

Package: splm (via r-universe)

October 16, 2024

Title Econometric Models for Spatial Panel Data

Version 1.6-5

Date 2023-12-11

Description ML and GM estimation and diagnostic testing of econometric models for spatial panel data.

Depends R (>= 2.12.0)

Imports plm, maxLik, MASS, bdsmatrix, nlme, Matrix, spam, methods, spatialreg (>= 1.2-1), spdep (>= 1.2-1), stringr

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LazyLoad yes

NeedsCompilation no

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Repository CRAN

Date/Publication 2023-12-20 16:20:05 UTC

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bsjktest	<i>Baltagi, Song, Jung and Koh LM test for spatial panels</i>
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Description

Baltagi, Song, Jung and Koh joint or conditional LM test for spatial error correlation or serial correlation sub spatial, serial correlation and random effects in panel models

Usage

```
bsjktest(x,...)
## S3 method for class 'formula'
bsjktest(x, data, index=NULL, listw,
test=c("C.1", "C.2", "C.3", "J"), ...)
```

Arguments

x	an object of class formula
data	a data.frame or pdata.frame containing the variables in the model
index	either NULL (default) or a character vector to identify the indexes among the columns of the data.frame
listw	either a matrix or a listw representing the spatial structure
test	one of c("C.1", "C.2", "C.3", "J"), the test to be performed.
...	additional arguments to be passed

Value

an object of class htest

Author(s)

Giovanni Millo

References

Baltagi, B.H., Song, S.H., Jung B. and Koh, W. (2007) Testing panel data regression models with spatial and serial error correlation. *Journal of Econometrics*, **140**, 5-51.

See Also

bsktest

Examples

```
data(Produc, package="plm")
data(usaww)
fm <- log(gsp)~log(pcap)+log(pc)+log(emp)+unemp
bsjktest(fm, data=Produc, listw = usaww, test="C.1")
```

bsktest

*Baltagi, Song and Koh LM test for spatial panels***Description**

Baltagi, Song and Koh marginal or conditional LM test for spatial error correlation or random effects in panel models

Usage

```
bsktest(x,...)
## S3 method for class 'formula'
bsktest(x, data, index=NULL, listw,
test=c("LMH", "LM1", "LM2", "CLMlambda", "CLMmu"),
standardize=FALSE, method = "eigen", ...)
```

Arguments

x	a formula
data	a data.frame or pdata.frame containing the variables in the model
index	either NULL (default) or a character vector to identify the indexes among the columns of the data.frame
listw	a listw representing the spatial structure
test	one of c("LMH", "LM1", "LM2", "CLMlambda", "CLMmu"), the test to be performed
standardize	whether to standardize the test statistic or not (applies only to LM1 and LM2)
method	select a method for ML in "CLMmu". the default is "eigen"
...	additional arguments to be passed

Value

an object of class htest

Author(s)

Gianfranco Piras

References

Baltagi, B.H., Song, S.H. and Koh, W. (2003) Testing panel data regression models with spatial error correlation. *Journal of Econometrics*, **117**, 123–150.

Millo, G., Piras, G. (2012) splm: Spatial Panel Data Models in R. *Journal of Statistical Software*, **47(1)**, 1–38. URL <http://www.jstatsoft.org/v47/i01/>.

See Also

sphtest

Examples

```
data(Produc, package="plm")
data(usaww)
fm <- log(gsp)~log(pcap)+log(pc)+log(emp)+unemp
bsktest(fm,data=Produc, listw = spdep::mat2listw(usaww),
        test="LM1")
```

effects.splm	<i>method for extracting fixed effects</i>
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Description

Methods used for extracting fixed effects from objects of class splm where type is one of "fixed effects lag" or "fixed effects error"

Usage

```
## S3 method for class 'splm'
effects(object,...)
```

Arguments

object	an object of class 'splm'
...	additional arguments to be passed over

Details

If the argument object is not of class splm the function will terminate with an error.

If the argument object is of class splm but type is not one of "fixed effects lag" or "fixed effects error", the function will terminate with an error.

Value

An object of class effects.splm

res	a list whose elements are various type of fixed effects and the intercept (when present)
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Author(s)

Gianfranco Piras

References

Elhorst, J.P. (2003) Specification and estimation of spatial panel data models, *International Regional Science Review*, **26**, pages 244–268.

Elhorst, J.P. (2009) Spatial panel data models, In Fischer, M.M. and Getis, A. (eds), *Handbook of Applied Spatial Analysis* Springer, Berlin.

See Also

spml summary.effects.splm

Examples

```
data(Produc, package = "plm")
data(usaww)
fm <- log(gsp) ~ log(pcap) + log(pc) + log(emp) + unemp
err <- spml(fm, data = Produc, listw = spdep::mat2listw(usaww), model="within")
summary(err)
eff <- effects(err)
print(eff)
```

Insurance

Insurance consumption across Italian provinces, 1998-2002

Description

A panel of 103 observations
number of observations : 515
observation : provinces
country : Italy

Usage

data(Insurance)

Format

A dataframe containing :

code the province code according to Istat**year** the year of observation**ppcd** real per capita premiums in 2000 euros, non-life insurance excluding mandatory motor third-party liability

rgdp real per-capita GDP
bank real per-capita bank deposits
den population density per square Km
rirs real interest rate on lending to families and small enterprises
agen density of insurance agencies per 1000 inhabitants
school share of people with second grade schooling or more
vaagr share of value added, agriculture
fam average number of family members
inef judicial inefficiency index: average years to settle first degree of civil case
trust survey result to the question "do you trust others?"
dXX year dummies
NorthWest macroregional dummy
NorthEast macroregional dummy
Centre macroregional dummy
South macroregional dummy
Islands macroregional dummy (Sicily and Sardinia)

Author(s)

Giovanni Millo

Source

Giovanni Millo and Gaetano Carmeci, (2011) "Non-life insurance consumption in Italy: a sub-regional panel data analysis", *Journal of Geographical Systems*, **13:273–298**.

itaww

Spatial weights matrix - Italian provinces

Description

Spatial weights matrix of the 103 Italian provinces as in 1992-2005.

Usage

data(itaww)

Format

A matrix with elements different from zero if province i and j are neighbors. Weights are row-standardized. Messina and Reggio Calabria, divided by the Messina Strait, are considered neighbours.

Author(s)

Giovanni Millo

listw2dgCMatrix	<i>Interface between Matrix class objects and weights list</i>
-----------------	--

Description

Interface between Matrix class objects and weights list

Usage

```
listw2dgCMatrix(listw, zero.policy = NULL)
```

Arguments

listw	a listw object created for example by <code>spdep::nb2listw</code>
zero.policy	See <code>lagsarlm</code> for details

Value

Matrix class object: a sparse Matrix

Author(s)

Gianfranco Piras

Examples

```
data(columbus, package="spdep")
listw<-spdep::nb2listw(col.gal.nb)
spW<-listw2dgCMatrix(listw)
```

print.splm	<i>print method for class splm</i>
------------	------------------------------------

Description

Method to print objects of class `summary.splm` and `splm`

Usage

```
## S3 method for class 'splm'
print(x, digits = max(3,getOption("digits") -3), ...)
```

Arguments

x	an object of class <code>splm</code>
digits	minimal number of significant digits, see <code>print.default</code>
...	additional arguments to be passed

Details

The summary function `summary.splm` returns an objects of class 'splm' organized in a coefficient matrix.

Also a matrix for the error components, or the spatial coefficients will be generated depending on the estimated model.

Author(s)

Giovanni Millo, Gianfranco Piras

See Also

`splm`, `spgm`

Examples

```
data(Produc, package = "plm")
data(usaww)
spremod<-splm(log(gsp)~log(pcap)+log(pc)+log(emp)+unemp, data=Produc,
listw = spdep::mat2listw(usaww), model="random", lag=TRUE, spatial.error="none")
summary(spremod)
```

RiceFarms

Production of Rice in India

Description

yearly observations of 171 farms

number of observations : 1026

country : Indonesia

economic topic : producer behavior

econometrics topic : error component

Usage

```
data(RiceFarms)
```

Format

A dataframe containing :

id the farm identifier

time the growing season

size the total area cultivated with rice, measured in hectares

status land status, on of 'owner' (non sharecroppers, owner operators or leasholders or both), 'share' (sharecroppers), 'mixed' (mixed of the two previous status)

- varieties** one of 'trad' (traditional varieties), 'high' (high yielding varieties) and 'mixed' (mixed varieties)
- bimas** bIMAS is an intensification program ; one of 'no' (non-bimas famer), 'yes' (bimas farmer) or 'mixed' (part but not all of farmer's land was registered to be in the bimas program)
- seed** seed in kilogram
- urea** urea in kilogram
- phosphate** phosphate in kilogram
- pesticide** pesticide cost in Rupiah
- pseed** price of seed in Rupiah per kg
- purea** price of urea in Rupiah per kg
- pphosph** price of phosphate in Rupiah per kg
- hiredlabor** hired labor in hours
- famlabor** family labor in hours
- totlabor** total labor (excluding harvest labor)
- wage** labor wage in Rupiah per hour
- goutput** gross output of rice in kg
- noutput** net output, gross output minus harvesting cost (paid in terms of rice)
- price** price of rough rice in Rupiah per Kg
- region** one of 'wargabinangun', 'langan', 'gunungwangi', 'malausma', 'sukaambit', 'ciwangi'

Source

Journal of Applied Econometrics Data Archive.

References

- Qu Feng and William C. Horrace, (2012) "Alternative Measures of Technical Efficiency: Skew, Bias and Scale", *Journal of Applied Econometrics*, **forthcoming**.
- Horrace, W.C. and P. Schmidt (1996) "Confidence statements for efficiency estimates from stochastic frontier models", *Journal of Productivity Analysis*, **7**, 257–282.

riceww

Spatial weights matrix of Indonesian rice farms

Description

Spatial weights matrix of the 171 farms in the Indonesian Rice Farming example. Farms in the same village (out of six) are considered contiguous.

Usage

data(riceww)

Format

A matrix with elements different from zero if farms i and j are neighbors. Farms are considered neighbors if in the same village. Weights are row-standardized.

Author(s)

Giovanni Millo, data provided by Yves Croissant

 rwtest

Randomization-based test of spatial dependence for panel models

Description

Randomization-based test of spatial dependence for panel models, robust to global dependence induced by common factors and to persistence (serial correlation) in the data

Usage

```
rwtest(x, ...)
## S3 method for class 'formula'
rwtest(x, data, w, index = NULL, model = NULL,
       replications = 99, seed=NULL, order=1,
       mc=1, test = c("rho", "cd", "sclm"),
       alternative=c("twosided", "onesided",
                    "symmetric"), ...)

## S3 method for class 'panelmodel'
rwtest(x, w, replications = 99, seed=NULL,
       order=1, mc=1,
       test = c("rho", "cd", "sclm"),
       alternative=c("twosided", "onesided",
                    "symmetric"), ...)

## S3 method for class 'pseries'
rwtest(x, w, replications = 99, seed=NULL,
       order=1, mc=1,
       test = c("rho", "cd", "sclm"),
       alternative=c("twosided", "onesided",
                    "symmetric"), ...)
```

Arguments

x	an object of class <code>formula</code> , <code>panelmodel</code> , or <code>pseries</code> (depending on the respective interface) describing the model to be tested
data	a <code>data.frame</code>

<code>w</code>	a $n \times n$ matrix describing proximity between individuals, with $w_{ij} = a$ where a is any number such that <code>as.logical(a)==TRUE</code> , if i, j are neighbours, 0 or any number b such that <code>as.logical(b)==FALSE</code> elsewhere. Only the lower triangular part (without diagonal) of w after coercing by <code>as.logical()</code> is evaluated for neighbouring information (but w can be symmetric). See also Details and Examples .
<code>index</code>	an optional numerical index, in case data has to be formatted by <code>plm.data</code>
<code>model</code>	an optional character string indicating which type of model to estimate; if left to <code>NULL</code> , the original heterogeneous specification of Pesaran is used
<code>replications</code>	the number of Monte Carlo randomizations of the neighbourhood matrix (default: 99),
<code>seed</code>	the optional random seed,
<code>order</code>	the order of neighbourhood to test for,
<code>mc</code>	the number of parallel threads to execute; defaults to 1 (serial execution); is limited to the number of execution cores actually available, and depends on operating system support.
<code>test</code>	the type of test statistic to be returned. One of <ul style="list-style-type: none"> • "rho" for the average correlation coefficient, • "cd" for Pesaran's CD statistic, or • "sclm" for the scaled version of Breusch and Pagan's LM statistic,
<code>alternative</code>	the alternative hypothesis for the test, defaulting to (asymmetric) twosided,
<code>...</code>	further arguments to be passed on to <code>plm</code> , such as e.g. <code>effect</code> or <code>random.method</code>

Details

This test is meant as a generalization of Pesaran's spatial dependence test "CD(p)" for robustness against global dependence (perhaps of the factor type) and persistence in the data, both of which the original test does not tolerate.

The procedure can be applied to model residuals as well as to individual `pseries`. See the comments in `pcdtest` as for the different methods.

Space is defined supplying a proximity matrix (elements coercible to `logical`) with argument `w` which provides information on whether any pair of individuals are neighbours or not. If `order=1`, only first-order neighbouring pairs will be used in computing the test; else, w will be transformed in the neighbourhood matrix of the appropriate order. The matrix need not be binary, so commonly used "row-standardized" matrices can be employed as well. `nb` objects from **spdep** must instead be transformed into matrices by **spdep**'s function `nb2mat` before using.

Notice that the "rho" and "cd" tests are permutationally equivalent.

The test is suitable also for unbalanced panels.

The test on a `pseries` is the same as a test on a pooled regression model of that variable on a constant, i.e. `rwtest(some_pseries)` is equivalent to `rwtest(plm(some_var ~ 1, data = some_pdata.frame, model = "pooling"))` and also equivalent to `rwtest(some_var ~ 1, data = some_data)`, where `some_var` is the variable name in the data which corresponds to `some_pseries`.

Value

An object of class "htest".

Author(s)

Giovanni Millo

References

Millo, G. (2016), A simple randomization test for spatial dependence in the presence of common factors and serial correlation, (*unpublished*), **xx(x)**, pp. xxx–xxx. Pesaran, M.H. (2004), General Diagnostic Tests for Cross Section Dependence in Panels, *CESifo Working Paper 1229*. Pesaran, M.H. (2015), Testing Weak Cross–Sectional Dependence in Large Panels, *Econometric Reviews*, **34**(6-10), pp. 1089–1117.

Examples

```
data(Produc, package = "plm")
data(usaww)
fm <- log(gsp) ~ log(pcap) + log(pc) + log(emp) + unemp
## test on heterogeneous model (separate time series regressions)
rwtest(fm, data = Produc, w=usaww, index = c("state", "year"))

## test on two-way fixed effects homogeneous model
rwtest(fm, data = Produc, w=usaww, index = c("state", "year"),
       model = "within", effect = "twoways")

## test on panelmodel object
library(plm)
g <- plm(fm, data = Produc)
rwtest(g, w=usaww)

## test on pseries, higher-order neighbourhood
pprod <- pdata.frame(Produc)
rwtest(pprod$gsp, w=usaww, order=3)
```

slag

Spatial lag operator

Description

Spatial lagging method for vectors or pseries objects.

Usage

```
## S3 method for class 'pseries'
slag(x, listw, maxlag, ...)
## Default S3 method:
slag(x, listw, maxlag, index, ...)
```

Arguments

x an object of class pseries
 listw an object of class listw
 maxlag the spatial lag order (including lower)
 index the default method needs a well-specified index
 ... additional arguments to be passed

Value

a pseries

Author(s)

Giovanni Millo

Examples

```
data(Produc, package="plm")
data(usaww)
usalw <- spdep::mat2listw(usaww)
fm <- log(gsp)~log(pcap)+log(pc)+log(emp)+unemp+slag(log(pcap),
  listw=usalw)
slxmod <- spreml(fm, data=Produc, w = usaww,
  model="pooling", lag=FALSE, errors="ols")
```

slmtest

Locally robust panel Lagrange Multiplier tests for spatial dependence

Description

Locally robust LM tests for spatial lag (error) correlation sub spatial error (lag) correlation in panel models

Usage

```
slmtest(x,...)
## S3 method for class 'formula'
slmtest(formula, data, listw, model="pooling",
  test=c("lme","lml","rlme","rlml"), index=NULL, ...)
## S3 method for class 'plm'
slmtest(x, listw,
  test=c("lme","lml","rlme","rlml"), ...)
```

Arguments

formula	an object of class formula
data	a data.frame or pdata.frame containing the variables in the model
x	an object of class plm
listw	either a matrix or a listw representing the spatial structure
model	a character value specifying the transformation to be applied to the data.
test	one of c("lme", "lml", "rlme", "rlml"), the test to be performed.
index	either NULL (default) or a character vector to identify the indexes among the columns of the data.frame
...	additional arguments to be passed

Details

This tests are panel versions of the locally robust LM tests of Anselin et al. (1996), based on a pooling assumption: i.e., they do not allow for any kind of individual effect. Therefore it is advisable to employ a within transformation whenever individual effects cannot be ruled out.

It must be kept in mind that these locally robust procedures have been designed for situations in which the "other" effect is not of substantial magnitude, and can behave suboptimally otherwise.

Four tests are available to be chosen through the test argument: "lml" for "LM lag" and, respectively, "lme" for "LM error" are the standard, non-robust versions, obtained simply pooling the cross-sectional versions; "rlml" and "rlme" are, respectively, the locally robust test for lag, allowing for a spatial error; and for error, allowing for a spatial lag.

The model argument, specified according to the standards of plm, is passed on internally and employed to determine the panel data transformation to be applied before calculating the test. Defaults to "pooling" (no transformation).

Value

an object of class htest

Author(s)

Giovanni Millo

References

Anselin, L., Bera, A.K., Florax, R. and Yoon, M.J. (1996) Simple diagnostic tests for spatial dependence. *Regional Science and Urban Economics*, **26**(1), 77-104. Elhorst, J.P. (2014) Spatial Panel data Models, in *Spatial Econometrics* (Springer) 37-93.

Examples

```
data(Produc, package="plm")
data(usaww)
fm <- log(gsp)~log(pcap)+log(pc)+log(emp)+unemp
## robust LM test for spatial error sub spatial lag
```

```
## model on original data, pooling hypothesis
slmtest(fm, data=Produc, listw = usaww, test="rlme")
## model on within-transformed (time-demeaned) data,
## eliminates individual effects
slmtest(fm, data=Produc, listw = usaww, test="rlme",
        model="within")
```

spgm

GM estimation of spatial panel data models

Description

GM estimation of panel data models with spatially correlated errors components of the form:

$$y_N(t) = \lambda W y + X_N(t)\beta + u_N(t)$$

$$u_N(t) = \rho W_N u_N(t) + \epsilon(t)$$

$$\epsilon_N = (e_T \otimes I_N)\mu_N + \nu_N$$

where ρ , and the variance components σ_μ^2 and σ_ν^2 are estimated by GM, and the model coefficients by a Feasible GLS estimator. The model can also include additional (other than the spatial lag) endogenous variables.

Usage

```
spgm(formula, data=list(), index=NULL, listw = NULL, listw2 = NULL, Durbin = FALSE,
      model = c("within", "random"), lag = FALSE, spatial.error = TRUE,
      moments = c("initial", "weights", "fullweights"), endog = NULL,
      instruments = NULL, lag.instruments = FALSE, verbose = FALSE,
      method = c("w2sls", "b2sls", "g2sls", "ec2sls"), control = list(),
      optim.method = "nlsminb", pars = NULL)
```

Arguments

formula	a description of the model to be fit. The details of model specification are given for <code>lm</code>
data	an object of class <code>data.frame</code> or <code>pdata.frame</code> . An optional data frame containing the variables in the model. When the object is a <code>data.frame</code> , the first two columns may contain the indexes. See <code>index</code>
index	if not <code>NULL</code> (default), a character vector to identify the indexes among the columns of the <code>data.frame</code>
listw	an object of class <code>listw</code> , <code>matrix</code> , or <code>Matrix</code>

<code>listw2</code>	an object of class <code>listw</code> , <code>matrix</code> , or <code>Matrix</code> . Only if both <code>lag</code> and <code>spatial.error</code> are both TRUE
<code>Durbin</code>	default FALSE. If TRUE all the regressors are also spatially lagged. If it is a formula (<code>Durbin = ~ xi + x2</code>), only the variables specified in the formula are lagged.
<code>model</code>	One of "within" or "random". The assumption made on the individual effects
<code>lag</code>	if TRUE a spatial lag of the dependent variable is added to the regression equation
<code>spatial.error</code>	a logic vector. If TRUE the spatial autoregressive error term is added to the model and an estimate for ρ is produced
<code>moments</code>	"initial" (default) defines the set of GM estimator to be used. Alternatives are "weights" and "fullweights" (See Details)
<code>endog</code>	additional endogenous variables. Default NULL. If not NULL should be specified as a formula with no dependent variable (<code>endog = ~ x1 + x2</code>). Note the ~ before the expression.
<code>instruments</code>	external instruments. Default NULL. If not NULL should be specified as a formula with no dependent variable (<code>instruments = ~ x1 + x2</code>). Note the ~ before the expression.
<code>lag.instruments</code>	should the external instruments be spatially lagged?
<code>verbose</code>	default FALSE, If TRUE reports function values during optimization
<code>method</code>	One of "w2s1s", "b2s1s", "g2s1s", "ec2s1s". (See Details)
<code>control</code>	a list of control parameters for the optimization
<code>optim.method</code>	default set to "nlnmb". or optionally a method passed to <code>optim</code> to use an alternative optimizer.
<code>pars</code>	initial values of the parameter <code>rho</code> and <code>sigmav</code> . The default for <code>rho</code> is to start from a regression of the spatially lagged residuals on the residuals (depending on the model). for <code>sigmav</code> the starting value is the variance of the residuals (again this depends on the model).

Details

The function is a very general interface to estimate various nested specifications of the general model including additional endogenous variables described above. When both `spatial.error` and `lag` are FALSE the model reduces to a panel data model with an additional endogeneous variable. The function then uses `ivsp1m` to perform the Instrumental Variables and two-stage least squares for panel data model. `method = "w2s1s"` corresponds to the fixed effects estimator, `method = "b2s1s"` to the between effects model, `method = "g2s1s"` to the GLS random effects model, and `method = "ec2s1s"` to the Baltagi's EC2SLS.

When `spatial.error` is TRUE and `lag` is FALSE the model is one with spatially autocorrelated error components. If `effects` is "random", the Kapoor et al. (2007) GM estimator is performed and the residuals in the first step come from an OLS regression. When `moments` is "initial", the initial estimator is calculated. This first set of GM estimators is based only on a subset of the moments conditions and assigns equal weights to each of them. When `moments` is "fullweights", the second set of GM estimators is calculated. This estimator is based on the full set of moments

conditions. It also involves the expression for the variance covariance matrix of the sample moments calculated under the assumption of normally distributed innovations. The calculation of the trace terms in the expression of the variance covariance matrix of the sample moments uses codes from the Matrix package. When moments is "weights", the third set of GM estimator is used. This is motivated by computational issues. The procedure is analogous to the second one but uses a simplified expression for the variance covariance matrix of the sample moments. If effects is "fixed", the initial estimator is a within estimator and the moments conditions of Kapoor et al. (2007) are modified accordingly.

Finally, when both `spatial.error` and `lag` are TRUE the complete model is estimated (with or without additional endogenous variables). OLS residuals are no longer consistent because of the spatially lagged dependent variable. If effects is "random", two initial estimators are computed: a within two-stage least squares and a between two stage least squares. The two sets of corresponding residuals are used in the spatial generalized moments estimator (GM) where the moments conditions of Kapoor et al. (2007) are again modified accordingly. If effects is "fixed", the initial estimator is a within two stage least squares estimator and the moments conditions of Kapoor et al. (2007) are modified accordingly.

Note that for the random effects models, σ_{μ}^2 is not reported. σ_1^2 is reported instead. However, a value for σ_{μ}^2 can easily be obtained from:

$$\sigma_1^2 = \sigma_{\nu}^2 + T\sigma_{\mu}^2$$

The function also produces an estimate for θ which is a function of the variance components.

Value

An object of class "splm".

<code>coefficients</code>	GLS coefficients estimate of the model parameters
<code>vcov</code>	the variance covariance matrix of the estimated coefficients
<code>residuals</code>	the GLS residuals
<code>fitted.values</code>	difference between response variable and residuals
<code>sigma2</code>	GLS residuals variance
<code>type</code>	'a description of the model estimated'
<code>rho</code>	a vector including the spatial parameter and the variance components (see Details)
<code>model</code>	the matrix of the data used
<code>call</code>	the call used to create the object

Author(s)

Gianfranco Piras

References

Kapoor, M., Kelejian, H.H. and Prucha, I.R. (2007) Panel data model with spatially correlated error components, *Journal of Econometrics*, **140**, pages 97–130.

Mutl, J., and Pfaffermayr, M. (2011) The Hausman test in a Cliff and Ord panel model, *Econometrics Journal*, **14**, pages 48–76.

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Millo, G., Piras, G. (2012) splm: Spatial Panel Data Models in R. *Journal of Statistical Software*, **47(1)**, 1–38. URL <http://www.jstatsoft.org/v47/i01/>.

Examples

```
library(splm)
library(plm)
data(Produc)
data(usaww)
#####SPATIAL LAG MODEL#####
#no space no endog: error use plm
## Not run: GM_lag_b2s1s <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
                             lag = FALSE, spatial.error = FALSE, method = "b2s1s",
                             listw = usaww)

## End(Not run)
#no space but endog
GM_b2s1s_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
                  lag = FALSE, spatial.error = FALSE, endog = ~ log(pcap),
                  method = "b2s1s",
                  instruments = ~log(hwy), listw = usaww)

summary(GM_b2s1s_e)
GM_g2s1s_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
                  lag = FALSE, spatial.error = FALSE, endog = ~ log(pcap),
                  method = "g2s1s",
                  instruments = ~log(hwy), listw = usaww)

summary(GM_g2s1s_e)
GM_ec2s1s_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
                  lag = FALSE, spatial.error = FALSE, endog = ~ log(pcap),
                  method = "ec2s1s",
                  instruments = ~log(hwy), listw = usaww)

summary(GM_ec2s1s_e)
GM_w2s1s_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
                  lag = FALSE, spatial.error = FALSE, endog = ~ log(pcap),
                  method = "w2s1s",
                  instruments = ~log(hwy), listw = usaww)

summary(GM_w2s1s_e)
#SPATIAL LAG MODEL
GM_lag_b2s1s <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
                   lag = TRUE, spatial.error = FALSE, method = "b2s1s",
                   listw = usaww)

summary(GM_lag_b2s1s)
GM_lag_g2s1s <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
                   lag = TRUE, spatial.error = FALSE, method = "g2s1s",
                   listw = usaww)
```

```

summary(GM_lag_g2s1s)
GM_lag_ec2s1s <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = TRUE, spatial.error = FALSE, method = "ec2s1s",
  listw = usaww)
summary(GM_lag_ec2s1s)
GM_lag_w2s1s <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = TRUE, spatial.error = FALSE,
  listw = usaww, method = "w2s1s")
summary(GM_lag_w2s1s)
#endogenous
GM_lag_b2s1s_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
  lag = TRUE,
  spatial.error = FALSE, endog = ~ log(pcap),
  method = "b2s1s",
  instruments = ~log(hwy), listw = usaww)
summary(GM_lag_b2s1s_e)
GM_lag_g2s1s_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
  lag = TRUE,
  spatial.error = FALSE, endog = ~ log(pcap),
  method = "g2s1s",
  instruments = ~log(hwy), listw = usaww)
summary(GM_lag_g2s1s_e)
GM_lag_ec2s1s_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
  lag = TRUE,
  spatial.error = FALSE, endog = ~ log(pcap),
  method = "ec2s1s",
  instruments = ~log(hwy), listw = usaww)
summary(GM_lag_ec2s1s_e)
GM_lag_w2s1s_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
  lag = TRUE,
  spatial.error = FALSE, endog = ~ log(pcap),
  method = "w2s1s",
  instruments = ~log(hwy), listw = usaww)
summary(GM_lag_w2s1s_e)

#####SPATIAL ERROR MODEL#####
GM_error_within <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = FALSE, spatial.error = TRUE, model = "within",
  listw = usaww)
summary(GM_error_within)
GM_error_random <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = FALSE, spatial.error = TRUE, model = "random",
  listw = usaww)
summary(GM_error_random)
GM_error_within_fw <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = FALSE, spatial.error = TRUE, model = "within",
  listw = usaww, moments = "fullweights")
summary(GM_error_within_fw)
GM_error_random_w <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = FALSE, spatial.error = TRUE, model = "random",
  listw = usaww, moments = "fullweights")
summary(GM_error_random_w)
#endogenous

```

```

GM_error_within_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
  lag = FALSE,
  spatial.error = TRUE, endog = ~ log(pcap),
  model = "within",
  instruments = ~log(hwy), listw = usaww)
summary(GM_error_within_e)
GM_error_random_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
  lag = FALSE,
  spatial.error = TRUE, endog = ~ log(pcap),
  model = "random",
  instruments = ~log(hwy), listw = usaww)
summary(GM_error_random_e)

#####SARAR MODEL#####
GM_sarar_within <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = TRUE, spatial.error = TRUE, model = "within",
  listw = usaww)
summary(GM_sarar_within)
GM_sarar_random <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = TRUE, spatial.error = TRUE, model = "random",
  listw = usaww)
summary(GM_sarar_random)
GM_sarar_within_fw <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = TRUE, spatial.error = TRUE, model = "within",
  listw = usaww, moments = "fullweights")
summary(GM_sarar_within_fw)
GM_sarar_random_fw <- spgm(log(gsp) ~ log(pcap) + log(pc) + unemp, data=Produc,
  lag = TRUE, spatial.error = TRUE, model = "random",
  listw = usaww, moments = "fullweights")
summary(GM_sarar_random_fw)
#endogenous
GM_sarar_within_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
  lag = TRUE,
  spatial.error = TRUE, endog = ~ log(pcap),
  model = "within",
  instruments = ~log(hwy), listw = usaww)
summary(GM_sarar_within_e)
GM_sarar_random_e <- spgm(log(gsp) ~ log(pc) + unemp, data=Produc,
  lag = TRUE,
  spatial.error = TRUE, endog = ~ log(pcap),
  model = "random",
  instruments = ~log(hwy), listw = usaww)
summary(GM_sarar_random_e)

```

sphtest

Hausman test for spatial panel data models

Description

Hausman specification test for spatial panel data models

Usage

```

sphtest(x, ...)
## S3 method for class 'formula'
sphtest(x, data, index = NULL, listw,
        spatial.model = c("lag", "error", "sarar"),
        method = c("ML", "GM"), errors = c("KPP", "BSK"),...)
## S3 method for class 'splm'
sphtest(x, x2, ...)

```

Arguments

<code>x</code>	an object of class <code>formula</code> or <code>splm</code>
<code>x2</code>	an object of class <code>splm</code>
<code>data</code>	an object of class <code>data.frame</code> or <code>pdata.frame</code> . An optional data frame containing the variables in the model. When the object is a <code>data.frame</code> , the first two columns may contain the indexes. See <code>index</code>
<code>index</code>	if not <code>NULL</code> (default), a character vector to identify the indexes among the columns of the <code>data.frame</code>
<code>listw</code>	an object of class <code>listw</code> created for example by <code>spdep::nb2listw</code>
<code>spatial.model</code>	one of <code>c("lag", "error", "sarar")</code> , the model to be estimated (only lag, only error, both lag and error dependence)
<code>method</code>	one of <code>c("ML", "GM")</code>
<code>errors</code>	one of <code>c("BSK", "KPP")</code> . When <code>method</code> is <code>"ML"</code> defines the specification of the innovations
<code>...</code>	additional arguments to be passed

Value

an object of class `hctest`

Author(s)

Gianfranco Piras

References

Millo, G., Piras, G. (2012) `splm`: Spatial Panel Data Models in R. *Journal of Statistical Software*, **47(1)**, 1–38. URL <http://www.jstatsoft.org/v47/i01/>.

See Also

`spgm`

Examples

```

data(Produc, package="plm")
data(usaww)
fm <- log(gsp)~log(pcap)+log(pc)+log(emp)+unemp
test1<-sphtest(fm,data=Produc, listw = spdep::mat2listw(usaww),
              spatial.model = "error", method="GM")

test1
mod1<- spgm(fm, data=Produc, listw = usaww, model = "random",
            spatial.error = TRUE, moments="fullweights")
mod2<- spgm(fm, data=Produc, listw = usaww, model = "within",
            spatial.error = TRUE)
test2<-sphtest(mod1, mod2)
test2

```

 spml

Spatial Panel Model by Maximum Likelihood

Description

Maximum likelihood (ML) estimation of spatial panel models, possibly with fixed or random effects.

Usage

```

spml(formula, data, index=NULL, listw, listw2=listw, na.action,
      model=c("within","random","pooling"),
      effect=c("individual","time","twoways"),
      lag=FALSE, spatial.error=c("b","kkp","none"),
      ...)
## S3 method for class 'splm_ML'
impacts(obj, listw = NULL,
        time = NULL, ...,
        tr = NULL, R = 200,
        type = "mult",
        empirical = FALSE, Q = NULL)
## S3 method for class 'splm_GM'
impacts(obj, ..., tr=NULL,
        R=NULL, listw=NULL,
        type = "mult",
        time = NULL,
        evalues=NULL, tol=1e-6,
        empirical=FALSE, Q=NULL,
        KPformula = FALSE, prt = TRUE)

```

Arguments

formula a symbolic description of the model to be estimated

<code>data</code>	an object of class <code>data.frame</code> or <code>pdata.frame</code> . A data frame containing the variables in the model. When the object is a <code>data.frame</code> , the first two columns shall contain the indexes, unless otherwise specified. See <code>index</code>
<code>index</code>	if not <code>NULL</code> (default), a character vector to identify the indexes among the columns of the <code>data.frame</code>
<code>listw</code>	an object of class <code>listw</code> or a matrix. It represents the spatial weights to be used in estimation.
<code>listw2</code>	an object of class <code>listw</code> or a matrix. Second of set spatial weights for estimation, if different from the first (e.g., in a 'sarar' model).
<code>na.action</code>	see spdep for more details.
<code>model</code>	one of <code>c("within", "random", "pooling")</code> .
<code>effect</code>	one of <code>c("individual", "time", "twoways")</code> ; the effects introduced in the model.
<code>lag</code>	default= <code>FALSE</code> . If <code>TRUE</code> , a spatial lag of the dependent variable is added.
<code>spatial.error</code>	one of <code>c("b", "kcp", "none")</code> . The type of spatial error in the specification, if any. See details.
<code>...</code>	additional argument to pass over to other functions
<code>obj</code>	fitted model object
<code>time</code>	??time??
<code>tr</code>	A vector of traces of powers of the spatial weights matrix created using 'trW', for approximate impact measures
<code>R</code>	If given, simulations are used to compute distributions for the impact measures, returned as 'mcmc' objects
<code>type</code>	Either "mult" (default) for powering a sparse matrix (with moderate or larger N, the matrix becomes dense, and may lead to swapping), or "MC" for Monte Carlo simulation of the traces (the first two simulated traces are replaced by their analytical equivalents), or "moments" to use the looping space saving algorithm proposed by Smirnov and Anselin (2009) - for "moments", 'W' must be symmetric, for row-standardised weights through a similarity transformation
<code>empirical</code>	Argument passed to 'mvrnorm' (default <code>FALSE</code>)
<code>Q</code>	default <code>NULL</code> , else an integer number of cumulative power series impacts to calculate if 'tr' is given
<code>evalues</code>	vector of eigenvalues of spatial weights matrix for impacts calculations
<code>tol</code>	Argument passed to 'mvrnorm'
<code>KPformula</code>	not yet implemented
<code>prt</code>	not yet implemented

Details

The models are estimated by two-step Maximum Likelihood. Further optional parameters to be passed on to the estimator may be: `pvar`: if `TRUE` the `pvar` function is called `hess`: if `TRUE` use numerical Hessian instead of GLS for the standard errors of the estimates `quiet`: if `FALSE` report

function and parameters values during optimization initial: one of c("zeros", "estimate"), the initial values for the parameters. If "zeros" a vector of zeros is used. if "estimate" the initial values are retrieved from the estimation of the nested specifications. Alternatively, a numeric vector can be specified. x.tol: Tolerance. See nlminb for details. rel.tol: Relative tolerance. See nlminb for details.

Value

An object of class "splm".

coefficients	coefficients estimate of the model parameters
arcoef	the coefficient for the spatial lag on y
errcomp	the estimates of the error variance components
vcov	the asymptotic variance covariance matrix of the estimated coefficients
vcov.arcoef	the asymptotic variance of the estimated spatial lag parameter
vcov.errcomp	the asymptotic variance covariance matrix of the estimated error covariance parameters
type	'random effects ML'
residuals	the model residuals
fitted.values	the fitted values, calculated as $\hat{y} = X\hat{\beta}$
sigma2	GLS residuals variance
model	the matrix of the data used
call	the call used to create the object
logLik	the value of the log likelihood function at the optimum
errors	the value of the errors argument

Author(s)

Giovanni Millo

References

- Baltagi, B.H., Song, S.H., Jung B. and Koh, W. (2007) Testing panel data regression models with spatial and serial error correlation. *Journal of Econometrics*, **140**, 5-51.
- Millo, G., Piras, G. (2012) splm: Spatial Panel Data Models in R. *Journal of Statistical Software*, **47(1)**, 1-38. URL <http://www.jstatsoft.org/v47/i01/>.

See Also

spgm

Examples

```

data(Produc, package = "plm")
data(usaww)
fm <- log(gsp) ~ log(pcap) + log(pc) + log(emp) + unemp
## the two standard specifications (SEM and SAR) one with FE
## and the other with RE:
## fixed effects panel with spatial errors
fespaterr <- spml(fm, data = Produc, listw = spdep::mat2listw(usaww),
                 model="within", spatial.error="b", Hess = FALSE)
summary(fespaterr)
## random effects panel with spatial lag
respatlag <- spml(fm, data = Produc, listw = spdep::mat2listw(usaww),
                 model="random", spatial.error="none", lag=TRUE)
summary(respatlag)
## calculate impact measures
#impac1 <- impacts.splm(respatlag, listw = spdep::mat2listw(usaww, #style = "W"), time = 17)
#summary(impac1, zstats=TRUE, short=TRUE)

```

```
spreml
```

Spatial Panel Model with Random Effects by Maximum Likelihood

Description

Maximum likelihood (ML) estimation of spatial panel models with random effects and serial error correlation.

Usage

```

spreml(formula, data, index = NULL, w, w2=w, lag = FALSE,
       errors = c("semsrre", "semsr", "srre", "semre",
                 "re", "sr", "sem", "ols", "sem2srre",
                 "sem2re", "semgre"),
       pvar = FALSE, hess = FALSE, quiet = TRUE,
       initval = c("zeros", "estimate"),
       x.tol = 1.5e-18, rel.tol = 1e-15, ...)

```

Arguments

formula	a symbolic description of the model to be estimated
data	an object of class <code>data.frame</code> or <code>pdata.frame</code> . A data frame containing the variables in the model. When the object is a <code>data.frame</code> , the first two columns shall contain the indexes, unless otherwise specified. See <code>index</code>
index	if not <code>NULL</code> (default), a character vector to identify the indexes among the columns of the <code>data.frame</code>
w	an object of class <code>listw</code> or a matrix. It represents the spatial weights to be used in estimation.

w2	an object of class <code>listw</code> or a matrix. Second set of spatial weights for estimation, if different from the first (e.g., in a 'sarar' model).
lag	default=FALSE. If TRUE, a spatial lag of the dependent variable is added.
errors	Specifies the error covariance structure. See details.
pvar	legacy parameter here only for compatibility.
hess	default=FALSE. If TRUE estimate the covariance for <code>beta_hat</code> by numerical Hessian instead of GLS at optimal values.
quiet	default=TRUE. If FALSE, report function and parameters values during optimization.
initval	one of <code>c("zeros", "estimate")</code> , the initial values for the parameters. If "zeros" a vector of zeros is used. if "estimate" the initial values are retrieved from the estimation of the nested specifications. Alternatively, a numeric vector can be specified.
x.tol	control parameter for tolerance. See <code>nlm</code> for details.
rel.tol	control parameter for relative tolerance. See <code>nlm</code> for details.
...	additional arguments to pass over to other functions, e.g. <code>method</code> .

Details

Second-level wrapper for estimation of random effects models with serial and spatial correlation. The specifications without serial correlation (no "sr" in errors) can be called through `spml`, the extended ones only through `spreml`. The models are estimated by two-step Maximum Likelihood. Abbreviations in errors correspond to: "sem" Anselin-Balagati type spatial autoregressive error; if present, random effects are not spatially correlated; "sem2" Kapoor, Kelejian and Prucha-type spatial autoregressive error model with spatially correlated random effects; "sr" serially correlated remainder errors; "re" random effects; "ols" spherical errors (usually combined with `lag=T`). The optimization method can be passed on as optional parameter. Default is "nlminb"; all constrained optimization methods from `maxLik` are allowed ("BFGS", "NM", "SANN") but the latter two are still experimental.

Value

An object of class "spml".

<code>coefficients</code>	coefficients estimate of the model parameters
<code>arcoef</code>	the coefficient for the spatial lag on <code>y</code>
<code>errcomp</code>	the estimates of the error variance components
<code>vcov</code>	the asymptotic variance covariance matrix of the estimated coefficients
<code>vcov.arcoef</code>	the asymptotic variance of the estimated spatial lag parameter
<code>vcov.errcomp</code>	the asymptotic variance covariance matrix of the estimated error covariance parameters
<code>type</code>	'random effects ML'
<code>residuals</code>	the model residuals
<code>fitted.values</code>	the fitted values, calculated as $\hat{y} = X\hat{\beta}$

sigma2	GLS residuals variance
model	the matrix of the data used
call	the call used to create the object
logLik	the value of the log likelihood function at the optimum
errors	the value of the errors argument

Author(s)

Giovanni Millo

References

Millo, G. (2014) Maximum likelihood estimation of spatially and serially correlated panels with random effects. *Computational Statistics and Data Analysis*, **71**, 914–933.

See Also

spml

Examples

```
data(Produc, package = "plm")
data(usaww)
fm <- log(gsp) ~ log(pcap) + log(pc) + log(emp) + unemp
## random effects panel with spatial lag and serial error correlation
## optimization method set to "BFGS"
sarsrmod <- spreml(fm, data = Produc, w = usaww, errors="sr", lag=TRUE,
                  method="BFGS")
summary(sarsrmod)
```

summary.splm

summary method for class splm

Description

Method for summarizing the results of objects of class 'splm'

Usage

```
## S3 method for class 'splm'
summary(object,...)
```

Arguments

object	an object of class 'splm'
...	additional arguments to be passed

Details

The summary function `summary.splm` returns an objects of class 'splm' organized in a coefficient matrix.

Also a matrix for the error components, or the spatial coefficients will be generated depending on the estimated model.

When the 'splm' is produced by the function 'spsegm', the summary will be generated looping over the number of equations in the system.

Value

An object of class 'summary.splm'

Author(s)

Giovanni Millo, Gianfranco Piras

See Also

`spml`, `spgm`

Examples

```
data(Produc, package = "plm")
data(usaww)
GM <- spgm(log(gsp)~log(pcap)+log(pc)+log(emp)+unemp, data=Produc,
           listw=usaww, moments = "fullweights", spatial.error = TRUE)
summary(GM)
```

usaww

Spatial weights matrix - US states

Description

Spatial weights matrix of the 48 continental US States based on the queen contiguity criterium.

Usage

```
data(usaww)
```

Format

A matrix with elements different from zero if state *i* and *j* are neighbors. Weights are row standardized. According to the queen contiguity criterium, Arizona and Colorado are considered neighbours.

Author(s)

Giovanni Millo

`vcov.splm`*Covariance extractor method for splm objects*

Description

Covariance extractor method for `splm` objects. Seldom used as such but needed, e.g., for interoperability with testing functions in `lmtest` and `car`.

Usage

```
## S3 method for class 'splm'  
vcov(object, ...)
```

Arguments

<code>object</code>	an object of class <code>splm</code>
<code>...</code>	additional arguments to be passed; currently not used

Value

a covariance matrix of beta coefficients

Author(s)

Giovanni Millo

References

Zeileis, A. (2006) Object-Oriented Computation of Sandwich Estimators. *Journal of Statistical Software*, **16(9)**, 1-16.

Examples

```
## not run:  
## data(Produc, package="plm")  
## data(usaww)  
## fm <- log(gsp)~log(pcap)+log(pc)+log(emp)+unemp  
## sarremod <- spml(fm, data=Produc, listw = spdep::mat2listw(usaww),  
## model="random", lag=TRUE, spatial.error="none")  
## ## compact representation of betas  
## library(lmtest)  
## coeftest(sarremod)  
## ## linear hypothesis test  
## library(car)  
## lht(sarremod, "log(pcap)=log(pc)")
```

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