

Package ‘wdnet’

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Title Weighted and Directed Networks

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Description Implementations of network analysis including

- (1) assortativity coefficient of weighted and directed networks, Yuan, Yan and Zhang (2021) <[doi:10.1093/comnet/cnab017](https://doi.org/10.1093/comnet/cnab017)>,
- (2) centrality measures for weighted and directed networks, Opsahl, Agneessens and Skvoretz (2010) <[doi:10.1016/j.socnet.2010.03.006](https://doi.org/10.1016/j.socnet.2010.03.006)>,
- Zhang, Wang and Yan (2022) <[doi:10.1016/j.physa.2021.126438](https://doi.org/10.1016/j.physa.2021.126438)>,
- (3) clustering coefficient of weighted and directed networks, Fagiolo (2007) <[doi:10.1103/PhysRevE.76.026107](https://doi.org/10.1103/PhysRevE.76.026107)> and Clemente and Grassi (2018) <[doi:10.1016/j.chaos.2017.12.007](https://doi.org/10.1016/j.chaos.2017.12.007)>,
- (4) rewiring networks with given assortativity coefficients, Wang, Yan, Yuan and Zhang (2022) <[doi:10.1007/s11222-022-10161-8](https://doi.org/10.1007/s11222-022-10161-8)>,
- (5) preferential attachment network generation.

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LinkingTo Rcpp, RcppArmadillo

BugReports <https://gitlab.com/wdnetwork/wdnet/-/issues>

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+.rpacontrol	<i>Add components to the control list</i>
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Description

'+' is used to combine components to control the PA network generation process. Available components are `rpa_control_scenario()`, `rpa_control_edgweight()`, `rpa_control_newedge()`, `rpa_control_preference()` and `rpa_control_reciprocal()`.

Usage

```
## S3 method for class 'rpacontrol'
e1 + e2
```

Arguments

e1	A list of class <code>rpacontrol</code> .
e2	A list of class <code>rpacontrol</code> .

Value

A list of class `rpacontrol` with components from e1 and e2.

Examples

```
control <- rpa_control_scenario(alpha = 0.5, beta = 0.5) +
  rpa_control_preference(ftype = "customized",
    spref = "pow(outs, 2) + 1",
    tpref = "pow(ins, 2) + 1")
```

```
control <- rpa_control_scenario(alpha = 1) +
  rpa_control_edgweight(distribution = rgamma,
    dparams = list(shape = 5, scale = 0.2),
    shift = 1)
```

assortcoef	<i>Assortativity coefficient</i>
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Description

Compute the assortativity coefficient of a network.

Usage

```
assortcoef(
  edgelist = NULL,
  edgweight = NULL,
  adj = NULL,
  directed = TRUE,
  f1 = NULL,
  f2 = NULL
)
```

Arguments

<code>edgelist</code>	A two column matrix represents edges. If <code>NULL</code> , <code>edgelist</code> and <code>edgweight</code> will be extracted from the adjacency matrix <code>adj</code> .
<code>edgweight</code>	A vector represents the weight of edges. If <code>edgelist</code> is provided and <code>edgweight</code> is <code>NULL</code> , all the edges will be considered have weight 1.
<code>adj</code>	An adjacency matrix.
<code>directed</code>	Logical. Whether the edges will be considered as directed.
<code>f1</code>	A vector, represents the first feature of existing nodes. Number of nodes = <code>length(f1) = length(f2)</code> . Defined for directed networks. If <code>NULL</code> , out-strength will be used.
<code>f2</code>	A vector, represents the second feature of existing nodes. Defined for directed networks. If <code>NULL</code> , in-strength will be used.

Value

Assortativity coefficient for undirected networks, or four assortativity coefficients for directed networks.

Note

When the adjacency matrix is binary (i.e., directed but unweighted networks), `assortcoef` returns the assortativity coefficient proposed in Foster et al. (2010).

References

- Foster, J.G., Foster, D.V., Grassberger, P. and Paczuski, M. (2010). Edge direction and the structure of networks. *Proceedings of the National Academy of Sciences of the United States*, 107(24), 10815–10820.
- Yuan, Y. Zhang, P. and Yan, J. (2021). Assortativity coefficients for weighted and directed networks. *Journal of Complex Networks*, 9(2), cnab017.

Examples

```
set.seed(123)
control <- rpa_control_edgweight(distribution = rgamma,
  dparams = list(shape = 5, scale = 0.2), shift = 0)
netwk <- rpanet(nstep = 10^4, control = control)
result <- assortcoef(netwk$edgelist, edgweight = netwk$edgweight, directed = TRUE)
```

centrality

Centrality measures

Description

Compute the centrality measures of the nodes in a weighted and directed network.

Usage

```
centrality(
  adj = NULL,
  edgelist = NULL,
  edgweight = NULL,
  measure = c("degree", "closeness", "wpr"),
  degree.control = list(alpha = 1, mode = "out"),
  closeness.control = list(alpha = 1, mode = "out", method = "harmonic", distance =
    FALSE),
  wpr.control = list(gamma = 0.85, theta = 1, prior.info = NULL)
)
```

Arguments

<code>adj</code>	An adjacency matrix of a weighted and directed network. If <code>NULL</code> , <code>edgelist</code> and <code>edgweight</code> will be used to construct the adjacency matrix.
<code>edgelist</code>	A two column matrix, each row represents a directed edge of the network. It will be ignored if <code>adj</code> is not <code>NULL</code> .
<code>edgweight</code>	A vector represents the weight of edges. If <code>edgelist</code> is provided and <code>edgweight</code> is <code>NULL</code> , all the edges will be considered have weight 1. It will be ignored if <code>adj</code> is not <code>NULL</code> .
<code>measure</code>	Which measure to use: "degree" (degree-based centrality), "closeness" (closeness centrality), or "wpr" (weighted PageRank centrality)?
<code>degree.control</code>	A list of parameters passed to the degree centrality measure. <ul style="list-style-type: none"> • <code>alpha</code> A tuning parameter. The value of <code>alpha</code> must be nonnegative. By convention, <code>alpha</code> takes a value from 0 to 1 (default). • <code>mode</code> Which mode to compute: "out" (default) or "in"? For undirected networks, this setting is irrelevant.
<code>closeness.control</code>	A list of parameters passed to the closeness centrality measure. <ul style="list-style-type: none"> • <code>alpha</code> A tuning parameter. The value of <code>alpha</code> must be nonnegative. By convention, <code>alpha</code> takes a value from 0 to 1 (default). • <code>mode</code> Which mode to compute: "out" (default) or "in"? For undirected networks, this setting is irrelevant. • <code>method</code> Which method to use: "harmonic" (default) or "standard"? • <code>distance</code> Whether to consider the entries in the adjacency matrix as distances or strong connections. The default setting is <code>FALSE</code>.
<code>wpr.control</code>	A list of parameters passed to the weighted PageRank centrality measure. <ul style="list-style-type: none"> • <code>gamma</code> The damping factor; it takes 0.85 (default) if not given. • <code>theta</code> A tuning parameter leveraging node degree and strength; <code>theta = 0</code> does not consider edge weight; <code>theta = 1</code> (default) fully considers edge weight. • <code>prior.info</code> Vertex-specific prior information for restarting when arriving at a sink. When it is not given (<code>NULL</code>), a random restart is implemented.

Value

A list of node names and associated centrality measures

Note

The degree-based centrality measure is an extension of function `strength` in package `igraph` and an alternative of function `degree_w` in package `tnet`.

The closeness centrality measure is an extension of function `closeness` in package `igraph` and function `closeness_w` in package `tnet`. The method of computing distances between vertices is the *Dijkstra's algorithm*.

The weighted PageRank centrality measure is an extension of function `page_rank` in package `igraph`.

References

- Dijkstra, E.W. (1959). A note on two problems in connexion with graphs. *Numerische Mathematik*, 1, 269–271.
- Newman, M.E.J. (2003). The structure and function of complex networks. *SIAM review*, 45(2), 167–256.
- Opsahl, T., Agneessens, F., Skvoretz, J. (2010). Node centrality in weighted networks: Generalizing degree and shortest paths. *Social Networks*, 32, 245–251.
- Zhang, P., Wang, T. and Yan, J. (2022) PageRank centrality and algorithms for weighted, directed networks with applications to World Input-Output Tables. *Physica A: Statistical Mechanics and its Applications*, 586, 126438.
- Zhang, P., Zhao, J. and Yan, J. (2020+) Centrality measures of networks with application to world input-output tables

Examples

```
## Generate a network according to the Erd\{o}s-Renyi model of order 20
## and parameter p = 0.3
edge_ER <- rbinom(400,1,0.3)
weight_ER <- sapply(edge_ER, function(x) x*sample(3,1))
adj_ER <- matrix(weight_ER,20,20)
mydegree <- centrality(adj_ER, measure = "degree", degree.control =
list(alpha = 0.8, mode = "in"))
myclose <- centrality(adj_ER, measure = "closeness", closeness.control =
list(alpha = 0.8, mode = "out", method = "harmonic", distance = FALSE))
mywpr <- centrality(adj_ER, measure = "wpr", wpr.control =
list(gamma = 0.85, theta = 0.75))
```

clustcoef	<i>Directed clustering coefficient</i>
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Description

Compute the clustering coefficient of a weighted and directed network.

Usage

```
clustcoef(adj, method = c("Clemente", "Fagiolo"), isolates = "zero")
```

Arguments

adj	is an adjacency matrix of an weighted and directed network.
method	which method used to compute clustering coefficients: Clemente and Grassi (2018) or Fagiolo (2007).
isolates	character, defines how to treat vertices with degree zero and one. If "zero", then their clustering coefficient is returned as 0 and are included in the averaging. Otherwise, their clustering coefficient is NaN and are excluded in the averaging. Default value is "zero".

Value

lists of local clustering coefficients (in terms of a vector), global clustering coefficient (in terms of a scalar) and number of weighted directed triangles (in terms of a vector) base on total, in, out, middleman (middle), or cycle triplets.

Note

Self-loops (if exist) are removed prior to the computation of clustering coefficient. When the adjacency matrix is symmetric (i.e., undirected but possibly unweighted networks), `clustcoef` returns local and global clustering coefficients proposed by Barrat et al. (2010).

References

- Barrat, A., Barthélemy, M., Pastor-Satorras, R. and Vespignani, A. (2004). The architecture of complex weighted networks. *Proceedings of National Academy of Sciences of the United States of America*, 101(11), 3747–3752.
- Clemente, G.P. and Grassi, R. (2018). Directed clustering in weighted networks: A new perspective. *Chaos, Solitons & Fractals*, 107, 26–38.
- Fagiolo, G. (2007). Clustering in complex directed networks. *Physical Review E*, 76, 026107.

Examples

```
## Generate a network according to the Erdős-Renyi model of order 20
## and parameter p = 0.3
edge_ER <- rbinom(400,1,0.3)
weight_ER <- sapply(edge_ER, function(x) x*sample(3,1))
adj_ER <- matrix(weight_ER,20,20)
mycc <- clustcoef(adj_ER, method = "Clemente")
system.time(mycc)
```

cvxr_control

Parameters passed to CVXR::solve().

Description

Defined for the convex optimization problems for solving eta.

Usage

```
cvxr_control(
  solver = "ECOS",
  ignore_dcp = FALSE,
  warm_start = FALSE,
  verbose = FALSE,
  parallel = FALSE,
  gp = FALSE,
```

```

    feastol = 1e-05,
    reltol = 1e-05,
    abstol = 1e-05,
    num_iter = NULL,
    ...
)

```

Arguments

<code>solver</code>	(Optional) A string indicating the solver to use. Defaults to "ECOS".
<code>ignore_dcp</code>	(Optional) A logical value indicating whether to override the DCP check for a problem.
<code>warm_start</code>	(Optional) A logical value indicating whether the previous solver result should be used to warm start.
<code>verbose</code>	(Optional) A logical value indicating whether to print additional solver output.
<code>parallel</code>	(Optional) A logical value indicating whether to solve in parallel if the problem is separable.
<code>gp</code>	(Optional) A logical value indicating whether the problem is a geometric program. Defaults to FALSE.
<code>feastol</code>	The feasible tolerance on the primal and dual residual. Defaults to 1e-5.
<code>reltol</code>	The relative tolerance on the duality gap. Defaults to 1e-5.
<code>abstol</code>	The absolute tolerance on the duality gap. Defaults to 1e-5.
<code>num_iter</code>	The maximum number of iterations.
<code>...</code>	Additional options that will be passed to the specific solver. In general, these options will override any default settings imposed by CVXR.

Value

A list containing the parameters.

Examples

```
control <- cvxr_control(solver = "OSQP", abstol = 1e-5)
```

dprewire

Degree preserving rewiring.

Description

Rewire a given network to have predetermined assortativity coefficients while preserving node degree.

Usage

```
dprewire(
  edgelist = NULL,
  directed = TRUE,
  adj = NULL,
  target.assortcoef = list(outout = NULL, outin = NULL, inout = NULL, inin = NULL),
  control = list(iteration = 200, nattempts = NULL, history = FALSE, cvxr_control =
    cvxr_control(), eta.obj = function(x) 0),
  eta = NULL
)
```

Arguments

<code>edgelist</code>	A two column matrix, each row represents an edge of the network.
<code>directed</code>	Logical, whether the network is directed or not.
<code>adj</code>	Adjacency matrix of an unweighted network. It will be ignored if <code>edgelist</code> is provided.
<code>target.assortcoef</code>	For directed networks, it is a list represents the predetermined value or range of assortativity coefficients. For undirected networks, it is a constant between -1 to 1. It will be ignored if <code>eta</code> is provided.
<code>control</code>	A list of parameters for controlling the rewiring process and the process for solving <code>eta</code> . <ul style="list-style-type: none"> <code>iteration</code> An integer, represents the number of rewiring iterations. Each iteration consists of <code>nattempts</code> rewiring attempts. The assortativity coefficient(s) of the network will be recorded after each iteration. <code>nattempts</code> An integer, number of rewiring attempts for each iteration. Default value equals the number of rows of <code>edgelist</code>. <code>history</code> Logical, whether the rewiring attempts should be recorded and returned. <code>eta.obj</code> A convex function of <code>eta</code> to be minimized when solving for <code>eta</code> with given <code>target.assortcoef</code>. Defaults to 0. It will be ignored if <code>eta</code> is provided. <code>cvxr_control</code> A list of parameters passed to <code>CVXR::solve()</code> for solving <code>eta</code> with given <code>target.assortcoef</code>. It will be ignored if <code>eta</code> is provided.
<code>eta</code>	An matrix represents the target network structure. If specified, <code>target.assortcoef</code> will be ignored. For directed networks, the element at row "i-j" and column "k-l" represents the proportion of directed edges linking a source node with out-degree i and in-degree j to a target node with out-degree k and in-degree l. For undirected networks, <code>eta</code> is symmetric, the summation of the elements at row "i", column "j" and row "j", column "i" represents the proportion of edges linking to a node with degree i and a node with degree j.

Details

There are two steps in this algorithm. It first solves for an appropriate `eta` using `target.assortcoef`, `eta.obj`, and `cvxr_control`, then proceeds to the rewiring process and rewire the network towards

the solved eta. If eta is given, the algorithm will skip the first step. The function only works for unweighted networks.

Each rewiring attempt samples two rows from edgelist, for example Edge1:(v_1, v_2) and Edge2:(v_3, v_4). For directed networks, if the rewiring attempt is accepted, the sampled edges are replaced as (v_1, v_4), (v_3, v_2); for undirected networks, the algorithm try to rewire the sampled edges as {v_1, v_4}, {v_3, v_2} (type 1) or {v_1, v_3}, {v_2, v_4} (type 2), each with probability 1/2.

Value

Rewired edgelist; assortativity coefficient(s) after each iteration; rewiring history (including the index of sampled edges and rewiring result) and solver results.

Examples

```
set.seed(123)
edgelist <- rpanet(1e4, control = rpa_control_scenario(
  alpha = 0.4, beta = 0.3, gamma = 0.3))$edgelist
## rewire a directed network to have predetermined assortativity coefficients
target.assortcoef <- list("outout" = -0.2, "outin" = 0.2)
ret1 <- dprewire(edgelist, directed = TRUE,
  target.assortcoef = target.assortcoef,
  control = list(iteration = 200))
plot(ret1$assortcoef$Iteration, ret1$assortcoef$"outout")
plot(ret1$assortcoef$Iteration, ret1$assortcoef$"outin")

edgelist <- rpanet(1e4, control = rpa_control_scenario(
  alpha = 0.3, beta = 0.1, gamma = 0.3, xi = 0.3),
  directed = FALSE)$edgelist
## rewire an undirected network to have predetermined assortativity coefficient
ret2 <- dprewire(edgelist, directed = FALSE, target.assortcoef = 0.3,
  control = list(iteration = 300, eta.obj = CVXR::norm2,
  history = TRUE))
plot(ret2$assortcoef$Iteration, ret2$assortcoef$Value)
```

dprewire.range

Range of assortativity coefficient.

Description

The assortativity coefficient of a given network may not achieve all the values within -1 and 1 via degree preserving rewiring. This function computes the range of assortativity coefficients that can be achieved through degree preserving rewiring. The algorithm is designed for unweighted networks.

Usage

```
dprewire.range(
  edgelist = NULL,
  directed = TRUE,
  adj = NULL,
  which.range = c("outout", "outin", "inout", "inin"),
  control = cvxr_control(),
  target.assortcoef = list(outout = NULL, outin = NULL, inout = NULL, inin = NULL)
)
```

Arguments

edgelist	A two column matrix, each row represents an edge of the network.
directed	Logical, whether the network is directed or not.
adj	Adjacency matrix of an unweighted network. It will be ignored if edgelist is provided.
which.range	The type of interested assortativity coefficient. For directed networks, it takes one of the values: "outout", "outin", "inout" and "inin". It will be ignored if the network is undirected.
control	A list of parameters passed to CVXR::solve() for solving an appropriate eta with the constraints target.assortcoef.
target.assortcoef	A list of constraints, it has the predetermined value or range imposed on assortativity coefficients other than which.range. It will be ignored if the network is undirected.

Details

The ranges are computed through convex optimization. The problems are defined and solved via the R package CVXR. For undirected networks, the function returns the range of the assortativity coefficient. For directed networks, the function computes the range of which.range while other assortativity coefficients are restricted through target.assortcoef.

Value

Range of the interested assortativity coefficient and solver results.

Examples

```
set.seed(123)
edgelist <- rpanet(5e3, control =
  rpa_control_scenario(alpha = 0.5, beta = 0.5))$edgelist
ret1 <- dprewire.range(edgelist, directed = TRUE, which.range = "outin",
  target.assortcoef = list("outout" = c(-0.3, 0.3), "inout" = 0.1))
ret1$range
```

rpanet *Generate PA networks.*

Description

Generate preferential attachment (PA) networks with linear or non-linear preference functions.

Usage

```
rpanet(
  nstep = 10^3,
  initial.network = list(edgelist = matrix(c(1, 2), nrow = 1)),
  control = list(),
  directed = TRUE,
  method = c("binary", "linear", "bagx", "bag")
)
```

Arguments

nstep	Number of steps.
initial.network	A list represents the seed network. By default, <code>initial.network</code> has one edge from node 1 to node 2 with weight 1. It consists of the following components: a two column matrix <code>edgelist</code> represents the edges; a vector <code>edgeweight</code> represents the weight of edges; an integer vector <code>nodegroup</code> represents the group of nodes. <code>nodegroup</code> is defined for directed networks, if <code>NULL</code> , all nodes from the seed network are considered from group 1.
control	A list of parameters controls the PA network generation process. Defaults to an empty list, i.e., all the control parameters are set to default. For more details about available controls, see <code>rpa_control_scenario()</code> , <code>rpa_control_newedge()</code> , <code>rpa_control_edgeweight()</code> , <code>rpa_control_preference</code> and <code>rpa_control_reciprocal()</code> . Under the default setup, in each step, a new edge of weight 1 is added from a new node A to an existing node B (alpha scenario), where B is chosen with probability proportional to its in-strength + 1.
directed	Logical, whether to generate a directed network. If <code>FALSE</code> , the edges are undirected.
method	Which method to use: <code>binary</code> , <code>linear</code> , <code>bagx</code> or <code>bag</code> . For <code>bag</code> and <code>bagx</code> methods, <code>beta.loop</code> must be <code>TRUE</code> ; default preference functions must be used and <code>sparams = c(1, 1, 0, 0, a)</code> , <code>tparams = c(0, 0, 1, 1, b)</code> , <code>param = c(1, c)</code> , where <code>a</code> , <code>b</code> and <code>c</code> are non-negative constants; reciprocal edges and sampling without replacement are not considered, i.e., option <code>rpa_control_reciprocal()</code> must be set as default, <code>snode.replace</code> , <code>tnode.replace</code> and <code>node.replace</code> must be <code>TRUE</code> . In addition, <code>nodelsit</code> method only works for unweighted networks and does not consider multiple edges, i.e., <code>rpa_control_edgeweight()</code> and <code>rpa_control_newedge()</code> must be set as default.

Value

A list with the following components: `edgelist`; `edgweight`; number of new edges at each step `newedge` (reciprocal edges are included); `node.attribute`, including node strengths, preference scores and node group (if applicable); control list `control`; edge scenario `scenario` (1~alpha, 2~beta, 3~gamma, 4~xi, 5~rho, 6~reciprocal). The edges from `initial.network` are denoted as scenario 0.

Note

The binary method implements binary search algorithm; linear represents linear search algorithm; bag method implements the algorithm from Wan et al. (2017); bagx puts all the edges into a bag, then samples edges and find the source/target node of the sampled edge.

References

- Wan P, Wang T, Davis RA, Resnick SI (2017). Fitting the Linear Preferential Attachment Model. *Electronic Journal of Statistics*, 11(2), 3738–3780.

Examples

```
# Control edge scenario and edge weight through rpa_control_scenario()
# and rpa_control_edgweight(), respectively, while keeping rpa_control_newedge(),
# rpa_control_preference() and rpa_control_reciprocal() as default.
set.seed(123)
control <- rpa_control_scenario(alpha = 0.5, beta = 0.5) +
  rpa_control_edgweight(distribution = rgamma,
    dparams = list(shape = 5, scale = 0.2), shift = 0)
ret1 <- rpanet(nstep = 1e3, control = control)

# In addition, set node groups and probability of creating reciprocal edges.
control <- control + rpa_control_reciprocal(group.prob = c(0.4, 0.6),
  recip.prob = matrix(runif(4), ncol = 2))
ret2 <- rpanet(nstep = 1e3, control = control)

# Further, set the number of new edges in each step as Poisson(2) + 1 and use
# ret2 as a seed network.
control <- control + rpa_control_newedge(distribution = rpois,
  dparams = list(lambda = 2), shift = 1)
ret3 <- rpanet(nstep = 1e3, initial.network = ret2, control = control)
```

rpa_control_edgweight

Control weight of new edges. Defined for rpanet.

Description

Control weight of new edges. Defined for rpanet.

Usage

```
rpa_control_edgweight(distribution = NA, dparams = list(), shift = 1)
```

Arguments

distribution	Distribution for drawing edge weights. Default is NA. If specified, its first argument must be the number of observations.
dparams	Additional parameters passed on to distribution. The name of parameters must be specified.
shift	A constant add to the values sampled from distribution. Default value is 1.

Value

A list of class `rpacontrol` with components `distribution`, `dparams`, and `shift` with meanings as explained under 'Arguments'.

Examples

```
# Edge weight follows Gamma(5, 0.2).
control <- rpa_control_edgweight(distribution = rgamma,
  dparams = list(shape = 5, scale = 0.2),
  shift = 0)

# Constant edge weight
control <- rpa_control_edgweight(shift = 2)
```

`rpa_control_newedge` *Control new edges in each step. Defined for rpanet.*

Description

Control new edges in each step. Defined for rpanet.

Usage

```
rpa_control_newedge(
  distribution = NA,
  dparams = list(),
  shift = 1,
  snode.replace = TRUE,
  tnode.replace = TRUE,
  node.replace = TRUE
)
```

Arguments

distribution	Distribution for drawing number of new edges. Default is NA. If specified, its first argument must be the number of observations.
dparams	Additional parameters passed on to distribution. The name of parameters must be specified.
shift	A constant add to the values sampled from distribution. Default value is 1.
snode.replace	Logical, whether the source nodes in the same step should be sampled with replacement. Defined for directed networks.
tnode.replace	Logical, whether the target nodes in the same step should be sampled with replacement. Defined for directed networks.
node.replace	Logical, whether the nodes in the same step should be sampled with replacement. Defined for undirected networks. If FALSE, self-loops will not be allowed under beta scenario.

Value

A list of class `rpacontrol` with components `distribution`, `dparams`, `shift`, `snode.replace`, `tnode.replace` and `node.replace` with meanings as explained under 'Arguments'.

Examples

```
control <- rpa_control_newedge(distribution = rpois,
  dparams = list(lambda = 2),
  shift = 1,
  node.replace = FALSE)
```

rpa_control_preference

Set preference function(s). Defined for rpanet.

Description

Set preference function(s). Defined for rpanet.

Usage

```
rpa_control_preference(
  ftype = c("default", "customized"),
  sparams = c(1, 1, 0, 0, 1),
  tparams = c(0, 0, 1, 1, 1),
  params = c(1, 1),
  spref = "outs + 1",
  tpref = "ins + 1",
  pref = "s + 1"
)
```

Arguments

f type	Preference function type. Either "default" or "customized". "customized" preference functions require "binary" or "linear" generation methods. If using default preference functions, sparams, tparams and params must be specified. If using customized preference functions, spref, tpref and pref must be specified.
sparams	A numerical vector of length 5 giving the parameters of the default source preference function. Defined for directed networks. Probability of choosing an existing node as the source node is proportional to $\text{sparams}[1] * \text{out-strength}^{\text{sparams}[2]} + \text{sparams}[3] * \text{in-strength}^{\text{sparams}[4]} + \text{sparams}[5]$.
tparams	A numerical vector of length 5 giving the parameters of the default target preference function. Defined for directed networks. Probability of choosing an existing node as the target node is proportional to $\text{tparams}[1] * \text{out-strength}^{\text{tparams}[2]} + \text{tparams}[3] * \text{in-strength}^{\text{tparams}[4]} + \text{tparams}[5]$.
params	A numerical vector of length 2 giving the parameters of the default preference function. Defined for undirected networks. Probability of choosing an existing node is proportional to $\text{strength}^{\text{params}[1]} + \text{params}[2]$.
spref	Character expression or an object of class XPtr giving the customized source preference function. Defined for directed networks. Default value is "outs + 1", i.e., node out-strength + 1. See Details and Examples for more information.
tpref	Character expression or an object of class XPtr giving the customized target preference function. Defined for directed networks. Default value is "ins + 1", i.e., node in-strength + 1.
pref	Character expression or an object of class XPtr giving the customized preference function. Defined for undirected networks. Default value is "s + 1", i.e., node strength + 1.

Details

If choosing customized preference functions, spref, tpref and pref will be used and the network generation method must be "binary" or "linear". spref (tpref) defines the source (target) preference function, it can be a character expression or an object of class XPtr.

- Character expression: it must be an one-line C++ style expression of outs (node out-strength) and ins (node-instrength). For example, "pow(outs, 2) + 1", "pow(outs, 2) + pow(ins, 2) + 1", etc. The expression will be used to define an XPtr via `RcppXPtrUtils::cppXPtr`. The XPtr will be passed to the network generation function. The expression must not have variables other than outs and ins.
- XPtr: an external pointer wrapped in an object of class XPtr defined via `RcppXPtrUtils::cppXPtr`. An example for defining an XPtr with C++ source code is included in Examples. For more information about passing function pointers, see <https://gallery.rcpp.org/articles/passing-cpp-function-pointers-rcppxptrutils/>. Please note the supplied C++ function takes two double arguments and returns a double. The first and second arguments represent node out- and in-strength, respectively.

pref is defined analogously. If using character expression, it must be a one-line C++ style expression of s (node strength). If using XPtr, the supplied C++ function takes only one double argument and returns a double.

Value

A list of class rpacontrol with components ftype, sparams, tparams, params or ftype, spref, tpref, pref with function pointers spref.pointer, tpref.pointer, pref.pointer.

Examples

```
# Set source preference as out-strength^2 + in-strength + 1,
# target preference as out-strength + in-strength^2 + 1.
# 1. use default preference functions
control1 <- rpa_control_preference(ftype = "default",
  sparams = c(1, 2, 1, 1, 1), tparams = c(1, 1, 1, 2, 1))
# 2. use character expressions
control2 <- rpa_control_preference(ftype = "customized",
  spref = "pow(outs, 2) + ins + 1", tpref = "outs + pow(ins, 2) + 1")
# 3. define XPtr's with C++ source code
spref.pointer <- RcppXPtrUtils::cppXPtr(code =
  "double spref(double outs, double ins) {return pow(outs, 2) + ins + 1;}")
tpref.pointer <- RcppXPtrUtils::cppXPtr(code =
  "double tpref(double outs, double ins) {return outs + pow(ins, 2) + 1;}")
control3 <- rpa_control_preference(ftype = "customized",
  spref = spref.pointer,
  tpref = tpref.pointer)
ret <- rpanet(1e5, control = control3)
```

rpa_control_reciprocal

Control reciprocal edges. Defined for rpanet.

Description

Control reciprocal edges. Defined for rpanet.

Usage

```
rpa_control_reciprocal(
  group.prob = NA,
  recip.prob = NA,
  selfloop.recip = FALSE
)
```

Arguments

group.prob A vector of probability weights for sampling the group of new nodes. Defined for directed networks. Groups are from 1 to length(group.prob). Its length must equal to the number of rows of recip.prob.

`recip.prob` A square matrix giving the probability of adding a reciprocal edge after a new edge is introduced. Defined for directed networks. Its element p_{ij} represents the probability of adding a reciprocal edge from node A, which belongs to group i, to node B, which belongs to group j, immediately after a directed edge from B to A is added.

`selfloop.recip` Logical, whether reciprocal edge of self-loops are allowed.

Value

A list of class `rpacontrol` with components `group.prob`, `recip.prob`, and `selfloop.recip` with meanings as explained under 'Arguments'.

Examples

```
control <- rpa_control_reciprocal(group.prob = c(0.4, 0.6),
  recip.prob = matrix(runif(4), ncol = 2))
```

`rpa_control_scenario` *Control edge scenarios. Defined for rpanet.*

Description

Control edge scenarios. Defined for rpanet.

Usage

```
rpa_control_scenario(
  alpha = 1,
  beta = 0,
  gamma = 0,
  xi = 0,
  rho = 0,
  beta.loop = TRUE,
  source.first = TRUE
)
```

Arguments

`alpha` Probability of adding an edge from a new node to an existing node.

`beta` Probability of adding an edge between existing nodes.

`gamma` Probability of adding an edge from an existing node to a new node.

`xi` Probability of adding an edge between two new nodes.

`rho` Probability of adding a new node with a self-loop.

`beta.loop` Logical, whether self-loops are allowed under beta scenario. Default value is TRUE.

`source.first` Logical. Defined for beta scenario edges of directed networks. If TRUE, the source node of a new edge is sampled from existing nodes before the target node is sampled; if FALSE, the target node is sampled from existing nodes before the source node is sampled. Default value is TRUE.

Value

A list of class `rpacontrol` with components `alpha`, `beta`, `gamma`, `xi`, `rho`, `beta.loop` and `source.first` with meanings as explained under 'Arguments'.

Examples

```
control <- rpa_control_scenario(alpha = 0.5, beta = 0.5, beta.loop = FALSE)
```

wdnet

wdnet: Weighted and Directed Networks

Description

This package provides functions to conduct network analysis

- Assortativity, centrality, clustering coefficient for weighted and directed networks
- Rewire an unweighted network with given assortativity coefficient(s)
- Preferential attachment (PA) network generation

Details

The development version of this package is available on Gitlab (<https://gitlab.com/wdnetwork/wdnet>).

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