

Package ‘simplegraph’

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Title Simple Graph Data Types and Basic Algorithms

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Description Simple classic graph algorithms for simple graph classes.

Graphs may possess vertex and edge attributes. 'simplegraph' has no dependencies and it is written entirely in R, so it is easy to install.

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URL <https://github.com/gaborcsardi/simplegraph>

BugReports <https://github.com/gaborcsardi/simplegraph/issues>

Suggests testthat

Imports methods, utils

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adjacent_vertices *Adjacent vertices for all vertices in a graph*

Description

A vertex is adjacent if it is either a successor, or a predecessor.

Usage

`adjacent_vertices(graph)`

Arguments

`graph` The graph.

Value

A named list of character vectors, the adjacent vertices for each vertex.

See Also

Other simple queries: `edges()`, `order()`, `vertices()`

Examples

```
G <- graph(list(A = c("B", "C"), B = "C", C = "A"))
adjacent_vertices(G)
```

bfs *Breadth-first search of a graph*

Description

Breadth-first search of a graph

Usage

```
bfs(graph, from = vertex_ids(graph))
```

Arguments

graph	Input graph.
from	Character vector, which vertices to start the search from. By default all vertices are attempted.

Value

Character vector of the names of the visited vertices, in the order of their visit.

Examples

```
funcs <- graph(list(
  drop_internal = character(0),
  get_deps = c("get_description", "parse_deps",
    "%||%", "drop_internal"),
  get_description = "pkg_from_filename",
  parse_deps = "str_trim",
  cran_file = c("get_pkg_type", "r_minor_version", "cran_file"),
  download_urls = c("split_pkg_names_versions", "cran_file"),
  filename_from_url = character(0),
  get_pkg_type = character(0),
  pkg_download = c("dir_exists", "download_urls",
    "filename_from_url", "try_download"),
  r_minor_version = character(0),
  try_download = character(0),
  drop_missing_deps = character(0),
  install_order = character(0),
  restore = c("pkg_download", "drop_missing_deps",
    "install_order", "get_deps"),
  snap = character(0),
  `%||%` = character(0),
  data_frame = character(0),
  dir_exists = character(0),
  pkg_from_filename = character(0),
  split_pkg_names_versions = "data_frame",
  str_trim = character(0)
))
bfs(funcs)
```

degree

*Degree of vertices***Description**

Degree of vertices

Usage

```
degree(graph, mode = c("out", "in", "total", "all"))
```

Arguments

graph

Input graph.

mode

Whether to calculate out-degree, in-degree, or the total degree.

Value

Named numeric vector of degrees.

Examples

```
G <- graph(list(A = c("B", "C"), B = "C", C = "A"))
degree(G, mode = "out")
degree(G, mode = "in")
degree(G, mode = "total")
```

edges

*Edges of a graph***Description**

Edges of a graph

Usage

```
edges(graph)
```

Arguments

graph

The graph

Value

Data frame of edge data and metadata. The tail and head vertices are in the first two columns. The rest of the columns are metadata.

See Also

Other simple queries: [adjacent_vertices\(\)](#), [order\(\)](#), [vertices\(\)](#)

Examples

```
bridges <- graph(list(
  "Altstadt-Loebenicht" = c(
    "Kneiphof",
    "Kneiphof",
    "Lomse"
  ),
  "Kneiphof" = c(
    "Altstadt-Loebenicht",
    "Altstadt-Loebenicht",
    "Vorstadt-Haberberg",
    "Vorstadt-Haberberg",
    "Lomse"
  ),
  "Vorstadt-Haberberg" = c(
    "Kneiphof",
    "Kneiphof",
    "Lomse"
  ),
  "Lomse" = c(
    "Altstadt-Loebenicht",
    "Kneiphof",
    "Vorstadt-Haberberg"
  )
))
edges(bridges)
```

graph*Create a graph*

Description

Graphs can be specified as adjacency lists or (two) data frames.

Usage

```
graph(x, ...)
```

Arguments

- | | |
|------------------|--|
| <code>x</code> | A data frame, or a named list of character vectors. See details below. |
| <code>...</code> | Additional arguments, see details below. |

Details

If the first argument is a data frame, then it is interpreted as vertex data, and a second data frame must be supplied as edge data. The first column of the vertex data must contain (character) vertex ids. The first two columns of the edge data frame must contain the directed edges of the graph, in the order of tail and head, as characters referring to the nodes ids. Other columns are kept as metadata.

If the first argument is not a data frame, but a list, then it is interpreted as an adjacency list. It must be named, and the names will be used as vertex ids. Each list element must be a character vector containing the successors of each vertex.

Value

A graph object.

Examples

```
funcs <- graph(list(
  drop_internal = character(0),
  get_deps = c("get_description", "parse_deps",
    "%||%", "drop_internal"),
  get_description = "pkg_from_filename",
  parse_deps = "str_trim",
  cran_file = c("get_pkg_type", "r_minor_version", "cran_file"),
  download_urls = c("split_pkg_names_versions", "cran_file"),
  filename_from_url = character(0),
  get_pkg_type = character(0),
  pkg_download = c("dir_exists", "download_urls",
    "filename_from_url", "try_download"),
  r_minor_version = character(0),
  try_download = character(0),
  drop_missing_deps = character(0),
  install_order = character(0),
  restore = c("pkg_download", "drop_missing_deps",
    "install_order", "get_deps"),
  snap = character(0),
  `%||%` = character(0),
  data_frame = character(0),
  dir_exists = character(0),
  pkg_from_filename = character(0),
  split_pkg_names_versions = "data_frame",
  str_trim = character(0)
))
funcs

vertices <- data.frame(
  stringsAsFactors = FALSE,
  name = c("Tom Hanks", "Cate Blanchett", "Matt Damon", "Kate Winslet",
    "Saving Private Ryan", "Contagion", "The Talented Mr. Ripley"),
  what = c("actor", "actor", "actor", "actor", "movie", "movie", "movie"),
  born = c("1956-07-09", "1966-05-26", "1970-10-08", "1975-10-05",
    NA, NA, NA),
```

```
gender = c("M", "F", "M", "F", NA, NA, NA),
year = c(NA, NA, NA, NA, 1998, 2011, 1999)
)

edges <- data.frame(
  stringsAsFactors = FALSE,
  actor = c("Tom Hanks", "Cate Blanchett", "Matt Damon", "Matt Damon",
            "Kate Winslet"),
  movie = c("Saving Private Ryan", "The Talented Mr. Ripley",
            "Saving Private Ryan", "The Talented Mr. Ripley", "Contagion")
)
actors <- graph(vertices, edges)
actors
```

incident_edges*Incident edges*

Description

Incident edges

Usage

```
incident_edges(graph, mode = c("out", "in", "all", "total"))
```

Arguments

graph	Input graph.
mode	Whether to use out edges, in edges or all edges.

Value

A list of data frames, each a set of edges.

Examples

```
G <- graph(list(A = c("B", "C"), B = "C", C = "A"))
incident_edges(G, mode = "out")
incident_edges(G, mode = "in")
incident_edges(G, mode = "all")
```

<code>is_loopy</code>	<i>Is this a loopy graph?</i>
-----------------------	-------------------------------

Description

A loopy graph has at least one loop edge: an edge from a vertex to itself.

Usage

```
is_loopy(graph)
```

Arguments

<code>graph</code>	The input graph.
--------------------	------------------

Value

Logical scalar.

See Also

Other multigraphs: [is_multigraph\(\)](#), [is_simple\(\)](#), [remove_loops\(\)](#), [remove_multiple\(\)](#), [simplify\(\)](#)

Examples

```
G <- graph(list(A = c("A", "B", "B"), B = c("A", "C"), C = "A"))
is_loopy(G)

G2 <- simplify(G)
is_loopy(G2)
```

<code>is_multigraph</code>	<i>Is this a multigraph?</i>
----------------------------	------------------------------

Description

A multigraph has at least one pair or multiple edges, edges connecting the same (ordered) pair of vertices.

Usage

```
is_multigraph(graph)
```

Arguments

<code>graph</code>	Input graph.
--------------------	--------------

Value

Logical scalar.

See Also

Other multigraphs: [is_loopy\(\)](#), [is_simple\(\)](#), [remove_loops\(\)](#), [remove_multiple\(\)](#), [simplify\(\)](#)

Examples

```
G <- graph(list(A = c("A", "B", "B"), B = c("A", "C"), C = "A"))
is_multigraph(G)

G2 <- simplify(G)
is_multigraph(G2)
```

is_simple

Is this a simple graph?

Description

A simple graph contains no loop and multiple edges.

Usage

```
is_simple(graph)
```

Arguments

graph The input graph.

Value

Logical scalar.

See Also

Other multigraphs: [is_loopy\(\)](#), [is_multigraph\(\)](#), [remove_loops\(\)](#), [remove_multiple\(\)](#), [simplify\(\)](#)

Examples

```
G <- graph(list(A = c("A", "B", "B"), B = c("A", "C"), C = "A"))
is_simple(G)

G2 <- simplify(G)
is_simple(G2)
```

<code>is_weighted</code>	<i>Is the graph weighted?</i>
--------------------------	-------------------------------

Description

Is the graph weighted?

Usage

```
is_weighted(graph)
```

Arguments

<code>graph</code>	The graph.
--------------------	------------

Examples

```
G <- graph(
  data.frame(
    stringsAsFactors = FALSE,
    id = c("a", "b", "c", "d")
  ),
  data.frame(
    stringsAsFactors = FALSE,
    from = c("a", "a", "b", "b", "c"),
    to = c("b", "d", "d", "c", "a"),
    weight = c( 1 , 2 , 1 , 3 , 2 )
  )
)
is_weighted(G)

G2 <- graph(
  data.frame(
    stringsAsFactors = FALSE,
    id = c("a", "b", "c", "d")
  ),
  data.frame(
    stringsAsFactors = FALSE,
    from = c("a", "a", "b", "b", "c"),
    to = c("b", "d", "d", "c", "a")
  )
)
is_weighted(G2)
```

order

Order of a graph

Description

The order of the graph is the number of vertices.

Usage

```
order(graph)
```

Arguments

graph The graph.

Value

Numeric scalar, the number of vertices.

See Also

Other simple queries: [adjacent_vertices\(\)](#), [edges\(\)](#), [vertices\(\)](#)

Examples

```
G <- graph(list(A = c("B", "C"), B = "C", C = "A"))
order(G)
```

predecessors

Predecessors and successors

Description

Predecessors and successors

Usage

```
predecessors(graph)
```

```
successors(graph)
```

Arguments

graph Input graph

Value

Named list of character vectors, the predecessors or the successors of each vertex.

Examples

```
G <- graph(list(A = c("B", "C"), B = "C", C = "A"))
predecessors(G)
successors(G)
```

remove_loops

Remove loop edges from a graph

Description

Remove loop edges from a graph

Usage

```
remove_loops(graph)
```

Arguments

graph	Input graph
-------	-------------

Value

Graph, with loop edges removed.

See Also

Other multigraphs: [is_loopy\(\)](#), [is_multigraph\(\)](#), [is_simple\(\)](#), [remove_multiple\(\)](#), [simplify\(\)](#)

Examples

```
G <- graph(list(A = c("A", "B", "B"), B = c("A", "C"), C = "A"))
is_loopy(G)
is_loopy(remove_loops(G))
```

remove_multiple	<i>Remove multiple edges from a graph</i>
-----------------	---

Description

Remove multiple edges from a graph

Usage

```
remove_multiple(graph)
```

Arguments

graph	Input graph.
-------	--------------

Value

Graph, without the multiple edges. (More precisely, from each set of multiple edges, only one, the first one, is kept.)

See Also

Other multigraphs: [is_loopy\(\)](#), [is_multigraph\(\)](#), [is_simple\(\)](#), [remove_loops\(\)](#), [simplify\(\)](#)

Examples

```
G <- graph(list(A = c("A", "B", "B"), B = c("A", "C"), C = "A"))
is_multigraph(G)
is_multigraph(remove_multiple(G))
```

sanitize	<i>Check the validity of a graph data structure</i>
----------	---

Description

This is mainly for internal checks, but occasionally it might be useful externally.

Usage

```
sanitize(x, ...)
```

Arguments

x	Graph.
...	Extra arguments are currently ignored.

Examples

```
G <- graph(list(A = c("B", "C"), B = "C", C = "A"))
sanitize(G)

G <- c(G, list("this is not good" = c(1, 2, 3)))
try(sanitize(G))
```

simplegraph

Simple Graph Data Types and Basic Algorithms

Description

Simple classic graph algorithms for simple graph classes. Graphs may possess vertex and edge attributes. 'simplegraph' has no dependencies and it is writing entirely in R, so it is easy to install.

See Also

Useful links:

- <https://github.com/gaborcsardi/simplegraph>
- Report bugs at <https://github.com/gaborcsardi/simplegraph/issues>

simplify

Remove multiple and loop edges from a graph

Description

Remove multiple and loop edges from a graph

Usage

```
simplify(graph)
```

Arguments

graph	Input graph.
-------	--------------

Value

Another graph, with the multiple and loop edges removed.

See Also

Other multigraphs: [is_loopy\(\)](#), [is_multigraph\(\)](#), [is_simple\(\)](#), [remove_loops\(\)](#), [remove_multiple\(\)](#)

Examples

```
G <- graph(list(A = c("A", "B", "B"), B = c("A", "C"), C = "A"))
is_simple(G)

G2 <- simplify(G)
is_simple(G2)
```

size

*The size of the graph is the number of edges***Description**

The size of the graph is the number of edges

Usage

```
size(graph)
```

Arguments

graph	The graph.
-------	------------

Value

Numeric scalar, the number of edges.

Examples

```
G <- graph(list(A = c("B", "C"), B = "C", C = "A"))
size(G)
```

strength

*Vertex strength: sum of weights of incident edges***Description**

This is also called weighed degree.

Usage

```
strength(graph, mode = c("out", "in", "total", "all"))
```

Arguments

graph	Input graph.
-------	--------------

mode	Whether to consider incoming (in), outgoing (out) or all (total) edges.
------	---

Details

For non-weighted graphs, the degree is returned as a fallback.

Value

Named numeric vector.

Examples

```
G <- graph(
  data.frame(
    stringsAsFactors = FALSE,
    id = c("a", "b", "c", "d")
  ),
  data.frame(
    stringsAsFactors = FALSE,
    from = c("a", "a", "b", "b", "c"),
    to = c("b", "d", "d", "c", "a"),
    weight = c( 1 , 2 , 1 , 3 , 2 )
  )
)
strength(G)

G2 <- graph(
  data.frame(
    stringsAsFactors = FALSE,
    id = c("a", "b", "c", "d")
  ),
  data.frame(
    stringsAsFactors = FALSE,
    from = c("a", "a", "b", "b", "c"),
    to = c("b", "d", "d", "c", "a")
  )
)
strength(G2)
```

topological_sort *Topological sorting of a graph*

Description

Topological sorting of a graph

Usage

```
topological_sort(graph)
```

Arguments

graph	Input graph.
-------	--------------

Value

Character vector of vertex ids, in topological order.

Examples

```
funcs <- graph(list(
  drop_internal = character(0),
  get_deps = c("get_description", "parse_deps",
    "%||%", "drop_internal"),
  get_description = "pkg_from_filename",
  parse_deps = "str_trim",
  cran_file = c("get_pkg_type", "r_minor_version", "cran_file"),
  download_urls = c("split_pkg_names_versions", "cran_file"),
  filename_from_url = character(0),
  get_pkg_type = character(0),
  pkg_download = c("dir_exists", "download_urls",
    "filename_from_url", "try_download"),
  r_minor_version = character(0),
  try_download = character(0),
  drop_missing_deps = character(0),
  install_order = character(0),
  restore = c("pkg_download", "drop_missing_deps",
    "install_order", "get_deps"),
  snap = character(0),
  `%||%` = character(0),
  data_frame = character(0),
  dir_exists = character(0),
  pkg_from_filename = character(0),
  split_pkg_names_versions = "data_frame",
  str_trim = character(0)
))
topological_sort(remove_loops(funcs))
```

transpose

*Transpose a graph***Description**

The transposed graph have the same vertices, and the same number of edges, but all edge directions are opposite compared to the original graph.

Usage

```
transpose(graph)
```

Arguments

graph	Input graph
-------	-------------

Value

Transposed graph.

Examples

```
funcs <- graph(list(
  drop_internal = character(0),
  get_deps = c("get_description", "parse_deps",
    "%||%", "drop_internal"),
  get_description = "pkg_from_filename",
  parse_deps = "str_trim",
  cran_file = c("get_pkg_type", "r_minor_version", "cran_file"),
  download_urls = c("split_pkg_names_versions", "cran_file"),
  filename_from_url = character(0),
  get_pkg_type = character(0),
  pkg_download = c("dir_exists", "download_urls",
    "filename_from_url", "try_download"),
  r_minor_version = character(0),
  try_download = character(0),
  drop_missing_deps = character(0),
  install_order = character(0),
  restore = c("pkg_download", "drop_missing_deps",
    "install_order", "get_deps"),
  snap = character(0),
  `%||%` = character(0),
  data_frame = character(0),
  dir_exists = character(0),
  pkg_from_filename = character(0),
  split_pkg_names_versions = "data_frame",
  str_trim = character(0)
))
edges(transpose(funcs))
```

vertex_ids*Vertex ids of a graph***Description**

Vertex ids of a graph

Usage

```
vertex_ids(graph)
```

Arguments

graph	The graph.
--------------	------------

Value

Character vector of vertex ids.

Examples

```
G <- graph(list(A = c("B", "C"), B = "C", C = "A"))
vertex_ids(G)
```

vertices

Vertices of a graph, with metadata

Description

Vertices of a graph, with metadata

Usage

```
vertices(graph)
```

Arguments

graph The graph.

Value

Character vector of vertex names.

See Also

Other simple queries: [adjacent_vertices\(\)](#), [edges\(\)](#), [order\(\)](#)

Examples

```
bridges <- graph(list(
  "Altstadt-Loebenicht" = c(
    "Kneiphof",
    "Kneiphof",
    "Lomse"
  ),
  "Kneiphof" = c(
    "Altstadt-Loebenicht",
    "Altstadt-Loebenicht",
    "Vorstadt-Haberberg",
    "Vorstadt-Haberberg",
    "Lomse"
  ),
  "Vorstadt-Haberberg" = c(
    "Kneiphof",
    "Kneiphof",
    "Kneiphof",
    "Kneiphof"
  )
))
```

```
"Lomse"
),
"Lomse" = c(
  "Altstadt-Loebenicht",
  "Kneiphof",
  "Vorstadt-Haberberg"
)
))
vertices(bridges)
```

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