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ntreatreg: A Stata module for estimation of treatment effects in the presence of neighborhood interactions

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Science is when we manage to pass from the theoretically fanciful to the empirically plausible

Outline of this presentation

- □ Statistical background and related studies
- □ The Rubin's *potential outcome model* with *neighborhood interactions*
- \Box Model's estimation
- □ Stata implementation via ntreatreg
- □ Application to real data
- □ Conclusions

In standard Econometrics of Program Evaluation (aimed at estimating the effect of a policy on supported individuals) it is assumed the so-called SUTVA (Rubin 1978):

SUTVA: Stable-Unit-Treatment-Value-Assumption



"treatment received by one unit do not affect outcomes for another unit"

It means that: only the treatment applied to the specific individual is assumed to potentially affect the outcome for that particular individual.

===> We would like to *relax* this assumption and understand what happens to the estimation of the effect of a "treatment" in the presence of potential *contagion* (or *neighborhood*, or *social*) effects.

SUTVA and NO-SUTVA setting



- ✓ Rubin (1978): calls this important assumption as Stable-Unit-Treatment-Value-Assumption (SUTVA)
- ✓ Manski (2011): refers to Individualistic-Treatment-Response (ITR) to emphasize that this poses a restriction in the form of the treatment response function that the analyst considers.

- AIM: estimating the "Average Treatment Effects" (ATEs) of a policy program in a *non-experimental* setup in the presence of *endogenous neighbourhood* (or *externality*) interactions (Manski, 1993), by assuming that *Conditional Mean Independence* (i.e., *selection-on-observables*) holds.
- **SETTING**: we consider a *binary* treatment variable *w* taking value 1 for treated and 0 for untreated units assumed to affect an *outcome* variable *y* that can take a variety of forms: binary, count, continuous, etc..
- NOTATION:
 - -N = number of units involved in the (social) experiment
 - $-N_1$ = number of treated units
 - $N_0 =$ number of untreated units
 - $-w_i$ = treatment variable assuming value "1" if the unit is treated and "0" if untreated
 - $y_{1i} =$ outcome of unit *i* when he is treated
 - $y_{0i} =$ outcome of unit *i* when he is untreated
 - $-\mathbf{x}_i = (x_{1i}, x_{2i}, x_{3i} \dots x_{Mi}) = \text{row vector of } M \text{ observable variables for unit } i.$

The notion of "endogenous" neighbourhood effects

Manski (1993) identifies three types of effects corresponding to three arguments of an individual outcome equation incorporating *social effects*:

- 1. *Endogenous effects*: the outcome of an individual depends on the outcomes of other individuals belonging to his neighbourhood.
- 2. *Exogenous (or contextual) effects*: the outcome of an individual is affected by the exogenous idiosyncratic characteristics of the individuals belonging to his neighbourhood.
- 3. *Correlated effects*: due to belonging to a specific group and thus sharing some institutional/normative condition (that one can loosely define as "environment").

Contextual and correlated effects are to be assumed as exogenous, as they clearly depend on pre-determined characteristics of the individuals in the neighbourhood (case 2) or of the neighbourhood itself (case 3).

Endogenous effects are of broader interest: they depend on the behaviour (measured as "outcome") of other individuals involved in the same neighbourhood.

Endogenous effects both comprise *direct* and *indirect* effects linked to a given external intervention on individuals.

The model presented here incorporates the presence of *endogenous neighbourhood effects* as defined by Manski within a traditional *binary counterfactual model* and provides both an identification and an estimation procedure of the **Average Treatment Effects (ATEs)** in a simple parametric case.

Some related literature

Rosenbaum (2007) discusses methods for testing *null hypotheses* on the presence of interference in trials where *random assignment occurs* within groups and interference *does not cross* group boundaries.

Hudgens and Halloran (2008) extend the previous work in the setting of a *two-stage randomized trial* in which some groups are randomly assigned to host treatments, and then treatments are assigned at random within the selected groups. Interference is presumed to operate only *within groups*.

Tchetgen-Tchetgen and VanderWeele (2010) extend Hudgens and Halloran's results, providing conservative variance estimators, a framework for finite sample inference and extensions to observational studies. Hierarchical treatment assignment and interference limited to groups greatly simplifies the estimation problem, as inference can proceed assuming *independence across groups*.

===> **Sobel (2006)** analyzes the potential for *bias* when <u>no-interference</u> is mistakenly assumed, and then defines a number of direct and indirect effects that may be identifiable.

He characterizes the usual estimators of treatment effects developing their form when *interference* is allowed.

"When interference is present, the difference between a **treatment group** mean and a **control group** mean (unadjusted or adjusted for covariates) estimates not an average treatment effect, but rather the difference between two effects defined on two distinct subpopulations. This result is of great importance, for a researcher who fails to recognize this could easily infer that a treatment is beneficial when in fact it is universally harmful" (p. 1398).

=== > Application: **social experiment** (with *randomization*)

=== > MTO program ("Move To Opportunity")

=== > LATE estimator ($\hat{a} \ la \ Angrist$)

Position of this paper within previous literature

Previous literature assumes:

- 1. Randomized assignment
- 2. Multiple treatment
- 3. Non-parametric form for the *potential outcome* and *interaction*

This paper assumes:

- 1. Non-randomized assignment
- 2. Binary treatment
- 3. Parametric form for the *potential outcome* and *interaction*

=== > **Therefore**: this paper suggests a *simpler* and *less general* way to relax SUTVA, but one that is easy to implement in many contexts of application.

DEFINITION OF (AVERAGE) TREATMENT EFFECTS (ATEs)

Unit *i* **Treatment Effect**:

 $TE_i = y_{1i} - y_{0i}$

we observe just *one* of the two quantities $(y_{1i}; y_{0i})$, but <u>*never*</u> both: missing observation problem (Holland, 1986).

What is *observable* to the analyst is the single status of unit *i*, that is:

 $y_i = y_{0i} + w_i (y_{1i} - y_{0i})$

called the **Potential Outcome Model**, and it links *unobservable* with *observable* outcomes.

Since recovering the entire distributions of y_{1i} and y_{0i} is too demanding, we focus on the population **Average Treatment Effects** (hereinafter **ATEs**) and on ATEs *conditional on* **x** (i.e., ATE(**x**)) of a policy intervention, defined as:

 $ATE = E(y_{i1} - y_{i0})$ $ATE(\mathbf{x}_i) = E(y_{i1} - y_{i0} | \mathbf{x}_i)$

 $ATET = E(y_{i1}-y_{i0} | w_i=1)$ $ATET(\mathbf{x}_i) = E(y_{i1} - y_{i0} | \mathbf{x}_i, w_i=1)$

 $ATENT = E(y_{i1}-y_{i0} | w_i=0)$ $ATENT(\mathbf{x}_i) = E(y_{i1} - y_{i0} | \mathbf{x}_i, w_i=0)$

where $E(\cdot)$ is the mean operator. These parameters are equal to the difference between the average of the target variable when the individual is treated (y_1), and the average of the target variable when the same individual is untreated (y_0). Observe that by LIE: ATE = E_x {ATE(x)}, ATET = E_x {ATET(x)}, ATENT = E_x {ATENT(x)}.

A NEIGHBORHOOD-EFFECT TREATMENT MODEL

 y_{0i} and y_{1i} need to have a representation including the **neighborhood effect** from *treated* to *untreated* units. We start by this **parametric model system**:

$y_{1i} = \boldsymbol{\mu}_1 + \mathbf{x}_i \boldsymbol{\beta}_1 + \boldsymbol{e}_{1i}$	Outcome equation for the treated status
$y_{0i} = \boldsymbol{\mu}_0 + \mathbf{x}_i \boldsymbol{\beta}_0 + \gamma s_i + e_{0i}$	Outcome equation for the non-treated status with <i>neighbourhood effect "s</i> "
$s_{i} = \begin{cases} \sum_{j=1}^{N_{1}} \omega_{j} y_{1j} & \text{if } i \in \{w = 0\} \\ 0 & \text{if } i \in \{w = 1\} \end{cases}$	Form of the <i>neighbourhood effect</i> of treated <i>j</i> s on unit <i>i</i> (<i>weighted mean</i>)
$y_i = y_{0i} + w_i (y_{1i} - y_{0i})$	Potential Outcome Equation (POM)
$\sum_{j=1}^{N_1} \boldsymbol{\omega}_{ij} = 1$	Weights add to one
$i = 1,, N$ and $j = 1,, N_1$	<i>i</i> : index for all units; <i>j</i> : index for treated units

and Conditional Mean Independence (CMI) holds:

 $E(y_{ig} | w_i, \mathbf{x}_i) = E(y_{ig} | \mathbf{x}_i) \text{ with } g = \{0, 1\}$

We need to solve the previous **SYSTEM** to recover an estimation of ATEs. By substitutions within the **previous system**, we eventually get that:

$$y_{0i} = \mu_0 + \mathbf{x}_i \boldsymbol{\beta}_0 + \gamma \sum_{j=1}^{N_1} \omega_{ij} y_{1j} + e_{0i}$$

Hence, ATE is equal to:

ATE = E(
$$y_{1i} - y_{0i}$$
) = E $\left[(\mu_1 + \mathbf{x}_i \boldsymbol{\beta}_1 + e_{1i}) - \left(\mu_0 + \mathbf{x}_i \boldsymbol{\beta}_0 + \gamma \sum_{j=1}^{N_1} \omega_{ij} y_{1j} + e_{0i} \right) \right]$

After some manipulations, we get that:

$$ATE = \boldsymbol{\mu} + \overline{\mathbf{x}}_i \boldsymbol{\delta} - \left(\sum_{j=1}^{N_1} \omega_{ij} \overline{\mathbf{x}}_j\right) \boldsymbol{\gamma} \boldsymbol{\beta}_1$$

where: $\overline{\mathbf{x}}_i = \mathbf{E}(\mathbf{x}_i)$.

We are also interested in estimating $ATE(\mathbf{x})$. Using the previous results, we finally get that:

$$ATE(\mathbf{x}_i) = ATE + (\mathbf{x}_i - \overline{\mathbf{x}})\mathbf{\delta} + \sum_{j=1}^{N_1} \omega_{ij}(\overline{\mathbf{x}} - \mathbf{x}_j)\gamma \beta_1$$

where it is clear that $ATE(\mathbf{x})$ depends on \mathbf{x} .

Once the formulas for ATE and $ATE(\mathbf{x})$ are available, it is also possible to recover the Average Treatment Effect on Treated (ATET) and on non-Treated (ATENT), that is:

ATET = ATE +
$$\frac{1}{\sum_{i=1}^{N} w_i} \sum_{i=1}^{N} w_i \left[(\mathbf{x}_i - \overline{\mathbf{x}}) \mathbf{\delta} + \sum_{j=1}^{N} \omega_{ij} (\overline{\mathbf{x}} - \mathbf{x}_j) \gamma \mathbf{\beta}_1 \right]$$

and:

ATENT = ATE +
$$\frac{1}{\sum_{i=1}^{N} (1-w_i)} \sum_{i=1}^{N} (1-w_i) \left[(\mathbf{x}_i - \overline{\mathbf{x}}) \boldsymbol{\delta} + \sum_{j=1}^{N} \omega_{jj} (\overline{\mathbf{x}} - \mathbf{x}_j) \boldsymbol{\gamma} \boldsymbol{\beta}_1 \right]$$

These quantities are functions of observable components and parameters to be firstly consistently estimated. Once these estimates are available, standard errors for ATET and ATENT can be obtained via bootstrapping (Wooldridge, 2010, Ch. 21).

How to get consistent estimation of ATEs ?

Using an *i.i.d.* sample of observed variables for each individual *i*:

 $\{y_i, w_i, \mathbf{x}_i\}$ with i = 1, ..., N

and by substitution into the POM, we get this *Switching Random Coefficient* Model:

$$y_{i} = \left(\mu_{0} + \mathbf{x}_{i}\boldsymbol{\beta}_{0} + \gamma \sum_{j=1}^{N_{1}} \omega_{ij} y_{1j} + e_{0i}\right) + w \left[\left(\mu_{1} + \mathbf{x}_{i}\boldsymbol{\beta}_{1} + e_{1i}\right) - \left(\mu_{0} + \mathbf{x}_{i}\boldsymbol{\beta}_{0} + \gamma \sum_{j=1}^{N_{1}} \omega_{ij} y_{1j} + e_{0i}\right) \right]$$

After sorting out previous formula, we finally get that:

$$y_i = \eta + w_i \cdot \mathbf{ATE} + \mathbf{x}_i \boldsymbol{\beta}_0 + w_i (\mathbf{x}_i - \overline{\mathbf{x}}) \boldsymbol{\delta} + w_i \sum_{j=1}^N \omega_{ij} w_j (\overline{\mathbf{x}} - \mathbf{x}_j) \gamma \boldsymbol{\beta}_1 + e_i$$

with:

$$\mu = \mu_1 - \mu_0 - \gamma \mu_1; \quad \eta = \mu_0 + \gamma \mu_1$$
$$e_i = \gamma \sum_{j=1}^{N_1} \omega_{ij} e_{1j} + e_{0i} + w_i (e_{1i} - e_{0i}) - w_i \gamma \sum_{j=1}^{N_1} \omega_{ij} e_{1j}$$

This is a usual regression model whose parameters – **under CMI** – can be estimated *consistently* by **Ordinary Least Squares (OLS)**. With an estimation of the parameters at hand we can estimate ATE (directly from the regression) and ATEs by plugging parameters into their formulas. Observe, however, that a **matrix of distance weights** $\Omega = [\omega_{ij}]$ needs beforehand to be provided by the analyst.

A PROTOCOL FOR ESTIMATING PARAMETRICALLY ATES UNDER "NEIGHBORHOOD INTERACTIONS"

1.Provide a matrix of distance weights $\Omega = [\omega_{ij}]$ between the generic unit *i* (untreated) and unit *j* (treated).

2.Estimate the regression model by an OLS of:

$$y_i$$
 on $\left\{1, w_i, \mathbf{x}_i, w_i(\mathbf{x}_i - \overline{\mathbf{x}}), w_i \sum_{j=1}^N \omega_{ij} w_j(\overline{\mathbf{x}} - \mathbf{x}_j)\right\}$

3.Obtain $\{\hat{\boldsymbol{\beta}}_{0}, \hat{\boldsymbol{\delta}}, \hat{\boldsymbol{\gamma}}, \hat{\boldsymbol{\beta}}_{1}\}$ and put them into the formulas for ATEs.

INTERPRETATION OF THE "NEIGHBOURHOOD BIAS"

By comparing the formula of ATE with ($\gamma \neq 0$) and without ($\gamma = 0$) neighbourhood effect, we get the so-called Neighbourhood Bias (Sobel, 2006):

$$\mathbf{Bias} = \left| \mathbf{ATE}_{\text{no-neigh}} - \mathbf{ATE}_{\text{with-neigh}} = \left(\sum_{j=1}^{N_1} \omega_{jj} \overline{\mathbf{x}}_j \right) \boldsymbol{\gamma} \boldsymbol{\beta}_1 \right|$$

This can also be seen as the *externality effect* produced by the policy: it depends on:

- 1. weights
- 2. mean of \mathbf{x}
- 3. magnitude and sign of coefficients γ and β_1 .

Observe that it can be *positive* as well as *negative*.

Observe that the **NEIGHBOURHOOD BIAS** can also be interpreted as a **SPECIFICATION ERROR** in the outcome equation arising when potential outcomes are modelled without taking into account externality effects.

Finally, by defining:

$$\boldsymbol{\gamma}\boldsymbol{\beta}_1 = \boldsymbol{\lambda}$$

one can (parametrically) test whether this *bias* is or is not statistically significant by testing this null:

$$\mathbf{H}_0: \quad \boldsymbol{\lambda}_1 = \boldsymbol{\lambda}_2 = \ldots = \boldsymbol{\lambda}_M = \mathbf{0}$$

STATA implementation: the "ntreatreg" command

The syntax of **ntreatreg** is a very common one for a STATA command:

```
ntreatreg outcome treatment varlist , hetero(varlist_h)
spill(matrix) graphic
```

where:

outcome: y
treatment: w
varlist: x
varlist_h: subset of x
matrix: distance matrix Ω

Stata help-file for **ntreatreg**

help ntreatreg

Title

ntreatreg - Stata module for estimation treatment effects in the presence of neighbourhood Interactions

Syntax

ntreatreg outcome treatment [varlist] [if] [in] [weight], [spill(matrix) hetero(varlist_h) conf(number) graphic vce(robust) const(noconstant) head(noheader)]

fweights, iweights, and pweights are allowed; see weight.

Description

ntreatreg estimates Average Treatment Effects (ATEs) under Conditional Mean Independence (CMI) when neighbourhood interactions may be present. It incorporates such externalities within the traditional Rubin's potential outcome model. As such, it provides an attempt to relax the Stable Unit Treatment Value Assumption (SUTVA) generally used in observational studies.

Options

- spill(matrix) specifies the adjacent (weighted) matrix used to define presence and strength of units' relationship.
 It could be a distance matrix, with distance loosely defined either as vector or spatial.
- hetero(varlist_h) specifies the variables over which to calculate the idyosincratic Average Treatment Effect ATE(x), ATET(x) and ATENT(x), where x=varlist_h. It is optional. When this option is not specified, the command estimates the specified model without heterogeneous average effect. Observe that varlist_h should be the same set or a subset of the variables specified in varlist.
- graphic allows for a graphical representation of the density distributions of ATE(x), ATET(x) and ATENT(x). It is optional for all models and gives an outcome only if variables into hetero() are specified.

vce(robust) allows for robust regression standard errors. It is optional for all models.

beta reports standardized beta coefficients. It is optional for all models.

const(noconstant) suppresses regression constant term. It is optional for all models.

conf(number) sets the confidence level equal to the specified number. The default is number=95.

_ws_varname_h are the additional regressors used in model's regression when hetero(varlist_h) is specified.

z_ws_varname_h are the spillover additional regressors used in model's regression when hetero(varlist_h) is specified.

ATE(x) is an estimate of the idiosyncratic Average Treatment Effect.

ATET(x) is an estimate of the idiosyncratic Average Treatment Effect on treated.

ATENT(x) is an estimate of the idiosyncratic Average Treatment Effect on Non-Treated.

ntreatreg returns the following scalars:

r(N_tot) is the total number of (used) observations.

r(N_treat) is the number of (used) treated units.

r(N_untreat) is the number of (used) untreated units.

r(ate) is the value of the Average Treatment Effect.

r(atet) is the value of the Average Treatment Effect on Treated.

r(atent) is the value of the Average Treatment Effect on Non-treated.

Remarks

The treatment has to be a 0/1 binary variable (1 = treated, 0 = untreated).

When option hetero is not specified, ATE(x), ATET(x) and ATENT(x) are one singleton number equal to ATE=ATET=ATENT.

Please remember to use the update query command before running this program to make sure you have an up-to-date version of Stata installed.

Example

- . ssc install ntreatreg
- . use "FERTIL2_200.DTA"
- . matrix dissimilarity dist = age agesq urban electric tv , corr
- . matewmf dist dist_abs, f(abs)
- . ntreatreg children educ7 age agesq evermarr electric tv , ///
 hetero(age agesq evermarr) spill(dist_abs) graphic
- . test z_ws_age1 = z_ws_agesq1 = z_ws_evermarr1 = 0

References

- Cerulli, G. 2014. Identification and Estimation of Treatment Effects in the Presence of Neighbourhood Interactions, Working Paper Cnr-Ceris, N° 04/2014.
- Wooldridge, J. M. 2010. Econometric Analysis of Cross Section and Panel Data, 2nd Edition. Chapter 21. The MIT Press, Cambridge.

Example 1: effect of *location* **on** *crime*

Dataset. "SPATIAL_COLUMBUS.DTA" provided by Anselin (1988) containing information (22 variables) on property crimes in 49 neighbourhoods in Columubus, Ohio, in 1980.

Objective. Evaluating the impact of *housing location* on *crimes*, i.e. the causal effect of the variable "cp" - taking value 1 if the neighbourhood is located in the "core" of the city and 0 if located in the "periphery" - on the number of residential burglaries and vehicle thefts per thousand households (i.e., the variable "crime").

Confounding observables. Only two main factors: the *household income* in \$1,000 ("inc") and the *housing value* in \$1,000 ("hoval").

===> We are interested in detecting the effect of housing location on the number of crimes in such a setting, by taking into account possible *interactions among neighbourhoods*.

STEP 0. INPUT DATA FOR THE REGRESSION MODEL

y: crime

W: cp

 $\boldsymbol{X}\mbox{:}\mbox{ inc hoval}$

Matrix Ω : w

STEP 1. LOAD THE STATA ROUTINE "NTREATREG" AND THE DATASET

- . ssc install ntreatreg
- . ssc install spatwmat // see package: sg162 from

http://www.stata.com/stb/stb60

. use "SPATIAL_COLUMBUS.DTA "

STEP 2. PROVIDE THE MATRIX "OMEGA" (HERE WE CALL IT "W")

. spatwmat, name(W) xcoord(\$xcoord) ycoord(\$ycoord) band(0 \$band) ///
standardize eigenval(E) // this generates the inverse distance matrix W

The following matrices have been created:

1. Inverse distance weights matrix W (row-standardized)

Dimension: 49x49

Distance band: 0 < d <= 10

Friction parameter: 1

Minimum distance: 0.7

1st quartile distance: 6.0

Median distance: 9.5

3rd quartile distance: 13.6

Maximum distance: 27.0

Largest minimum distance: 3.37

Smallest maximum distance: 14.51

2. Eigenvalues matrix E

Dimension: 49x1

STEP 3. ESTIMATE THE MODEL USING "NTREATREG" TO GET THE "ATE" WITH NEIGHBORHOOD-INTERACTIONS

- . set more off
- . xi: ntreatreg crime cp inc hoval , hetero(inc hoval) spill(W) graphic

Source	SS	df	MS		Number of obs	= 49	
+-					F(7, 41)	= 15.74	AIE
Model	9793.37437	7 1399	.05348		Prob > F	= 0.0000	
Residual	3644.84518	41 88.8	986629		R-squared	= 0.7288	
+-					Adj R-squared	= 0.6825	
Total	13438.2195	48 279.	962907		Root MSE	= 9.4286	
crime	Coef.	Std. Err.	t	P> t	[95% Conf.	Interval]	
+-	K						
cp	9.492458	4.816401	1.97	0.056	2344611	19.21938	
 inc	9.492498 4968051	4.816401 .3653732	1.97 -1.36	0.056 0.181	2344611 -1.234691	19.21938 .241081	
 inc hoval	9.492498 4968051 2133293	4.816401 .3653732 .101395	1.97 -1.36 -2.10	0.056 0.181 0.042	2344611 -1.234691 4181006	19.21938 .241081 008558	
 cp inc hoval ws_inc	9.492498 4968051 2133293 -1.19053	4.816401 .3653732 .101395 .9911119	1.97 -1.36 -2.10 -1.20	0.056 0.181 0.042 0.237	2344611 -1.234691 4181006 -3.192121	19.21938 .241081 008558 .8110612	
cp inc hoval _ws_inc _ws_hoval	9.492498 4968051 2133293 -1.19053 .1440651	4.816401 .3653732 .101395 .9911119 .2268815	1.97 -1.36 -2.10 -1.20 0.63	0.056 0.181 0.042 0.237 0.529	2344611 -1.234691 4181006 -3.192121 3141313	19.21938 .241081 008558 .8110612 .6022616	
cp inc hoval _ws_inc _ws_hoval z_ws_inc1	9.492498 4968051 2133293 -1.19053 .1440651 -5.719737	4.816401 .3653732 .101395 .9911119 .2268815 2.934276	1.97 -1.36 -2.10 -1.20 0.63 -1.95	0.056 0.181 0.042 0.237 0.529 0.058	2344611 -1.234691 4181006 -3.192121 3141313 -11.64563	19.21938 .241081 008558 .8110612 .6022616 .2061538	
cp inc hoval _ws_inc _ws_hoval z_ws_inc1 z_ws_hoval1	9.492498 4968051 2133293 -1.19053 .1440651 -5.719737 .3889889	4.816401 .3653732 .101395 .9911119 .2268815 2.934276 .9016162	1.97 -1.36 -2.10 -1.20 0.63 -1.95 0.43	0.056 0.181 0.042 0.237 0.529 0.058 0.668	2344611 -1.234691 4181006 -3.192121 3141313 -11.64563 -1.431862	19.21938 .241081 008558 .8110612 .6022616 .2061538 2.20984	
cp inc hoval _ws_inc z_ws_hoval z_ws_hoval1 _cons	9.492498 4968051 2133293 -1.19053 .1440651 -5.719737 .3889889 34.78312	4.816401 .3653732 .101395 .9911119 .2268815 2.934276 .9016162 8.655264	1.97 -1.36 -2.10 -1.20 0.63 -1.95 0.43 4.02	0.056 0.181 0.042 0.237 0.529 0.058 0.668 0.000	2344611 -1.234691 4181006 -3.192121 3141313 -11.64563 -1.431862 17.30346	19.21938 .241081 008558 .8110612 .6022616 .2061538 2.20984 52.26279	

- . scalar ate_neigh = _b[cp] // put ATE into a scalar
- . rename ATE_x _ATE_x_spill // rename ATE_x as _ATE_x_spill
- . rename ATET_x _ATET_x_spill
- . rename ATENT_x _ATENT_x_spill

STEP 4. DO A TEST TO SEE IF THE COEFFICIENTS OF THE NEIGHBOURHOOD-EFFECT ARE JOINTLY ZERO

4.1. if one accepts the null Ho: $\gamma \beta_0 = 0 \implies$ the neighbourhood-effect is negligible;

4.2. if one does not accept the null => the neighbourhood-effect effect is relevant.

```
. test z_ws_inc1 = z_ws_hoval1 = 0
```

```
(1) z_ws_incl - z_ws_hovall = 0
```

```
(2) z_ws_inc1 = 0
```

F(2, 41) = 2.35
Prob > F = 0.1078 // externality effect seems not significant

STEP 5. ESTIMATE THE MODEL USING "IVTREATREG" (TO GET ATE "WITHOUT" NEIGHBOURHOOD-INTERACTIONS)

. xi: ivtreatreg crime cp inc hoval , hetero(inc hoval) model(cf-ols) graphic

Source	SS	df	MS		Number of obs	: =	49	ΓΛ	Ъ
+					F(5, 43)	=	19.84	AI	
Model	9375.05895	5 187	5.01179		Prob > F	=	0.0000		
Residual	4063.1606	43 94.	4921069		R-squared	=	0.6976		
+					Adj R-squared	-	0.6625		
Total	13438.2195	48 279	.962907		Root MSE	=	9.7207		
crime	Coef.	Std. Err.	t	P> t	[95% Conf.	Int	erval]		
+									
Cp	13.59008	4.119155	3.30	0.002	5.283016	21	.89715		
inc	8335211	.3384488	-2.46	0.018	-1.516068	1	509741		
hoval	1885477	.1036879	-1.82	0.076	3976543	.0	205588		
_ws_inc	-1.26008	1.004873	-1.25	0.217	-3.286599	.7	664396		
_ws_hoval	.2021829	.2300834	0.88	0.384	2618246	.6	661904		
_cons	46.52524	6.948544	6.70	0.000	32.51217	60	.53832		

. scalar ate_no_neigh = _b[educ7] // put ATE into a scalar

. di ate_no_neigh

STEP 6. SEE THE MAGNITUDE OF THE NEIGHBORHOOD-INTERACTIONS BIAS

. scalar bias= ate_no_neigh - ate_neigh // in level

. di bias

4.09 // the difference in level is around four crimes

- . scalar bias_perc=(bias/ate_no_neigh)*100 // in percentage
- . di bias_perc

30.15 // there is a 30% of bias due to neighbourhood interaction

STEP 7. COMPARE GRAPHICALLY THE DISTRIBUTION OF ATE(**x**), ATET(**x**) and ATENT(**x**) WITH AND WITHOUT NEIGHBOURHOOD-INTERACTION

```
* ATE
twoway kdensity ATE_x , ///
|| ///
kdensity _ATE_x_spill ,lpattern(longdash_dot) xtitle() ///
ytitle(Kernel density) legend(order(1 "ATE(x)" 2 "ATE_spill(x)")) ///
title("Model `model': Comparison of ATE(x) and ATE_spill(x)", size(medlarge))
```



```
* ATET
twoway kdensity ATET_x , ///
|| ///
kdensity _ATET_x_spill ,lpattern(longdash_dot) xtitle() ///
ytitle(Kernel density) legend(order(1 "ATET(x)" 2 "ATET_spill(x)")) ///
title("Model `model': Comparison of ATE(x) and ATE_spill(x)", size(medlarge))
```



```
* ATENT
twoway kdensity ATENT_x , ///
|| ///
kdensity _ATENT_x_spill ,lpattern(longdash_dot) xtitle() ///
ytitle(Kernel density) legend(order(1 "ATENT(x)" 2 "ATENT_spill(x)")) ///
title("Model `model': Comparison of ATE(x) and ATE_spill(x)", size(medlarge))
```



STEP 8. COMPARING UNCONSTRAINED (i.e., WITH SPILLOVER) VS. UNCONSTRAINED (i.e., WITHOUT SPILLOVER) PREDICTIONS

We write a program, "_marg", returning the difference between the constrained and the unconstrained prediction, when cp=1:

```
cap prog drop _marg
program _marg , rclass
qui ntreatreg crime cp inc hoval , hetero(inc hoval) spill(W)
* uncontrained prediction
margins , at(cp= 1)
mat A=r(table)
mat B=A["b","_cons"]
return scalar _marg1=B[1,1]
* contrained prediction
margins , at(cp= 1 z_ws_inc1=0 z_ws_hoval1=0)
mat A=r(table)
mat B=A["b","_cons"]
return scalar _marg2=B[1,1]
end
```

We test:

H₀: $E(y_1|$ with spillover) - $E(y_1|$ without spillover) = 0

We can use "__marg" to test whether predictions are different by bootstrap:

The average difference in prediction is around -10 and it is significant. This entails that, in terms of prediction, the neighbourhood effect accounts for **10 fewer burglaries**.

===> **Conclusion**: not considering "neighbourhood effects" leads to "overestimate" the actual effect of housing location on crime of around a 30%. Although, the Wald-test seems to show that the neighbourhood effect is not significant, if we accept the model with spillovers as the actual one, the average difference in prediction without and with spillovers is around -10 and it is also significant.

Limits and further developments

- Extending the model to "multiple" or "continuous" treatment (i.e., *w* no more binary, but **multi-valued** or **continuous**), by still holding CMI.
- Identifying the model when *w* is **endogenous** (i.e., CMI does not hold), by implementing some GMM-IV estimation.
- So far we have assumed the weighting matrix Ω to be "exogenous". But: what happens if individuals **strategically** modify their "distance weights" to better profit of others' treatment? In this case weights become *endogenous*. It poses severe identification problems.
- Providing Monte Carlo studies to see how the model is robust under different specificationerrors in the weighting matrix Ω provided.
- Going towards a **semi-parametric** approach

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