

Package ‘SNAGEEdata’

May 29, 2025

Version 1.44.0

Date 2012-01-26

Title SNAGEE data

Author David Venet <davenet@ulb.ac.be>

Maintainer David Venet <davenet@ulb.ac.be>

Depends R (>= 2.6.0)

Suggests ALL, hgu95av2.db, SNAGEE

Description SNAGEE data - gene list and correlation matrix

License Artistic-2.0

biocViews MicroarrayData

URL <http://fleming.ulb.ac.be/SNAGEE>

git_url <https://git.bioconductor.org/packages/SNAGEEdata>

git_branch RELEASE_3_21

git_last_commit 27e11a7

git_last_commit_date 2025-04-15

Repository Bioconductor 3.21

Date/Publication 2025-05-29

Contents

SNAGEEdata-package	2
getCC	2
Index	4

SNAGEEdata-package	<i>SNAGEE - data</i>
--------------------	----------------------

Description

Signal-to-Noise applied to Gene Expression Experiments - database of gene correlations.

Details

Package:	SNAGEEdata
Version:	0.99.0
Date:	2012-01-26
Depends:	R (>= 2.6.0)
Suggests:	SNAGEE
License:	Artistic-2.0
URL:	http://fleming.ulb.ac.be/SNAGEE

Index:

getCC	Gene-gene correlations and list of genes
-------	--

Author(s)

David Venet <davenet@ulb.ac.be>

Maintainer: David Venet <davenet@ulb.ac.be>

Examples

```
# the gene-gene correlations
cc = getCC();
```

getCC	<i>Gene-gene correlations</i>
-------	-------------------------------

Description

Get the gene-gene correlations and the list of genes.

Usage

```
getCC(mode="complete")
```

Arguments

mode Which correlations should be recovered. complete: calculated with all platforms; woAffy: calculated without the Affymetrix platforms.

Value

A list with two elements: g is the list of gene IDs, cc is the upper triangular part of the correlation matrix.

Examples

```
# Get the list of genes
geneList = getCC()$g;
```

Index

getCC, [2](#)

SNAGEEdata (SNAGEEdata-package), [2](#)

SNAGEEdata-package, [2](#)