{bmatrix} y_{t-2} \\ y_{t-3} \end{bmatrix} + \begin{bmatrix} a_{t-1} \\ 0 \end{bmatrix}$$

$$y_t = \phi_1 y_{t-1} + \phi_2 y_{t-2} + a_t.$$

WAY(1) Define  $\tilde{x}_t = \begin{bmatrix} y_t \\ y_{t-1} \end{bmatrix}$ .

$$\text{Then } y_t = [1 \ 0] \begin{bmatrix} y_t \\ y_{t-1} \end{bmatrix},$$

$$\begin{bmatrix} y_t \\ y_{t-1} \end{bmatrix} = [\phi_1 \phi_2] \begin{bmatrix} y_{t-1} \\ y_{t-2} \end{bmatrix} + \begin{bmatrix} a_t \\ 0 \end{bmatrix}$$

WAY(2) Define  $\tilde{x}_t = \begin{bmatrix} y_t \\ \phi_2 y_{t-1} \end{bmatrix}$

$$y_t = [1 \ 0] \begin{bmatrix} y_t \\ \phi_2 y_{t-1} \end{bmatrix},$$

$$\begin{bmatrix} y_t \\ \phi_2 y_{t-1} \end{bmatrix} = [\phi_1 \ 1] \begin{bmatrix} y_{t-1} \\ y_{t-2} \end{bmatrix} + \begin{bmatrix} a_t \\ 0 \end{bmatrix}.$$

e.g.  $y_t \sim \text{ARIMA}(1,1)$ .

$$y_t - \phi_1 y_{t-1} = a_t + \theta_1 a_{t-1}$$

$$\Rightarrow y_t = \phi_1 y_{t-1} + a_t + \theta_1 a_{t-1}$$

Define  $\tilde{x}_t = \begin{bmatrix} y_t \\ y_{t-1} \end{bmatrix}$ .

~~$$\text{Then } y_t = [\phi_1 \ 0] \begin{bmatrix} y_t \\ y_{t-1} \end{bmatrix} + a_t + \theta_1 a_{t-1},$$~~

~~$$\begin{bmatrix} y_t \\ y_{t-1} \end{bmatrix} = [\phi_1 \ 0] \begin{bmatrix} y_{t-1} \\ y_{t-2} \end{bmatrix} + \begin{bmatrix} a_t \\ 0 \end{bmatrix}$$~~

$y_t \sim ARIMA(1, 1)$

$$y_t = \phi_1 y_{t-1} + \theta_1 a_{t-1} + a_t$$

$$\begin{bmatrix} \tilde{x}_t \\ \tilde{a}_{t-1} \end{bmatrix} = \begin{bmatrix} y_t \\ a_{t-1} \end{bmatrix}, \quad \begin{bmatrix} \tilde{w}_t \\ \tilde{a}_t \end{bmatrix} = \begin{bmatrix} a_t \\ a_t \end{bmatrix}$$

$$y_t = [1 \ 0] \begin{bmatrix} \tilde{x}_t \\ \tilde{a}_{t-1} \end{bmatrix}$$

$$\begin{bmatrix} y_t \\ a_{t-1} \end{bmatrix} = \begin{bmatrix} \phi_1 & 1 \\ 0 & 0 \end{bmatrix} \begin{bmatrix} y_{t-1} \\ a_{t-1} \end{bmatrix} + \begin{bmatrix} a_t \\ a_t \end{bmatrix}$$

Combining two SSM

$$y_t = F_1 \tilde{x}_t + F_2 \tilde{z}_t + \varepsilon_t,$$

$$\tilde{x}_t = G_1 \tilde{x}_{t-1} + \tilde{w}_t$$

$$\tilde{z}_t = G_2 \tilde{z}_{t-1} + \tilde{u}_t$$

$$\text{Combined SSM: } y_t = [F_1 \ F_2] \begin{bmatrix} \tilde{x}_t \\ \tilde{z}_t \end{bmatrix} + \varepsilon_t,$$

$$\begin{bmatrix} \tilde{x}_t \\ \tilde{z}_t \end{bmatrix} = \begin{bmatrix} G_1 & 0 \\ 0 & G_2 \end{bmatrix} \begin{bmatrix} \tilde{x}_{t-1} \\ \tilde{z}_{t-1} \end{bmatrix} + \begin{bmatrix} \tilde{w}_t \\ \tilde{u}_t \end{bmatrix}$$

SSM for reg. model with AR( $\varepsilon$ ) error.

$$y_t = \alpha + \beta f_t + \eta_t, \quad \eta_t \sim AR(\varepsilon),$$

$$(\eta_t = \phi_1 \eta_{t-1} + \phi_2 \eta_{t-2} + a_t)$$

Let  $\tilde{x}_t = \begin{bmatrix} \alpha \\ f_t \end{bmatrix}$ . Then  $\tilde{x}_t = \tilde{z}_t$ .

$$\text{So } y_t = [1 \ f_t] \begin{bmatrix} \alpha \\ f_t \end{bmatrix} + \eta_t.$$

$$\text{Also, } \tilde{z}_t = \begin{bmatrix} \eta_t \\ \phi_1 \eta_{t-1} \end{bmatrix} \Rightarrow \eta_t = [1 \ 0] \begin{bmatrix} \eta_t \\ \phi_1 \eta_{t-1} \end{bmatrix}$$

$$\text{and } \tilde{z}_t = \begin{bmatrix} \phi_1 & 0 \\ 0 & 0 \end{bmatrix} \tilde{z}_{t-1} + \begin{bmatrix} a_t \\ 0 \end{bmatrix}$$

$$y_t = [1 \ f_t] \begin{bmatrix} \alpha \\ \beta \end{bmatrix} + [1 \ 0] \begin{bmatrix} \eta_t \\ \phi_1 \eta_{t-1} \end{bmatrix}$$

$$\tilde{x}_t = \begin{bmatrix} \alpha \\ \beta \end{bmatrix} = \begin{bmatrix} 1 & 0 \\ 0 & 1 \end{bmatrix} \begin{bmatrix} \alpha \\ \beta \end{bmatrix} + \begin{bmatrix} 0 \\ 0 \end{bmatrix},$$

$$\tilde{z}_t = \begin{bmatrix} \eta_t \\ \phi_1 \eta_{t-1} \end{bmatrix} = \begin{bmatrix} \phi_1 & 1 \\ 0 & 0 \end{bmatrix} \begin{bmatrix} \eta_{t-1} \\ \phi_2 \eta_{t-2} \end{bmatrix} + \begin{bmatrix} a_t \\ 0 \end{bmatrix}$$

$$\Rightarrow y_t = [F_1' \ F_2'] \begin{bmatrix} \tilde{x}_t \\ \tilde{z}_t \end{bmatrix} = [1 \ f_t \ 1 \ 0] \begin{bmatrix} \alpha \\ \beta \\ \eta_t \\ \phi_1 \eta_{t-1} \end{bmatrix}$$

$$\begin{bmatrix} \tilde{x}_t \\ \tilde{z}_t \end{bmatrix} = \begin{bmatrix} G_1 & 0 \\ 0 & G_2 \end{bmatrix} \begin{bmatrix} \tilde{x}_{t-1} \\ \tilde{z}_{t-1} \end{bmatrix} + \begin{bmatrix} \vec{0} \\ a_t \end{bmatrix}$$

$$\begin{bmatrix} \alpha \\ \beta \\ \eta_t \\ \phi_1 \eta_{t-1} \end{bmatrix} = \begin{bmatrix} 1 & 0 & 0 & 0 \\ 0 & 1 & 0 & 0 \\ 0 & 0 & \phi_1 & 1 \\ 0 & 0 & \phi_2 & 0 \end{bmatrix} \begin{bmatrix} \alpha \\ \beta \\ \eta_{t-1} \\ \phi_2 \eta_{t-2} \end{bmatrix} + \begin{bmatrix} 0 \\ 0 \\ a_t \\ 0 \end{bmatrix}$$

STA355:  
 Estimate  
 the sampling  
 dist. of  $\hat{\theta}$   
 by sampling  
 w/ replacement

### \* Bootstrap

- Replicating an experiment by resampling from observed data
- Used to estimate bias, se, percentiles, or other properties of a statistic

from the data and then computing estimates from these samples.

### General approach

- Draw repeated samples w/ replacement
- Since TS obs aren't iid, the method needs to be modified.

e.g. library(boot), bigcity

→ population (in 1000s) of 49 U.S. cities in 1920 (u) and 1930 (x).

→ 49 cities are among the biggest 196 cities in the U.S. in 1920. Randomly chosen.

$$\begin{aligned} X &= U = \text{pop. in 1920} \\ U &= \text{pop. in 1930} \end{aligned}$$

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→ parameter of interest:  $\theta = E(X)/E(U)$ .

$$\text{Let } \theta = \bar{X}/\bar{U}.$$

So how "uncertain" or "certain" are we about  $\theta$ ?

> `dat <- bigcity[1:4, ]`

> `dat`

|   | U   | X   |
|---|-----|-----|
| 1 | 138 | 143 |
| 2 | 93  | 104 |
| 3 | 61  | 69  |
| 4 | 179 | 260 |

> `set.seed(1234)`

> `i <- sample(1:4, size=4, replace=T)`

> `i`

[1] 1 3 3 3

> `dat[i, ]`

|   | U   | X   |
|---|-----|-----|
| 1 | 138 | 143 |
| 3 | 61  | 69  |
| 3 | 61  | 69  |
| 3 | 61  | 69  |

> `ratio <- function(X) { mean(X[2])/mean(X[1]) }`

> `n <- nrow(bigcity)`

> `n`

[1] 49

> `t0 <- ratio(bigcity)`

> `t0`

[1] 1.23909

```

> set.seed(1234)
> t.replicates <- replicate(1000, expr = {
+ i <- sample(1:n, size=n, replace=T)
+ boots.obs <- bigcity[i,]
+ ratio(boots.obs)
+ })
> library(MASS)
> triwheelise(t.replicates) →
> points(t0, 0, pch=17, col=2) ←
    t.replicates
    ↓ has appeared.

```

$$\widehat{\text{Var}}(\hat{\theta}) = \frac{1}{R-1} \sum_j^R (\hat{\theta}_j - \bar{\theta}^*)^2$$

$\bar{\theta}^*$  is the arithmetic mean of the replicates.

```

> mean(t.replicates)
[1] 1.241886
> se.hat <- sd(t.replicates)
> se.hat
[1] 0.03635716

```

$$\text{Bias}(\hat{\theta}) = E(\hat{\theta}) - \theta. \quad \text{And } \widehat{\text{Bias}}(\hat{\theta}) = \bar{\theta}^* - \hat{\theta}.$$

```

> bias.hat <- mean(t.replicates) - t0
> bias.hat
[1] 0.002867242

```

- boot function (in boot package)

boot(data, statistic, R)

matrix or  
data.frame

name of a function to compute the statistic of interest # of replicates to generate

```

> ratio.boot <- function(X, i) {
+   y <- X[i, ]
+   mean(y[2]) / mean(y[1])
+ }
> out <- boot(bigcity, ratio.boot, 999)
> out

```

## ORDINARY NONPARAMETRIC BOOTSTRAP

Call:

```
boot(data = bigcity, statistic = ratio.boot, R = 999)
```

Bootstrap Statistics:

|     | original | bias        | std. error |
|-----|----------|-------------|------------|
| t1* | 1.239019 | 0.001209301 | 0.03499868 |

```

> truehist(out$t, main = "By boot function")
> points(t0, 0, pch=17, col=2)
> out$t0
[1] 1.239019 # explicit computation
> mean(out$t)
[1] 1.243021

```

$\hat{F}(x) = \frac{1}{n} \sum_{i=1}^n I(X_i \leq x)$

$\hat{F}(x) = \sum_{i=1}^n I(X_i \leq x)$

$\hat{F}(x) = \sum_{i=1}^n I(X_i \leq x)$



$$\hat{F}(x) = \sum_{i=1}^n I(X_i \leq x)$$

"N" (norm)  
 Percentile CI (perc)  
 "t" (sead)  
 Basic Bootstrap (basic)  
 BC<sub>a</sub> CI (bca) ← "better" bootstrap (I)

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> boot.ci(out, type = c("norm", "perc", "basic", "bca"))

| level | Normal         | Basic          |
|-------|----------------|----------------|
| 95%   | (1.166, 1.304) | (1.160, 1.299) |

| Level | Percentile     | BC <sub>a</sub> |
|-------|----------------|-----------------|
| 95%   | (1.179, 1.318) | (1.179, 1.318)  |

- Bootstrapping regression

$$y_t = \hat{X}_t \beta + u_t, \quad E(u_t | X_t) = 0, \quad u_t \sim \text{iid}(0, \sigma^2),$$

n obs, k regressors.

Assumptions we make:

- ① independent errors?
- ② identically distributed errors?

Types:

- Residual bootstrap
- Parametric bootstrap
- Wild bootstrap
- Pair bootstrap

What about bootstrap for dependent data?

→ Types

- Parametric bootstrap
- Sieve bootstrap
- Block bootstrap

popular approaches

Carlo Stein (nonoverlapping)  
Künsch (overlapping)

Comments on block bootstrap methods

$$\hat{F}(x) = \frac{1}{n} \sum_{i=1}^n I(X_i \leq x)$$

## — Bootstrap regression

n obs.  
k regressors

$$y_t = X_t \beta + u_t, \quad E(u_t | X_t) = 0, \quad u_t \sim \text{iid}(0, \sigma^2).$$

Is  $u_t$  i.i.d.? Is  $u_t$  id?

Types:

residual

parametric

with

pair

### • Residual bootstrap

→ requires errors to be indep. of contemporaneous regressors and iid, but w/ minimal distributional assumptions

→ Steps:

① Use OLS to obtain  $\hat{\beta}$  and  $\hat{u}_t$ .

② (Optional) Rescale residuals so that they have correct variance.

$$\text{e.g. } \tilde{u}_t = (\frac{n}{n-k})^{1/2} \hat{u}_t.$$

③ Generate a typical observation of the bootstrap sample as

$$y_t^* = X_t \hat{\beta} + u_t^*, \quad u_t^* \sim \text{edf}(\tilde{u}_t).$$

$u_t^*$ 's are often said to be resampled from  $\tilde{u}_t$

### • Parametric bootstrap

→ assume  $u_t$  follow a specific distribution, e.g.  $N$ .

→ Steps:

① Use OLS to obtain  $\hat{\beta}$  and  $\hat{u}_t$ .

② Generate a typical observation using

$$y_t^* = X_t \hat{\beta} + u_t^*, \quad u_t^* \sim N(0, S^2).$$

$S^2$  = sample var of  $\hat{u}_t$ .

$$\begin{aligned} Syy &= Y'(I - H)Y \\ SSReg &= Y'(H - H)Y \\ RSS &= Y'(I - H)Y \\ &= \hat{e}'\hat{e} \end{aligned}$$

$$S^2 = \frac{RSS}{n-p-1}$$

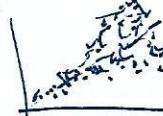
$$\begin{aligned} \widehat{\text{Var}}(\hat{\beta}) \\ = S^2(X'X)^{-1} \end{aligned}$$

$$\hat{\beta} = Y - \hat{Y} = (I - H)Y.$$

$$(3) \quad \hat{\beta} = X'(X'X)^{-1}X$$

$$\begin{aligned} \hat{\beta} &= (X'X)^{-1}XY \\ H &= X(X'X)^{-1}X \end{aligned}$$

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e.g. 

non-constant variance among residuals

### • Wild bootstrap

- is designed to handle heteroskedasticity in reg. models.
- data generating process (DGP):

$$y_t^* = X_t \hat{\beta} + f(u_t) v_t^* \quad v_t^* \sim \text{dist}(0, 1),$$

e.g.  $v_t^* \sim \text{Bern}(0.5)$  (-1 or 1).

$f(u_t)$  is a transformation of the  $t$ -th residual  $\hat{u}_t$ .

$$\text{e.g. } f(\hat{u}_t) = \frac{\hat{u}_t}{\sqrt{1-h_{tt}}}.$$

### • Pair bootstrap

- resample from the matrix with typical row  $[y_t \ X_t]$ .

We no longer condition on the  $X_t$  since each bootstrap sample now has a different  $X$  matrix.

A typical observation of the bootstrap sample is  $[y_t^* \ X_t^*]$ .

→ Comments:

- ① the pairs of bootstrap is valid even when the errors display heteroskedasticity of unknown form
- ② it works even for dynamic models.
- ③ pairs of bootstrap can be applied to an enormous range of models.
- ④ in the case of multivariate models, we can combine the pairs and residual bootstraps: organize residuals as a matrix and apply the pairs bootstrap to its rows. This preserves cross-equation correlations.

→ Deficiencies:

- (1) The bootstrap DGP does not impose  $H_0$ 's restrictions on  $\beta$ .
- (2) Compared to residual bootstrap (under validity) and wild bootstrap, pairs bootstrap does not yield very accurate results.

$$\begin{aligned} x_t &= \phi_1 x_{t-1} + \alpha_t \\ x_t &= \phi_1 x_{t-1} + \phi_1 \alpha_{t-1} + \alpha_t \\ x_t &= \phi_1 x_{t-1} + \sigma_a^2 + \alpha_t \\ x_t &= \phi_1 x_{t-1} + \alpha_t \end{aligned}$$

## Bootstrap for dependent data

- Types: → Resampling breaks up any dependence and is therefore inappropriate for dependent data.
- Parametric Sieve Block → Two popular approaches: sieve and block

### • Parametric bootstrapping

Consider stationary  $x_t \sim AR(1)$ .

$$x_t = \phi_1 x_{t-1} + \alpha_t, \alpha_t \sim N(0, \sigma_a^2) \rightarrow (1)$$

$$(Unconditional) x_t \sim N\left(\frac{\mu}{1-\phi_1}, \frac{\sigma_a^2}{1-\phi_1^2}\right) \rightarrow (2)$$

Why

$$x_t = \alpha_t + \phi_1 x_{t-1} + \alpha_{t-1}$$

$$x_t x_{t-k} = \alpha_t x_{t-k} + \phi_1 x_{t-1} x_{t-k} + \alpha_{t-1} x_{t-k}$$

$$\begin{aligned} x_k &= \alpha_t E(x_{t-k}) + \phi_1 x_{t-1} + E(\alpha_{t-1} x_{t-k}) \\ &= 0 \end{aligned}$$

$$k=0 \Rightarrow x_k = \phi_1 x_{k-1} + E(\alpha_t x_{t-k}).$$

$$k=0 \Rightarrow E(\alpha_t x_{t-k})$$

$$= E(\alpha_t (\phi_1 x_{t-1} + \alpha_{t-1}))$$

$$= 0 + 0 + \sigma_a^2 = \sigma_a^2.$$

$$\Rightarrow x_0 = \phi_1 x_1 + \sigma_a^2$$

$$k=1 \Rightarrow x_1 = \phi_1 x_0$$

$$\Rightarrow x_0 = \phi_1 (\phi_1 x_0) + \sigma_a^2$$

$$= \phi_1^2 x_0 + \sigma_a^2$$

$$\Rightarrow (1 - \phi_1^2) x_0 = \sigma_a^2$$

$$\Rightarrow x_0 = \frac{\sigma_a^2}{1 - \phi_1^2} \dots$$

$$x_t = \phi_1 x_{t-1} + \alpha_t$$

$$\frac{\alpha_t}{1 - \phi_1} = \phi_1 x_{t-1} + \frac{\alpha_t}{1 - \phi_1}$$

$$(Conditional) x_t | X_{t-1} \sim N(\phi_1 x_{t-1}, \sigma_a^2) \rightarrow (3)$$

→ The (unconditional) simulation procedure may be summarized as!

① Simulate  $X_0$  by drawing a random # from (2)

② Simulate  $X_1 = \alpha_t + \phi_1 X_0 + \alpha_{t-1}$

③ Simulate  $X_t = \alpha_t + \phi_1 X_{t-1} + \alpha_{t-2}$  recursively.

$$y_t = \tilde{X}_t \tilde{\beta} + u_t$$

- The Sieve Bootstrap.

→ Assumptions:

$u_t$  in a reg model follows an unknown, stationary process w/ homoscedastic,  $\text{iid}$ , innovations

→ Approximate such process by  $\text{AR}(p)$  where  $p$  is chosen by some sort of model selection criteria like AIC or BIC, or by sequential testing ( $p=1$  vs.  $p$ )

→ Steps:

① Estimate the model to obtain residuals  $\hat{u}_t$

② Estimate  $\text{AR}(p)$  model:  $\hat{u}_t = \sum_{i=1}^p \phi_i \hat{u}_{t-i} + \hat{\epsilon}_t$ . (1)

③ Generate bootstrap error terms

$$\hat{u}_t^* = \sum_{i=1}^p \hat{\phi}_i \hat{u}_{t-i}^* + \hat{\epsilon}_t^*, \quad (2)$$

where  $\hat{\epsilon}_t^*$ 's are resampled from the (rescaled) residuals from (1).

④ Generate the bootstrap data according to

$$\hat{y}_t^* = \tilde{X}_t \tilde{\beta} + \hat{u}_t^*, \text{ where } \hat{u}_t^* \text{ is from (2).}$$

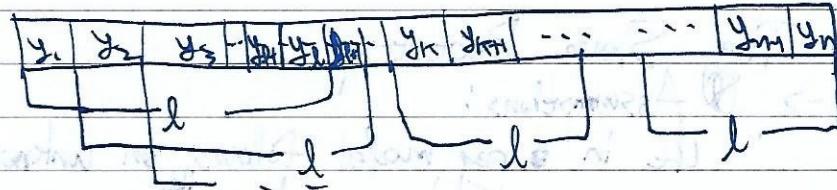
- MBB (the moving block bootstrap)

→ Carlstein: bootstrapping blocks of observations, rather than individual observations; nonoverlapping blocks

→ Künsch: MBB, applicable to stationary time series data; blocks of observations are overlapping.

Idea

Divide the data ~~into~~ of  $n$  observations into blocks of length  $l$  and select  $b$  of these blocks ~~with replacement~~ by resampling with replacement all the possible blocks.



④ Carlstein

Block 1:  $(y_1, y_2, \dots, y_l)$ Block 2:  $(y_{l+1}, \dots, y_{2l})$ Block k:  $(y_{k(l-1)+1}, \dots, y_{kl})$ Block last(b):  
 $(y_{n-l+1}, \dots, y_n)$ 

$$\Rightarrow \# \text{ of blocks: } b = \frac{n}{l}$$

So  $n = lb$ .

(l = size of a block, b = # of blocks).

e.g.:  $\Delta^t = \{ \boxed{3} | \boxed{6} | \boxed{n} | \boxed{2} | \boxed{1} | \boxed{5} \}$ ,  
l=3, n=6.Carlstein  $\Rightarrow b=2$ 

$$\boxed{1} \boxed{3} \boxed{6} \boxed{7} \quad \boxed{2} \boxed{1} \boxed{5}$$
  
~~⊗⊗⊗~~ = ♦2
Draw a sample of two blocks w/ replacement;  
possible outcomes:
~~⊗⊗⊗, ⊗⊗⊗, ⊗⊗⊗, ⊗⊗⊗~~  
 $\Omega = \{ (\star_1, \star_1), (\star_1, \star_2), (\star_2, \star_1), (\star_2, \star_2) \}$ 

Künsch

Block 1:  $(y_1, \dots, y_l)$ Block 2:  $(y_2, \dots, y_{l+1})$ Block k:  $(y_k, \dots, y_{k+l-1})$ Block last(b):  
 $(y_b, \dots, y_n)$ 

$$\Rightarrow \# \text{ of blocks: } b = n - l + 1.$$

So  $n = l + b - 1$ .

Künsch: b=4

$$\boxed{1} \boxed{3} \boxed{6} \boxed{7} \quad \boxed{1} \boxed{6} \boxed{1} \boxed{2} \quad \boxed{1} \boxed{7} \boxed{2} \quad \boxed{1} \boxed{2} \boxed{5}$$
  
~~⊗⊗⊗~~ || || ||  
 $\square_1 \quad \square_2 \quad \square_3 \quad \square_4$ 
Draw a sample of two blocks w/ replacement;  
possible outcomes:
 $\Omega = \{ (\square_1, \square_1), \dots, (\square_4, \square_4) \}$ ,  
 $|\Omega| = 16$ .

$$\begin{array}{l|l} P(\text{1st in the second row}) & P_k(\text{1st in 2nd row} \mid \text{1st was 1st}) \\ P_k(\text{1st in 2nd row} \mid \text{1st was 1st}) & = 1/4 = 0.25. \\ = 1/2 = 0.5. & \end{array}$$

### Problems w/ MBB

① Even if  $\{x_t\}$  is stationary, the pseudo time series generated from  $\{x_t\}$  by MBB is not stationary.

Possible solution: let  $l \sim \text{Geom}(p) \neq p$ .

② The mean of MBB,  $\bar{x}_n^*$ , is biased:

$$E(\bar{x}_n^* \mid x_1, \dots, x_n) = \bar{x}_n \neq 0.$$

③ MBB estimator of the variance of  $\sqrt{n}(\bar{x}_n)$  is also biased; instead of  $\hat{\sigma}^2 = \frac{1}{n} \sum_{i=1}^n (x_i - \bar{x}_n)^2$ , we should use  $\tilde{\sigma}^2 = \frac{1}{n} \sum_{i=1}^n [(x_i - \bar{x}_i)^2 + \sum_{k=1}^{i-1} \sum_{j=1}^{n-k} (x_j - \bar{x}_n)(x_{i+k} - \bar{x}_n)]$ .

### Optimal $l$

① Carlstein < Kunsch

⑦  $l \sim \text{Geom}(p)$ , or  $l = \frac{1}{p} (E(l))$ .

→ less sensitive to the choice of  $p$  than the application of MBB is to the choice of  $l$ .

### Comments

- Block bootstrap methods divide the quantities that are being resampled, which might be either rescaled residuals or  $[y, X]$  pairs, into blocks of ~~con-~~ consecutive observations. We then resample the blocks.

- Overlapping blocks are better.

- $l$  fixed is better than  $l$  variable.

- Choice of  $l$  is critical; it must  $\uparrow$  as  $n \uparrow$ . Often,  $l \propto n^{1/3}$ .

- $l$  too small  $\Rightarrow$  cannot mimic original.  
Dependence is broken.
- $l$  too long  $\Rightarrow$  bootstrap samples are not random enough.
- Block-of-blocks of bootstrap is the analog of the pairs bootstrap for dynamic models.

Steps

Consider  $y_t = X_t \beta + \gamma y_{t-1} + u_t$ ,  $u_t \stackrel{\text{iid}}{\sim} \text{dist}(0, \sigma^2)$ .  
Define  $\tilde{z}_t^* = [y_t, y_{t-1}, X_t]$ .

Construct  $\textcircled{B}$  length  $l$  overlapping blocks:

Block 1 :  $(\tilde{z}_1, \dots, \tilde{z}_l)$

Block 2 :  $(\tilde{z}_2, \dots, \tilde{z}_{l+1})$

⋮ ⋮

Block  $k$  :  $(\tilde{z}_k, \dots, \tilde{z}_{k+l-1})$

⋮ ⋮

Block  $b$  (last) :  $(\tilde{z}_b, \dots, \tilde{z}_n)$ .

( $n = l+b-1 \Rightarrow b = n-l+1$  blocks in total).

$\rightarrow$  ~~B~~ BofB bootstrap works w/ nonconseant variance & feature as well as serial correlation.

$\rightarrow$  Generally, Block bootstrap offer higher-order accuracy than asymptotic methods, but only by a modest extent.

$\rightarrow$  Block bootstraps can yield more reliable standard errors.