Package 'NCIgraphData'

April 14, 2020

Title Data for the NCIgraph	software package		
Version 1.22.0 Created 2011-04-01 Author Laurent Jacob Maintainer Laurent Jacob < laurent.jacob@gmail.com>			
		Description Provides pathwa	ays from the NCI Pathways Database as R graph objects
		License GPL-3	
		LazyData yes	
Depends R (>= 2.10.0) Suggests Rgraphviz biocViews NCI git_url https://git.bioconductor.org/packages/NCIgraphData			
		git_branch RELEASE_3_10)
		git_last_commit 7b2f9f7	
		git_last_commit_date 2019-	-10-29
Date/Publication 2020-04-1	4		
R topics documented:			
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NCI.cyList	Raw NCI networks from Nature curated pathways and BioCarta imported as graphNEL objects		

Description

A list of graphNEL objects read from Cytoscape using the CytoscapeRPC Cytoscape plugin in combination with the RCytoscape bioconductor package. This list contains 460 of the NCI-Nature curated and BioCarta imported pathways of the NCI PID.

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Usage

```
NCI.cyList
```

Format

A list of 460 graphNEL objects.

Author(s)

Laurent Jacob

Examples

```
data("NCI-cyList")
length(NCI.cyList)
library(Rgraphviz)
plot(NCI.cyList[[1]])
```

reactome.cyList

Raw NCI networks from reactome as graphNEL objects

Description

A list of graphNEL objects read from Cytoscape using the CytoscapeRPC Cytoscape plugin in combination with the RCytoscape bioconductor package. This list contains 487 of the Reactome pathways of the NCI PID.

Usage

```
reactome.cyList
```

Format

A list of 460 graphNEL objects.

Author(s)

Laurent Jacob

Examples

```
data("reactome-cyList")
length(reactome.cyList)

library(Rgraphviz)
plot(reactome.cyList[[1]])
```

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