## Package 'hyperdraw'

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**Version** 1.57.0 **Depends** R (>= 2.9.0) Imports methods, grid, graph, hypergraph, Rgraphviz, stats4 SystemRequirements graphviz Title Visualizing Hypergaphs Author Paul Murrell Maintainer Paul Murrell <p.murrell@auckland.ac.nz> Description Functions for visualizing hypergraphs. License GPL (>= 2) Collate AllClasses.R affine.R draw.R graphBPH.R grid.R hypergraph.R legacy.R node.R RagraphBPH.R biocViews Visualization, GraphAndNetwork git\_url https://git.bioconductor.org/packages/hyperdraw git\_branch devel git\_last\_commit ea89127 git\_last\_commit\_date 2024-04-30 Repository Bioconductor 3.20 Date/Publication 2024-05-24

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graphBPH

#### Description

A convenience constructor for graphBPH-class objects. This is a generic function.

#### Usage

```
graphBPH(graph, edgeNodePattern, ...)
```

## Arguments

graphSome form of graph that is to be converted into a graphBPH object.edgeNodePatternA regular expression used to distinguish between normal nodes and edge nodes.

... Potential arguments to other methods.

## Value

An object of class graphBPH-class

### Methods

- graphBPH signature(graph = "graphNEL", edgeNodePattern = "character"): create a graphBPH
   object from a (directed) graphNEL object.
- graphBPH signature(graph = "Hypergraph", edgeNodePattern = "missing"): create a graphBPH
   object from a Hypergraph object (where all Hyperedges are DirectedHyperedges).

#### Author(s)

Paul Murrell

#### References

Falcon, S. and Gentleman, R. hypergraph: A package providing hypergraph data structures.

Gentleman, R. and Whalen, E. and Huber, W. and Falcon, S. graph: A package to handle graph data structures.

#### See Also

graphBPH-class

graphBPH-class Class "graphBPH"

#### Description

A bipartite representation of a hypergraph. The purpose of this class is to support visualization of the hypergraph; it is not intended for analysis or manipulation of the hypergraph.

#### **Objects from the Class**

Objects can be created by calls of the form new("graphBPH", graph, edgeNodePattern, ...). There is also a convenience function graphBPH().

A graphBPH object consists of a graphNEL object, which must obey some strict rules:

- nodes in the graph are divided into two sets: normal nodes and edge-nodes,
- all edges in the graph must connect a normal node to an edge node,
- the graph must be a directed graph.

The edgeNodePattern is a regular expression that is used to define the set of edge-nodes.

#### Slots

graph: Object of class graphNEL. This graph must obey the constraints described above.

edgeNodePattern: Object of class character. The regular expression used to define edge-nodes.

nodes: Object of class character. Records which nodes in the graph are normal nodes.

- edgeNodes: Object of class character. Records which nodes in the graph are edge-nodes.
- edgeNodeIO: Object of class list. Records information about which edges enter and exit each edge-node.

#### Methods

- plot signature(x = "graphBPH", y = "ANY"): draw a representation of the hypergraph where edges between normal nodes in the graph pass through an intermediate edge-node in a nice smooth curve.
- graphLayout signature(graph = "graphBPH", layoutType = "missing"): convert the graphBPH
   object to a RagraphBPH object (using a default layout method).
- graphLayout signature(graph = "graphBPH", layoutType = "character"): convert the graphBPH
  object to a RagraphBPH object (using the specified layout method).

#### Author(s)

Paul Murrell

#### References

Gansner, E.R. and and North, S.C. (1999) An open graph visualization system and its applications to software engineering, *Software - Practice and Experience*, 30:1203–1233.

Gentleman, R. and Whalen, E. and Huber, W. and Falcon, S. graph: A package to handle graph data structures.

Gentry, J. and Long, L. and Gentleman, R. and Falcon, S. and Hahne, F. and Sarkar, D. and Hansen, K. **Rgraphviz**: Provides plotting capabilities for R graph objects.

#### See Also

agopen, graphLayout and graphNEL RagraphBPH

### Examples

```
nodes <- c(LETTERS[1:5], paste("R", 1:3, sep=""))</pre>
testgnel <- new("graphNEL",</pre>
                 nodes=nodes,
                 edgeL=list(
                   A=list(edges=c("R1", "R2")),
                   B=list(edges="R2"),
                   C=list(),
                   D=list(edges="R3"),
                   E=list(),
                   R1=list(edges="B"),
                   R2=list(edges=c("C", "D")),
                   R3=list(edges="E")),
                 edgemode="directed")
testbph <- graphBPH(testgnel, "^R")</pre>
plot(testbph)
# A Hypergraph equivalent
require(hypergraph)
dh1 <- DirectedHyperedge("A", "B", "R1")</pre>
dh2 <- DirectedHyperedge(c("A", "B"), c("C", "D"), "R2")</pre>
dh3 <- DirectedHyperedge("D", "E", "R3")</pre>
hg <- Hypergraph(LETTERS[1:5], list(dh1, dh2, dh3))</pre>
plot(graphBPH(hg))
```

graphLayout

Layout a graph.

#### Description

This function is designed to layout a graph using the **Rgraphviz** package. The **hyperdraw** package makes this a generic function with a method for graphBPH objects. The function of the same name in the **Rgraphviz** package is used as a method for Ragraph objects.

## graphLayout

## Usage

```
graphLayout(graph, layoutType, ...)
```

## Arguments

graph	An graphBPH object, which is to be laid out.
layoutType	The layout method (e.g., dot or neato).
	These arguments will be passed to the agopen() function.

## Value

An RagraphBPH object.

## Author(s)

Paul Murrell

## References

Gansner, E.R. and and North, S.C. (1999) An open graph visualization system and its applications to software engineering, *Software - Practice and Experience*, 30:1203–1233.

Gentry, J. and Long, L. and Gentleman, R. and Falcon, S. and Hahne, F. and Sarkar, D. and Hansen, K. **Rgraphviz**: Provides plotting capabilities for R graph objects.

### See Also

agopen and GraphvizLayouts

#### Examples

RagraphBPH-class Class "RagraphBPH"

#### Description

The purpose of this class is to represent a laid out version of a graphBPH object. The laying out is performed by the **Rgraphviz** package. This is an intermediate step in the process of drawing a graphBPH object.

#### **Objects from the Class**

Objects of this class should be created via the graphLayout() function.

#### Slots

graph: Object of class Ragraph. The laid out graph.

allNodes: Object of class character. The names of all nodes in the graph.

nodes: Object of class character. Records normal nodes in the graph.

edgeNodes: Object of class character. Records edge-nodes in the graph.

edgeNodeIO: Object of class list. Records which edges enter and exit each edge-node.

#### Methods

- plot signature(x = "RagraphBPH", y = "ANY"): draw a representation of the hypergraph where edges between normal nodes in the graph pass through an intermediate edge-node in a nice smooth curve.
- edgeDataDefaults<- signature(self = "RagraphBPH", attr = "character", value = "ANY"):
   set the default drawing attributes for all edges.</pre>
- nodeDataDefaults<- signature(self = "RagraphBPH", attr = "character", value = "ANY"):
   set the default drawing attributes for all nodes.</pre>
- graphDataDefaults<- signature(self = "RagraphBPH", attr = "character", value = "ANY"):
   set the default drawing attributes for the graph.</pre>

#### Author(s)

Paul Murrell

## RagraphBPH-class

## See Also

graphLayout, graphBPH, and Ragraph

#### Examples

```
nodes <- c(LETTERS[1:5], paste("R", 1:3, sep=""))</pre>
testgnel <- new("graphNEL",</pre>
                  nodes=nodes,
                  edgeL=list(
                    A=list(edges=c("R1", "R2")),
                    B=list(edges="R2"),
                    C=list(),
                    D=list(edges="R3"),
                    E=list(),
                    R1=list(edges="B"),
                    R2=list(edges=c("C", "D")),
                    R3=list(edges="E")),
                  edgemode="directed")
testbph <- graphBPH(testgnel, "^R")</pre>
testrabph <- graphLayout(testbph)</pre>
edgeDataDefaults(testrabph, "lwd") <- 1
edgeData(testrabph, c("A", "R1"), c("R1", "B"), "lwd") <- c("3", 5)</pre>
edgeDataDefaults(testrabph, "color") <- "black"</pre>
edgeData(testrabph, c("A", "R1"), c("R1", "B"), "color") <- "red"</pre>
nodeDataDefaults(testrabph, "margin") <- 'unit(2, "mm")'</pre>
nodeDataDefaults(testrabph, "shape") <- "circle"</pre>
plot(testrabph)
graphDataDefaults(testrabph, "arrowLoc") <- "middle"</pre>
graphData(testrabph, "arrowLoc") <- "end"</pre>
plot(testrabph)
graphData(testrabph, "arrowLoc") <- "none"</pre>
plot(testrabph)
```

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