# Package 'NCIgraphData'

June 25, 2024

Title Data for the NCIgraph software package

**Version** 1.41.0

<b>Created</b> 2011-04-01		
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Description Provides pathways from the NCI Pathways Database as R graph objects		
License GPL-3		
LazyData yes		
<b>Depends</b> R ( $>= 2.10.0$ )		
Suggests Rgraphviz		
biocViews NCI		
git_url https://git.bioconductor.org/packages/NCIgraphData		
git_branch devel		
git_last_commit 00f4ca3		
git_last_commit_date 2024-04-30		
Repository Bioconductor 3.20		
Date/Publication 2024-06-25		
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NCI.cyList	Raw NCI networks from Nature curated pathways and BioCarta im-
	ported 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.

#### 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.

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### Author(s)

Laurent Jacob

## Examples

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

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

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