Package ‘ergm.ego’

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  ergm (>= 4.5.0),
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  RColorBrewer (>= 1.1.2),
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  tibble (>= 2.1.1),
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  survey,
  stats,
  methods
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Description Utilities for managing egocentrically sampled network data and a wrapper around the ‘ergm’ package to facilitate ERGM inference and simulation from such data. See Krivitsky and Morris (2017) <doi:10.1214/16-AOAS1010>.

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*.svystat

A scalar multiplication method for svystat

Description

Multiply the values of survey statistics by a specified vector elementwise, adjusting the variance.

Usage

```r
## S3 method for class 'svystat'
x * y
```

Arguments

- `x`: an object of class [svystat][survey::svymean].
- `y`: a numeric vector equal in length to `x`; shorter vectors will be recycled.

Value

A [svystat][survey::svymean] object with the updated statistics and variance-covariance matrix.
Examples

```r
library(survey)
data(api)
# From example(svymean):
dclus1 <- svydesign(id=~dnum, weights=~pw, data=apiclus1, fpc=~fpc)

(m1 <- svymean(~api99, dclus1))
(v1 <- vcov(m1))

# Scale the survey stat object by a factor of two:
(m2 <- m1 * 2)
(v2 <- vcov(m2))
```

---

**as.egor.egodata** — *Convert (deprecated) egodata Objects to egor Objects*

**Description**

Convert (deprecated) egodata Objects to egor Objects

**Usage**

```r
## S3 method for class 'egodata'
as.egor(x, ...)
as_egor.egodata(x, ...)
```

**Arguments**

- `x` — a egodata object
- `...` — additional arguments, currently unused.

**Value**

An egor object.

**Author(s)**

Pavel N. Krivitsky
as.egor.network  
Construct an Egocentric View of a network Object

Description

Given a network object, construct an egor object representing a census of all the actors in the network. Used mainly for testing.

Usage

```r
## S3 method for class 'network'
as.egor(x, special.cols = c("na"), ...)
```

Arguments

- `x`  A network object.
- `special.cols`  Vertex attributes that should not be copied to the egos and alters tables. Defaults to attributes special to the network objects.
- `...`  Additional arguments, currently unused.

Value

An egor object.

Author(s)

Pavel N. Krivitsky

See Also

template_network, which performs the inverse operation (though drops the ties).

Examples

```r
# See example(ergm.ego) and example(template_network).
```
control.ergm.ego

Control parameters for ergm.ego.

Description

Constructs and checks the list of control parameters for estimation by ergm.ego.

Usage

control.ergm.ego(
  ppopsize = c("auto", "samp", "pop"),
  ppopsize.mul = 1,
  ppop.wt = c("round", "sample"),
  stats.wt = c("data", "ppop"),
  stats.est = c("survey", "asymptotic", "bootstrap", "jackknife", "naive"),
  boot.R = 10000,
  ignore.max.alters = TRUE,
  ergm = control.ergm(),
  ...
)

Arguments

ppopsize, ppopsize.mul

Parameters to determine the size $|N'|$ of the pseudopopulation network. ppopsize can be

"auto" If the ppopsize ($|N|$) argument is specified and is different from 1, as if "pop"; otherwise, as "samp".
"samp" set $|N'|$ based on the sample size: $|N'| = |S| \times \text{ppopsize.mul}$
"pop" set $|N'|$ based on the population size: $|N'| = |N| \times \text{ppopsize.mul}$
a number set $|N'|$ directly (ppopsize.mul ignored)
a network object use the specified network as the pseudo-population network directly; use at your own risk
a data frame use the specified data frame as the pseudo-population; use at your own risk

The default is to use the same pseudopopulation size as the sample size, but, particularly if there are sampling weights in the data, it should be bigger.

Note that depending on ppop.wt, this may only be an approximate target specification, with the actual constructed pseudopopulation network being slightly bigger or smaller.

ppop.wt

Because each ego must be represented in the pseudopopulation network an integral number of times, if the sample is weighted (or the target $|N'|$ calculated from ppopsize and ppopsize.mul is not a multiple of the sample size), it may not be possible, for a finite $|N'|$ to represent each ego exactly according to its relative weight, and ppop.wt controls how the fractional egos are allocated:
"round" (default) Rather than treating ppopsize as a hard setting, calculate \(|N'|w_i/w_i\) for each ego \(i\) and round it to the nearest integer. Then, the \(|N'|\) actually used will be the sum of these rounded frequencies.

"sample" Resample in proportion to \(w_i\).

\texttt{stats.wt} Weight assigned to each ego's contribution to the ERGM's sufficient statistic:

"data" (default) Use weights \(|N'|w_i/w_i\) for each ego \(i\) as in the data.

"ppop" Use weights ultimately used in the pseudopopulation network.

\texttt{stats.est, boot.R} Method to be used to estimate the ERGM's sufficient statistics and their variance:

"survey" Variance estimator returned by \texttt{survey::svymean()}, appropriate to the design of the dataset.

"asymptotic" Delta method, as derived by Krivitsky and Morris (2017), assuming the ego weights are sampled alongside the egos. (default) Delta method, as derived by Krivitsky and Morris (2017), assuming the ego weights are sampled alongside the egos.

"bootstrap" Nonparametric bootstrap with bias correction, resampling egos, using \(R\) replications.

"jackknife" Jackknife with bias correction.

"naive" "Naive" estimator, assuming that weights are fixed.

\texttt{ignore.max.alters} if \texttt{TRUE}, ignores any constraints on the number of nominations. Used to be \texttt{FALSE}, now \texttt{TRUE} in light of the findings of Krivitsky et. al (2020).

\texttt{ergm} Control parameters for the \texttt{ergm()} call to fit the model, constructed by \texttt{control.ergm()}.

... Not used at this time.

\textbf{Value}

A list with arguments as components.

\textbf{Author(s)}

Pavel N. Krivitsky

\textbf{References}


See Also

control.ergm()

---

control.simulate.ergm.ego

Control parameters for `simulate.ergm.ego`.

**Description**

Constructs and checks the list of control parameters for simulation by `simulate.ergm.ego`.

**Usage**

```r
control.simulate.ergm.ego(
  ppop.wt = c("round", "sample"),
  SAN = control.san(),
  simulate = control.simulate(),
  ...
)
```

**Arguments**

- `ppop.wt`  
  Because each ego must be represented in the pseudopopulation network an integral number of times, if the sample is weighted (or the target $|N'|$ calculated from `ppopsize` and `ppopsize.mul` is not a multiple of the sample size), it may not be possible, for a finite $|N'|$ to represent each ego exactly according to its relative weight, and `ppop.wt` controls how the fractional egos are allocated:
  - "round" (default) Rather than treating `ppopsize` as a hard setting, calculate $|N'|_i w_i / w$ for each ego $i$ and round it to the nearest integer. Then, the $|N'|$ actually used will be the sum of these rounded frequencies.
  - "sample" Resample in proportion to $w_i$.

- `SAN`  
  A list of control parameters for `san` constructed by `control.ergm`, called to construct a pseudopopulation network consistent with the data.

- `simulate`  
  A list of control parameters for `simulate.formula` constructed by `control.simulate`, called to simulate from the model fit.

  ...  
  Not used at this time.

**Value**

A list with arguments as components.
Author(s)

Pavel N. Krivitsky

See Also

control.simulate, control.san

degreedist.egor  
Plotting the degree distribution of an egocentric dataset

Description

A `degreedist()` method for `egodata` objects: plot a histogram of the degree distribution of actors in the egocentric dataset, optionally broken down by group and/or compared with a Bernoulli graph.

Usage

```r
## S3 method for class 'egor'
degreedist(
  object,
  freq = FALSE,
  prob = !freq,
  by = NULL,
  brgmod = FALSE,
  main = NULL,
  plot = brgmod,
  weight = TRUE,
  ...
)
```

Arguments

- **object**: A `egor` object.
- **freq, prob**: Whether to plot the raw frequencies or the conditional proportions of the degree values. Defaults to the latter.
- **by**: A character vector giving the name of a vertex attribute; if given, plots the frequencies broken down by that attribute.
- **brgmod**: Plot the range of predicted frequencies/probabilities according to a Bernoulli graph having the same expected density as the observed.
- **main**: Main title of the plot.
- **plot**: Whether to plot the histogram; defaults to the same value as `brgmod`, i.e., `FALSE`.
- **weight**: Whether sampling weights should be incorporated into the calculation (`TRUE`, the default) or ignored (`FALSE`).
- **...**: Additional arguments to `simulate.ergm.ego()`.
ergm.ego

Value

Returns either a vector of degree frequencies/proportions if by=NULL or a matrix with a row for each category if not. If plot==TRUE returns invisibly.

See Also
degreedist, summary

Examples

data(faux.mesa.high)
fmh.ego <- as.egor(faux.mesa.high)

degreedist(fmh.ego,by="Grade",brgmod=TRUE)
# Compare:
degreedist(faux.mesa.high)

Description

A wrapper around the ergm to fit an ERGM to an egor.

Usage

ergm.ego(
  formula,
  popsize = 1,
  offset.coef = NULL,
  constraints = ~.,
  ...,
  basis = eval_lhs.formula(formula),
  control = control.ergm.ego(),
  na.action = na.fail,
  na.rm = FALSE,
  do.fit = TRUE
)

Arguments

formula An formula object, of the form e ~ <model terms>, where e is a egor object. See ergm for details and examples. For a list of currently implemented egocentric terms for the RHS, see ergm.ego-terms.
The size $|N|$ of the finite population network from which the egocentric sample was taken; only affects the shift in the coefficients of the terms modeling the overall propensity to have ties. Setting it to 1 (the default) essentially uses the $-\log |N'|$ offset on the edges term. Passing 0 disables network size adjustment and uses the egocentric sample size; passing $I(N)$ uses the specified size $N$ (though can be overridden by the ppop control.ergm.ego() option) and disables network size adjustment.

A vector of coefficients for the offset terms.

A one-sided formula formula giving the sample space constraints. See ergm for details and examples.

Additional arguments passed to ergm.

A value (usually an egor) to override the LHS of the formula.

A control.ergm.ego control list.

How to handle missing actor attributes in egos or alters, when the terms need them for models that scale.

How to handle missing actor attributes in egos or alters, when the terms need them for models that do not scale.

Whether to actually call ergm

An object of class ergm.ego inheriting from ergm, with the following additional or overridden elements:

Variance-covariance matrix of the estimate of the sufficient statistics

Estimate of the sufficient statistics

The egor object passed

Population network size used

Pseudopopulation size used, see control.ergm.ego

The coefficients, along with the network size adjustment netsize.adj coefficient.

Pseudo-MLE estimate of the variance-covariance matrix of the parameter estimates under repeated egocentric sampling

The variance-covariance matrix of parameter estimates under the ERGM superpopulation process (without incorporating sampling).

Estimated Jacobian of the expectation of the sufficient statistics with respect to the model parameters

Pavel N. Krivitsky
References


See Also

ergm()

Examples

data(faux.mesa.high)
fmh.ego <- as.egor(faux.mesa.high)

head(fmh.ego)

egofit <- ergm.ego(fmh.ego~edges+degree(0:3)+nodefactor("Race") +nodefactor("Sex") +nodematch("Sex") +absdiff("Grade") +gwesp(0,fix=TRUE),
popsize=network.size(faux.mesa.high))

# Run convergence diagnostics
mcmc.diagnostics(egofit)

# Estimates and standard errors
summary(egofit)

--

ergm.ego-terms

---

This page describes the **ergm** terms (and hence network statistics) for which inference based on egocentrically sampled data is implemented in **ergm.ego** package. Other packages may add their own terms. These functions should not be called by the end-user.
Details

The current recommendation for any package implementing additional egocentric calculator terms is to create a help file with a name or alias `ergm.ego-terms`, so that `help("ergm.ego-terms")` will list egocentric ERGM terms available from all loaded packages.

Currently implemented egocentric statistics

For each of these, please see their respective package’s `ergm-terms` help for meaning and parameters. The simplest way to do this is usually via `? TERM`.

Special-purpose terms: `netsize.adj(edges=+1, mutual=0, transitiveties=0, cyclicalties=0)` A special-purpose term equivalent to a linear combination of `edges-ergmTerm`, `mutual-ergmTerm`, `transitiveties-ergmTerm`, and `cyclicalties-ergmTerm`, to house the network-size adjustment offset. This term is added to the model automatically and should not be used in the model formula directly.

**ergm:**
- `offset`
- `edges`
- `nodecov`
- `nodefactor`
- `nodematch`
- `nodemix`
- `absdiff`
- `degree`
- `degrange`
- `concurrent`
- `concurrentties`
- `degree1.5`
- `transitiveties`
- `cyclicalties`
- `esp`
- `gwesp`
- `gwdegree`
- `mm`
- `meandeg*`

**tergm:**
- `mean.age*`

Starred terms are *nonscaling*, in that while they can be evaluated, some inferential results and standard error calculation methods may not be applicable.

See Also

`ergm-terms`
fmhfit

Fitted ergm.ego model object

Description
This is an object with a fitted model to faux.mesa.high data using the code shown below in the Examples section.

Format
An object of class ergm.ego.

Examples
```r
## Not run:
data(faux.mesa.high)
fmh.ego <- egor::as.egor(faux.mesa.high)
fmhfit <- ergm.ego(
  fhm.ego ~ edges + degree(0:3) +
  nodefactor("Race") + nodematch("Race")
  + nodefactor("Sex") + nodematch("Sex")
  + absdiff("Grade") + gwesp(0, fix=TRUE),
popsize = network.size(faux.mesa.high),
  control = control.ergm.ego(
    ergm = control.ergm(parallel=2)
  )
)
## End(Not run)
```

gof.ergm.ego

Conduct Goodness-of-Fit Diagnostics on a Exponential Family Random Graph Model fit to Egocentrically Sampled Data

Description
gof.ergm.ego implements the gof method for ergm.ego fit objects. An enhanced plotting method is also provided, giving uncertainty bars for the observed statistics as well.

Usage
```r
## S3 method for class 'ergm.ego'
gof(
  object,
  ..., 
)```
GOF = c("model", "degree", "espartners"),
control = control.gof.ergm(),
verbose = FALSE
)

## S3 method for class 'gof.ergm.ego'
plot(x, ..., ego.conf.level = 0.95)

Arguments

object
An \texttt{ergm.ego} fit.

\ldots
Additional arguments. Unused by \texttt{gof.ergm.ego()}, passed to \texttt{ergm::plot.gof()} by \texttt{plot.gof.ergm.ego()}.

GOF
A string specifying the statistics whose goodness of fit is to be evaluated. Currently, only “degree”, “espartners” and “model” are implemented; see \texttt{gof} documentation for details.

control
A list to control parameters, constructed using \texttt{control.gof.formula} or \texttt{control.gof.ergm} (which have different defaults).

verbose
Provide verbose information on the progress of the simulation.

x
an object returned by \texttt{gof.ergm.ego()}. 

ego.conf.level
confidence level for the observed statistic estimates as well.

Value

An object of class \texttt{gof.ergm.ego}, inheriting from \texttt{gof.ergm}.

Author(s)

Pavel N. Krivitsky

References


See Also

For examples, see \texttt{ergm.ego}.

Examples

data(faux.mesa.high)
fmh.ego <- as.egor(faux.mesa.high)

head(fmh.ego)

egofit <- ergm.ego(fmh.ego~edges+degree(0:3)+nodefactor("Race")+nodematch("Race"))
mixingmatrix.egor

+nodefactor("Sex")+nodematch("Sex")+absdiff("Grade"),
popsize=network.size(faux.mesa.high))

# Check whether the model "converged":
(modelgof <- gof(egofit, GOF="model"))
plot(modelgof)

# Check whether the model reconstructs the degree distribution:
(deggof <- gof(egofit, GOF="degree"))
plot(deggof)

mixingmatrix.egor  

Summarizing the mixing among groups in an egocentric dataset

Description

A mixingmatrix method for egor objects, to return counts of how often a ego of each group
nominates an alter of each group.

Usage

## S3 method for class 'egor'
mixingmatrix(object, attrname, rowprob = FALSE, weight = TRUE, ...)

Arguments

object

A egor object.

attrname

A character vector containing the name of the network attribute whose mixing
matrix is wanted.

rowprob

Whether the counts should be normalized by row sums. That is, whether they
should be proportions conditional on the ego's group.

weight

Whether sampling weights should be incorporated into the calculation (TRUE, the default) or ignored (FALSE).

... Additional arguments, currently unused.

Value

A matrix with a row and a column for each level of attrname.

Note that, unlike mixingmatrix, what is counted are nominations, not ties. This means that
under an egocentric census, the diagonal of mixingmatrix.egor will be twice that returned by
mixingmatrix for the original undirected network.

See Also

mixingmatrix, nodemix-ergmTerm, summary method for egocentric data
**Examples**

```r
data(faux.mesa.high)
fmh.ego <- as.egor(faux.mesa.high)

(mm <- mixingmatrix(faux.mesa.high,"Grade"))
(mm.ego <- mixingmatrix(fmh.ego,"Grade"))
```

**Description**

These functions are meant to be used in EgoStat and other implementations to provide the user with a way to extract nodal attributes and select their levels in standardized and flexible ways. They are intended to parallel `ergm::nodal_attributes-API` of `ergm` package.

`ergm.ego_get_vattr` extracts and processes the specified nodal attribute vector. It is strongly recommended that `check.ErgmTerm()`'s corresponding vartype="function,formula,character" (using the ERGM_VATTR_SPEC constant).

`ergm.ego_attr_levels` filters the levels of the attribute. It is strongly recommended that `check.ErgmTerm()`'s corresponding vartype="function,formula,character,numeric,logical,AsIs,NULL" (using the ERGM_LEVELS_SPEC constant).

**Usage**

```r
ergm.ego_get_vattr(
  object,
  df,
  accept = "character",
  multiple = if (accept == "character") "paste" else "stop",
  ...
)

## S3 method for class 'character'
ergm.ego_get_vattr(
  object,
  df,
  accept = "character",
  multiple = if (accept == "character") "paste" else "stop",
  ...
)

## S3 method for class ``function`'
ergm.ego_get_vattr(
  object,
  df,
  accept = "character",
  multiple = if (accept == "character") "paste" else "stop",
  ...
)
```
accept = "character",
multiple = if (accept == "character") "paste" else "stop",
...
)

## S3 method for class 'formula'
ergm.ego_get_vattr(
  object,
  df,
  accept = "character",
  multiple = if (accept == "character") "paste" else "stop",
  ...
)

ergm.ego_attr_levels(object, attr, egor, levels = sort(unique(attr)), ...)

## S3 method for class 'numeric'
ergm.ego_attr_levels(object, attr, egor, levels = sort(unique(attr)), ...)

## S3 method for class 'logical'
ergm.ego_attr_levels(object, attr, egor, levels = sort(unique(attr)), ...)

## S3 method for class 'AsIs'
ergm.ego_attr_levels(object, attr, egor, levels = sort(unique(attr)), ...)

## S3 method for class 'character'
ergm.ego_attr_levels(object, attr, egor, levels = sort(unique(attr)), ...)

## S3 method for class `NULL`
ergm.ego_attr_levels(object, attr, egor, levels = sort(unique(attr)), ...)

## S3 method for class 'matrix'
ergm.ego_attr_levels(object, attr, egor, levels = sort(unique(attr)), ...)

## S3 method for class `function`
ergm.ego_attr_levels(object, attr, egor, levels = sort(unique(attr)), ...)

## S3 method for class 'formula'
ergm.ego_attr_levels(object, attr, egor, levels = sort(unique(attr)), ...)

COLLAPSE_SMALLEST(object, n, into)

Arguments

object An argument specifying the nodal attribute to select or which levels to include.
df Table of egos or of alters.
accept A character vector listing permitted data types for the output. See the Details section for the specification.
multiple Handling of multiple attributes or matrix or data frame output. See the Details section for the specification.

... Additional argument to the functions of network or to the formula's environment.

attr A vector of length equal to the number of nodes, specifying the attribute vector.

egor An egor object.

levels Starting set of levels to use; defaults to the sorted list of unique attributes.

n, into see `ergm::COLLAPSE_SMALLEST()`.

Details

The accept argument is meant to allow the user to quickly check whether the output is of an acceptable class or mode. Typically, if a term accepts a character (i.e., categorical) attribute, it will also accept a numeric one, treating each number as a category label. For this reason, the following outputs are defined:

"character" Accept any mode or class (since it can be converted to character).

"numeric" Accept real, integer, or logical.

"logical" Accept logical.

"integer" Accept integer or logical.

"natural" Accept a strictly positive integer.

"natural" Accept a nonnegative integer or logical.

"nonnegative" Accept a nonnegative number or logical.

"positive" Accept a strictly positive number or logical.

"paste" Paste together with dot as the separator.

"stop" Fail with an error message.

"matrix" Construct and/or return a matrix whose rows correspond to vertices.

Value

`ergm.ego_get_vattr` returns a vector of length equal to the number of nodes giving the selected attribute function. It may also have an attribute "name", which controls the suggested name of the attribute combination.

`ergm.ego_attr_levels` returns a vector of levels to use and their order.

Functions

- `COLLAPSE_SMALLEST()`: A version of `ergm::COLLAPSE_SMALLEST()` that can handle both `network` and `egodata` objects.
Examples

data(florentine)
flomego <- as.egor(flomarriage)
ergm.ego_get_vattr("priorates", flomego)
ergm.ego_get_vattr(~priorates, flomego)
ergm.ego_get_vattr(c("wealth","priorates"), flomego)
ergm.ego_get_vattr(~priorates>30, flomego)
(a <- ergm.ego_get_vattr(~cut(priorates,c(-Inf,0,20,40,60,Inf),label=FALSE)-1, flomego))
ergm.ego_attr_levels(NULL, a, flomego)
ergm.ego_attr_levels(-1, a, flomego)
ergm.ego_attr_levels(1:2, a, flomego)
ergm.ego_attr_levels(I(1:2), a, flomego)

predict.ergm.ego

---

**predict.ergm.ego**

**ERGM-based predicted tie probabilities for the pseudo-population network**

Description

ERGM-based predicted tie probabilities for the pseudo-population network

Usage

```r
## S3 method for class 'ergm.ego'
predict(object, ...)
```

Arguments

- `object`: model fit as returned by `ergm.ego()`
- `...`: other arguments passed to/from other methods

Value

See `ergm::predict.ergm()`

---

**sample**

**Draw random egocentric subsamples**

Description

Implementations of the `base::sample()` function for `egor::egor()` data.
Usage

```r
sample(x, size, replace = FALSE, prob = NULL, ...)
```

## Default S3 method:
sample(x, ...)

## S3 method for class 'egor'
sample(x, size, replace = FALSE, prob = NULL, ...)

Arguments

- `x`, `size`, `replace`, `prob`
  - see `base::sample()`.
- `...` extra arguments, currently unused.

Value

An `egor::egor()` object whose egos have been resampled in accordance with the arguments. Note that its `egor::ego_design()` information is overwritten in favor of the selection probabilities used in the sampling.

Note

A reimplementation of `sample` as a generic was necessary because `base::sample()` is not a generic and cannot take data-frame-alikes as arguments.

Examples

data(faux.mesa.high)
fmh.ego <- as.egor(faux.mesa.high)

# Create a tiny weighted sample:
(s3 <- sample(fmh.ego, 3, replace=TRUE, prob=1:nrow(fmh.ego$ego)))
# Resampling with prob=weights(egor) creates a self-weighted
# sample:
(sample(s3, 3, replace=TRUE, prob=weights(s3)))

# Create a large weighted sample, oversampling 12th-graders:
p <- ifelse(as_tibble(fmh.ego$ego)$Grade==12, 2, 1)
s2000 <- sample(fmh.ego, 2000, replace=TRUE, prob=p)

# Summary function adjusts for weights:
(summ.net <- summary(faux.mesa.high ~ edges + nodematch("Grade") +
    nodefactor("Race") + gwesp(0,fix=TRUE)))
(summ.ego <- summary(s2000 ~ edges + nodematch("Grade") +
    nodefactor("Race") + gwesp(0,fix=TRUE),
    scaleto=network.size(faux.mesa.high)))
**simulate.ergm.ego**

Simulate from a **ergm.ego** fit.

### Description

A wrapper around `simulate.formula` to simulate networks from an ERGM fit using `ergm.ego`.

### Usage

```r
define simulate
  ## S3 method for class 'ergm.ego'
simulate(object, nsim = 1, seed = NULL, constraints = object$constraints, popsize = if (object$popsize == 1 || object$popsize == 0 || is(object$popsize, "AsIs")) object$ppopsize else object$popsize, control = control.simulate.ergm.ego(), output = c("network", "stats", "edgelist", "pending_update_network", "ergm_state"), ... basis = NULL, verbose = FALSE )
```

### Arguments

- **object**: An `ergm.ego` fit.
- **nsim**: Number of realizations to simulate.
- **seed**: Seed value (integer) for the random number generator. See `set.seed`.
- **constraints, ...**: Additional arguments passed to `san` and `simulate.formula`.
- **popsize, basis**: A network size to which to scale the model for simulation; a `data.frame` with at least those ego attributes used to estimate the model to simulate over a specific set of actors; or a `network` object to use as is. `basis` is provided for consistency with `ergm()`, `ergm.ego()`, `simulate.ergm()` and others. If both are specified, `popsize` overrules.
- **control**: A `control.simulate.ergm.ego` control list.
- **output**: one of "network", "stats", "edgelist", "pending_update_network", or, for future compatibility, "ergm_state". See help for `simulate.ergm()` for explanation.
- **verbose**: A logical or an integer to control the amount of progress and diagnostic information to be printed. FALSE/0 produces minimal output, with higher values producing more detail. Note that very high values (5+) may significantly slow down processing.
Value

The output has the same format (with the same options) as `simulate.formula`. If `output="stats"` is passed, an additional attribute, "ppopsize" is set, giving the actual size of the network reconstructed, when the `pop.wt` control parameter is set to "round" and "popsiz"e is not a multiple of the egocentric sample size or the sampling weights.

Author(s)

Pavel N. Krivitsky

References


See Also

`simulate.formula`, `simulate.ergm`

Examples

data(faux.mesa.high)
data(fmhfit)
colMeans(egosim <- simulate(fmhfit, popsiz=300,nsim=50,
        output="stats", control=control.simulate.ergm.ego(
        simulate=control.simulate.formula(MCMC.burnin=2e6))))
colMeans(egosim)/attr(egosim,"ppopsize")*network.size(faux.mesa.high)
summary(faux.mesa.high~edges+degree(0:3)+nodefactor("Race")+nodefactor("Race")
+nodefactor("Sex")+nodefactor("Sex")+absdiff("Grade"))

---

**snctrl** Statnet Control

Description

A utility to facilitate argument completion of control lists, reexported from `statnet.common`.
Currently recognised control parameters

This list is updated as packages are loaded and unloaded.

See Also

`statnet.common::snctrl()`

---

## S3 method for class `egor`

```r
summary_formula(object, ..., basis = NULL, individual = FALSE, scaleto = NULL)
```

## S3 method for class `ergm.ego_svystat`

```r
x * y
```

### Arguments

- **object**: An `ergm`-style formula with an `egor` object as the LHS.
  - For a list of currently implemented egocentric terms for the RHS, see `ergm.ego-terms`.
- **...**: Not used at this time.
- **basis**: An optional `egor` object relative to which the statistics should be calculated.
- **individual**: If `FALSE` (the default), calculate the estimated per-capita statistics, weighted according to the ego weights, then scale them up to a network of size `scaleto`. If `TRUE`, calculate each ego’s individual contribution to the specified network statistics.
- **scaleto**: Size of a hypothetical network to which to scale the statistics. Defaults to the number of egos in the dataset.
- **x, y**: see `*.svystat`.

### Value

- If `individual==FALSE`, an `ergm.ego_svystat` object, which is a subclass of `svystat`—effectively a named vector of statistics. If `individual==TRUE`, a matrix with a row for each ego, giving that ego’s contribution to the network statistic.

### Functions

- `*`: A multiplication method that takes into account which statistics are scalable.
**template_network**

**Construct an Empty “Template” Network Consistent with an Egocentric Sample**

**Description**

Taking an object with ego information, constructs a network object with no edges whose vertices have the attributes of the egos in the dataset, replicating the egos as needed, and taking into accounts their sampling weights.

**Author(s)**

Pavel N. Krivitsky

**References**


**See Also**

`summary_formula`, `summary_formula.ergm`

**Examples**

```r
data(faux.mesa.high)
fmh.ego <- as.egor(faux.mesa.high)
(nw.summ <- summary(faux.mesa.high ~ edges + degree(0:3) + nodematch("Race") + nodematch("Sex") + absdiff("Grade") + nodemix("Grade")))

(ego.summ <- summary(fmh.ego ~ edges + degree(0:3) + nodematch("Race") + nodematch("Sex") + absdiff("Grade") + nodemix("Grade", scaleto = network.size(faux.mesa.high))))

stopifnot(isTRUE(all.equal(as.vector(nw.summ), as.vector(ego.summ))))

(ego.summ2 <- summary(fmh.ego ~ edges + meandeg + degree(0:2)))
vcov(ego.summ2)

ego.summ2 * 2 # edges and degrees scales, meandeg doesn't
vcov(ego.summ2 * 2)
```
template_network

Usage

```r
template_network(x, ...)
```

## S3 method for class 'data.frame'
template_network(x, ...)

## S3 method for class 'egor'
template_network(x, N, scaling = c("round", "sample"), ...)

Arguments

- **x**: A `egor` object.
- **...**: Additional arguments, currently unused.
- **N**: The target number of vertices the output network should have.
- **scaling**: If `egor` contains weights or `N` is not a multiple of number of egos in the sample, it may not be possible, for a finite `N` to represent each ego exactly according to its relative weight, and scaling controls how the fractional egos are allocated:
  - **"round"**: (the default) Rather than treating `N` as a hard setting, calculate \( \frac{Nw_i}{w} \) for each ego \( i \) and round it to the nearest integer. Then, the `N` actually used will be the sum of these rounded frequencies.
  - **"sample"**: Resample in proportion to \( w_i \).

Value

A `network` object.

Methods (by class)

- `template_network(data.frame)`: method for `data.frames` and `tibbles`, specifying ego composition directly.
- `template_network(egor)`: method for `egor` objects; weights, if any, are obtained from the `egor`'s design information.

Author(s)

Pavel N. Krivitsky

See Also

`as.egor.network`, which performs the inverse operation.

Examples

```r
data(faux.mesa.high)
summary(faux.mesa.high, print.adj = FALSE)

fmh.ego <- as.egor(faux.mesa.high)
```
# Same actor attributes
fmh.template <- template_network(fmh.ego, N=network.size(faux.mesa.high))
summary(fmh.template, print.adj = FALSE)

# Twice the actors, same distribution
fmh2.template <- template_network(fmh.ego, N=2*network.size(faux.mesa.high))
summary(fmh2.template, print.adj = FALSE)
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