

Package: spldv (via r-universe)

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Type Package

Title Spatial Models for Limited Dependent Variables

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Description The current version of this package estimates spatial autoregressive models for binary dependent variables using GMM estimators <[doi:10.18637/jss.v107.i08](https://doi.org/10.18637/jss.v107.i08)> and RIS estimator <[doi:10.1007/978-3-662-05617-2_8](https://doi.org/10.1007/978-3-662-05617-2_8)>. It supports one-step (Pinkse and Slade, 1998) <[doi:10.1016/S0304-4076\(97\)00097-3](https://doi.org/10.1016/S0304-4076(97)00097-3)> and two-step GMM estimator along with the linearized GMM estimator proposed by Klier and McMillen (2008) <[doi:10.1198/073500107000000188](https://doi.org/10.1198/073500107000000188)>. It also allows for either Probit or Logit model and compute the average marginal effects. The RIS estimator allows to estimate the SAR and SEM model. All these models are presented in Sarrias and Piras (2023) <[doi:10.1016/j.jocm.2023.100432](https://doi.org/10.1016/j.jocm.2023.100432)>.

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BugReports <https://github.com/gpiras/spldv/issues>

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getSummary.bingmm *Get Model Summaries for use with "mtable" for objects of class bingmm*

Description

A generic function to collect coefficients and summary statistics from a *bingmm* object. It is used in *mtable*

Usage

```
## S3 method for class 'bingmm'
getSummary(obj, alpha = 0.05, ...)
```

Arguments

<i>obj</i>	a <i>bingmm</i> object,
<i>alpha</i>	level of the confidence intervals,
...	further arguments,

Details

For more details see package **memisc**.

Value

A list with an array with coefficient estimates and a vector containing the model summary statistics.

getSummary.binlgmm	<i>Get Model Summaries for use with "mtable" for objects of class binlgmm</i>
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Description

A generic function to collect coefficients and summary statistics from a `binlgmm` object. It is used in `mtable`

Usage

```
## S3 method for class 'binlgmm'  
getSummary(obj, alpha = 0.05, ...)
```

Arguments

obj	a <code>binlgmm</code> object,
alpha	level of the confidence intervals,
...	further arguments,

Details

For more details see package **memisc**.

Value

A list with an array with coefficient estimates and a vector containing the model summary statistics.

getSummary.binris	<i>Get Model Summaries for use with "mtable" for objects of class binris</i>
-------------------	--

Description

A generic function to collect coefficients and summary statistics from a `binris` object. It is used in `mtable`

Usage

```
## S3 method for class 'binris'  
getSummary(obj, alpha = 0.05, ...)
```

Arguments

obj	a <code>binris</code> object,
alpha	level of the confidence intervals,
...	further arguments,

Details

For more details see package **memisc**.

Value

A list with an array with coefficient estimates and a vector containing the model summary statistics.

impacts.bingmm	<i>Estimation of the average marginal effects for SARB models.</i>
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Description

Obtain the average marginal effects from **bingmm**, **binlgmm** or **binris** class model.

Usage

```
## S3 method for class 'bingmm'
impacts(
  obj,
  vcov = NULL,
  vce = c("robust", "efficient", "ml"),
  het = TRUE,
  atmeans = FALSE,
  type = c("mc", "delta"),
  R = 100,
  approximation = FALSE,
  pw = 5,
  tol = 1e-06,
  empirical = FALSE,
  ...
)

## S3 method for class 'binlgmm'
impacts(
  obj,
  vcov = NULL,
  het = TRUE,
  atmeans = FALSE,
  type = c("mc", "delta"),
  R = 100,
  approximation = FALSE,
  pw = 5,
  tol = 1e-06,
  empirical = FALSE,
  ...
)
```

```

## S3 method for class 'binris'
impacts(
  obj,
  vcov = NULL,
  het = TRUE,
  atmeans = FALSE,
  type = c("mc", "delta"),
  R = 100,
  approximation = FALSE,
  pw = 5,
  tol = 1e-06,
  empirical = FALSE,
  ...
)

## S3 method for class 'impacts.bingmm'
print(x, ...)

## S3 method for class 'impacts.bingmm'
summary(object, ...)

## S3 method for class 'summary.impacts.bingmm'
print(x, digits = max(3,getOption("digits") - 3), ...)

```

Arguments

obj	an object of class <code>bingmm</code> , <code>binlgmm</code> or <code>binris</code> .
vcov	an estimate of the asymptotic variance-covariance matrix of the parameters for a <code>bingmm</code> or <code>binlgmm</code> object.
vce	string indicating what kind of variance-covariance matrix of the estimate should be computed when using <code>effect.bingmm</code> . For the one-step GMM estimator, the options are <code>"robust"</code> and <code>"m1"</code> . For the two-step GMM estimator, the options are <code>"robust"</code> , <code>"efficient"</code> and <code>"m1"</code> . The option <code>"vce = m1"</code> is an exploratory method that evaluates the VC of the RIS estimator using the GMM estimates.
het	logical. If <code>TRUE</code> (the default), then the heteroskedasticity is taken into account when computing the average marginal effects.
atmeans	logical. If <code>FALSE</code> (the default), then the average marginal effects are computed at the unit level.
type	string indicating which method is used to compute the standard errors of the average marginal effects. If <code>"mc"</code> , then the Monte Carlo approximation is used. If <code>"delta"</code> , then the Delta Method is used.
R	numerical. Indicates the number of draws used in the Monte Carlo approximation if <code>type = "mc"</code> .
approximation	logical. If <code>TRUE</code> then $(I - \lambda W)^{-1}$ is approximated as $I + \lambda W + \lambda^2 W^2 + \lambda^3 W^3 + \dots + \lambda^q W^q$. The default is <code>FALSE</code> .
pw	numeric. The power used for the approximation $I + \lambda W + \lambda^2 W^2 + \lambda^3 W^3 + \dots + \lambda^q W^q$. The default is 5.

tol	Argument passed to <code>mvrnorm</code> : tolerance (relative to largest variance) for numerical lack of positive-definiteness in the coefficient covariance matrix.
empirical	logical. Argument passed to <code>mvrnorm</code> (default FALSE): if TRUE, the coefficients and their covariance matrix specify the empirical not population mean and covariance matrix
...	further arguments. Ignored.
x	an object of class <code>impacts.bingmm</code> .
object	an object of class <code>impacts.bingmm</code> for summary methods.
digits	the number of digits.

Details

Let the model be:

$$y^* = X\beta + WX\gamma + \lambda Wy^* + \epsilon = Z\delta + \lambda Wy^* + \epsilon$$

where $y = 1$ if $y^* > 0$ and 0 otherwise; $\epsilon \sim N(0, 1)$ if `link = "probit"` or $\epsilon \sim L(0, \pi^2/3)$ if `link = "logit"`. The RIS estimator assumes that $\epsilon \sim N(0, 1)$.

The marginal effects respect to variable x_r can be computed as

$$\text{diag}(f(a))D_\lambda^{-1}A_\lambda^{-1}(I_n\beta_r + W\gamma_r) = C_r(\theta)$$

where $f()$ is the pdf, which depends on the assumption of the error terms; `diag` is the operator that creates a $n \times n$ diagonal matrix; $A_\lambda = (I - \lambda W)$; and D_λ is a diagonal matrix whose elements represent the square root of the diagonal elements of the variance-covariance matrix of $u = A_\lambda^{-1}\epsilon$.

We implement these three summary measures: (1) The average total effects, $ATE_r = n^{-1}i'_n C_r i_n$, (2) The average direct effects, $ADE_r = n^{-1}tr(C_r)$, and (3) the average indirect effects, $ATE_r - ADE_r$.

The standard errors of the average total, direct and indirect effects can be estimated using either Monte Carlo (MC) approximation, which takes into account the sampling distribution of θ , or Delta Method.

Value

An object of class `impacts.bingmm`.

Author(s)

Mauricio Sarrias and Gianfranco Piras.

See Also

[sbinaryGMM](#), [sbinaryLGMM](#).

Examples

```
# Data set
data(olcld, package = "spdep")

# Create dependent (dummy) variable
COL.OLD$CRIMED <- as.numeric(COL.OLD$CRIME > 35)

# Two-step (Probit) GMM estimator
ts <- sbinaryGMM(CRIMED ~ INC + HOVAL | HOVAL,
                   link = "probit",
                   listw = spdep::nb2listw(COL.nb, style = "W"),
                   data = COL.OLD,
                   type = "twostep")

# Marginal effects using Delta Method
summary(impacts(ts, type = "delta"))

# Marginal effects using MC with 100 draws
summary(impacts(ts, type = "mc", R = 100))

# Marginal effects using efficient VC matrix
summary(impacts(ts, type = "delta", vce = "efficient"))

# Marginal effects using efficient VC matrix and ignoring the heteroskedasticity
summary(impacts(ts, type = "delta", vce = "efficient", het = FALSE))

# Marginal effects using RIS estimator
ris_sar <- sbinaryRis(CRIMED ~ INC + HOVAL, data = COL.OLD,
                       R = 50,
                       listw = spdep::nb2listw(COL.nb, style = "W"))
summary(impacts(ris_sar, method = "delta"))
summary(impacts(ris_sar, method = "mc", R = 100))
```

make.instruments

Make instruments for spatial models

Description

Make instruments for spatial models

Usage

```
make.instruments(listw, x, q)
```

Arguments

- | | |
|-------|--|
| listw | object. An object of class <code>listw</code> , <code>matrix</code> , or <code>Matrix</code> . |
| x | variable(s) to be lagged |
| q | number of lags |

Author(s)

Mauricio Sarrias and Gianfranco Piras.

sbinaryGMM

Estimation of SAR for binary dependent models using GMM

Description

Estimation of SAR model for binary dependent variables (either Probit or Logit), using one- or two-step GMM estimator. The type of model supported has the following structure:

$$y^* = X\beta + WX\gamma + \lambda Wy^* + \epsilon = Z\delta + \lambda Wy^* + \epsilon$$

where $y = 1$ if $y^* > 0$ and 0 otherwise; $\epsilon \sim N(0, 1)$ if `link = "probit"` or $\epsilon \sim L(0, \pi^2/3)$ if `link = "logit"`.

Usage

```
sbinaryGMM(
  formula,
  data,
  listw = NULL,
  nins = 2,
  link = c("probit", "logit"),
  winitial = c("optimal", "identity"),
  s.matrix = c("robust", "iid"),
  type = c("onestep", "twostep"),
  gradient = TRUE,
  start = NULL,
  cons.opt = FALSE,
  approximation = FALSE,
  verbose = TRUE,
  print.init = FALSE,
  pw = 5,
  tol.solve = .Machine$double.eps,
  ...
)

## S3 method for class 'bingmm'
coef(object, ...)

## S3 method for class 'bingmm'
vcov(
  object,
  vce = c("robust", "efficient", "ml"),
  method = "bhhh",
  R = 1000,
```

```

tol.solve = .Machine$double.eps,
...
)

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

## S3 method for class 'bingmm'
summary(
  object,
  vce = c("robust", "efficient", "ml"),
  method = "bhhh",
  R = 1000,
  tol.solve = .Machine$double.eps,
  ...
)

## S3 method for class 'summary.bingmm'
print(x, digits = max(5, getOption("digits") - 3), ...)

```

Arguments

formula	a symbolic description of the model of the form $y \sim x wx$ where y is the binary dependent variable, x are the independent variables. The variables after $ $ are those variables that enter spatially lagged: WX . The variables in the second part of formula must also appear in the first part. This rules out situations in which one of the regressors can be specified only in lagged form.
data	the data of class <code>data.frame</code> .
listw	object. An object of class <code>listw</code> , <code>matrix</code> , or <code>Matrix</code> .
nins	numerical. Order of instrumental-variable approximation; as default <code>nins</code> = 2, such that $H = (Z, WZ, W^2Z)$ are used as instruments.
link	string. The assumption of the distribution of the error term; it can be either <code>link</code> = "probit" (the default) or <code>link</code> = "logit".
winitial	string. A string indicating the initial moment-weighting matrix Ψ ; it can be either <code>winitial</code> = "optimal" (the default) or <code>winitial</code> = "identity".
s.matrix	string. Only valid of type = "twostep" is used. This is a string indicating the type of variance-covariance matrix \hat{S} to be used in the second-step procedure; it can be <code>s.matrix</code> = "robust" (the default) or <code>s.matrix</code> = "iid".
type	string. A string indicating whether the one-step (<code>type</code> = "onestep"), or two-step GMM (<code>type</code> = "twostep") should be computed.
gradient	logical. Only for testing procedures. Should the analytic gradient be used in the GMM optimization procedure? TRUE as default. If FALSE, then the numerical gradient is used.
start	if not NULL, the user must provide a vector of initial parameters for the optimization procedure. When <code>start</code> = NULL, <code>sbinaryGMM</code> uses the traditional Probit or Logit estimates as initial values for the parameters, and the correlation between y and Wy as initial value for λ .

<code>cons.opt</code>	logical. Should a constrained optimization procedure for λ be used? FALSE as default.
<code>approximation</code>	logical. If TRUE then $(I - \lambda W)^{-1}$ is approximated as $I + \lambda W + \lambda^2 W^2 + \lambda^3 W^3 + \dots + \lambda^q W^q$. The default is FALSE.
<code>verbose</code>	logical. If TRUE, the code reports messages and some values during optimization.
<code>print.init</code>	logical. If TRUE the initial parameters used in the optimization of the first step are printed.
<code>pw</code>	numeric. The power used for the approximation $I + \lambda W + \lambda^2 W^2 + \lambda^3 W^3 + \dots + \lambda^q W^q$. The default is 5.
<code>tol.solve</code>	Tolerance for <code>solve()</code> .
<code>...</code>	additional arguments passed to <code>maxLik</code> .
<code>vce</code>	string. A string indicating what kind of standard errors should be computed when using <code>summary</code> . For the one-step GMM estimator, the options are "robust" and "ml". For the two-step GMM estimator, the options are "robust", "efficient" and "ml". The option "vce = ml" is an exploratory method that evaluates the VC of the RIS estimator using the GMM estimates.
<code>method</code>	string. Only valid if <code>vce = "ml"</code> . It indicates the algorithm used to compute the Hessian matrix of the RIS estimator. The defult is "bhhh".
<code>R</code>	numeric. Only valid if <code>vce = "ml"</code> . It indicates the number of draws used to compute the simulated probability in the RIS estimator.
<code>x, object</code>	an object of class <code>bingmm</code>
<code>digits</code>	the number of digits

Details

The data generating process is:

$$y^* = X\beta + WX\gamma + \lambda Wy^* + \epsilon = Z\delta + \lambda Wy^* + \epsilon$$

where $y = 1$ if $y^* > 0$ and 0 otherwise; $\epsilon \sim N(0, 1)$ if `link = "probit"` or $\epsilon \sim L(0, \pi^2/3)$ if `link = "logit"`. The general GMM estimator minimizes

$$J(\theta) = g'(\theta)\hat{\Psi}g(\theta)$$

where $\theta = (\beta, \gamma, \lambda)$ and

$$g = n^{-1}H'v$$

where v is the generalized residuals. Let $Z = (X, WX)$, then the instrument matrix H contains the linearly independent columns of $H = (Z, WZ, \dots, W^q Z)$. The one-step GMM estimator minimizes $J(\theta)$ setting either $\hat{\Psi} = I_p$ if `winitial = "identity"` or $\hat{\Psi} = (H'H/n)^{-1}$ if `winitial = "optimal"`. The two-step GMM estimator uses an additional step to achieve higher efficiency by computing the variance-covariance matrix of the moments \hat{S} to weight the sample moments. This matrix is computed using the residuals or generalized residuals from the first-step, which are consistent. This matrix is computed as $\hat{S} = n^{-1} \sum_{i=1}^n h_i(f^2/(F(1-F)))h_i'$ if `s.matrix = "robust"` or $\hat{S} = n^{-1} \sum_{i=1}^n \hat{v}_i h_i h_i'$, where \hat{v} are the first-step generalized residuals.

Value

An object of class “bingmm”, a list with elements:

<code>coefficients</code>	the estimated coefficients,
<code>call</code>	the matched call,
<code>callF</code>	the full matched call,
<code>X</code>	the X matrix, which contains also WX if the second part of the <code>formula</code> is used,
<code>H</code>	the H matrix of instruments used,
<code>y</code>	the dependent variable,
<code>listw</code>	the spatial weight matrix,
<code>link</code>	the string indicating the distribution of the error term,
<code>Psi</code>	the moment-weighting matrix used in the last round,
<code>type</code>	type of model that was fitted,
<code>s.matrix</code>	the type of S matrix used in the second round,
<code>winitial</code>	the moment-weighting matrix used for the first step procedure
<code>opt</code>	object of class <code>maxLik</code> ,
<code>approximation</code>	a logical value indicating whether approximation was used to compute the inverse matrix,
<code>pw</code>	the powers for the approximation,
<code>formula</code>	the formula.

Author(s)

Mauricio Sarrias and Gianfranco Piras.

References

- Pinkse, J., & Slade, M. E. (1998). Contracting in space: An application of spatial statistics to discrete-choice models. *Journal of Econometrics*, 85(1), 125-154.
- Fleming, M. M. (2004). Techniques for estimating spatially dependent discrete choice models. In *Advances in spatial econometrics* (pp. 145-168). Springer, Berlin, Heidelberg.
- Klier, T., & McMillen, D. P. (2008). Clustering of auto supplier plants in the United States: generalized method of moments spatial logit for large samples. *Journal of Business & Economic Statistics*, 26(4), 460-471.
- LeSage, J. P., Kelley Pace, R., Lam, N., Campanella, R., & Liu, X. (2011). New Orleans business recovery in the aftermath of Hurricane Katrina. *Journal of the Royal Statistical Society: Series A (Statistics in Society)*, 174(4), 1007-1027.
- Piras, G., & Sarrias, M. (2023). One or Two-Step? Evaluating GMM Efficiency for Spatial Binary Probit Models. *Journal of choice modelling*, 48, 100432.
- Piras, G., & Sarrias, M. (2023). GMM Estimators for Binary Spatial Models in R. *Journal of Statistical Software*, 107(8), 1-33.

See Also

[sbinaryLGMM](#), [impacts.bingmm](#).

Examples

```
# Data set
data(olddol, package = "spdep")

# Create dependent (dummy) variable
COL.OLD$CRIMED <- as.numeric(COL.OLD$CRIME > 35)

# Two-step (Probit) GMM estimator
ts <- sbinaryGMM(CRIMED ~ INC + HOVAL,
                  link = "probit",
                  listw = spdep::nb2listw(COL.nb, style = "W"),
                  data = COL.OLD,
                  type = "twostep",
                  verbose = TRUE)

# Robust standard errors
summary(ts)
# Efficient standard errors
summary(ts, vce = "efficient")

# One-step (Probit) GMM estimator
os <- sbinaryGMM(CRIMED ~ INC + HOVAL,
                  link = "probit",
                  listw = spdep::nb2listw(COL.nb, style = "W"),
                  data = COL.OLD,
                  type = "onestep",
                  verbose = TRUE)
summary(os)

# One-step (Logit) GMM estimator with identity matrix as initial weight matrix
os_l <- sbinaryGMM(CRIMED ~ INC + HOVAL,
                     link = "logit",
                     listw = spdep::nb2listw(COL.nb, style = "W"),
                     data = COL.OLD,
                     type = "onestep",
                     winitial = "identity",
                     verbose = TRUE)
summary(os_l)

# Two-step (Probit) GMM estimator with WX
ts_wx <- sbinaryGMM(CRIMED ~ INC + HOVAL | INC + HOVAL,
                      link = "probit",
                      listw = spdep::nb2listw(COL.nb, style = "W"),
                      data = COL.OLD,
                      type = "twostep",
                      verbose = FALSE)
summary(ts_wx)
```

```
# Constrained two-step (Probit) GMM estimator
ts_c <- sbinaryGMM(CRIMED ~ INC + HOVAL,
                     link = "probit",
                     listw = spdep::nb2listw(COL.nb, style = "W"),
                     data = COL.OLD,
                     type = "twostep",
                     verbose = TRUE,
                     cons.opt = TRUE)
summary(ts_c)
```

sbinaryLGMM*Estimation of SAR for binary models using Linearized GMM.***Description**

Estimation of SAR model for binary dependent variables (either Probit or Logit), using Linearized GMM estimator suggested by Klier and McMillen (2008). The model is:

$$y^* = X\beta + WX\gamma + \lambda Wy^* + \epsilon = Z\delta + \lambda Wy^* + \epsilon$$

where $y = 1$ if $y^* > 0$ and 0 otherwise; $\epsilon \sim N(0, 1)$ if `link = "probit"` or $\epsilon \sim L(0, \pi^2/3)$ `link = "logit"`.

Usage

```
sbinaryLGMM(
  formula,
  data,
  listw = NULL,
  nins = 2,
  link = c("logit", "probit"),
  ...
)

## S3 method for class 'binlgmm'
coef(object, ...)

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

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

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

## S3 method for class 'summary.binlgmm'
print(x, digits = max(3, getOption("digits") - 2), ...)
```

Arguments

<code>formula</code>	a symbolic description of the model of the form $y \sim x wx$ where y is the binary dependent variable, x are the independent variables. The variables after $ $ are those variables that enter spatially lagged: WX . The variables in the second part of <code>formula</code> must also appear in the first part.
<code>data</code>	the data of class <code>data.frame</code> .
<code>listw</code>	object. An object of class <code>listw</code> , <code>matrix</code> , or <code>Matrix</code> .
<code>nins</code>	numerical. Order of instrumental-variable approximation; as default <code>nins</code> = 2, such that $H = (Z, WZ, W^2Z)$ are used as instruments.
<code>link</code>	string. The assumption of the distribution of the error term; it can be either <code>link</code> = "probit" (the default) or <code>link</code> = "logit".
<code>...</code>	additional arguments.
<code>x, object</code>	an object of class <code>binlgmm</code> .
<code>digits</code>	the number of digits

Details

The steps for the linearized spatial Probit/Logit model are the following:

1. Estimate the model by standard Probit/Logit model, in which spatial autocorrelation and heteroskedasticity are ignored. The estimated values are $\hat{\beta}_0$. Calculate the generalized residuals assuming that $\lambda = 0$ and the gradient terms G_β and G_λ .
2. The second step is a two-stage least squares estimator of the linearized model. Thus regress G_β and G_λ on $H = (Z, WZ, W^2Z, \dots, W^qZ)$ and obtain the predicted values \hat{G} . Then regress $u_0 + G'_\beta \hat{\beta}_0$ on \hat{G} . The coefficients are the estimated values of β and λ .

The variance-covariance matrix can be computed using the traditional White-corrected coefficient covariance matrix from the last two-stage least squares estimator of the linearized model.

Value

An object of class "bingmm", a list with elements:

<code>coefficients</code>	the estimated coefficients,
<code>call</code>	the matched call,
<code>X</code>	the X matrix, which contains also WX if the second part of the formula is used,
<code>H</code>	the H matrix of instruments used,
<code>y</code>	the dependent variable,
<code>listw</code>	the spatial weight matrix,
<code>link</code>	the string indicating the distribution of the error term,
<code>fit</code>	an object of <code>lm</code> representing the T2SLS,
<code>formula</code>	the formula.

Author(s)

Mauricio Sarrias and Gianfranco Piras.

References

- Klier, T., & McMillen, D. P. (2008). Clustering of auto supplier plants in the United States: generalized method of moments spatial logit for large samples. *Journal of Business & Economic Statistics*, 26(4), 460-471.
- Piras, G., & Sarrias, M. (2023). One or Two-Step? Evaluating GMM Efficiency for Spatial Binary Probit Models. *Journal of choice modelling*, 48, 100432.
- Piras, G., & Sarrias, M. (2023). GMM Estimators for Binary Spatial Models in R. *Journal of Statistical Software*, 107(8), 1-33.

See Also

`sbinaryGMM`, `impacts.bingmm`.

Examples

```
# Data set
data(olddol, package = "spdep")

# Create dependent (dummy) variable
COL.OLD$CRIMED <- as.numeric(COL.OLD$CRIME > 35)

# LGMM for probit using q = 3 for instruments
lgmm <- sbinaryLGMM(CRIMED ~ INC + HOVAL | INC,
                      link = "probit",
                      listw = spdep::nb2listw(COL.nb, style = "W"),
                      nins = 3,
                      data = COL.OLD)
summary(lgmm)
```

sbinaryRis

*Estimation of spatial probit model for binary outcomes using RIS
(GHK) simulator*

Description

Estimation of spatial probit model using RIS-normal (a.k.a GHK) simulator. The models can be the SAR or SEM probit model model. The SAR probit model has the following structure:

$$y^* = X\beta + WX\gamma + \lambda Wy^* + \epsilon = Z\delta + \lambda Wy^* + \epsilon = A_\lambda^{-1}Z\delta + u,$$

where $y = 1$ if $y^* > 0$ and 0 otherwise, $Z = (X, WX)$, $\delta = (\beta', \gamma')'$, $u = A_\lambda^{-1}\epsilon$ with $A_\lambda = (I - \lambda W)$, and $\epsilon \sim N(0, I)$. The SEM probit model has the following structure:

$$y^* = X\beta + WX\gamma + u = Z\delta + u$$

where $y = 1$ if $y^* > 0$ and 0 otherwise, $Z = (X, WX)$, $\delta = (\beta', \gamma')'$, $u = \rho Wu + \epsilon$ such that $u = A_\rho^{-1}\epsilon$, and $\epsilon \sim N(0, I)$,

Usage

```

sbinaryRis(
  formula,
  data,
  subset,
  na.action,
  listw = NULL,
  R = 20,
  model = c("SAR", "SEM"),
  varcov = c("invsigma", "sigma"),
  approximation = TRUE,
  pw = 5,
  start = NULL,
  Qneg = FALSE,
  print.init = FALSE,
  ...
)

## S3 method for class 'binris'
terms(x, ...)

## S3 method for class 'binris'
estfun(x, ...)

## S3 method for class 'binris'
bread(x, ...)

## S3 method for class 'binris'
df.residual(object, ...)

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

## S3 method for class 'binris'
coef(object, ...)

## S3 method for class 'binris'
logLik(object, ...)

## S3 method for class 'binris'
print(x, ...)

## S3 method for class 'binris'
summary(object, eigentol = 1e-12, ...)

## S3 method for class 'summary.binris'
print(x, digits = max(3,getOption("digits") - 2), ...)

```

Arguments

formula	a symbolic description of the model of the form $y \sim x wx$ where y is the binary dependent variable, x are the independent variables. The variables after $ $ are those variables that enter spatially lagged: WX . The variables in the second part of formula must also appear in the first part. This rules out situations in which one of the regressors can be specified only in lagged form.
data	the data of class <code>data.frame</code> .
subset	an optional vector specifying a subset of observations to be used in the fitting process.
na.action	a function which indicates what should happen when the data contains NAs.
listw	object. An object of class <code>listw</code> , <code>matrix</code> , or <code>Matrix</code> .
R	numerical. The number of draws used in RIS (GHK) simulator.
model	string. A string indicating which model to estimate. It can be "SAR" for the spatial autoregressive spatial model or "SEM" for the spatial error model.
varcov	string. A string indicating over which variance-covariance matrix to apply the Cholesky factorization.
approximation	logical. If TRUE (the default) then $(I - \lambda W)^{-1}$ or $(I - \rho W)^{-1}$ is approximated as $I + \lambda W + \lambda^2 W^2 + \lambda^3 W^3 + \dots + \lambda^q W^q$. If FALSE, then the inverse is computed without approximations.
pw	numeric. The power used for the approximation $I + \lambda W + \lambda^2 W^2 + \lambda^3 W^3 + \dots + \lambda^q W^q$. The default is 5.
start	if not NULL, the user must provide a vector of initial parameters for the optimization procedure. When <code>start = NULL</code> , <code>sbinaryRis</code> uses the traditional Probit estimates as initial values for the parameters, and the correlation between y and Wy as initial value for λ or ρ .
Qneg	logical. Whether to construct the diagonal elements of Q . If <code>Qneg = FALSE</code> , then $q_{ii} = 2y_i - 1$. If <code>Qneg = TRUE</code> , then $q_{ii} = 1 - 2y_i$.
print.init	logical. If TRUE the initial parameters used in the optimization of the first step are printed.
...	additional arguments passed to <code>maxLik</code> .
x, object	an object of class <code>bingmm</code> .
eigentol	the standard errors are only calculated if the ratio of the smallest and largest eigenvalue of the Hessian matrix is less than <code>eigentol</code> . Otherwise the Hessian is treated as singular.
digits	the number of digits

Details

The models are estimated by simulating the probabilities using the RIS-normal (GHK) simulator. The aim is to evaluate the multivariate density function $P(v = Qu < s)$, where Q is a diagonal matrix with entries $q_{ii} = 2y_i - 1$ and the $n \times 1$ vector s depends on whether the model is SAR or SEM. If `model = "SAR"`, then $s = QA_\lambda^{-1}Z\delta$ where $A_\lambda = (I - \lambda W)$; if `model = "SEM"`, then $s = QZ\delta$.

Let $\Sigma_v = QVar(u)Q'$ be the variance-covariance model of the transformed model. If `model = "SAR"` $\Sigma_v = Q\Sigma_\lambda Q'$, where $\Sigma_\lambda = (A'_\lambda A_\lambda)^{-1}$. If `model = "SEM"`, then $\Sigma_v = Q\Sigma_\rho Q'$, where $\Sigma_\rho = (A'_\rho A_\rho)^{-1}$.

Since Σ_v is positive definite, there exists a Cholesky decomposition such that $C'C = \Sigma_v^{-1}$, where C is the upper triangular Cholesky matrix and Σ_v^{-1} is the precision matrix. Let $B = C^{-1}$. Then the random vector Qu can be replaced by $Qu = C^{-1}\eta = B\eta$, where η is a vector of standard normal variables. Then, the upper limit of the integration becomes $Qu = B\eta < s$, which can be written as $\eta < B^{-1}s = \nu$.

The RIS simulator is implemented by drawing a large number R of random vector η and computing η_{rj} recursively for $j = 1, \dots, n$. The parameters are estimated using the simulated maximum likelihood (SML) function:

$$\ln \tilde{L}(\theta) = \ln \left(\frac{1}{R} \sum_{r=1}^R \tilde{p}_r \right),$$

where:

$$\tilde{p}_r = \prod_{j=1}^n \Phi(\hat{\eta}_{jr}),$$

and $\Phi(\cdot)$ is the univariate CDF of the standard normal density.

By default, `sbinaryRis` compute the SML using the Cholesky transformation on Σ_v^{-1} , `varcov = "invsigma"`. The transformation can also be applied to Σ_v using `varcov = "sigma"`, which is slower than the previous option.

This estimator can take several minutes for large datasets. Thus, by default the inverse matrices A_λ^{-1} and A_ρ^{-1} are approximated using the Leontief expansion.

Value

An object of class "`binris`", a list with elements:

<code>varcov</code>	the matrix over which the Cholesky factorization is applied,
<code>model</code>	type of model that was fitted,
<code>Qneg</code>	matrix <code>Q</code> used,
<code>approximation</code>	a logical value indicating whether approximation was used to compute the inverse matrix,
<code>pw</code>	the powers for the approximation,
<code>call</code>	the matched call,
<code>X</code>	the <code>X</code> matrix, which contains also <code>WX</code> if the second part of the <code>formula</code> is used,
<code>y</code>	the dependent variable,
<code>listw</code>	the spatial weight matrix,
<code>formula</code>	the formula,
<code>R</code>	number of draws,
<code>mf</code>	model frame.

Author(s)

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References

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- Fleming, M. M. (2004). Techniques for estimating spatially dependent discrete choice models. In Advances in spatial econometrics (pp. 145-168). Springer, Berlin, Heidelberg.
- Pace, R. K., & LeSage, J. P. (2016). Fast simulated maximum likelihood estimation of the spatial probit model capable of handling large samples. In Spatial Econometrics: Qualitative and Limited Dependent Variables (pp. 3-34). Emerald Group Publishing Limited.
- Piras, G., & Sarrias, M. (2023). One or Two-Step? Evaluating GMM Efficiency for Spatial Binary Probit Models. Journal of choice modelling, 48, 100432.

See Also

[sbinaryGMM](#), [impacts.binris](#).

Examples

```
data(olddol, package = "spdep")

# Create dependent (dummy) variable
COL.OLD$CRIMED <- as.numeric(COL.OLD$CRIME > 35)

# Estimate SAR probit model using RIS simulator using Sigma_v^{-1}
ris_sar <- sbinaryRis(CRIMED ~ INC + HOVAL,
                       data = COL.OLD,
                       R = 50,
                       listw = spdep::nb2listw(COL.nb, style = "W"),
                       print.level = 2)
summary(ris_sar)

# Estimate SAR probit model using RIS simulator using Sigma_v
ris_sar_2 <- sbinaryRis(CRIMED ~ INC + HOVAL,
                        data = COL.OLD,
                        R = 50,
                        listw = spdep::nb2listw(COL.nb, style = "W"),
                        varcov = "sigma",
                        print.level = 2)
summary(ris_sar_2)

# Estimate SDM probit model using RIS simulator
ris_sdm <- sbinaryRis(CRIMED ~ INC + HOVAL | INC + HOVAL,
                       data = COL.OLD,
                       R = 50,
                       listw = spdep::nb2listw(COL.nb, style = "W"),
                       print.level = 2)
```

```
summary(ris_sdm)

# Estimate SEM probit model using RIS simulator
ris_sem <- sbinaryRis(CRIMED ~ INC + HOVAL | INC,
                       data = COL.OLD,
                       R = 50,
                       listw = spdep::nb2listw(COL.nb, style = "W"),
                       model = "SEM")
summary(ris_sem)
```

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