

# Package: causalnet (via r-universe)

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**Title** Directed Causal Network Enumeration and Simulation

**Version** 0.1.0

**Description** Enumerate orientation-consistent directed networks from an undirected or partially directed skeleton, detect feedback loops, summarize topology, and simulate node dynamics via stochastic differential equations.

**License** GPL-3

**URL** <https://github.com/KyuriP/causalnet>

**BugReports** <https://github.com/KyuriP/causalnet/issues>

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detect\_feedback\_loops *Detect Unique Feedback Loops in a Directed Network*

---

### Description

Detects all directed cycles (including 2-node loops) and returns each as a unique set of nodes (ignores order and entry point).

### Usage

```
detect_feedback_loops(adj_matrix, include_self_loops = FALSE, use_names = TRUE)
```

### Arguments

adj\_matrix      Square directed adjacency (non-zero = edge).  
include\_self\_loops  
                  Logical; include 1-node self-loops (default FALSE).  
use\_names        Logical; return node names if available (default TRUE).

### Value

List of unique loops, each as a sorted vector (names if available, else indices).

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generate\_directed\_networks  
*Generate Directed Networks Consistent with Constraints*

---

### Description

Enumerate all directed adjacency matrices that are consistent with a given undirected skeleton and optional direction constraints. Enumeration can optionally include bidirected edges and display a simple progress bar.

### Usage

```
generate_directed_networks(  
  adj_matrix,  
  allow_bidirectional = TRUE,  
  fixed_edges = NULL,  
  max_networks = Inf,  
  show_progress = interactive()  
)
```

**Arguments**

adj_matrix	Symmetric binary (0/1) adjacency matrix giving the undirected skeleton. Only pairs with <code>adj_matrix[i, j] = 1</code> are considered for orientation; all other pairs remain 0.
allow_bidirectional	Logical. If TRUE, bidirected edges ( $i \leftrightarrow j$ ) are allowed during enumeration. Default: TRUE.
fixed_edges	Numeric matrix the same size as <code>adj_matrix</code> that encodes per-edge constraints (interpreted on the directed $i \rightarrow j$ entry): <ul style="list-style-type: none"> <li>• 1: force <math>i \rightarrow j</math></li> <li>• 2: force <math>i \leftrightarrow j</math> (both <math>i \rightarrow j</math> and <math>j \rightarrow i</math>)</li> <li>• 0 or NA: unconstrained</li> </ul> To exclude an edge entirely, set <code>adj_matrix[i, j] = adj_matrix[j, i] = 0</code> in the skeleton; <code>fixed_edges</code> does not remove edges, it only constrains orientations of skeleton edges.
max_networks	Integer. Maximum number of networks to return. Use to cap output size when constraints are loose and the search space is large. Default: Inf.
show_progress	Logical. Show a text progress bar during enumeration. Default: <code>interactive()</code> .

**Details**

If the skeleton has  $m$  undirected edges, the number of orientation-consistent digraphs is at most  $2^m$  when `allow_bidirectional = FALSE` and  $3^m$  when TRUE (before applying constraints). Consider setting `max_networks` for exploratory use.

**Value**

A list of unique directed 0/1 adjacency matrices, each with the same dimensions and dimnames as `adj_matrix`.

**See Also**

[detect\\_feedback\\_loops](#), [summarize\\_network\\_metrics](#)

**Examples**

```
skel <- matrix(0, 3, 3); skel[upper.tri(skel)] <- 1; skel <- skel + t(skel)
colnames(skel) <- rownames(skel) <- paste0("X", 1:3)
out <- generate_directed_networks(skel, allow_bidirectional = TRUE)
length(out)

# Force X1 -> X2 and X2 <-> X3:
F <- matrix(NA_real_, 3, 3, dimnames = dimnames(skel))
F["X1", "X2"] <- 1
F["X2", "X3"] <- 2
out2 <- generate_directed_networks(skel, fixed_edges = F)
length(out2)
```

---

get\_sample\_parameters *Generate Sample Parameters for Node Dynamics*

---

### Description

Returns a list of simulation parameters (domain-agnostic: nodes can be any variables). If a parameter vector is not supplied, values are sampled i.i.d. from a uniform range.

### Usage

```
get_sample_parameters(
  n_nodes,
  beta_range = c(-1.5, -1),
  alpha_range = c(0.05, 0.3),
  delta_range = c(1, 5),
  sigma_range = c(0.01, 0.1),
  beta = NULL,
  alpha_self = NULL,
  delta = NULL,
  sigma = NULL,
  nodes = NULL
)
```

### Arguments

n_nodes	Integer number of nodes.
beta_range	Length-2 numeric range for baseline/exogenous drive (used if beta is NULL).
alpha_range	Length-2 numeric range for self-activation (used if alpha_self is NULL).
delta_range	Length-2 numeric range for nonlinear amplification (used if delta is NULL).
sigma_range	Length-2 numeric range for noise SD (used if sigma is NULL).
beta, alpha_self, delta, sigma	Optional fixed numeric vectors. If length-1, recycled to n_nodes.
nodes	Optional character vector of node names (length n_nodes) used to name outputs.

### Value

A named list with elements beta, alpha\_self, delta, sigma (each length n\_nodes).

**Description**

Visualizes node/variable dynamics over time using ggplot2, with optional stress intervals and customizable styling.

**Usage**

```
plot_dynamics(
  S,
  stress_windows = NULL,
  title = "Dynamics",
  colors = NULL,
  legend_labels = NULL,
  show_lines = FALSE,
  line_width = 0.8,
  line_alpha = 1,
  base_size = 14,
  label_stress = TRUE,
  stress_label = "Stress Period",
  stress_fill = "gray60",
  stress_alpha = 0.2,
  stress_line_color = "gray40",
  y_label = "Level",
  legend_position = "right",
  y_limits = NULL
)
```

**Arguments**

<code>S</code>	Matrix (time x variables) of simulated states. If <code>attr(S,"time")</code> exists, it is used for the x-axis (continuous time). Otherwise the x-axis is step index <code>1:nrow(S)</code> .
<code>stress_windows</code>	Optional list of numeric <code>c(start, end)</code> intervals, or a 2-column matrix/data.frame with <code>start, end</code> . Units must match the x-axis (i.e., the "Time" used for plotting).
<code>title</code>	Plot title.
<code>colors</code>	Optional vector of line colors (length = #variables).
<code>legend_labels</code>	Optional vector of legend labels (length = #variables).
<code>show_lines</code>	If TRUE, draw dashed vertical lines instead of shaded rectangles.
<code>line_width</code>	Line width for trajectories.
<code>line_alpha</code>	Line transparency (0–1).
<code>base_size</code>	Base font size for theme.

label_stress	If TRUE and using shading, label each stress window.
stress_label	Text label (length 1 or length = #windows).
stress_fill	Fill color for shaded windows.
stress_alpha	Alpha for shaded windows.
stress_line_color	Color for dashed lines (if show_lines = TRUE).
y_label	Y-axis label.
legend_position	Legend position (e.g., "right", "bottom", "none").
y_limits	Optional numeric length-2 vector for y-axis limits.

**Value**

A ggplot object.

---

plot\_network\_metrics *Generate ggplot objects summarizing network metrics*

---

**Description**

Produces a list of ggplot2 objects visualizing summary metrics across a list of directed networks.

**Usage**

```
plot_network_metrics(
  summary_df,
  n_bins = 6,
  fill_colors = c("skyblue", "darkgreen", "orange", "lightcoral"),
  base_size = 14,
  return_grid = TRUE,
  p2_style = c("auto", "jitter", "point", "beeswarm"),
  auto_threshold = 200
)
```

**Arguments**

summary_df	Data frame from summarize_network_metrics().
n_bins	Number of histogram bins (default = 6).
fill_colors	Optional vector of 4 fill colors.
base_size	Base font size for plots (default = 14).
return_grid	If TRUE, returns cowplot grid; otherwise, returns list of plots.
p2_style	How to plot panel (b) (sigma_total by loop count). One of "auto" (default), "jitter", "point", or "beeswarm". If "auto", uses beeswarm when nrow(summary_df) < auto_threshold, otherwise jitter.
auto_threshold	Threshold for switching when p2_style = "auto" (default = 200).

**Details**

The "beeswarm" option uses `ggbeeswarm::geom_beeswarm()` if the `ggbeeswarm` package is installed. If not installed, the function falls back to jitter and emits a warning.

**Value**

A cowplot grid or a named list of `ggplot2` objects.

**Examples**

```
## Not run:
summary_df <- summarize_network_metrics(nets)

# automatic switching
plot_network_metrics(summary_df, p2_style = "auto", auto_threshold = 200)

# force beeswarm
plot_network_metrics(summary_df, p2_style = "beeswarm")

# force jitter
plot_network_metrics(summary_df, p2_style = "jitter")

## End(Not run)
```

---

simulate_dynamics	<i>Simulate network state dynamics via SDEs (nonlinear, linear, or custom)</i>
-------------------	--

---

**Description**

Simulates the evolution of node states in a directed network using an Euler–Maruyama discretization of stochastic differential equations (SDEs). Choose the built-in nonlinear model, a linear alternative, or provide a custom update function.

**Usage**

```
simulate_dynamics(
  adj_matrix,
  params,
  t_max = 100,
  dt = 0.1,
  S0 = NULL,
  model_type = "nonlinear",
  model_fn = NULL,
  stress_event = NULL,
  boundary = c("auto", "reflect", "clamp", "none"),
  clamp = NULL
)
```

**Arguments**

adj_matrix	Numeric matrix (square; directed adjacency). Interpreted as $i \rightarrow j$ .
params	Named list of model parameters. For model_type = "nonlinear", requires vectors (length = n nodes): <ul style="list-style-type: none"> <li>• beta: baseline/exogenous drive per node.</li> <li>• alpha_self: self-activation per node.</li> <li>• delta: nonlinear amplification of incoming effects.</li> <li>• sigma: noise SD per node.</li> </ul> For model_type = "linear", requires beta, alpha_self, sigma. For a custom model, include whatever your model_fn expects.
t_max	Total simulated time (must be > 0).
dt	Time step (must be > 0). The output has $\text{floor}(t\_max/dt) + 1$ rows.
S0	Optional numeric vector of initial states (length = n). Defaults to 0.01.
model_type	One of "nonlinear" (default), "linear", or NULL when using a custom model_fn.
model_fn	Optional function with signature <code>function(current, interaction, dt, ...)</code> returning a numeric vector of increments dS. Additional args are taken from params.
stress_event	Optional function <code>f(time, state) -&gt; numeric(n)</code> that returns an exogenous input vector added each step (e.g., shocks/perturbations).
boundary	One of "auto", "reflect", "clamp", "none". <ul style="list-style-type: none"> <li>• "reflect": mirror overshoot back into <code>[clamp[1], clamp[2]]</code>.</li> <li>• "clamp": hard-box to <code>[clamp[1], clamp[2]]</code>.</li> <li>• "none": no bounding.</li> <li>• "auto": pick a sensible default based on the model and clamp: nonlinear -&gt; boundary = "reflect" (and if clamp is NULL, use <code>c(0, 1)</code>); linear/custom -&gt; boundary = "none" unless a clamp range is supplied, in which case use "clamp".</li> </ul>
clamp	Either NULL (no numeric range) or a length-2 numeric vector <code>c(min, max)</code> used by "reflect" or "clamp" to keep states within bounds.

**Details**

**Direction convention.** By default `adj[i, j] = 1` encodes a directed edge  $i \rightarrow j$ . Under this convention, the *incoming input* to node  $j$  is the dot product of column  $j$  with the current state; in vector form `t(adj) %*% state`. If your internal convention differs, transpose accordingly.

Integration uses Euler–Maruyama. The per-step diffusion term is added as  $\sigma\sqrt{dt}Z$  with  $Z \sim \mathcal{N}(0, I)$  (component-wise), i.e., `sigma * sqrt(dt) * rnorm(n)`.

**Value**

Numeric matrix of states over time (rows = time steps, cols = nodes). The time vector is attached as `attr(result, "time")`.

### Boundary handling

- **Reflecting** avoids “sticky” edges by bouncing trajectories back inside the range, which is useful for bounded variables on  $[0, 1]$ .
- **Clamping** is numerically simple but can create artificial absorbing states at the limits.
- For smoothly bounded dynamics, consider modeling on an unbounded latent scale and applying a link (e.g., logistic) instead of hard post-step bounds.

### Examples

```
set.seed(1)
net <- matrix(c(0,1,0,0,
               0,0,1,0,
               0,0,0,1,
               1,0,0,0), 4, byrow = TRUE)

# Linear model, automatic boundary selection ("none" because no clamp supplied)
p_lin <- list(beta = rep(0.8, 4), alpha_self = rep(0.2, 4), sigma = rep(0.05, 4))
S1 <- simulate_dynamics(net, p_lin, model_type = "linear", boundary = "auto", t_max = 5, dt = 0.01)

# Linear model with a finite box -> "auto" switches to clamp on [0, 5]
S2 <- simulate_dynamics(net, p_lin, model_type = "linear",
                       boundary = "auto", clamp = c(0, 5), t_max = 5, dt = 0.01)

# Nonlinear model -> "auto" uses reflecting boundaries on [0,1]
p_nl <- list(beta = rep(0.2, 4), alpha_self = rep(0.2, 4),
            delta = rep(0.5, 4), sigma = rep(0.05, 4))
S3 <- simulate_dynamics(
  net, p_nl, model_type = "nonlinear",
  boundary = "auto", t_max = 5, dt = 0.01
)
```

---

```
summarize_network_metrics
```

*Summarize Directed Network List*

---

### Description

Compute feedback loop and topology metrics for a list of directed networks.

### Usage

```
summarize_network_metrics(net_list)
```

### Arguments

`net_list`            A list of directed adjacency matrices (can be from any source).

**Value**

A data frame with one row per network and the following columns:

- net\_id
- n\_nodes
- n\_edges
- num\_loops (number of unique feedback loops)
- sigma\_total (sum of SDs of in/out degrees)
- node\_overlap\_score
- avg\_loop\_size

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