

# Package ‘classGraph’

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**Type** Package

**Title** Construct Graphs of S4 Class Hierarchies

**Version** 0.7-4

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**Depends** methods, graph, Rgraphviz

**Imports** methods, graph, Rgraphviz

**Suggests** Matrix

**Description** Construct directed graphs of S4 class hierarchies and visualize them. In general, these graphs typically are DAGs (directed acyclic graphs), often simple trees in practice.

**License** GPL

**Repository** CRAN

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classGraph-package      *The R Package 'classGraph'*

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### Description

The package **classGraph** is package using graph and graph visualization methods to visualize inheritance graphs of S4 classes.

### Details

Package: classGraph  
Type: Package  
Version: 0.7-0  
Date: 2007-02-10  
License: GPL

### Author(s)

Martin Maechler

### See Also

[classTree\(\)](#) is the main function of this package.

---

bGraph      *Create a "Branch Graph" (Simple Tree with Root and Leaves)*

---

### Description

Create a "Branch Graph", i.e., a simple tree with root and  $n$  (simple) branches or leaves.

### Usage

```
bGraph(n, root = "Mom",  
       leaves = paste(l.prefix, seq(length = n), sep = ""),  
       l.prefix = "D", weights = NULL,  
       mode = c("undirected", "directed"))
```

**Arguments**

|          |  |
|----------|--|
| n        | integer specifying the number of leave branches. |
| root     | the node on which to root the tree.              |
| leaves   | the nodes to be used as leaves.                  |
| l.prefix | a string specifying .....                        |
| weights  | .....  |
| mode     | string indicating which mode is to be used.      |

**Value**

a graph object of class [graphNEL](#).

**Author(s)**

Martin Maechler, Aug.2005

**See Also**

class [graphNEL](#); [ftM2graphNEL](#).

**Examples**

```
(bg7 <- bGraph(7)) # 8 nodes {Mom, D1..D7}; 7 edges
plot(bg7) # draws the graph

(bgD3 <- bGraph(3, mode="directed"))
plot(bgD3) # directed: using arrows

(bgw2 <- bGraph(2, weights = c(10,1)))
plot(bgw2) # {maybe use lwd for weights in the future?}
if(require("Matrix"))
  show(as(bgw2, "sparseMatrix")) # shows the weights
```

---

class2Graph

*Build the Graph of Super Classes from an S4 Class Definition*


---

**Description**

From an S4 class definition `class`, computes the graph of all super classes, i.e., of all classes that `class` extends.

**Usage**

```
class2Graph(class, fullNames = TRUE, simpleOnly = FALSE,
            bottomUp = FALSE, package = class@package)
```

**Arguments**

|            |   |
|------------|---|
| class      | class name  |
| fullNames  | logical indicating if full name should be applied....               |
| simpleOnly | logical, simply passed to <a href="#">getAllSuperClasses(...)</a> . |
| bottomUp   | logical indicating the <i>direction</i> of the graph.               |
| package    | package where the super classes should be gotten from.              |

**Value**

an R object inheriting from class [graph](#).

**Author(s)**

Robert Gentleman (original code) and Martin Maechler

**See Also**

[classTree](#) which builds the graph of all *subclasses*.

**Examples**

```
require("graph")
cg <- class2Graph("graphNEL") # simple : graphNEL |-> graph
plot(cg)

if(require("Matrix")) {
  cg2 <- class2Graph("dgCMatrix")
  as(cg2, "sparseMatrix")
  plot(cg2)
  ## alternative: don't show the initial "Matrix:"
  cg2. <- class2Graph("dgCMatrix", fullNames=FALSE)
  plot(cg2.)
  ## 'simpleOnly' does not change anything here :
  stopifnot(identical(cg2.,
    class2Graph("dgCMatrix", fullNames=FALSE, simpleOnly = TRUE)))

  ## very simple, since "sparseMatrix" only extends "Matrix" :
  cg3 <- class2Graph("sparseMatrix")
  plot(cg3)
}
```

---

|           |  |
|-----------|--|
| classTree | <i>builds a directed graph, typically a tree from a class Object</i> |
|-----------|--|

---

### Description

From an S4 class, by investigating all subclasses, an inheritance graph is built, a directed graph, often a tree.

### Usage

```
classTree(Cl, all = FALSE, ...)
```

### Arguments

|     |  |
|-----|--|
| Cl  | class name ...   |
| all | logical indicating if all instead of just direct sub-classes should be used. |
| ... | ....   |

### Value

an R object inheriting from class [graph](#).

### Author(s)

Martin Maechler

### See Also

[class2Graph](#), ...

### Examples

```
trGclass <- classTree("graph")
as(trGclass, "matrix")
plot(trGclass) # using package 'Rgraphviz'
```

**Description**

My constructor of an [Ragraph](#) object, a kind of “laid-out” graph, from package **Rgraphviz**. This allows more customization in plotting than just calling `plot(gr, ...)` for a [graph](#) object from package **graph**.

**Usage**

```
mRagraph(gr, lType, fixedsize = FALSE,
         fill = c("lightblue", "gray90"),
         color = c("blue3", "gray60"),
         labcol = c("blue3", "green4", "purple"))
```

**Arguments**

|                        |   |
|------------------------|---|
| <code>gr</code>        | an R object of class <a href="#">graph</a> (from package <b>graph</b> ), in our case typically the result of <code>classTree()</code> . |
| <code>lType</code>     | a string specifying the layout type, see <code>agopen()</code> in package <b>Rgraphviz</b> for the possibilities.                       |
| <code>fixedsize</code> | logical indicating if the ellipses should all get the same size – or should rather adapt to the situation.                              |
| <code>fill</code>      | character vector of length 2....  |
| <code>color</code>     | character vector of length 2....  |
| <code>labcol</code>    | vector of labels to be used ....  |

**Value**

an object of class [Ragraph](#), produced by an appropriate call to `agopen`.

**Author(s)**

Martin Maechler

**See Also**

the customized plotting function `plotRag`.

### Examples

```
if(require("Matrix")) {
  trMatrix <- classTree("Matrix")
  trMatrix
  RtrM <- mRagraph(trMatrix)
  RtrM # (the show method will hopefully improve)
  str(RtrM, max=2) # shows a bit more

  plot(RtrM)
}
```

---

numOutEdges

*For each Node of a Directed Graph give the Number Outgoing Edges*

---

### Description

In a directed or undirected graph, for each node count the number of edges “leaving” that nodes.

### Usage

```
numOutEdges(g)
```

### Arguments

`g` an R object of class `graph` (from package `graph`).

### Value

an `integer` vector the same length as `nodes(g)` giving the number of edges that “go out” from each node.

### Author(s)

Martin Maechler

### See Also

[edgeL](#) on which this function is built, and [leaves](#), both from package `graph`.

### Examples

```
## Simplistic leaves() definition for *directed graphs* :
## { compare with graph::leaves() }
is.leaf <- function(g) numOutEdges(g) == 0 ## (also exists hiddenly..)
Leaves <- function(g) nodes(g)[is.leaf(g)]
Leaves(bGraph(4, mode = "directed"))
```

---

plotRag

*Plot an Ragraph (using Rgraphviz)*


---

### Description

Plot an [Ragraph](#) object (a kind of “laid-out” graph, from package **Rgraphviz**). This simply uses the `plot` method from package **Rgraphviz** (i.e., `selectMethod(plot, "Ragraph")`) and additionally adds a “footnote”-like subtitle.

### Usage

```
plotRag(ragr, sub, subArgs = .optRagargs(), ...)

.optRagargs(side = 1, adj = 0.05, cex = 0.75, line = 3)
```

### Arguments

|                                   |   |
|-----------------------------------|---|
| <code>ragr</code>                 | an object of class <a href="#">Ragraph</a> (as defined in the <b>Rgraphviz</b> package).                                  |
| <code>sub</code>                  | a “footnote” or subtitle to be added to <code>plot(ragr, ...)</code> . By default gives the number of nodes and edges.    |
| <code>subArgs</code>              | a <a href="#">list</a> of arguments to <code>mtext</code> , typically from calling <code>.optRagargs()</code> .           |
| <code>...</code>                  | further arguments passed to <code>plot(.)</code> , i.e., the plot method for <a href="#">Ragraph</a> objects.             |
| <code>side, adj, cex, line</code> | arguments passed to <code>mtext()</code> with non-standard defaults in order to place <code>sub</code> , the “sub title”. |

### Author(s)

Martin Maechler

### See Also

[mRagraph](#), [Ragraph](#).

### Examples

```
if(require("Matrix")) {
  trMatrix <- classTree("Matrix")
  trMatrix
  RtrM <- mRagraph(trMatrix)
  RtrM # (the show method will hopefully improve)
  str(RtrM, max=2) # shows a bit more

  plot(RtrM) ## almost the same as
  plotRag(RtrM, subArgs=.optRagargs(adj = 0.5))
  ## which just gives "<n> nodes with <m> edges"
}
```

---

`subClasses`*All Subclasses of a Given S4 Class*

---

**Description**

Return all subclasses of a given S4 class; either only the direct sub classes are also those “further away” ( $distance > 1$ ).

**Usage**

```
subClasses(Cl, directOnly = TRUE, complete = TRUE, ...)
```

**Arguments**

|                         |   |
|-------------------------|---|
| <code>Cl</code>         | a class representation or a class name ( <a href="#">character</a> ).   |
| <code>directOnly</code> | logical indicating if you <i>direct</i> subclasses are desired (or also the ones with <i>distance &gt; 1</i> ). |
| <code>complete</code>   | logical,.. as in....  |
| <code>...</code>        | .....   |

**Value**

a [character](#) vector of class names.

**Author(s)**

Martin Maechler

**See Also**

[superClasses](#); [Classes](#) in general.

**Examples**

```
subClasses("graph") # -> the direct ones
subClasses("graph", directOnly = FALSE) # the same: has only direct subclasses
if(require("Matrix")) {
  print( subClasses("sparseMatrix") )
  print( subClasses("sparseMatrix", directOnly = FALSE) )# much more
}
```

---

`superClasses`*List of Super Classes of a Given S4 Class*

---

**Description**

Give a [list](#) of all super classes of a specific S4 class (definition).

**Usage**

```
superClasses(x)
```

**Arguments**

`x` a class representation as returned by [getClassDef](#).

**Value**

a list of length-1 [character](#) strings, typically with a "package" attribute each.

**Author(s)**

Robert Gentleman and Martin Maechler

**See Also**

[subClasses](#), ...

**Examples**

```
superClasses(getClassDef("graphNEL"))

if(require("Matrix")) {
  scl <- superClasses(getClassDef("dgeMatrix"))
  str(scl) # a list of two
} # 'Matrix'
```

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