

# Package ‘ergm.ego’

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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.

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as.egodata.network	<i>Construct an Egocentric View of a Network</i>
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### Description

Given a [network](#) object, construct an [egodata](#) object representing a census of all the actors in the network. Used mainly for testing.

### Usage

```
## S3 method for class 'network'
as.egodata(
  object,
  special.cols = c("na", "vertex.names"),
  ...,
  egoIDcol = "vertex.names"
)
```

### Arguments

object	A <a href="#">network</a> object.
special.cols	Vertex attributes that should not be copied to the egos and alters tables. Defaults to attributes special to the <a href="#">network</a> objects.
...	Additional arguments, currently unused.
egoIDcol	The name of the vertex attribute containing unique ego IDs. Defaults to "vertex.names".

**Value**

An `egodata` object.

**Author(s)**

Pavel N. Krivitsky

**See Also**

`as.network.egodata`, which performs the inverse operation (though drops the ties).

**Examples**

```
# See example(ergm.ego) and example(as.network.egodata).
```

---

<code>as.network.egodata</code>	<i>Construct an Empty “Template” Network Consistent with an Egocentric Sample</i>
---------------------------------	---

---

**Description**

Taking a `egodata` object, 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.

**Usage**

```
## S3 method for class 'egodata'
as.network(x, N, scaling = c("round", "sample"), ...)
```

**Arguments**

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

**Value**

A [network](#) object. It has an additional vertex attribute `.ego.ind` giving the *row index* (not ego ID) of the ego from which a particular node was generated.

**Author(s)**

Pavel N. Krivitsky

**See Also**

[as.egodata.network](#), which performs the inverse operation.

**Examples**

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

fmh.ego <- as.egodata(faux.mesa.high)

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

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

---

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("asymptotic", "bootstrap", "jackknife", "naive"),
  boot.R = 10000,
  ergm.control = control.ergm(),
  ...
)
```

## Arguments

ppopsize, ppopsize.mul	<p>Parameters to determine the size <math> N' </math> of the pseudopopulation network. ppopsize can be</p> <p><b>"auto"</b> If the popsize (<math> N </math>) argument is specified and is different from 1, as if "pop"; otherwise, as "samp".</p> <p><b>"samp"</b> set <math> N' </math> based on the sample size: <math> N'  =  S  \times \text{ppopsize.mul}</math></p> <p><b>"pop"</b> set <math> N' </math> based on the population size: <math> N'  =  N  \times \text{ppopsize.mul}</math></p> <p><b>a number</b> set <math> N' </math> directly (ppopsize.mul ignored)</p> <p><b>a network object</b> use the specified network as the pseudo-population network directly; use at your own risk</p> <p><b>a data frame</b> use the specified data frame as the pseudo-population; use at your own risk</p> <p>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.</p>
ppop.wt	<p>Because each ego must be represented in the pseudopopulation network an integral number of times, if the sample is weighted (or the target <math> N' </math> calculated from ppopsize and ppopsize.mul is not a multiple of the sample size), it may not be possible, for a finite <math> N' </math> to represent each ego exactly according to its relative weight, and ppop.wt controls how the fractional egos are allocated:</p> <p><b>"round"</b> (default) Rather than treating ppopsize as a hard setting, calculate <math> N' w_i/w</math> for each ego <math>i</math> and round it to the nearest integer. Then, the <math> N' </math> actually used will be the sum of these rounded frequencies.</p> <p><b>"sample"</b> Resample in proportion to <math>w_i</math>.</p>
stats.wt	<p>Weight assigned to each ego's contribution to the ERGM's sufficient statistic:</p> <p><b>"data"</b> (default) Use weights <math> N' w_i/w</math> for each ego <math>i</math> as in the data.</p> <p><b>"ppop"</b> Use weights ultimately used in the pseudopopulation network.</p>
stats.est, boot.R	<p>Method to be used to estimate the ERGM's sufficient statistics and their variance:</p> <p><b>"asymptotic"</b> Delta method, as derived by Krivitsky and Morris (2015), assuming the ego weights are sampled alongside the egos.</p> <p><b>(default)</b> Delta method, as derived by Krivitsky and Morris (2015), assuming the ego weights are sampled alongside the egos.</p> <p><b>"bootstrap"</b> Nonparametric bootstrap with bias correction, resampling egos, using R replications.</p> <p><b>"jackknife"</b> Jackknife with bias correction.</p> <p><b>"naive"</b> "Naive" estimator, assuming that weights are fixed.</p>
ergm.control	Control parameters for the <code>ergm</code> call to fit the model, constructed by <code>control.ergm</code> .
...	Not used at this time.

**Value**

A list with arguments as components.

**Author(s)**

Pavel N. Krivitsky

**References**

Pavel N. Krivitsky and Martina Morris. Inference for Social Network Models from Egocentrically-Sampled Data, with Application to Understanding Persistent Racial Disparities in HIV Prevalence in the US. Technical Report. National Institute for Applied Statistics Research Australia, University of Wollongong, 2015(05-15). <http://niasra.uow.edu.au/publications/UOW190187.html>

**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**

```
control.simulate.ergm.ego(
  ppop.wt = c("round", "sample"),
  SAN.control = control.san(),
  simulate.control = 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  $|N'|w_i/w$  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  $w_i$ .

<code>SAN.control</code>	A list of control parameters for <code>san</code> constructed by <code>control.ergm</code> , called to construct a pseudopopulation network consistent with the data.
<code>simulate.control</code>	A list of control parameters for <code>simulate.formula</code> constructed by <code>control.simulate</code> , called to simulate from the model fit.
<code>...</code>	Not used at this time.

**Value**

A list with arguments as components.

**Author(s)**

Pavel N. Krivitsky

**See Also**

`control.simulate`, `control.san`

---

`degreedist.egodata`      *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**

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

**Arguments**

object	A <a href="#">egodata</a> 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; if FALSE, graphical parameters and brgmod have no effect.
weight	Whether sampling weights should be incorporated into the calculation (TRUE, the default) or ignored (FALSE).
...	Additional arguments to <a href="#">simulate.ergm.ego()</a> .

**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.egodata(faux.mesa.high)

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

---

egodata

---

*Convert to or Construct [egodata](#) Objects*


---

**Description**

[as.egodata](#) is a generic function to construct [egodata](#) objects from a variety of sources. [egodata](#) function is the standard constructor, taking two data frames. For other methods for this class, see the Miscellaneous Methods section.



**Usage**

```
egodata(egos, alters, egoWt = 1, ..., egoIDcol = "egoID")

as.egodata(object, ..., egoIDcol = "egoID")

## S3 method for class 'data.frame'
as.egodata(
  object,
  alters,
  egoWt = 1,
  ...,
  egoIDcol = "egoID",
  alterIcol = "alterInd",
  alterIDcol = "alterID"
)
```

**Arguments**

alters	<p>The <code>data.frame</code> containing at least the column named in <code>egoIDcol</code>, whose values do not have to be unique, and not every ego has to be represented. Other columns contain information about the alters.</p> <p>For the <code>data.frame</code> method in which the <code>object</code> argument also contains alter information in ‘wide’ format, a <code>list</code> with the following information:</p> <p><code>columns</code> A character, integer, or logical vector identifying which columns contain the alters’ information.</p> <p><code>count</code> The name of the column containing the number of alters nominated by that ego.</p> <p><code>name.sep</code> A one-character string or an empty string (defaulting to <code>"."</code>) specifying the character, if any, used to separate alter attribute name from alter’s index within the ego. If an empty string (<code>""</code>), attribute name is assumed to be made of letters, with any numbers being the alter index.</p> <p><code>egoIDcol</code>, whose values do not have to be unique, and not every ego must be represented. Other columns contain information about the alters.</p>
egoWt	A vector of the same length as number of rows in <code>egos</code> or <code>object</code> , containing the relative sampling weight of each ego.
...	Additional arguments; currently unused.
egoIDcol	Name of the column in the ego table containing the unique ego identifier.
object, egos	<p>The object from which the egocentric data should be constructed. For the <code>data.frame</code> methods and <code>egodata</code> itself, a data frame containing at least the column named in <code>egoIDcol</code>, whose values must all be unique. Other columns contain information about the egos.</p> <p>For the <code>data.frame</code> method, it may also contain the information about the alters in a ‘wide’ format, in the form of additional columns with names like <code>ATTRNAME1</code>, <code>ATTRNAME2</code>, etc. for attribute <code>ATTRNAME</code> of alter 1, 2, etc., as well as a column containing the number of alters nominated by that ego.</p>
alterIcol	Column name to use for the within-ego index of the alter.

`alterIDcol` Column name to use for the unique ID of each alter, constructed by concatenating the ID of the ego that nominated them and their index within that ego.

### Value

An `egodata` object. The object is a list containing the following elements:

<code>egos</code>	A data frame with one row for each ego, containing at least the column named in <code>egoIDcol</code> , and other columns containing attributes of the egos.
<code>alters</code>	A data frame containing at least the column named in <code>egoIDcol</code> , and other columns containing attributes of the alters.
<code>egoWt</code>	A vector of the same length as the number of egos, containing the relative sampling weight of each ego.
<code>egoIDcol</code>	Name of the column in the ego table containing the unique ego identifier.

### Miscellaneous Methods

The following “standard” methods have also been implemented for `egodata`:

**"dim.egodata"** A vector with three elements containing the “dimensions” of the `egodata` object: number of egos, number of columns in the egos table, and number of columns in the alters table, inclusive of the ego identifier column. As a corollary, `nrow` returns the number of egos in the dataset.

**"dimnames.egodata"** A list with three elements containing the “dimension names” of the `egodata` object: ego IDs, column names of the egos table, and column names of the alters table, inclusive of the ego identifier column.

**"sample.egodata"** As `sample`, but takes and returns a simulated `egodata` dataset by resampling egos, adjusting ego weights as necessary, if weighted sampling was used.

**"head.egodata"** As `head`, but returns the first `n` rows of egos, alters, and weights.

**"na.omit.egodata"** As `na.omit.data.frame`, but takes and returns an `egodata` dataset, with egos with NA in their rows or in their alters’ rows. An optional argument `relevant`, defaulting to all columns, can be used to select (by index or name) based on which columns an ego may be dropped. (I.e., NAs in those not “relevant” are ignored.)

### Author(s)

Pavel N. Krivitsky

### See Also

`ergm.ego` for examples, `as.network.egodata`, `as.egodata.network`, `subset.egodata`, `[.egodata`

---

ergm.ego	<i>Inference for Exponential-Family Random Graph Models based on Egocentrically Sampled Data</i>
----------	--

---

### Description

A wrapper around the [ergm](#) to fit an ERGM to an [egodata](#) object.

### Usage

```
ergm.ego(
  formula,
  popsize = 1,
  offset.coef = NULL,
  ...,
  control = control.ergm.ego(),
  na.action = na.fail,
  do.fit = TRUE
)
```

### Arguments

formula	An <a href="#">formula</a> object, of the form $e \sim \langle \text{model terms} \rangle$ , where $e$ is a <a href="#">egodata</a> object. See <a href="#">ergm</a> for details and examples. For a list of currently implemented egocentric terms for the RHS, see <a href="#">ergm.ego-terms</a> .
popsize	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.
offset.coef	A vector of coefficients for the offset terms.
...	Additional arguments passed to <a href="#">ergm</a> .
control	A <a href="#">control.ergm.ego</a> control list.
na.action	How to handle missing actor attributes in egos or alters, when the terms need them.
do.fit	Whether to actually call <a href="#">ergm</a>

### Value

An object of class `ergm.ego` inheriting from [ergm](#), with the following additional or overridden elements:

"v"	Variance-covariance matrix of the estimate of the sufficient statistics
"m"	Estimate of the sufficient statistics
"egodata"	The <code>egodata</code> object passed
"popsize"	Population network size used

"ppopsize"	Pseudopopulation size used, see <a href="#">control.ergm.ego</a>
"coef"	The coefficients, along with the network size adjustment <code>net.size.adj</code> coefficient.
"covar"	Pseudo-MLE estimate of the variance-covariance matrix of the parameter estimates under repeated egocentric sampling
"ergm.covar"	The variance-covariance matrix of parameter estimates under the ERGM superpopulation process (without incorporating sampling).
"DtDe"	Estimated Jacobian of the expectation of the sufficient statistics with respect to the model parameters

### Author(s)

Pavel N. Krivitsky

### References

Pavel N. Krivitsky and Martina Morris. Inference for Social Network Models from Egocentrically-Sampled Data, with Application to Understanding Persistent Racial Disparities in HIV Prevalence in the US. Technical Report. National Institute for Applied Statistics Research Australia, University of Wollongong, 2015(05-15). <http://niasra.uow.edu.au/publications/UOW190187.html>

### Examples

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

head(fmh.ego)

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

# Run convergence diagnostics
mcmc.diagnostics(egofit)

# Estimates and standard errors
summary(egofit)

# Note that we recover the ergm() parameters
## Not run:
coef(ergm(faux.mesa.high~edges+degree(0:3)+nodefactor("Race")+nodematch("Race")
         +nodefactor("Sex")+nodematch("Sex")+absdiff("Grade"),
         eval.loglik=FALSE))

## End(Not run)
rbind(c(0, -0.8407, 2.3393, 1.4686, 0.6323, 0.5287, -1.3603, -1.0454,
        -2.4998, -0.7207, 0.833, -0.1823, 0.6357, -1.3513),
      coef(egofit))
```

---

ergm.ego-terms

[ergm Terms Implemented for egodata](#)

---

## Description

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.

## Details

The current recommendation for any package implementing additional egocentric calculator terms is to create a help file with a name or alias `ergm.egodata-terms`, so that `help("ergm.egodata-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`** A special-purpose term equivalent to [edges](#), 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`
  - `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](#)

---

`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.

## Usage

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

## Arguments

<code>object</code>	An <code>ergm.ego</code> fit.
<code>...</code>	Additional arguments, currently unused.
<code>GOF</code>	A string specifying the statistics whose goodness of fit is to be evaluated. Currently, only “degree” and “model” are implemented; see <code>gof</code> documentation for details.
<code>control</code>	A list to control parameters, constructed using <code>control.gof.formula</code> or <code>control.gof.ergm</code> (which have different defaults).
<code>verbose</code>	Provide verbose information on the progress of the simulation.

## Value

An object of class `gofobject`.

## Author(s)

Pavel N. Krivitsky

## See Also

For examples, see `ergm.ego`.

## Examples

```

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

head(fmh.ego)

egofit <- ergm.ego(fmh.ego~edges+degree(0:3)+nodefactor("Race")+nodematch("Race")
                  +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.egodata    *Summarizing the mixing among groups in an egocentric dataset*

---

## Description

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

## Usage

```

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

```

## Arguments

<code>object</code>	A <code>egodata</code> object.
<code>attrname</code>	A character vector containing the name of the network attribute whose mixing matrix is wanted.
<code>rowprob</code>	Whether the counts should be normalized by row sums. That is, whether they should be proportions conditional on the ego's group.
<code>weight</code>	Whether sampling weights should be incorporated into the calculation (TRUE, the default) or ignored (FALSE).
<code>...</code>	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.egodata` will be twice that returned by `mixingmatrix` for the original undirected network.

**See Also**

`mixingmatrix`, `nodemix`, `summary` method for egocentric data

**Examples**

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

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

stopifnot(isTRUE(all.equal({tmp<-unclass(mm$matrix); diag(tmp) <- diag(tmp)*2;
tmp}, mm.ego, check.attributes=FALSE)))
```

---

nodal\_attributes-API    *Helper functions for specifying nodal attribute levels*

---

**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**

```
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",
  ...
)

## 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, egodata, levels = sort(unique(attr)), ...)

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

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

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

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

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

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

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

### 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.
egodata	An <a href="#">egodata</a> object.
levels	Starting set of levels to use; defaults to the sorted list of unique attributes.

### 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 beconverted to character).

"numeric" Accept real, integer, or logical.

"logical" Accept logical.

"integer" Accept integer or logical.

"natural" Accept a strictly positive integer.

"0natural" 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.

**Examples**

```

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

```

---

predict.ergm.ego	<i>ERGM-based predicted tie probabilities for the pseudo-population network</i>
------------------	---

---

**Description**

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

**Usage**

```

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

```

**Arguments**

object	model fit as returned by <a href="#">ergm.ego()</a>
...	other arguments passed to/from other methods

**Value**

See [ergm::predict.ergm\(\)](#)

---

simulate.ergm.ego	<i>Simulate from a <a href="#">ergm.ego</a> fit.</i>
-------------------	--

---

**Description**

A wrapper around [simulate.formula](#) to simulate networks from an ERGM fit using [ergm.ego](#).

**Usage**

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

**Arguments**

object	An <a href="#">ergm.ego</a> fit.
nsim	Number of realizations to simulate.
seed	Random seed.
constraints, ...	Additional arguments passed to <a href="#">san</a> and <a href="#">simulate.formula</a> .
popsize	Either network size to which to scale the model for simulation or a <a href="#">data.frame</a> with at least those ego attributes required to estimate the model, to simulate over a specific set of actors.
control	A <a href="#">control.simulate.ergm.ego</a> control list.
output	one of "network", "stats", "edgelist", "pending_update_network", or, for future compatibility, "ergm_state". See help for <a href="#">simulate.ergm()</a> for explanation.
verbose	Verbosity of output.

**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 "popsize" is not a multiple of the egocentric sample size or the sampling weights.

**Author(s)**

Pavel N. Krivitsky

**References**

Pavel N. Krivitsky and Martina Morris. Inference for Social Network Models from Egocentrically-Sampled Data, with Application to Understanding Persistent Racial Disparities in HIV Prevalence in the US. Technical Report. National Institute for Applied Statistics Research Australia, University of Wollongong, 2015(05-15). doi: [10.1214/16AOAS1010](https://doi.org/10.1214/16AOAS1010)

Pavel N. Krivitsky, Mark S. Handcock, and Martina Morris. Adjusting for Network Size and Composition Effects in Exponential-Family Random Graph Models. *Statistical Methodology*, 2011, 8(4), 319-339. doi: [10.1016/j.stamet.2011.01.005](https://doi.org/10.1016/j.stamet.2011.01.005)

### See Also

[simulate.formula](#), [simulate.ergm](#)

### Examples

```
data(faux.mesa.high)
fmh.ego <- as.egodata(faux.mesa.high)
egofit <- ergm.ego(fmh.ego~edges+degree(0:3)+nodefactor("Race")+nodematch("Race")
  +nodefactor("Sex")+nodematch("Sex")+absdiff("Grade"),
  popsize=network.size(faux.mesa.high))
colMeans(egosim <- simulate(egofit, popsize=300,nsim=50,
  output="stats", control=control.simulate.ergm.ego(
  simulate.control=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")+nodematch("Race")
  +nodefactor("Sex")+nodematch("Sex")+absdiff("Grade"))
```

---

summary\_formula.egodata

*Calculation of ERGM-style summary statistics for [egodata](#) objects.*

---

### Description

Used to calculate the specified network statistics inferred from a [egodata](#) object.

### Usage

```
## S3 method for class 'egodata'
summary_formula(object, ..., basis = NULL, individual = FALSE, scaleto = NULL)
```

### Arguments

object	An <a href="#">ergm</a> -style formula with a <a href="#">egodata</a> object as the LHS. For a list of currently implemented egocentric terms for the RHS, see <a href="#">ergm.ego-terms</a> .
...	Not used at this time.
basis	An optional <a href="#">egodata</a> 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 <code>scaleto</code> . 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.

**Value**

If `individual==FALSE`, 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.

**Author(s)**

Pavel N. Krivitsky

**References**

Pavel N. Krivitsky and Martina Morris. Inference for Social Network Models from Egocentrically-Sampled Data, with Application to Understanding Persistent Racial Disparities in HIV Prevalence in the US. Technical Report. National Institute for Applied Statistics Research Australia, University of Wollongong, 2015(05-15). doi: [10.1214/16AOAS1010](https://doi.org/10.1214/16AOAS1010)

Pavel N. Krivitsky, Mark S. Handcock, and Martina Morris. Adjusting for Network Size and Composition Effects in Exponential-Family Random Graph Models. *Statistical Methodology*, 2011, 8(4), 319-339. doi: [10.1016/j.stamet.2011.01.005](https://doi.org/10.1016/j.stamet.2011.01.005)

**See Also**

[summary\\_formula](#), [summary\\_formula.ergm](#)

**Examples**

```
data(faux.mesa.high)
fmh.ego <- as.egodata(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(nw.summ, ego.summ)))
```

**Description**

Returns subsets of `egodata` objects that meet conditions.

**Usage**

```
## S3 method for class 'egodata'  
x[i, j, ..., dup.action = c("make.unique", "fail", "number")]  
  
## S3 method for class 'egodata'  
subset(x, subset, select, ..., dup.action = c("make.unique", "fail", "number"))
```

**Arguments**

x	An <a href="#">egodata</a> object.
...	Additional arguments, currently unused.
dup.action	What to do when an ego is referenced multiple times: <b>"make.unique"</b> Construct new unique ego IDs using the <a href="#">make.unique</a> function <b>"fail"</b> Exit with an error. <b>"number"</b> Number the egos consecutively in the order they were selected
subset, i	An expression (evaluated in the context of the egos table of x producing a logical, integer, or character vector indicating which egos to select (and, for the latter two, how many times).
select, j	A numeric or character vector specifying the columns of egos and alters to select.

**Value**

An [egodata](#) object.

**Author(s)**

Pavel N. Krivitsky

**See Also**

[sample.egodata](#)

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