Package 'NCIgraphData'

October 16, 2021

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biocViews NCI git_url https://git.bioconductor.org/packages/NCIgraphData git_branch RELEASE_3_13 git_last_commit 3b4dd42 git_last_commit_date 2021-05-19 Date/Publication 2021-10-16						
				Suggests Rgraphviz		
			Depends R (>= 2.10.0)			
			License GPL-3 LazyData yes			
					Created 2011-04-01 Author Laurent Jacob Maintainer Laurent Jacob < laurent.jacob@gmail.com> Description Provides pathways from the NCI Pathways Database as R graph objects	
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Version	1 20 0					

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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