

hyperdraw

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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"): 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.

edgeData<- signature(`self` = "RagraphBPH", `from` = "character", `to` = "character", `attr` = "character", `value` = "ANY"): set a specific drawing attribute for one or more edges.

nodeDataDefaults<- signature(`self` = "RagraphBPH", `attr` = "character", `value` = "ANY"): set the default drawing attributes for all nodes.

nodeData<- signature(`self` = "RagraphBPH", `n` = "character", `attr` = "character", `value` = "ANY"): set a specific attribute for one or more nodes.

Author(s)

Paul Murrell

See Also[graphLayout](#), [graphBPH](#), and [Ragraph](#)**Examples**

```

nodes <- c(LETTERS[1:5], paste("R", 1:3, sep=""))
testgnel <- new("graphNEL",
  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")
testrabph <- graphLayout(testbph)
edgeDataDefaults(testrabph, "lwd") <- 1
edgeData(testrabph, c("A", "R1"), c("R1", "B"), "lwd") <- c("3", 5)
edgeDataDefaults(testrabph, "color") <- "black"
edgeData(testrabph, c("A", "R1"), c("R1", "B"), "color") <- "red"
nodeDataDefaults(testrabph, "margin") <- 'unit(2, "mm")'
nodeDataDefaults(testrabph, "shape") <- "circle"
plot(testrabph)
graphDataDefaults(testrabph, "arrowLoc") <- "middle"
graphData(testrabph, "arrowLoc") <- "end"
plot(testrabph)
graphData(testrabph, "arrowLoc") <- "none"
plot(testrabph)

```

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"`): 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 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=""))
testgnel <- new("graphNEL",
               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")
plot(testbph)

# A Hypergraph equivalent
dh1 <- DirectedHyperedge("A", "B", "R1")
dh2 <- DirectedHyperedge(c("A", "B"), c("C", "D"), "R2")
dh3 <- DirectedHyperedge("D", "E", "R3")
hg <- Hypergraph(LETTERS[1:5], list(dh1, dh2, dh3))
plot(graphBPH(hg))

```

graphBPH

Constructor for graphBPH objects

Description

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

Usage

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

Arguments

<code>graph</code>	Some form of graph that is to be converted into a <code>graphBPH</code> object.
<code>edgeNodePattern</code>	A regular expression used to distinguish between normal nodes and edge nodes.
<code>...</code>	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](#)

graphLayout	<i>Layout a graph.</i>
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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.

Usage

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

Arguments

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

Value

An `RagraphBPH` object.

Author(s)

Paul Murrell

References

Gansner, E.R. 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**

```
nodes <- c(LETTERS[1:5], paste("R", 1:3, sep=""))
testgnel <- new("graphNEL",
              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 <- new("graphBPH", testgnet, "^R")
testrabph <- graphLayout(testbph)
```

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